

UNIVERSIDADE DA BEIRA INTERIOR Engineering

Biologically Motivated Keypoint Detection for RGB-D Data

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Biologically Motivated Keypoint Detection for RGB-D Data

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Dedicatory

Dedicated to the persons who love me.

Biologically Motivated Keypoint Detection for RGB-D Data

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"There is a real danger that computers will develop intelligence and take over. We urgently need to develop direct connections to the brain so that computers can add to human intelligence rather than be in opposition."

Stephen Hawking

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List of Publications

Journal Papers

1. BIK-BUS: Biologically Motivated 3D Keypoint based on Bottom-Up Saliency Filipe, S., Itti, L., Alexandre, L.A., IEEE Transactions on Image Processing, vol. 24, pp. 163-175, January 2015.

International Conference Papers

- 1. **PFBIK-Tracking: Particle Filter with Bio-Inspired Keypoints Tracking** Filipe, S., Alexandre, L.A., in IEEE Symposium on Computational Intelligence for Multimedia, Signal and Vision Processing, 2014, December 9-12, Orlando, Florida, USA.
- 2. A Biological Motivated Multi-Scale Keypoint Detector for local 3D Descriptors Filipe, S., Alexandre, L.A., in 10th International Symposium on Visual Computing, 2014, pp. 218-227, December 8-10, Las Vegas, Nevada, USA.
- 3. A Comparative Evaluation of 3D Keypoint Detectors in a RGB-D Object Dataset Filipe, S., Alexandre, L.A., in 9th International Conference on Computer Vision Theory and Applications, 2014, pp. 476-483, January 5-8, Lisbon, Portugal.

Portuguese Conference Papers

- A 3D Keypoint Detector based on Biologically Motivated Bottom-Up Saliency Map Filipe, S., Alexandre, L.A., in 20th edition of the Portuguese Conference on Pattern Recognition, 2014, October 31, Covilhã, Portugal, 2014.
- A Comparative Evaluation of 3D Keypoint Detectors
 Filipe, S., Alexandre, L.A., in 9th Conference on Telecommunications, 2013, pp. 145-148, May 8-10, Castelo Branco, Portugal, 2013.

Resumo

Com o interesse emergente na visão ativa, os investigadores de visão computacional têm estado cada vez mais preocupados com os mecanismos de atenção. Por isso, uma série de modelos computacionais de atenção visual, inspirado no sistema visual humano, têm sido de-senvolvidos. Esses modelos têm como objetivo detetar regiões de interesse nas imagens.

Esta tese está focada na atenção visual seletiva, que fornece um mecanismo para que o cérebro concentre os recursos computacionais num objeto de cada vez, guiado pelas propriedades de baixo nível da imagem (atenção *Bottom-Up*). A tarefa de reconhecimento de objetos em diferentes locais é conseguida através da concentração em diferentes locais, um de cada vez. Dados os requisitos computacionais dos modelos propostos, a investigação nesta área tem sido principalmente de interesse teórico. Mais recentemente, psicólogos, neurobiólogos e engenheiros desenvolveram cooperações e isso resultou em benefícios consideráveis. No início deste trabalho, o objetivo é reunir os conceitos e ideias a partir dessas diferentes áreas de investigação. Desta forma, é fornecido o estudo sobre a investigação da biologia do sistema visual humano e uma discussão sobre o conhecimento interdisciplinar da matéria, bem como um estado de arte dos modelos computacionais de atenção visual (bottom-up). Normalmente, a atenção visual é denominada pelos engenheiros como saliência, se as pessoas fixam o olhar numa determinada região da imagem é porque esta região é saliente. Neste trabalho de investigação, os métodos saliência são apresentados em função da sua classificação (biologicamente plausível, computacional ou híbrido) e numa ordem cronológica.

Algumas estruturas salientes podem ser usadas, em vez do objeto todo, em aplicações tais como registo de objetos, recuperação ou simplificação de dados. É possível considerar estas poucas estruturas salientes como pontos-chave, com o objetivo de executar o reconhecimento de objetos. De um modo geral, os algoritmos de reconhecimento de objetos utilizam um grande número de descritores extraídos num denso conjunto de pontos. Com isso, estes têm um custo computacional muito elevado, impedindo que o processamento seja realizado em tempo real. A fim de evitar o problema da complexidade computacional requerido, as características devem ser extraídas a partir de um pequeno conjunto de pontos, geralmente chamados pontoschave. O uso de detetores de pontos-chave permite a redução do tempo de processamento e a quantidade de redundância dos dados. Os descritores locais extraídos a partir das imagens têm sido amplamente reportados na literatura de visão por computador. Uma vez que existe um grande conjunto de detetores de pontos-chave, sugere a necessidade de uma avaliação comparativa entre eles. Desta forma, propomos a fazer uma descrição dos detetores de pontos-chave 2D e 3D, dos descritores 3D e uma avaliação dos detetores de pontos-chave 3D existentes numa biblioteca de pública disponível e com objetos 3D reais. A invariância dos detetores de pontoschave 3D foi avaliada de acordo com variações nas rotações, mudanças de escala e translações. Essa avaliação retrata a robustez de um determinado detetor no que diz respeito às mudanças de ponto-de-vista e os critérios utilizados são as taxas de repetibilidade absoluta e relativa. Nas experiências realizadas, o método que apresentou melhor taxa de repetibilidade foi o método ISS3D.

Com a análise do sistema visual humano e dos detetores de mapas de saliência com inspiração biológica, surgiu a ideia de se fazer uma extensão para um detetor de ponto-chave com base na informação de cor na retina. A proposta produziu um detetor de ponto-chave 2D inspirado pelo comportamento do sistema visual. O nosso método é uma extensão com base na cor do detetor de ponto-chave BIMP, onde se incluem os canais de cor e de intensidade de uma imagem. A informação de cor é incluída de forma biológica plausível e as características multi-escala da imagem são combinadas num único mapas de pontos-chave. Este detetor é comparado com os detetores de estado-da-arte e é particularmente adequado para tarefas como o reconhecimento de categorias e de objetos. O processo de reconhecimento é realizado comparando os descritores 3D extraídos nos locais indicados pelos pontos-chave. Para isso, as localizações do pontos-chave 2D têm de ser convertido para o espaço 3D. Isto foi possível porque o conjunto de dados usado contém a localização de cada ponto de no espaço 2D e 3D. A avaliação permitiu-nos obter o melhor par detetor de ponto-chave/descritor num *RGB-D object dataset*. Usando o nosso detetor de ponto-chave e o descritor SHOTCOLOR, obtemos uma noa taxa de reconhecimento de categorias e para o reconhecimento de objetos é com o descritor PFHRGB que obtemos os melhores resultados.

Um sistema de reconhecimento 3D envolve a escolha de detetor de ponto-chave e descritor, por isso é apresentado um novo método para a deteção de pontos-chave em nuvens de pontos 3D e uma análise comparativa é realizada entre cada par de detetor de ponto-chave 3D e descritor 3D para avaliar o desempenho no reconhecimento de categorias e de objetos. Estas avaliações são feitas numa base de dados pública de objetos 3D reais. O nosso detetor de ponto-chave é inspirado no comportamento e na arquitetura neural do sistema visual dos primatas. Os pontos-chave 3D são extraídas com base num mapa de saliências 3D bottom-up, ou seja, um mapa que codifica a saliência dos objetos no ambiente visual. O mapa de saliência é determinada pelo cálculo dos mapas de conspicuidade (uma combinação entre diferentes modalidades) da orientação, intensidade e informações de cor de forma bottom-up e puramente orientada para o estímulo. Estes três mapas de conspicuidade são fundidos num mapa de saliência 3D e, finalmente, o foco de atenção (ou "localização do ponto-chave") está sequencialmente direcionado para os pontos mais salientes deste mapa. Inibir este local permite que o sistema automaticamente orientado para próximo local mais saliente. As principais conclusões são: com um número médio similar de pontos-chave, o nosso detetor de ponto-chave 3D supera os outros oito detetores de pontos-chave 3D avaliados, obtendo o melhor resultado em 32 das métricas avaliadas nas experiências do reconhecimento das categorias e dos objetos, quando o segundo melhor detetor obteve apenas o melhor resultado em 8 dessas métricas. A única desvantagem é o tempo computacional, uma vez que BIK-BUS é mais lento do que os outros detetores. Dado que existem grandes diferenças em termos de desempenho no reconhecimento, de tamanho e de tempo, a seleção do detetor de ponto-chave e descritor tem de ser interligada com a tarefa desejada e nós damos algumas orientações para facilitar esta escolha neste trabalho de investigação.

Depois de propor um detetor de ponto-chave 3D, a investigação incidiu sobre um método robusto de deteção e *tracking* de objetos 3D usando as informações dos pontos-chave num filtro de partículas. Este método consiste em três etapas distintas: Segmentação, Inicialização do *Tracking* e *Tracking*. A segmentação é feita de modo a remover toda a informação de fundo, a fim de reduzir o número de pontos para processamento futuro. Na inicialização, usamos um detetor de ponto-chave com inspiração biológica. A informação do objeto que queremos seguir é dada pelos pontos-chave extraídos. O filtro de partículas faz o acompanhamento dos pontos-chave, de modo a se poder prever onde os pontos-chave estarão no próximo *frame*. As experiências com método PFBIK-Tracking são feitas no interior, num ambiente de escritório/casa, onde se espera que robôs pessoais possam operar. Também avaliado quantitativamente este método utilizando um "*Tracking Error*". A avaliação passa pelo cálculo das centróides dos pontos-chave e das partículas. Comparando o nosso sistema com o método de *tracking* que existe na biblioteca

usada no desenvolvimento, nós obtemos melhores resultados, com um número muito menor de pontos e custo computacional. O nosso método é mais rápido e mais robusto em termos de oclusão, quando comparado com o *OpenniTracker*.

Palavras-chave

Atenção Visual; Sistema Visual Humano; Saliência; Região de Interesse; Visão Computacional Biologicamente Motivada; Detetores de Ponto-Chave; Pontos de Interesse; Reconhecimento de Objetos 3D; Extração de Características; Avaliação da Performance; Tracking; Filtro de Particulas; Aprendizagem Automática.

Resumo Alargado

Este capítulo resume, de forma alargada e em Língua Portuguesa, o trabalho de investigação descrito na tese de doutoramento intitulada "*Biologically Motivated Keypoint Detection for RGB-D Data*". A parte inicial deste capítulo descreve o enquadramento da tese, o problema abordado, os objetivos do doutoramento, o argumento da tese, e descreve as suas principais contribuições. De seguida, é abordado o tópico de investigação sobre a deteção de pontos-chave e são apresentados com maior detalhe os trabalhos de investigação e as principais contribuições da tese. O capítulo termina com a discussão breve das principais conclusões e a apresentação de algumas linhas de investigação futura.

Introdução

Esta tese aborda o tema da deteção de ponto-chave, propondo novos métodos com inspiração biológica e uma avaliação contra os métodos do estado-da-arte num sistema de reconhecimento de objetos. O contexto e o foco da tese são ainda descritos neste capítulo, em conjunto com a definição do problema, motivação e objetivos, o estado da tese, as principais contribuições, bem como a organização de tese.

Motivação e Objetivos

Vivemos num mundo cheio de dados visuais. O fluxo contínuo de dados visuais está constantemente a bombardear as nossas retinas e precisa de ser processado de forma a extrair apenas a informação que é importante para as nossas ações. Para selecionar as informações importantes a partir da grande quantidade de dados recebidos, o cérebro deve filtrar as suas entradas. O mesmo problema é enfrentado por muitos sistemas técnicos modernos. Os sistemas de visão computacional precisam de lidar com um grande número de pixels em cada imagem, bem como com a elevada complexidade computacional das muitas abordagens relacionadas com a interpretação dos dados numa imagem [1]. A tarefa torna-se especialmente difícil, se um sistema tem de funcionar em tempo real.

A atenção visual seletiva fornece um mecanismo para que o cérebro seja capaz de concentrar os recursos computacionais num único objeto de cada vez, guiados pelas propriedades da imagem de baixo nível (atenção *Bottom-Up*) ou com base numa tarefa específica (atenção *Top-Down*). O reconhecimento dos objetos em diferentes localizações é conseguido através da concentração da atenção em diferentes locais, um de cada vez. Durante muitos anos, a investigação nesta área teve principalmente um interesse teórico, dadas as exigências computacionais dos modelos apresentados. Por exemplo, Koch e Ullman [2] apresentaram o primeiro modelo teórico de atenção seletiva em macacos, mas foi só Itti et al. [3] que conseguiram reproduzir este modelo num computador. Desde então, o poder computacional aumentou substancialmente, permitindo o aparecimento de mais implementações de sistemas computacionais de atenção, que são úteis em aplicações práticas.

No início, esta tese pretende apresentar ambas as faces dos sistemas de atenção visual, da neurociência aos sistemas computacionais. Para os investigadores interessados em sistemas computacionais de atenção, o conhecimento da neurociência sobre a atenção visual humana é dada no capítulo 2. Enquanto que para os neurocientistas são apresentados os vários tipos de abordagens computacionais disponíveis para a simulação de atenção visual humana baseada na atenção *Bottom-Up* (capítulo 3). Este trabalho apresenta não só abordagens biologicamente plausíveis, mas também discute as abordagens computacionais e híbridas (uma mistura de conceitos biológicos e computacionais). Heinke e Humphreys [4] realizaram uma revisão dos modelos de atenção computacionais com um propósito psicológico. Por outro lado, um estudo sobre modelos computacionais inspirados na neurobiologia e psicofísica da atenção são apresentados por Rothenstein e Tsotsos [1]. Finalmente, Bundesen e Habekost [5], e mais recentemente Borji e Itti [6], apresentam uma revisão abrangente dos modelos de atenção psicológica em geral.

Uma outra área que tem atraído muita atenção na comunidade de visão computacional tem sido a deteção de ponto-chave, o desenvolvimento de uma série de métodos que são estáveis a uma ampla gama de transformações [7]. Os pontos-chave são pontos de interesse e podem ser considerados pontos que ajudam os humanos a reconhecer os objetos de uma forma computacional. Alguns deles são desenvolvidos com base em características gerais [8], mais específicas [7,9,10] ou uma mistura delas [11]. Dado o número de detetores de pontos-chave, é surpreendente como é que muitos dos melhores sistemas de reconhecimento não usam estes detetores. Em vez disso, eles processam a imagem inteira, quer pelo pré-processamento de imagem inteira de forma obter vetores de características [12], por sub-amostragem dos descritores numa grelha [13] ou pelo processamento de imagens inteiras de forma hierárquica e detetando características salientes durante processo [14]. Estas abordagens fornecem uma série de dados que ajudam a classificação, mas também introduzem uma grande quantidade de redundância [15] ou alto custo computacional [13]. Normalmente, o maior custo computacional destes sistemas está na fase de processar as características (ou descritores em 3D), por isso, faz sentido usar apenas um subconjunto não redundante de pontos obtidos a partir da imagem de entrada ou da nuvem de pontos. O custo computacional dos descritores é geralmente elevado, por isso não faz sentido extrair descritores em todos os pontos. Assim, os detetores de pontos-chave são usados para selecionar pontos de interesse sobre os quais descritores serão então computados. A finalidade dos detetores de pontos-chave é a de determinar os pontos que são diferentes, a fim de permitir que uma descrição eficiente do objeto e que continue a existir uma correspondência mesmo com variações no ponto-de-vista do objeto [16].

Motivado pela necessidade de comparar quantitativamente as diferentes abordagens de deteção de ponto-chave, num processo experimentalmente comum e bem estabelecido, dado o grande número de detetores de pontos-chave disponíveis e inspirado pelo trabalho apresentado para 2D em [17, 18] e para 3D em [19], uma comparação entre vários detetores de pontos-chave 3D é feita neste trabalho. Em relação ao trabalho apresentado em [17, 19], a novidade consiste no uso de um conjunto de dados real em vez de um artificial, maior número de nuvens de pontos 3D e detetores de pontos-chave diferentes. A vantagem de usar nuvens de pontos 3D reais é que estas refletem o que acontece na vida real, como com a visão do robô. Estes nunca "vêem" um objeto perfeito ou completo, como os apresentados por objetos artificiais. Para avaliar a invariância dos métodos de deteção de ponto-chave, os pontos-chave são extraídos diretamente da nuvem inicial. Além disso, uma transformação é aplicada à nuvem de pontos 3D original antes de extrair um outro conjunto de pontos-chave. Uma vez obtidos os pontos-chave da nuvem de pontos transformada, é possível aplicar uma transformação inversa, de modo que possam ser comparados com os pontos-chave extraídos a partir da nuvem inicial. Se um dado método for invariante à transformação aplicada, os pontos-chave extraídos diretamente da nuvem original devem corresponder aos pontos-chave extraídos a partir da nuvem onde a transformação foi aplicada.

O interesse sobre o uso de informações da profundidade em aplicações de visão computacional vem crescendo recentemente, devido à diminuição dos preços das câmaras 3D, como a Kinect ou a Asus Xtion. Com este tipo de câmaras, é possível fazer uma análise 2D e 3D dos objetos capturados. As informações de profundidade melhora a perceção do objeto, uma vez que permite determinar de sua forma ou geometria. As câmaras podem retornar diretamente a imagem 2D e a nuvem de pontos correspondente, a qual é composta pela informação RGB e profundidade. Informações de profundidade melhoram a percepção do objeto, uma vez que permite a determinação de sua forma ou geometria. Um recurso útil para os utilizadores deste tipo de sensores é a biblioteca PCL [20] que contém muitos algoritmos que lidam com dados das nuvens de pontos, desde a segmentação ao reconhecimento. Esta biblioteca é utilizada para lidar com dados reais em 3D e também para avaliar a robustez dos detetores com variações no ponto-de-vista dos dados reais em 3D.

Nesta tese, é apresentado um novo detetor de ponto-chave em 2D. O método possui motivação biológica e multi-escala, que usa os canais da cor e da intensidade de uma imagem. Este tem por base o método Biologically Inspired keyPoints (BIMP) [7], o qual é um detetor rápido de ponto-chave com base na biologia do córtex visual humano. A extensão deste método é feita introduzindo a análise da cor de forma similar ao que é feito na retina humana. A avaliação comparativa é feita no conjunto de dados *RGB-D Object Dataset* [21], composto por 300 objetos reais e divididos em 51 categorias. A avaliação do método apresentado e dos detetores de pontos-chave do estado-da-arte é feita com base no reconhecimento do próprio objeto e da sua categoria utilizando descritores 3D. Este conjunto de dados contém a localização de cada ponto no espaço 2D, o que nos permite usar detetores de ponto-chave 2D nestas nuvens de pontos.

Aqui também é proposto um detetor de ponto-chave para 3D derivado de um modelo de deteção de saliências baseado na atenção espacial numa arquitetura biologicamente plausível proposta em [2, 3]. Este utiliza os três canais de características: a cor, a intensidade e a orientação. O algoritmo computacional deste modelo de saliência foi apresentado em [3] e continua a ser a base de muitos modelos posteriores e o detetor de saliência padrão nas imagens 2D. Neste trabalho é apresentado a versão 3D deste detetor de saliência e foi demonstrado como podem ser extraídos os pontos-chave a partir de um mapa de saliência. Os detetores de pontos-chave 3D e os descritores comparados podem ser encontrados na versão 1.7 da PCL [20]. Com isso, é possível encontrar o qual é o melhor par de detetor ponto-chave/descritor para objetos em nuvem de pontos 3D. Isto é feito a fim de superar as dificuldades que surgem quando se pretende escolher o par mais adequado para uso numa determinada tarefa. Este trabalho propõe-se a responder a esta pergunta com base no conjunto de dados *RGB-D Object Dataset*.

Em [22], o Alexandre foca-se nos descritores disponíveis na PCL, explicando como eles funcionam e fez uma avaliação comparativa sobre o mesmo conjunto de dados. Ele compara descritores com base em dois métodos de extração de ponto-chave: o primeiro é um detetor de ponto-chave e a segunda abordagem consiste apenas numa sub-amostragem da nuvem de pontos de entrada com dois tamanhos diferentes, usando uma *voxelgrid* com 1 e 2 cm. Os pontos sub-amostradas são considerados pontos-chave. Uma das conclusões deste trabalho é que o aumento do número de pontos-chave melhora os resultados de reconhecimento à custa do aumento do tamanho e custo computacional. O mesmo autor estuda a precisão das distâncias, tanto para os reconhecimento dos objetos bem como das suas categorias [23].

As propostas desta tese terminam com um sistema de *tracking* de pontos-chave. O *tracking* é o processo de seguir objetos em movimento ao longo do tempo usando uma câmara. Existe uma vasta gama de aplicações para estes sistemas, tais como, aviso de colisão de veículos, robótica móvel, localização de um orador, seguimento de pessoas e de animais, o acompanhamento de um alvo militar e imagens médicas. Para realizar o seguimento, o algoritmo analisa as sequências de imagens de vídeo e emite a localização dos alvos em cada uma das imagens.

Existem duas componentes principais num sistema de seguimento visual: representação do alvo e a sua filtragem. A representação do alvo é principalmente um processo *bottom-up*, ao passo que a filtragem é principalmente um processo *top-down*. Estes métodos fornecem uma variedade de ferramentas para identificar o objeto em movimento. Alguns algoritmos de seguimento mais comuns são: *Blob tracking, Kernel-based* ou *mean-shift tracking* e *contour tracking*. A filtragem envolve a incorporação de informação prévia sobre a cena ou sobre o objeto, tem de lidar com a dinâmica de objetos e realizar uma avaliação das diferentes hipóteses. Estes métodos permitem o seguimento de objetos complexos juntamente com a interação objeto mais complexo como seguir objetos em movimento atrás de obstáculos [24]. Nesta tese, a informação é fornecida diretamente por uma câmara Kinect. Com esta câmara, não é necessário gastar recursos computacionais para produzir o mapa de profundidade, uma vez que este é fornecido diretamente pela câmara. Na visão estéreo tradicional, com duas câmaras, colocadas horizontalmente uma da outra são utilizadas para obter dois pontos de vista diferentes de uma cena, de uma maneira semelhante à visão binocular humana.

Principais Contribuições

Esta secção descreve brevemente as quatro principais contribuições científicas resultantes do trabalho de investigação apresentado nesta tese.

A primeira contribuição é a descrição e a avaliação da invariância de detetores de pontoschave 3D que estão disponíveis publicamente na biblioteca PCL. A invariância de detetores de pontos-chave 3D é avaliada de acordo com várias rotações, mudanças de escala e translações. Os critérios de avaliação utilizados são a taxa de repetibilidade absoluta e a relativa. Usando estes critérios, a robustez dos detetores é avaliada em relação às mudanças do ponto-de-vista. Este estudo é parte do capítulo 4, que consiste num artigo publicado na 9th Conference on Telecommunications (Conftele'13) [25] e estendido para a 9th International Joint Conference on Computer Vision, Imaging and Computer Graphics Theory and Applications (VISAPP'14) [26].

A segunda contribuição desta tese é a proposta de um detector de ponto-chave 2D, que contém inspiração biológica. O método é uma extensão da colorimétrica do detetor de pontoschave BIMP [7], onde a informação de cor está incluído em uma maneira plausível biológica e reproduz a informação como a cor é analisada na retina. Características da imagens em várias escalas são combinadas num único mapa pontos-chave. O detetor é comparado com os detetores do estado-da-arte e é particularmente adequado para tarefas como o reconhecimento de categorias e de objetos. Com base nesta comparação, foi obtido o melhor par de detetor de ponto-chave 2D/descritor 3D no conjunto de dados *RGB-D Object Dataset*. Este detetor de ponto-chave 2D é apresentado no capítulo 5 e foi publicado na 10th International Symposium on *Visual Computing (ISVC'14)* [27].

A terceira contribuição desta tese consiste num detetor de ponto-chave 3D baseado na saliência e inspirado pelo comportamento e arquitetura neural do sistema visual dos primatas. Os pontos-chave são extraídos com base num *bottom-up* mapa de saliências 3D, ou seja, um mapa que codifica a saliência dos objetos no ambiente visual. O mapa de saliência é determinado pelo cálculo de mapas conspicuidade (uma combinação entre diferentes modalidades) da orientação, intensidade e informações de cor num processo *bottom-up* e de uma maneira puramente orientada para o estímulo. Estes três mapas de conspicuidade são fundidos num único mapa de saliência 3D e, finalmente, o foco de atenção (ou "localização dos pontos-chave") é

sequencialmente direcionado para os pontos mais salientes deste mapa. A inibição de este local permite que o sistema seja capaz de automaticamente mudar o foco de atenção para o próximo local mais saliente. A análise comparativa entre cada par de detetores de ponto-chave 3D e descritores 3D é realizada, a fim de avaliar o seu desempenho no reconhecimento de objetos e categorias. Este detetor de ponto-chave 3D é descrito no capítulo 6, que consiste num artigo publicado na 20th Portuguese Conference on Pattern Recognition (RecPad'14) [28] e estendido para a IEEE Transaction on Image Processing (IEEE TIP) [29].

A última contribuição desta tese é a proposta de um sistema robusto de deteção e de seguimento (*tracking*) de objetos 3D usando informações dos pontos-chave num filtro de partículas. O método é composto por três etapas distintas: Segmentação, Inicialização do *Tracking* e *Tracking*. Um passo da segmentação é realizado para remover toda a informação de fundo, a fim de reduzir o número de pontos para os processamentos posteriores. A informação inicial do objeto a ser seguido é dado pelos pontos-chave extraídos. O filtro de partículas faz o acompanhamento dos pontos-chave, de modo a que se possa prever onde é que este se encontrará na próximo imagem. Este *tracker* é apresentado no capítulo 7 e publicado na 10th IEEE Symposium Series on Computational Intelligence (IEEE SSCI'14) [30].

Organização de Tese

Esta tese está organizada em oito capítulos principais. O primeiro capítulo descreve o contexto, foco e o problema abordado no trabalho de investigação, bem como a motivação da tese, objetivos, declaração e a abordagem adotada para resolver o problema. Também está incluído um resumo das principais contribuições desta tese, seguido da descrição da organização e estrutura da tese. Os temas e a organização dos principais capítulos restantes desta tese são apresentados a seguir.

O capítulo 2 fornece uma visão geral sobre o sistema visual humano, descrevendo como é feito o processamento dos sinais visuais captadas pelos nossos olhos. A descrição é baseada na opinião de neurocientistas e psicólogos, sendo mais focado numa área de atenção visual humana. Este capítulo é adicionado nesta tese, a fim de dar suporte à análise das diferenças entre as aplicações com inspiração biológica e as computacionais apresentadas no capítulo 3.

No capítulo 3 é apresentado o estado-da-arte de métodos *bottom-up* de deteção de saliência. Estes são classificados com base na sua inspiração: biologicamente plausível, puramente computacional, ou híbrido. Quando um método é classificado como biologicamente plausível, significa que resulta do conhecimento sobre o sistema visual humano. Outros métodos são puramente computacionais e não com base em qualquer dos princípios biológicos da visão. Os métodos classificados como híbridos são aqueles que incorporam ideias parcialmente baseados em modelos biológicos.

O capítulo 4 é composto por três partes: 1) descrição dos detetores de ponto-chave 2D e 3D que serão usados em capítulos posteriores; 2) descrição de descritores 3D que serão usados para avaliar os detetores de ponto-chave e obter o melhor par de detetor de ponto-chave/descritor no reconhecimento de objetos; 3) por fim, uma avaliação da repetibilidade dos detetores de ponto-chave 3D, a fim de medir a invariância dos métodos relativamente à rotação, mudança de escala e translação.

No capítulo 5 é apresentado um novo método de deteção de ponto-chave com inspiração biológica e compara com os métodos do estado-da-arte numa perspetiva de reconhecimento de objetos. Uma extensão de cor com base na retina foi desenvolvido para um detetor de ponto-chave existente na literatura e inspirado no sistema visual humano.

Enquanto que no capítulo 6 é proposto um detetor de ponto-chave 3D baseado num método *bottom-up* de deteção de saliência e avaliados da mesma forma como apresentado anteriormente. Os mapas de conspicuidade obtidos a partir da intensidade e orientação são então fundidas a fim de produzir o mapa de saliência. Com isso, a atenção pode ser direcionada para o ponto mais saliente e considerá-lo um ponto-chave.

O capítulo 7 apresenta um método de *tracking* de pontos-chave 3D que usa um filtro de partículas e é composto por três etapas principais: Segmentação, Inicialização do *Tracking* e *Tracking*. Este método é comparado com um outr que está presente na biblioteca utilizada. As experiências são feitas no interior, num ambiente de escritório/casa, onde se espera que robôs pessoais possam operar.

Por fim, o capítulo 8 apresenta as conclusões e contribuições desta tese e discute direções para um futuro trabalho de investigação.

Atenção Visual Humana

O capítulo 2 aborda o tema da atenção visual humana por parte dos neurocientistas e psicólogos, a fim de facilitar a compreensão de como é feito o processamento de informações no sistema visual humano. A maioria da informação vem de uma área normalmente referida como *Computational Neuroscience* e definida por Trappenberg como: *"o estudo teórico do cérebro usado para descobrir os princípios e mecanismos que orientam as capacidades de desenvolvimento, organização, processamento das informações mentais e do sistema nervoso"* [31].

Sistema Visual

Nesta secção é apresentada uma introdução sobre a anatomia e fisiologia do sistema visual. Informações mais detalhadas podem ser encontrados em, por exemplo, Hubel [32] e Kolb et al. [33].

Retina

A retina é uma parte do cérebro responsável pela formação de imagens, isto é, o sentido da visão [32]. Em cada retina há cerca de 120 milhões de fotoreceptores (cones e bastonetes) que libertam moléculas neurotransmissoras a uma taxa que é máxima na escuridão e diminui, de forma logarítmica, com o aumento da intensidade da luz. Este sinal é então transmitido para uma rede local de células bipolares e células ganglionares.

Há cerca de 1 milhão de células ganglionares na retina e nos seus *axons* que formam o nervo ótico (ver figura 2.1). Há, portanto, cerca de 100 fotoreceptores por célula ganglionar; no entanto, cada uma das células do gânglio recebe sinais de um campo recetivo na retina, uma área mais ou menos circular que cobre milhares de fotoreceptores.

Uma imagem é produzida pela excitação dos bastonetes e cones da retina. A excitação é processada por várias partes do cérebro que funcionam em paralelo, para formar uma representação do ambiente externo no cérebro.

Os bastonetes, que são muito mais numerosos do que os cones e são responsáveis pela nossa visão com pouca luz, sendo que com luz do dia estes não contribuem para a formação da imagem [34,35]. Por outro lado, os cones não respondem à luz fraca, mas são responsáveis pela nossa capacidade de ver detalhes finos e para a nossa visão a cores [32].

A retina, ao contrário de uma câmara, não envia apenas uma imagem para o cérebro. Esta codifica espacial (comprime) a imagem para a ajustar à capacidade limitada do nervo ótico. A compressão é necessária porque há 100 vezes mais células fotoreceptoras do que ganglionares. Na retina, a codificação espacial é realizada pelas estruturas *center-surround* implementadas pelas células bipolares e ganglionares. Existem dois tipos de estruturas *center-surround* na retina (ver figura 2.2): *ON-Center* e *OFF-Center*. As *ON-Center* utilizam um centro de com peso positivo e um peso negativo na vizinhança. As *OFF-Center* usam exatamente o oposto. A pesagem positiva é mais conhecida como excitadora e pela negativa o inibidora [32].

Estas estruturas *center-surround* não são físicas, no sentido em que podem ser vistas através da coloração de tecidos e a análise das amostras anatómicas da retina. As estruturas *center-surround* são apenas lógicas (isto é, matematicamente abstratas) no sentido em que elas dependem da força da conexão entre as células bipolares e ganglionares. Acredita-se que a força de ligação entre as células depende do número e tipos de canais de iões incorporados nas sinapses entre as células bipolares. Kuffler, na década de 1950, foi o primeiro a começar a entender as estruturas *center-surround* na retina dos gatos

As estruturas *center-surround* são matematicamente equivalentes aos algoritmos de deteção de arestas utilizados por programadores de computador para extrair ou reforçar os contornos de uma imagem. Assim, a retina realiza operações sobre as arestas dos objetos dentro do campo visual. Depois da imagem ser espacialmente codificada pelas estruturas *center-surround*, o sinal é enviado através do nervo ótico (isto é, dos *axons* das células do gânglio) para o quiasma através do *Lateral Geniculate Nucleus*, como apresentado na figura 2.1.

Lateral Geniculate Nucleus

O Lateral Geniculate Nucleus é o centro de transmissão primária para informações visuais recebidas da retina e encontra-se no interior do tálamo. Este recebe informações diretamente das células ganglionares da retina através do nervo ótico e do sistema de ativação reticular. O sistema de ativação reticular é uma área do cérebro responsável pela regulação da excitação (estado fisiológico e psicológico de estar acordado ou recetivo a estímulos). Os neurónios no LGN enviam seus *axons* através da radiação ótica, uma via direta para o córtex visual primário, como mostrado na figura 2.3. Nos mamíferos, os dois caminhos mais fortes que ligam o olho ao cérebro são aqueles que são projetados para a parte dorsal do LGN no tálamo e para o *superior colliculus* [36].

Tanto o LGN do hemisfério direito e como o esquerdo recebem entradas de cada um dos olhos. No entanto, cada um recebe apenas informação de uma metade do campo visual. Isto é devido aos *axons* das células do gânglio da metade interna da retina (lado nasal), atravessando para o outro lado do cérebro através do quiasma, como é apresentado na figura 2.1. Os *axons* das células ganglionares da metade exterior dos lados da retina (temporais) permanecem no mesmo lado do cérebro. Por isso, o hemisfério direito recebe informações visuais do campo visual esquerdo, e o hemisfério esquerdo recebe a informação visual do campo visual direito [37].

Córtex Visual

O córtex visual do cérebro é a parte responsável pelo processamento de informação visual. Ele está localizado no lobo occipital, na parte de trás do cérebro (ver figura 2.4). O termo córtex visual refere-se ao córtex visual primário (também conhecida como córtex estriado ou V1) e as áreas do córtex extra-estriado compreende as áreas visuais V2, V3, V4 e V5.

Biologically Motivated Keypoint Detection for RGB-D Data

Os neurónios no córtex visual permitem o desenvolvimento de uma ação quando os estímulos visuais aparecem dentro de seu campo recetivo. Por definição, o campo recetivo é a região dentro de todo o campo visual que provoca uma "potencial de ação" (na fisiologia, um potencial de ação é um evento de curta duração em que o potencial elétrico da membrana de uma célula rapidamente sobe e desce, seguindo uma trajetória consistente). Mas um determinado neurónio pode responder melhor a um subconjunto de estímulos dentro de seu campo recetivo. Por exemplo, um neurónio no V1 pode disparar a qualquer estímulo vertical no seu campo recetivo e ignorar outros tipos de estímulo. Nas áreas visuais anteriores, como no córtex *inferotemporal* (ver figura 2.3), um neurónio pode disparar apenas quando uma determinada face aparece em seu campo recetivo.

Córtex Primário (V1) O córtex visual primário é a área mais bem estudados do sistema visual. Em todos os mamíferos estudados, está localizada no pólo posterior do córtex occipital (responsável por processar os estímulos visuais), como apresentado nas figuras 2.3 e 2.4. É altamente especializada no processamento de informações sobre objetos estáticos e em movimento, e é excelente em reconhecimento de padrões.

O V1 tem um mapa bem definido de informação espacial visual. Por exemplo, nos seres humanos todo o topo do *calcarine sulcus* responde fortemente para a metade inferior do campo visual, e a parte inferior para a metade superior do campo visual. Concetualmente, o mapeamento *retinotopic* é uma transformação da imagem visual da retina para V1. A correspondência entre um determinado local em V1 no campo subjetivo da visão é muito precisa: até mesmo os pontos cegos são mapeados para o V1.

O consenso atual parece ser que as respostas iniciais de neurónios do V1 são compostas por uns conjuntos de filtros espaço-temporais seletivos. No espaço, o funcionamento do V1 pode ser pensado como sendo semelhante a muitas funções espaciais locais, transformadas de Fourier, ou mais precisamente filtros de Gabor. Teoricamente, esses filtros juntos podem realizar o processamento neural das frequências espaciais, orientações, movimentos, direções, velocidades (frequência temporal), e muitas outras características espaço-temporais.

Os neurónios do V1 também são sensíveis à organização global de uma cena [38]. Estas propriedades provavelmente resultam da repetição do processamento e conexões laterais nas pirâmides dos neurónios [39]. As conexões *feedforward* são na sua maioria de condução, e as conexões de *feedback* são as que na sua maioria modulam em seus efeitos [40, 41].

As teorias computacionais da atenção espacial no sistema visual propõem que a modulação da atenção aumenta as respostas dos neurónios em muitas áreas do córtex visual [42--44]. O lugar natural onde é possível prever um aumento precoce deste tipo é no V1 e recente as evidências do functional Magnetic Resonance Imaging (fMRI) mostram que o córtex estriado pode ser modulado pela atenção de uma maneira consistente com esta teoria [45].

Área Visual V2 A área visual V2, também denominada por córtex *prestriate* [46], é a segunda maior área do córtex visual e a primeira região dentro da área de associação visual. Esta recebe fortes conexões *feedforward* do V1 e envia fortes ligações para o V3, V4 e V5. Ele também envia forte conexões de *feedback* para o V1. Funcionalmente, o V2 tem muitas propriedades em comum com V1. Investigações recentes têm mostrado que as células no V2 mostram uma pequena quantidade de modulação à atenção (mais do que em V1, menos do que em V4), sendo definidas como padrões moderadamente complexos, e pode ser acionado por várias orientações em diferentes sub-regiões dentro de um único recetivo campo [47, 48].

Argumenta-se que todo o fluxo ventral (ver figura 2.4) é importante para a memória visual [49]. Esta teoria prevê que a memória relacionada com o reconhecimento de objetos sofre alterações e que pode resultar na manipulação do V2. Um estudo recente revelou que certas células do V2 desempenham um papel muito importante no armazenamento da informação relacionada com o reconhecimento de objetos e na conversão de memórias de curto prazo em memórias de longo prazo [50]. A maioria dos neurónios nesta área respondem a características visuais simples, como a orientação, frequência espacial, tamanho, cor e forma [51--53]. As células do V2 também responder a várias características de formas complexas, tais como a orientação dos contornos ilusórios [51] e se o estímulo provém do *foreground* ou do *background* [54, 55].

Área Visual V3 A região do córtex V3 está localizado imediatamente à frente do V2. Há ainda uma certa controvérsia sobre a extensão exata da área V3, alguns investigadores propõem que o córtex está localizado à frente do V2 e podem incluir duas ou três subdivisões funcionais. Por exemplo, Felleman et al. [56] propõem a existência de um V3 dorsal no hemisfério superior, que é distinto do V3 ventral localizado na parte inferior do cérebro. A região dorsal e ventral do V3 têm ligações distintas com outras partes do cérebro e possuem neurónios que respondem a diferentes combinações de estímulos visuais.

Área Visual V4 A área visual V4 está localizada antes do V2 e depois da área *Posterior Inferotemporal*, como se mostra na figura 2.3. V4 é a terceira área cortical no fluxo ventral e recebe fortes entradas *feedforward* do V2 e envia fortes ligações para o *Posterior Inferotemporal*. V4 é a primeira área ventral na corrente que tem uma forte modulação da atenção. A maioria dos estudos indicam que a atenção seletiva pode alterar as taxas de disparo dos neurónios do V4 em cerca de 20%. Moran e Desimone [57] caracterizaram estes efeitos, e este foi o primeiro estudo a encontrar efeitos de atenção em qualquer lugar no córtex visual [58]. Ao contrário do V1, o V4 é ajustado de forma a extrair características dos objetos de média complexidade, como formas geométricas simples, embora ninguém consiga ainda apresentar uma descrição completa dos parâmetros do V4.

Área Visual V5 ou MT A área visual V5, também conhecida como área visual Middle Temporal (MT), é uma região do córtex visual que se pensa ter um papel importante na perceção do movimento e orientações globais de alguns movimentos oculares [59]. As suas entradas incluem das áreas visuais V1, V2 e da parte dorsal do V3 [60, 61], regiões *koniocellulare* do *Lateral Geniculate Nucleus* [62]. As projeções para o MT variam um pouco, dependendo do campo visual periférico [63]. DeAmgelis e Newsome [64] argumentam que os neurónios no MT estão organizados com base em seu ajustes na disparidade binocular.

De uma forma global, o V1 é a área que fornece a entrada mais importante para o MT [59] (ver figura 2.3). No entanto, vários estudos têm demonstrado que os neurónios do MT são capazes de responder às informações visuais muitas vezes de forma seletiva [65]. Além disso, a investigação realizada por Zeki [66] sugere que certos tipos de informações visuais podem chegar MT antes mesmo de chegarem ao V1.

Atenção Visual

Nesta secção são discutidos vários conceitos sobre a atenção visual. Informações mais detalhadas podem ser encontrados em, por exemplo, Pashler [67,68], Style [69], e Johnson and

Proctor [70].

De um modo geral, embora parece que estamos a manter uma representação rica do nosso mundo visual, apenas uma pequena região da cena é analisada em detalhe, em cada momento: foco da atenção. Esta é, geralmente, mas nem sempre, a mesma região que é capturada pelos olhos [71, 72]. A ordem pela qual uma cena é analisada é determinada pelos mecanismos de atenção seletiva. Corbetta propôs a seguinte definição de atenção: "define a capacidade mental para selecionar estímulos, respostas, memórias, ou pensamentos que são comportamentalmente relevantes entre muitos outros que são comportamentalmente irrelevantes" cite Corbetta1998.

Existem duas categorias de fatores que motivam a atenção: os fatores *bottom-up* e os fatores *top-down* [73]. Corbetta e Shulman [74] analisar as evidências em redes parcialmente segregadas de áreas do cérebro que desempenham diferentes funções da atenção. A preparação e aplicação de uma meta direcionada (*top-down*) de seleção de estímulos é realizada por um sistema que inclui partes do córtex *intraparietal* e do córtex frontal superior, o que também é modulado pela deteção de estímulos. Um outro sistema, onde a seleção *top-down* não está incluída, é em grande parte lateralizado para o hemisfério direito, onde se inclui o córtex *temporoparietal* e o córtex frontal inferior. Este sistema é especializado na deteção de estímulos comportamentalmente relevantes, particularmente quando eles são salientes ou inesperados. Assim, é possível indicar que existem duas áreas separadas do cérebro que estão envolvidos na atenção. De acordo com Theeuwes [75], a influência *bottom-up* não é voluntariamente supressivo: uma região altamente salientes captura o foco de atenção, independentemente da tarefa.

Os fatores *bottom-up* derivam exclusivamente da cena visual [76]. As regiões de interesse que atraem a atenção de um modo *bottom-up* são denominadas por salientes e as características responsáveis por estas reações devem ser suficientemente discriminantes em relação às características circundantes. Além da atenção *bottom-up*, este mecanismo é também chamado a atenção exógena, automática, reflexiva, ou atenção periférica dirigida [77].

Em contraste, a atenção *top-down* é estimulada por fatores cognitivos como as expectativas de conhecimento e objetivos atuais [74]. Por exemplo, os condutores de automóveis são mais propensos a ver postos de gasolina numa rua e os ciclistas a notar a existência de ciclovias [78].

Os mecanismos de atenção *bottom-up* foram mais cuidadosamente investigados do que os mecanismos de atenção *top-down*. Uma razão é que os dados que impulsionam os estímulos são mais fáceis de controlar do que os fatores cognitivos, como o conhecimento e as expectativas, embora pouco se sabe sobre a interação entre os dois processos.

Os mecanismos de atenção seletiva no cérebro humano ainda permanecem em aberto no campo da investigação da perceção. A inexistência de uma área do cérebro exclusivamente orientada para atenção visual [79--81] é uma das descobertas mais importantes da neurofisiologia, mas a seleção visual parece estar presente em quase todas as áreas do cérebro associadas com o processamento visual [82]. Além disso, as novas descobertas indicam que muitas áreas do cérebro partilham o processamento da informações através dos diferentes sentidos e há cada vez mais evidências de que grandes partes do córtex são multi-sensoriais [83]. A rede das áreas anatómicas executa os mecanismos de atenção [74]. As opiniões divergem sobre a questão: quais são as áreas que executam determinadas tarefas.

Saliência, Modelos Computacionais para a Atenção Visual

A atenção visual seletiva, inicialmente proposta por Koch e Ullman [2], é usada por muitos modelos computacionais de atenção visual. Mapa de saliência é o termo introduzido por Itti et al. [3] no seu trabalho sobre *rapid scene analysis*, e por Tsotsos et al. [84] e Olshausen et al. [85] nos seus trabalhos sobre "atenção visual". Em alguns estudos, como por exemplo em [84, 86], o termo saliência aparece referido como "atenção visual" ou em [87, 88] como "imprevisibilidade, raridade ou surpresa". Os mapas de saliência são utilizados como sendo um mapa escalar bidimensional que representam a localização da saliência visual, independentemente do estímulo particular que faz com que a localização seja saliente [1].

Com o interesse emergente na "visão ativa", os investigadores da área da visão por computador têm-se preocupado cada vez mais com os mecanismos de atenção e propuseram numerosos modelos computacionais de atenção. Um sistema de visão ativo é um sistema que pode manipular o ponto de vista da(s) câmara(s), a fim de analisar o seu meio ambiente circundante e de forma obter uma melhor informação a partir dele.

Os métodos de deteção de saliências podem ser classificados em: biologicamente plausíveis, puramente computacionais, ou híbridos [89]. Outros tipos de categorias são descritas em [6]. Em geral, todos os métodos utilizam uma abordagem de baixo nível para determinar o contraste das regiões na imagem em relação ao seu ambiente, utilizando uma ou mais características de intensidade, cor ou orientação. Quando um método é dito biologicamente plausível que significa que este resulta do conhecimento do sistema visual humano. Geralmente, há uma tentativa de combinar elementos conhecidos, extraídos pela retina, *Lateral Geniculate Nucleus*, córtex visual primário (V1), ou por outros campos visuais (tais como V2, V3, V4 e V5). Itti et al. [3], por exemplo, a base de seu método é uma arquitetura biologicamente plausível proposta em [2], onde eles determinam o contraste *center-surround* com o abordagem Difference of Gaussians (DoG). Frintrop et al. [90] apresentam um método inspirado no método do Itti et al., mas as diferenças no *center-surround* são obtidas recorrendo a filtros quadrados e imagens integrais de forma a reduzir o tempo de processamento.

Os métodos são puramente computacionais e não possuem qualquer tipo de base nos princípios biológicos da visão. Ma and Zhang [86] and Achanta et al. [91] estimam a saliência usando as distâncias do *center-surround*. Enquanto Hu et al. [92] estimam a saliência através da aplicação de medidas heurísticas sobre medidas de saliência iniciais obtidas pelo *thresholding* do histograma dos mapas de características. A maximização da informação mútua entre as distribuições das características do centro e da vizinhança de uma imagem é feita em [93]. Hou e Zhang [94] executam o processamento no domínio das frequências.

Os métodos classificados como híbridos são aqueles que incorporam ideias que são parcialmente baseadas nos modelos biológicos. Aqui, o método de Itti et al. [3] é usado por Harrel et al. [95] de forma a gerar os mapas de características e a normalização é feita através de uma abordagem em grafos. Outros métodos utilizam abordagens computacionais como a maximização da informação [96] que representam modelos plausíveis biológicos de deteção de saliências.

Exemplos da Deteção de Saliências

Nesta secção são apresentados alguns resultados obtidos por vários métodos de deteção de saliência. A avaliação foi realizada em duas bases de dados, que vamos descrever.

A primeira base de dados, denominada por "*Toronto*", foi apresentado em [96]. Esta contém 120 imagens capturadas em ambientes fechados e ao ar livre com uma resolução de $681 \times$

511 pixels. Para o *eye tracking*, as imagens foram apresentadas aleatoriamente a 20 pessoas e entre cada imagem era apresentada uma tela cinzenta durante 2 segundos num monitor CRT de 21 polegadas e as pessoas estavam a uma distância de 0.75 metros do monitor. Os estímulos eram imagens a cores e a tarefa passava pela visualização da mesma, de forma a registarem quais eram as zonas para onde as pessoas olhavam mais.

A segunda base de dados é denominada por "*MIT*" e foi apresentada em [97]. As imagens foram coletadas a partir da "*Flicker: creative commons*" e do conjunto de dados "*LabelMe*". Nesta base de dados contém 1007 imagens, estas foram vistas livremente e a tela cinza aparecia durante 1 segundo entre cada imagem e o sistema de *eye tracking* era reajustado após cada 50 imagens.

A tabela 3.1 apresenta os mapas de saliência produzidos por 13 métodos em três imagens de cada uma das base de dados, aqui também são apresentados os tempos médios que cada um dos métodos demorou a produzir o mapa.

Aplicações

Até agora, a atenção concentrou-se nos conceitos da atenção visual humana e apresentam teorias psicológicas e neurológicas sobre o que se sabe sobre o sistema visual humano que tem influenciado os modelos de atenção computacionais. Também foi feita uma descrição da estrutura geral e das características dos modelos computacionais de atenção, dando uma visão geral do estado-da-arte nesta área. Há, no entanto, muitas aplicações tecnológicas destes modelos que foram desenvolvidos ao longo dos anos e que têm aumentado ainda mais o interesse na modelação da atenção. As aplicações que modelam a atenção estão organizadas em quatro categorias: imagem, objeto, robótica e vídeo, como mostra a tabela 3.2.

A categoria das imagens foi dividida em cinco sub-categorias: *assembling*, compressão, avaliação da qualidade, resolução e *target*. Há também algumas aplicações adaptadas para funcionarem com vídeos. A diferença entre um método que só pode funciona com imagens estáticas e outro que funciona nos vídeos está ligado à sua complexidade computacional, porque se eles pretenderem analisar todo o vídeo, o método tem de ser extremamente rápido. Além disso, as operações realizadas num vídeo utilizando mapas de saliência são muito semelhantes às usadas nas imagens.

A divisão feita para a categoria objeto é a seguinte: deteção, reconhecimento, segmentação e *tracking*. A deteção de objetos é um passo muito importante na visão de computador e isso pode ser feito através de mapas de saliência, como demonstrado por vários autores. Os métodos apresentados que se focam na segmentação utilizando os mapas de saliência são métodos que dão mais importância para às arestas dos objetos.

Detetores de Ponto-Chave, Descritores e Avaliação

Aqui é feita uma descrição de alguns detetores de pontos-chave 2D e 3D (mais focado no 3D), e também dos descritores 3D. Finalmente, uma avaliação de detetores de pontos-chave 3D (disponíveis na biblioteca PCL) são feitos com objetos reais em nuvens de pontos 3D. A invariância dos detetores de pontos chave 3D é avaliada de acordo com a rotação, mudança de escala e translação. Os critérios de avaliação utilizados são a taxa de repetibilidade absoluta e a relativa. Usando estes critérios, a robustez dos detetores é avaliada em relação às mudanças de ponto-de-vista.

Detetores de Ponto-Chave

Harris 3D, Lowe and Noble Methods O método de Harris [98] é baseado na deteção de arestas e estes tipos de métodos são caracterizados pelas variações nas intensidades. Na biblioteca PCL estão disponíveis duas variantes do detetor de pontos-chave Harris3D: estes são denominados por Lowe [99] e Noble [100]. A diferença entre eles é a função que define a resposta dos pontos-chave.

Kanade-Lucas-Tomasi (KLT) Este detetor [98] foi proposto alguns anos após o detetor Harris e possui a mesma base que o detetor Harris3D. A principal diferença é que a matriz de covariância é calculada usando os valores das intensidades, em vez dos normais da superfície.

Curvature O método de curvatura calcula as curvaturas principais da superfície em cada ponto usando os normais. A resposta dos pontos-chave utilizada para suprimir os pontos-chave mais fracos em torno dos mais fortes é o mesmo que no detetor Harris3D.

Smallest Univalue Segment Assimilating Nucleus (SUSAN) Este é um método genérico de baixo nível no processamento de imagem que, para além da deteção de cantos, também tem sido utilizado para deteção e de supressão de ruído [101].

Scale Invariant Feature Transform (SIFT) Este foi proposto em [9] e a versão 3D em [102], sendo que partilha propriedades semelhantes às dos neurónios no córtex temporal inferior que são usados no reconhecimento de objetos na visão dos primatas.

Speeded-Up Robust Features (SURF) Os autores deste método inspiraram-se no método SIFT para o desenvolver [10]. Este é baseado na soma das respostas das 2D Haar wavelet e fizeram uma utilização eficiente das imagens integrais.

Intrinsic Shape Signatures 3D (ISS3D) O ISS3D [103] é um método relacionado com a medição da qualidade das regiões. Este método utiliza a magnitude do menor valor próprio (para incluir apenas os pontos com grandes variações ao longo de cada direção principal) e a relação entre dois valores próprios sucessivos (para excluir pontos similares ao longo de direções principais).

Biologically Inspired keyPoints (BIMP) O BIMP [7] é um detetor de ponto-chave com base no córtex visual e visa resolver o do problema computacional do método apresentado em [104].

Descritores 3D

3D Shape Context O descritor 3DSC [105] é a versão 3D do descritor *Shape Context* [106] e é baseado numa grelha esférica centrada em cada ponto-chave.

Point Feature Histograms Este descritor é representado pelas normais da superfície, as estimativas da curvatura e as distâncias entre os pares de pontos [107]. Este possui uma versão que usa a informação da cor denominado por PFHRGB. **Fast Point Feature Histograms** O descritor FPFH [108, 109] é uma simplificação do PFH (definido mais à frente) e neste caso os ângulos das orientações das normais não são calculadas para todos os pares de pontos e seu vizinhos.

Viewpoint Feature Histogram Em [110], os autores propõem uma extensão do descritor FPFH, denominada por VFH (definido mais à frente). A principal diferença é que a superfície da normal é centrada na centróide e não num ponto.

Clustered Viewpoint Feature Histogram O descritor CVFH [111] é uma extensão do VFH e a ideia por trás deste é que os objetos possuem regiões estáveis.

Oriented, Unique and Repeatable Clustered Viewpoint Feature Histogram O descritor OUR-CVFH [112] é um descritor semi-global baseado no *Semi-Global Unique Reference Frames* e no CVFH, sendo que este explora a orientação fornecida pelo *reference frame* para codificar as propriedades geométricas da superfície do objeto.

Point Pair Feature O descritor PPF [113] assume que tanto a cena e como o modelo são representados como um conjunto finito de pontos orientados, onde uma normal é associada a cada ponto, este também possui uma versão que usa informação da cor denominada por PPFRGB.

Signature of Histograms of OrienTations O descritor SHOT [114] baseia-se numa assinatura de histogramas que representam características topológicas, de forma a torná-lo invariante à translação e à rotação. Em [115], eles propõem duas variantes: a primeira é uma versão que usa informação da cor, no espaço CIELab, (SHOTCOLOR); no segundo (SHOTLRF), eles codificam apenas a informação referencial local, descartando os *bins* do histograma provenientes da forma e das informações esféricas.

Unique Shape Context Uma atualização do descritor 3DSC é proposto em [116], denominado por USC. Os autores relataram que um dos problemas encontrados no 3DSC reside nas múltiplas descrições para o mesmo ponto-chave, com base na necessidade de obter tantas versões do descritor como o número de *azimuth bins*.

Ensemble of Shape Functions Em [117], eles introduziram o descritor ESF, que se baseia na forma para descrever as propriedades do objeto. Isto é feito recorrendo às três funções de forma apresentadas em [118]: o ângulo, a distância entre pontos e área.

Point Curvature Estimation O descritor PCE calcula as direções e magnitudes das principais curvaturas da superfície em cada ponto-chave.

Características dos Descritores Na tabela 4.1 são apresentadas algumas características dos descritores apresentados e é baseada naquela que é apresentada em [22]. A segunda coluna contém o número de pontos gerados por cada descritor dado um ponto da nuvem de entrada com *n* pontos Neste trabalho, a nuvem de entrada serão apenas os pontos-chave. A terceira coluna mostra o comprimento de cada ponto. A quarta coluna indica se o descritor requer o cálculo dos normais de superfície em cada ponto. A coluna 5 mostra se o método é um descritor global ou apenas local. Descritores globais requerem a noção de objeto completo, enquanto que os

descritores locais são processados localmente em torno de cada ponto-chave e trabalham sem esse pressuposto. A sexta coluna indica se o descritor é baseado na geometria ou da forma do objeto, e se a análise de um ponto é feita usando uma esfera.

Conjunto de Dados

Neste trabalho de investigação foi utilizado o conjunto de dados RGB-D Object Dataset¹ [21]. Este conjunto de dados foi coletado por meio de uma câmara RGB-D e contém um total de 207621 nuvens segmentadas. O conjunto de dados contém 300 objetos distintos capturados numa plataforma giratória em 4 diferentes poses e os objetos são organizados em 51 categorias. Exemplos de alguns objetos são apresentados na figura 4.1. É possível ver que existem alguns erros nas nuvens de pontos, isto deve-se a erros de segmentação ou ruído do sensor de profundidade (alguns materiais não refletem o infravermelho usado para obter informações de profundidade). Os objetos escolhidos são normalmente encontrados em residências e escritórios, onde se espera que robôs pessoais possam operar.

Avaliação dos Detetores de Ponto-Chave

Este trabalho é motivado pela necessidade de comparar quantitativamente diferentes abordagens para a deteção de pontos-chave numa *framework* experimental, dado o grande número de detetores de pontos-chave disponíveis. Inspirado pelos trabalhos em 2D apresentados em [17, 18] e para 3D em [19] é feita uma comparação de vários detetores de pontos-chave 3D. Em relação aos trabalhos em [17, 19], a novidade é que foi usado um conjunto de dados real em vez de um artificial, o grande número de nuvens de pontos 3D e diferentes detetores de pontoschave. A vantagem de usar nuvens de pontos 3D riais é que estas refletem o que acontece na vida real, como na visão do robô. Estes nunca "veem" um objeto perfeito ou completo, como os representados por objetos artificiais.

O sistema de avaliação dos detetores de pontos-chave utilizado é apresentado na figura 4.2. De forma a avaliar a invariância destes métodos, os pontos-chave são extraídos diretamente da nuvem inicial. Além disso, a transformação é aplicada na nuvem 3D original antes de extrair um novo conjunto de pontos-chave. Obtendo estes pontos-chave da nuvem transformada, a transformação inversa é aplicada, de modo a compará-los com os pontos-chave extraídos a partir da nuvem inicial. Se um método particular é invariante para uma determinada transformação aplicada, os pontos-chave extraídos diretamente da nuvem original devem corresponder aos pontos-chave extraídos a partir da nuvem onde a transformação foi aplicada.

A característica mais importante de um detetor de ponto-chave é a sua repetibilidade. Esta característica leva em conta a capacidade do detetor conseguir encontrar o mesmo conjunto de pontos-chave em diferentes aparições do mesmo modelo. As diferenças no modelos podem ser devido ao ruído, mudança de ponto de vista, oclusão ou por uma combinação dos anteriores. A medida repetibilidade usada nesta neste trabalho é baseada na medida utilizada em [17] para pontos chave 2D e em [19] para os pontos-chave em 3D, que são repetibilidade absoluta e relativa.

A invariância dos métodos é avaliada em relação à rotação, translação e mudança de escala. Para isto, a rotação é realizada de acordo com os três eixos (X, Y e Z). As rotações aplicadas variaram entre os 5° e os 45°, com saltos de 10°. A translação é realizada simultaneamente nos três eixos e o deslocamento da nuvem de pontos é aplicado em cada eixo e

¹Conjunto de dados público e disponível em http://www.cs.washington.edu/rgbd-dataset.

obtido aleatoriamente. Por fim, as mudanças de escala são aplicadas de forma aleatória (entre $]1 \times, 5 \times$]).

Na tabela 4.2 são apresentados alguns resultados em relação a cada detetor de pontochave aplicado às nuvens originais. A percentagem de nuvens onde os detetores de pontos-chave são extraídos com sucesso (mais do que um ponto-chave) é apresentado na coluna 2. A coluna 3 representa o número médio de pontos-chave extraídos em cada nuvem. E finalmente, o tempo médio gasto na deteção dos pontos-chave (em segundos) por cada método.

Para fazer uma comparação justa entre os detetores, todas as etapas são iguais (ver figura 4.2). As figuras 4.4, 4.5 e 4.6 mostram os resultados da avaliação dos diferentes métodos aplicados com as várias transformações. O *threshold* das distâncias analisadas variam entre [0, 2] cm, com pequenas variações entre elas e foram calculadas para 33 distâncias identicamente espaçadas. Conforme apresentado na secção 4.1, os métodos têm um conjunto relativamente grande de parâmetros a serem ajustado: os valores utilizados foram os estabelecidos por padrão na biblioteca PCL.

Extensão Colorimétrica Inspirada na Retina para um Detetor de

Ponto-Chave 2D

O BMMSKD usa a informação da cor de forma a criar uma extensão do método BIMP. A maneira pela qual se adiciona a informação de cor é baseada numa arquitetura neural do sistema visual primata [3,119]. A figura 5.1 apresenta o diagrama de blocos deste novo detetor de ponto-chave.

Para uma dada imagem a cores, são criadas três novas imagens a partir dos canais RGB, que são: RG, BY e a imagem em escala de cinza / (apresentadas na coluna da esquerda da figura 5.2). Os canais r, $g \in b$ são normalizados por / a fim de dissociar a tonalidade da intensidade. No entanto, as variações de tonalidade não são percetíveis a muito baixa luminância (e, portanto, não são salientes), logo a normalização é aplicada somente nos locais onde / é maior do que 1/10 de seu máximo ao longo de toda a imagem. Quatro canais de cores são criados: R para o canal vermelho, G para o verde, B para o azul e Y para amarelo. Em cada um dos canais de cor RG, BY e I, o detetor de ponto-chave BIMP é aplicado e são fundidos os locais dos pontos-chave.

Dada a aplicação do método BIMP em cada canal, são obtidos três conjuntos de pontoschave k_{RG} , k_{BY} e k_I e apresentados na coluna da direita entre a segunda e quarta linha da figura 5.2. A localização é considerada um ponto-chave, se existe um outro canal de cor na sua vizinhança que indica que existe um ponto-chave na região. Um exemplo do resultado da fusão é apresentada no fundo da primeira coluna na figura 5.2.

Resultados e Discussão

O processo de captura das imagens/nuvens de pontos e a segmentação são simulados pelo conjunto de dados *RGB-D Object Dataset* [21]. Foi selecionado de modo aleatório um conjunto de 5 imagens/nuvens de pontos de cada objeto distinto, num total de 1500 imagens. Deste conjunto de dados foram selecionadas 1500 imagens e com estas foi possível gerar mais de 2 milhões de comparações para cada par detetor de ponto-chave/descritor. Neste trabalho de investigação foram avaliados 60 pares (4 detetores de ponto-chave \times 15 descritores). Nesta parte da tese existe a particularidade que os detetores de ponto-chave funcionam com imagens

2D e os descritores em 3D, sendo para isso necessário fazer uma projeção dos pontos-chave para o espaço 3D.

Uma das etapas no reconhecimento é a correspondência entre um descritor de entrada (objeto a ser reconhecido) e um descritor que esteja armazenado na base de dados. A correspondência é tipicamente feita recorrendo uma função de distância entre os dois conjuntos de descritores. Em [23], são estudadas várias funções de distância, sendo que neste trabalho foi usada a medida D_6 .

A fim de realizar a avaliação do reconhecimento serão utilizados três medidas, que são as curvas ROC, AUC e DEC. Os valores para a AUC e DEC obtidos no reconhecimento das categorias e dos objetos são apresentados nas tabelas 5.3 e 5.4, e as curvas ROCs são apresentadas nas figuras 5.5 e 5.6.

Como mostra a tabela 5.3 e 5.4, o método aqui apresentado melhora os resultados do reconhecimento, tanto a nível da categoria do objeto como do próprio objeto. Comparando esta com a abordagem com a original, é possível verificar que a informação de cor apresentou uma melhoria significativa em ambos os tipos de reconhecimento.

Para o reconhecimento da categoria (tabela 5.3), o método BMMSKD, aqui apresentado, mostra piores resultados em apenas três casos para a medida AUC e em seis casos para o DEC. Nos outros pares existem melhorias significativas em comparação com os outros três métodos de deteção de pontos-chave, que também são visíveis nos gráficos da figura 5.5. O melhor resultado para o reconhecimento da categoria foi obtido pelo par BMMSKD/PFHRGB, sendo o único em que o índice DEC ultrapassou o limiar de 1.000. Além disso, o BMMSKD só apresenta uma menor AUC em comparação com o valor médio (no caso do descritor ESF), mas em termos do valor médio do DEC este já é inferior em cinco casos.

Os resultados do reconhecimento de objetos são apresentados na tabela 5.4 e nos gráficos da figura 5.6. Comparando os resultados globais do detetor de pontos-chave SIFT para o reconhecimento de categorias com os do detetor de pontos-chave SURF no reconhecimento de objetos, é possível verificar que existe uma inversão entre os resultados. Ou seja, enquanto o método SIFT apresentou melhores resultados em vários casos e o SURF não, aqui é o oposto. De forma geral, existe uma melhoria nos resultados do reconhecimento de objetos para todos os métodos, porque não existem tantas variações nos dados.

Detetor de Ponto-Chave 3D com Inspiração Biológica

O BIK-BUS é um detetor de pontos-chave baseado nos mapas de saliência. Os mapas de saliência são determinados pelo cálculo de mapas de conspicuidade da intensidade e orientação de forma *bottom-up*. Estes mapas de conspicuidade são fundidos num mapa de saliência e, por fim, o foco de atenção é sequencialmente direcionado para os pontos mais salientes neste mapa [120]. Usando esta teoria e seguindo os passos apresentados em [3, 119] é apresentado este novo detetor de pontos-chave (ver diagrama na figura 6.1).

Filtragem Linear

A parte inicial deste método é semelhante à extensão colorimétrica inspirada na retina apresentada anteriormente. Aqui, os quatro canais de cor (R, G, B and Y) e o canal da intensidade / também são usados. As pirâmides Gaussianas [121] são usadas nas escalas espaciais, que progressivamente reduzem a nuvem de pontos. Cinco pirâmides Gaussianas $R(\sigma)$, $G(\sigma)$, $B(\sigma)$, $Y(\sigma)$ and $I(\sigma)$ são criadas a partir dos canais da cor e da intensidade, onde o σ representa o desvio padrão do *kernel* Gaussiano.

As pirâmides das orientações $O(\sigma, \theta)$ são obtidas recorrendo às normais extraídas a partir do canal da intensidade *I*, onde $\theta \in \{0^o, 45^o, 90^o, 135^o\}$ são as orientações preferenciais [121]. No córtex visual primário, a resposta aos impulsos nos neurónios da orientação seletiva é aproximada por filtros de Gabor [122]. As pirâmides de orientação são criadas de uma forma semelhante aos canais de cor, mas aplicando filtros Gabor 3D com diferentes orientações θ .

Diferenças Center-Surround

Na retina, as células bipolares e ganglionares codificam a informação espacial, utilizando estruturas *center-surround*. As estruturas *center-surround* na retina podem ser descritas como *on-center* e *off-center*. O *on-center* usam um centro pesado positivamente e os vizinhos negativamente, sendo que o *off-center* usam exatamente o oposto. A pesagem positiva é mais conhecida como excitadora e a negativa como inibidora [123].

O primeiro conjunto de mapas de características está preocupado com o contraste das intensidades. Nos mamíferos, este é detetado pelos neurónios sensíveis aos centros escuros e vizinhanças brilhantes (*off-center*) ou aos centros brilhantes e vizinhanças escuras (*on-center*) [3, 122].

Para os canais de cor, o processo é semelhante e no córtex é normalmente denominado por um sistema "color double-opponent" [3]. No centro dos seus campos recetivos, os neurónios são excitados por uma cor e inibida por outra, enquanto que o inverso é verdadeiro na vizinhança. A existência de um oponente espacial e cromático entre pares de cores no córtex visual primário humano é descrito em [124]. Dado um oponente cromático são criados os mapas RG(c, s) e BY(c, s) de forma a ter em conta o oponente cromático vermelho/verde e verde/vermelho, e azul/amarelo e amarelo/azul.

Normalização

Um passo de normalização é realizado visto que não podemos combinar diretamente os diferentes mapas de características, isto porque representam diferentes dinâmicas e mecanismos de extração. Alguns objetos salientes aparecem apenas em alguns mapas, que podem ser mascarados pelo ruído ou por outro objetos menos salientes presentes num maior número de mapas. De forma a resolver este problema é utilizado um operador de normalização $\mathcal{N}(.)$. Isto promove os mapas que contêm um pequeno número de fortes atividades, e suprime os picos no mapas que possuem muitos [3].

Combinação Escalar

Os mapas de conspicuidade são a combinação dos mapas de características, para a intensidade, cor e orientação. Eles são obtidos através da redução de cada mapa para a escala quatro e uma adição ponto-a-ponto ` \oplus '. O mapa de conspicuidade para a intensidade é definido por \overline{I} e para os canais de cor por \overline{C} . Para orientação são criados inicialmente quatro mapas intermediários, que são uma combinação dos seis mapas de características para um determinado θ . Finalmente, eles são combinados num único mapa. Os três canais separados ($\overline{I}, \overline{C} \in \overline{O}$) têm uma contribuição independente no mapa de saliência e onde as características semelhantes entre eles terão um forte impacto no saliência.

Combinação Linear

O mapa final da saliência é obtido pela normalização e por uma combinação linear entre eles:

$$S = \frac{1}{3} \left(\mathcal{N}(\overline{l}) + \mathcal{N}(\overline{C}) + \mathcal{N}(\overline{O}) \right).$$
(1)

Inhibition-Of-Return

O Inhibition-Of-Return faz parte do método e é responsável pela seleção de pontos-chave. Ele deteta a localização mais saliente (máximo global) e dirige a atenção para ele, considerandoo a localização de um ponto-chave. Depois disso, o mecanismo Inhibition-Of-Return suprime este local no mapa de saliências e as suas vizinhanças num pequeno raio, de tal forma a que a atenção seja dirigida autonomamente para o próximo local mais saliente na imagem. A supressão é conseguida substituindo valores mapa de saliência com zero. O seguinte iteração vai encontrar o ponto mais saliente (novo máximo) num local diferente. Este processo iterativo é interrompido quando o máximo do mapa de saliências atinge um determinado valor mínimo, o qual é definido por um limiar. Computacionalmente, o Inhibition-Of-Return executa um processo semelhante ao de selecionar os máximos globais e locais.

Avaliação Experimental e Discussão

Porções desta avaliação, bem como as nuvens de pontos selecionadas, são as mesmas que as apresentadas no método anterior. As principais diferenças entre estas duas avaliações são relativas ao número de pares de detetores de pontos-chave/descritores avaliados e ao fato de que estes detetores de pontos-chave funcionarem diretamente sobre as nuvens de pontos e não nas imagens 2D. Aqui, é avaliado um total de 135 pares (9 detetores de ponto-chave \times 15 descritores).

As medidas de AUC e DEC são apresentadas na tabela 6.3, enquanto os ROCs para o reconhecimento das categorias e de objetos são apresentados nas figuras 6.4 e 6.5, respetivamente. A tabela 6.4 apresenta as informações sobre o número de vezes em que cada detetor de pontoschave conseguiu obter o melhor resultado no reconhecimento de categorias e de objetos, e as somas dessas contagens (na coluna total).

Em termos de tempo computacional e espaço, os requisitos dos descritores variam muito. Se a aplicação desejada necessita de desempenho em tempo real ou de usar dispositivos embebidos com recursos limitados existem alguns descritores que não podem ser consideradas.

Considerando apenas a precisão, a melhor combinação para o reconhecimento das categorias é o par BIK-BUS/PFHRGB, seguido de perto do BIK-BUS/SHOTCOLOR, ISS3D/PFHRGB e ISS3D/SHOTCOLOR, tanto em termos de AUC e DEC. Os pares BIK-BUS/PFHRGB e BIK-BUS/SHOTCOLOR têm exatamente a mesma AUC, a diferença está no DEC onde é ligeiramente superior no caso de PFHRGB. Em relação aos descritores 3DSC e SHOTLRF, o detetor de pontos-chave proposto obtém o melhor DEC, enquanto que a AUC é melhor quando se utiliza o detetor *Curvature* em ambos os descritores.

Em termos de reconhecimento de objetos, o melhor par é o BIK-BUS/PFHRGB, mas só bate a segunda melhor combinação, ISS3D/PFHRGB, porque apresenta um melhor DEC. Para os descritores SHOT e SHOTCOLOR se compararmos o detetor de ponto-chave aqui apresentado com o ISS3D obtemos melhorias de 1.5% no caso de reconhecimento de categorias, e de 1.1%

e 1.4% no reconhecimento de objetos, respetivamente. O único ponto contra é em relação ao tempo de processamento, uma vez que é de aproximadamente 6 vezes mais lento do que ISS3D. O tempo de processamento pode ser reduzido através de uma paralelização ou por uma implementação em GPGPU. A arquitetura do BIK-BUS, apresentada na figura 6.1, mostra que a paralelização seria uma boa estratégia para resolver este problema.

Aplicação de Pontos-Chave 3D no Tracking

Esta parte da tese foca-se numa aplicação dos detetores de pontos-chave no *tracking*. O método PFBIK-Tracking é apresentado na figura 7.1. Este é composto por duas etapas principais: Segmentação e *Tracking*, que serão descritas em detalhe a seguir.

A segmentação começa com o *Pass Through Filter*. Este filtro remove regiões de profundidade que não estão contidas nas distâncias de trabalho desejadas $[d_{min}, d_{max}]$, onde d_{min} é a distância mínima em que o sistema deve funcionar e d_{max} a distância máxima. As regiões com profundidades que não estejam incluídas entre essas distâncias são considerados de fundo (*background*) e são descartadas pelo nosso sistema de *tracking*. Ao retirar estas regiões (apresentado na figura 7.2(b)), que não tem informações interessantes para o sistema de *tracking* de objetos, é obtida uma considerável redução no tempo de processamento.

A segunda etapa da segmentação é uma segmentação planar, que se baseia no algoritmo RANSAC [125]. Este é um método iterativo para estimar os parâmetros de um modelo matemático do conjunto de dados observados. A distribuição dos dados *inliers* pode ser explicada por um conjunto de parâmetros do modelo, mas podem estar sujeitos a ruído e *outliers* que são dados que não se encaixam no modelo. Os *outliers* podem vir de valores extremos do ruído, de medições erradas ou hipóteses incorretas sobre a interpretação dos dados. Dada a região plana estimada por algoritmo RANSAC, é possível remover as regiões planas a partir da nuvem, mantendo apenas os restantes objetos (como apresentado na figura 7.2(c)).

Inicialização do Tracking

Na primeira nuvem de pontos capturada, de forma inicializar o *tracking*, é aplicado uma terceira etapa da segmentação que passa pela extração de um *cluster*. O objetivo é que os pontos do mesmo *cluster* tenham uma pequena distância entre eles, enquanto que os pontos em diferentes agrupamentos estejam a uma grande distância uns dos outros. Esta etapa irá retornar uma lista com os *clusters* (apresentado na figura 7.2(d)) e onde cada um contém as informações de um objeto presente na cena da nuvem.

Como mencionado anteriormente, em [26], foi apresentada uma avaliação dos detetores de pontos-chave disponíveis na biblioteca PCL. O detetor SIFT foi proposto em [9] e é representado por vetores de medições locais nas nuvens de pontos. A implementação 3D d detetor de pontos-chave SIFT (SIFT3D) foi apresentado em [102]. Ele usa uma versão 3D da Hessiana para selecionar esses pontos de interesse.

Tracking

Como apresentado na figura 7.1, o *tracking* é realizado por um filtro de partículas adaptativo apresentado em [126, 127]. Eles apresentaram uma abordagem estatística para adaptar o tamanho do conjunto de amostras dos filtros de partículas *on-the-fly*. O número de partículas
adapta-se com base na distância *Kullback-Leibler* [128], onde é interligado o erro introduzido pela representação à base da amostra do filtro de partículas. Este método irá escolher diferentes números de amostras, dependendo da densidade da nuvem de pontos 3D.

Resultados

Para avaliar o desempenho do método é calculada a distância euclidiana entre a centróide dos pontos-chave e a centróide da particulas do método. O objetivo de realizar esta comparação é verificar se um sistema é capaz de seguir os pontos-chave de um objeto. Isto é feito, a fim de não ser necessário aplicar um detetor de ponto-chave em todas as *frames*. Num sistema de tempo real, não é possível aplicar um detetor de ponto-chave em cada *frame*, devido ao custo computacional do seu cálculo.

A fim de avaliar adequadamente o desempenho do método, este será comparado com um método que realiza a sub-amostragem dos pontos na nuvem, denominado por *OpenniTracker* disponível na biblioteca PCL. É aplicado o processo de segmentação neste *tracker*, onde o resultado desta etapa é apresentado na figura 7.3. Assim, os dados de entrada são exatamente os mesmos para ambos os métodos.

A diferença entre os dois métodos é a inicialização do filtro de partículas. Considerando que um é inicializado com os resultados do detetor de ponto-chave e o *OpenniTracker* com uma sub-amostragem. Isto é uma diferença muito importante no reconhecimento de objetos, porque a sub-amostragem só reduz o número de pontos de uma forma linear, enquanto o detetor de ponto-chave faz uma redução do número de pontos com base nas características do objeto.

Os resultados são apresentados nas tabelas 7.1, 7.2 e 7.3, e foram obtidos com o conjunto de dados capturado por nós e apresentado na figura 7.3. Este conjunto de dados contém 10 objetos diferentes em movimento num total de 3300 nuvens de pontos.

Principais Conclusões

Esta tese foi focada em sistemas baseados na atenção visual humana. Os sistemas desenvolvidos tem características que foram obtidas a partir de estudos no campo da neurociência e da psicologia. Para entender essas características, uma visão geral do sistema visual humano (capítulo 2) e uma revisão dos métodos computacionais que tentam modelar atenção visual (capítulo 3) foi fornecida. O foco foi principalmente nos modelos de atenção *bottom-up*, embora alguns modelos *top-down* também foram discutidos em [129--133].

A atenção visual é um campo altamente interdisciplinar e os investigadores nesta área provêm de diferentes origens. Para os psicólogos, as investigações realizadas na área do comportamento humano é feita através do isolamento de determinadas funções específicas, a fim de compreender os processos internos do cérebro, muitas vezes resultando em teorias ou modelos psicofísicos [134]. Enquanto que os neurocientistas observam a resposta do cérebro em relação a determinados estímulos [135], usando técnicas como o fMRI, tendo portanto, uma visão direta das áreas do cérebro que estão ativas sob certas condições [45, 136, 137]. Finalmente, os engenheiros utilizam as descobertas feitas nessas áreas, e tentam reproduzi-las em modelos computacionais, de modo a que possam reduzir o tempo de processamento em algumas aplicações [42--44].

Nos últimos anos, estas diferentes áreas têm lucrado consideravelmente umas das outras. Os psicólogos usam pesquisa realizada por neurocientistas, a fim de melhorar os seus modelos de atenção, enquanto os neurocientistas usam as experiências feitas pelos psicólogos para interpretar seus dados [134]. Além disso, os psicólogos começaram a implementar modelos computacionais ou usam modelos computacionais desenvolvidos anteriormente, para verificar se eles têm um comportamento semelhante ao da perceção humana. Assim, os psicólogos tendem a melhorar a compreensão dos mecanismos e ajudam no desenvolvimento de melhores modelos computacionais.

A atenção computacional ganhou uma popularidade significativa na última década. Um dos fatores que contribuiu para o aumento na popularidade foi a melhoria dos recursos computacionais. Outra contribuição, foi os ganhos de desempenho obtidos a partir da inclusão de módulos de atenção visual (ou deteção saliência) em sistemas de reconhecimento de objetos [131, 138, 139].

A maioria das investigações apresentadas, centrou-se na componente *bottom-up* da atenção visual. Enquanto os esforços anteriores são apreciados, o campo da atenção visual ainda carece de princípios computacionais para a atenção dirigida a uma determinada tarefa. A direção promissora para investigações futuras é o desenvolvimento de modelos que levem em conta o custo computacional dependendo das exigências da tarefa, especialmente em ambientes interativos, complexos e dinâmicos. Além disso, ainda não há um entendimento nos princípios computacionais baseados na atenção visual. A solução está além do escopo de uma única área. A fim de se obter uma solução, é necessário que exista a cooperação entre as várias áreas, a partir da comunidade de aprendizagem automática, de visão por computador e também as áreas biológicas, assim como neurologia e psicologia.

A tabela 3.2 mostra algumas áreas onde foram aplicados os mapas de saliência, mas não houve referências ao fato de estes serem usados para extrair diretamente de pontos-chave, os que mais se aproximaram foram o Rodrigues e du Buf [104]. O trabalho de Ardizzone et al. [140] compara se um determinado método extrai os pontos-chave nas regiões mais salientes. Com isso, foi feita uma análise dos detetores de pontos-chave mais populares e apresentados no capítulo 4, especialmente para os que utilizam informações RGB-D. Além de uma descrição dos descritores 3D e foi feita uma avaliação de detetores de pontos-chave 3D, em dados públicos disponíveis com objetos 3D reais. A comparação experimental proposta neste trabalho delineou aspetos dos métodos do estado-da-arte para os detetores de pontos-chave 3D. Este trabalho permitiu assim avaliar qual dos métodos apresenta o melhor desempenho em termos de várias transformações (rotação, mudança de escala e de translação).

A novidade deste trabalho em comparação com os trabalhos apresentados em [17] e [19] são: é uso de um conjunto de dados real em vez de um artificial, um grande número de nuvens de pontos e diferentes detetores de pontos-chave. A vantagem de utilizar uma base de dados real é que os nossos objetos possuem "oclusões", obtidos por algum tipo de falha no sensor de infravermelhos da câmara ou do método de segmentação. Nos objetos artificiais isso não acontece, desta forma os métodos de pontos-chave podem gerar resultados melhores, mas menos realistas. Pelo contrário, as experiências realizadas refletem o que pode acontecer na vida real, como, com a visão de um robô. Em geral, SIFT3D e ISS3D produziram os melhores resultados em termos de repetibilidade e o ISS3D demonstrou ser o mais invariante.

Uma outra parte deste trabalho de investigação é descrita no capítulo 5 e abrangeu o estudo de um detetor de ponto-chave 2D num sistema de reconhecimento. Aqui também é feita a proposta de um novo método de deteção de ponto-chave que usa as informações de cor da retina, chamado BMMSKD. A informação da cor da retina foi aplicada como uma extensão ao método BIMP, a fim deste suportar a utilização das informações provenientes da cor. A avaliação da abordagem proposta foi feita em dados públicos disponíveis com objetos 3D reais. Para esta

avaliação, os detetores de ponto-chave utilizados foram desenvolvidos recorrendo à biblioteca OpenCV e os locais dos pontos em 2D foram projetados para o espaço 3D de forma a se poder usar os descritores 3D disponíveis na biblioteca PCL.

Com este trabalho, foi possível verificar que os locais de pontos chave podem ajudar ou prejudicar o processo de reconhecimento e os descritores que usam informações de cor devem ser usados em vez de um similar que use apenas informações da forma. Uma vez que existem grandes diferenças em termos de resultados no reconhecimento, de tamanho dos descritores e de custo computacional, o descritor a ser usado deve ser ajustado dependendo da tarefa desejada. Se pretender realizar o reconhecimento da categoria de um objeto ou um sistema em tempo real, a recomendação passa por usar o método SHOTCOLOR, isto porque mesmo apresentando uma taxa de reconhecimento de 7% abaixo do PFHRGB o seu custo computacional é muito menor. Por outro lado, para fazer o reconhecimento de 12.9% superior ao descritor SHOTCOLOR.

Um novo detetor de ponto-chave 3D biologicamente motivado pelo comportamento e a arquitetura neuronal do sistema visual dos primatas foi apresentado no capítulo 6. Da mesma forma que no capítulo 5, uma avaliação comparativa foi realizada entre vários detetores de pontos-chave e descritores num conjunto de dados públicos disponíveis com objetos 3D reais. O BIK-BUS é um detetor de ponto-chave que usa uma técnica computacional para determinar a atenção visual, que também são conhecidos como mapas de saliência. Os mapas de saliência são determinados por um conjunto de características *bottom-up*. A fusão desses conjuntos produziram o mapa de saliência e o foco de atenção é sequencialmente direcionado para os pontos mais salientes neste mapa, o que representa um ponto-chave.

Na avaliação, os detetores de pontos-chave 3D e os descritores 3D estão disponíveis na biblioteca PCL. Com um número médio similar de pontos-chave, o detetor de ponto-chave 3D proposto supera os outros oito detetores de pontos-chave 3D avaliados por obtendo o melhor resultado em 32 das métricas avaliadas nas experiências de reconhecimento por categoria e objeto, quando o segundo melhor detetor só obteve o melhor resultado em 8 dessas métricas (ver tabela 6.4), num total de 60 testes. A única desvantagem é o tempo computacional, uma vez que BIK-BUS é mais lento do que os outros detetores. Para um sistema em tempo real, os detetores ISS3D ou Curvatura são boas escolhas, uma vez que têm um desempenho que só é superado pelo BIK-BUS e são mais rápidos. Finalmente, em termos dos descritores, a recomendação passa pela utilização de PFHRGB ou SHOTCOLOR. PFHRGB deve ser usado se pretender um sistema de reconhecimento mais preciso e em tempo real uma boa escolha é o SHOTCOLOR porque apresenta um bom equilíbrio entre as taxas de reconhecimento e complexidade temporal.

Neste trabalho de investigação, também foi apresentado uma aplicação os detetores de pontos-chave 3D, denominado por PFBIK-Tracking, que era um sistema para realizar o seguimento dos pontos-chave. O objetivo era eliminar a necessidade de aplicação de um detetor de ponto-chave em todas as sequências de imagens que queríamos analisar. Isso porque, se os detetores de pontos-chave fossem aplicados a todos as imagens, o sistema não seria capaz de operar em tempo real. Para resolver este problema, um *tracker* de ponto-chave foi desenvolvido a fim de simular a aplicação dos detetores de pontos-chave em todos as imagens do vídeo, uma vez que o principal objetivo seria para extrair os descritores de um objeto particular na cena, de modo a executar o reconhecimento. Para isso, várias etapas de segmentação foram apresentadas, de forma a remover todo o fundo e os objetos ficarem isolados. Com os objetos segmentados, um método de *clustering* e o detetor de ponto-chave SIFT3D são aplicados, o qual foi utilizado para inicializar o filtro de partículas. O detetor de ponto-chave SIFT3D foi usada porque tem características semelhantes às do IT [99]. Depois deste ser inicializado com o objeto pretendido, só será necessário ter como entrada a saída da segmentação. Esta abordagem obteve melhores resultados do que usar o OpenniTracker disponível na biblioteca PCL, sendo um método mais rápido e robusto.

Os principais objetivos desta tese foram cumpridos mediante a apresentação dos três métodos. Juntos, os métodos propostos permitem a incorporação de características com inspiração biológica em sistemas de reconhecimento. Neste caso, as experiências foram realizadas apenas num sistema de reconhecimento de objetos, mas pode ser aplicada a outros tipos, tais como sinais biométricos 3D (um exemplo seria o uso da face em 3D).

Direções Para Trabalho Futuro

Como trabalho futuro destacamos três direções principais. A primeira seria a redução do custo computacional dos dois detetores ponto-chave apresentados. É possível considerar a paralelização do código ou uma implementação do GPGPU, a fim de reduzir o tempo computacional de BMMSKD e BIK-BUS. Esta paralelização é possível por causa da arquitetura dos métodos, mostrado nas figuras 5.1 e 6.1.

Em segundo lugar, seria uma boa ideia fornecer algumas dicas sobre o porquê de um detetor de ponto-chave ou uma combinação de um detetor de ponto-chave e descritor funcionam melhor do que os outros por um determinado teste específico. Isto pode ser feito selecionando um pequeno número dos melhores detetores de pontos-chave e descritores (com base nos resultados apresentados neste trabalho de investigação), a fim de analisar quais são as melhores pares para fazerem o reconhecimento de um tipo particular de categoria ou um objeto. Neste trabalho, uma análise foi feita a fim de abranger o conjunto de dados completo e não se concentrou num casos específicos. Esta análise não foi realizada por dois motivos: 1) o conjunto de dados usado neste trabalho é muito grande, sendo composto por 300 objetos e estes estão divididos em 51 categorias; 2) também foram avaliar 135 pares de detetor de ponto-chave/descritor e esta análise é inviável usando todos esses métodos.

Por fim, o trabalho futuro incidirá sobre o sistema de *tracking* de pontos-chave proposto e aqui há várias possibilidades ainda em aberto e que podem ser exploradas. A primeira possibilidade centra-se na a substituição do método de detetor de ponto-chave, a fim de utilizar o BIK-BUS em vez de SIFT3D. Isso deve ser feito visto que o BIK-BUS apresentou melhores resultados do que SIFT3D no *framework* de reconhecimento de objetos e categorias.

Outro ponto a explorar e melhorar neste trabalho de investigação será o conjunto de dados. Os dados a serem adicionados a este conjunto de dados passam pela adição de mais objetos e a forma como os objetos se movem na cena. Este novo conjunto de dados já foi capturado, ele contém 46 objetos diferentes, que são organizados em 24 categorias e têm vários longos períodos de oclusão (como por exemplo, fora do alcance da câmara ou atrás de caixas). A captura dos objetos foi feita usando uma câmara Kinect colocada em vários locais de uma sala e os objetos deslocavam-se sobre um carro telecomandado, a fim de ser capaz de se moverem ao longo da sala. Para utilizar este conjunto de dados ainda é necessário segmentar os objetos que estão em movimento na cena. A segmentação vai permitir a realização de várias experiência e comparações neste conjunto de dados. As primeiras experiências incluem uma avaliação similar à que foi feita nos capítulos 5 e 6, isso irá permitir consolidar os resultados apresentados nesses capítulos. Com os objetos segmentados, é possível propor uma extensão para o CLEAR MOT Metric [141]. Esta medida só está disponível para métodos 2D e não considera a profundidade

do objeto, desta forma a extensão para o 3D deverá incluir a informação da profundidade e não apenas a largura e altura dos objetos, como é feito no 2D. A diferença com a medida proposta no capítulo 7 é que ela também faz a avaliação da sobreposição entre a posição do objeto real e o estimado pelo método de *tracking*. Por fim, deverá ser feito o reconhecimento das categorias e dos objetos usando as partículas do método de *tracking* como pontos-chave, a fim de comparar com o processo de reconhecimento utilizando detetores de pontos-chave.

Abstract

With the emerging interest in active vision, computer vision researchers have been increasingly concerned with the mechanisms of attention. Therefore, several visual attention computational models inspired by the human visual system, have been developed, aiming at the detection of regions of interest in images.

This thesis is focused on selective visual attention, which provides a mechanism for the brain to focus computational resources on an object at a time, guided by low-level image properties (*Bottom-Up* attention). The task of recognizing objects in different locations is achieved by focusing on different locations, one at a time. Given the computational requirements of the models proposed, the research in this area has been mainly of theoretical interest. More recently, psychologists, neurobiologists and engineers have developed cooperation's and this has resulted in considerable benefits. The first objective of this doctoral work is to bring together concepts and ideas from these different research areas, providing a study of the biological research on human visual system and a discussion of the interdisciplinary knowledge in this area, as well as the state-of-art on computational models of visual attention (bottom-up). Normally, the visual attention is referred by engineers as saliency: when people fix their look in a particular region of the image, that's because that region is salient. In this research work, saliency methods are presented based on their classification (biological plausible, computational or hybrid) and in a chronological order.

A few salient structures can be used for applications like object registration, retrieval or data simplification, being possible to consider these few salient structures as keypoints when aiming at performing object recognition. Generally, object recognition algorithms use a large number of descriptors extracted in a dense set of points, which comes along with very high computational cost, preventing real-time processing. To avoid the problem of the computational complexity required, the features have to be extracted from a small set of points, usually called keypoints. The use of keypoint-based detectors allows the reduction of the processing time and the redundancy in the data. Local descriptors extracted from images have been extensively reported in the computer vision literature. Since there is a large set of keypoint detectors, this suggests the need of a comparative evaluation between them. In this way, we propose to do a description of 2D and 3D keypoint detectors, 3D descriptors and an evaluation of existing 3D keypoint detectors in a public available point cloud library with 3D real objects. The invariance of the 3D keypoint detectors was evaluated according to rotations, scale changes and translations. This evaluation reports the robustness of a particular detector for changes of point-of-view and the criteria used are the absolute and the relative repeatability rate. In our experiments, the method that achieved better repeatability rate was the ISS3D method.

The analysis of the human visual system and saliency maps detectors with biological inspiration led to the idea of making an extension for a keypoint detector based on the color information in the retina. Such proposal produced a 2D keypoint detector inspired by the behavior of the early visual system. Our method is a color extension of the BIMP keypoint detector, where we include both color and intensity channels of an image: color information is included in a biological plausible way and multi-scale image features are combined into a single keypoints map. This detector is compared against state-of-art detectors and found particularly well-suited for tasks such as category and object recognition. The recognition process is performed by comparing the extracted 3D descriptors in the locations indicated by the keypoints after mapping the 2D keypoints locations to the 3D space. The evaluation allowed us to obtain the best pair keypoint detector/descriptor on a RGB-D object dataset. Using our keypoint detector and the SHOTCOLOR descriptor a good category recognition rate and object recognition rate were obtained, and it is with the PFHRGB descriptor that we obtain the best results.

A 3D recognition system involves the choice of keypoint detector and descriptor. A new method for the detection of 3D keypoints on point clouds is presented and a benchmarking is performed between each pair of 3D keypoint detector and 3D descriptor to evaluate their performance on object and category recognition. These evaluations are done in a public database of real 3D objects. Our keypoint detector is inspired by the behavior and neural architecture of the primate visual system: the 3D keypoints are extracted based on a bottom-up 3D saliency map, which is a map that encodes the saliency of objects in the visual environment. The saliency map is determined by computing conspicuity maps (a combination across different modalities) of the orientation, intensity and color information, in a bottom-up and in a purely stimulusdriven manner. These three conspicuity maps are fused into a 3D saliency map and, finally, the focus of attention (or "keypoint location") is sequentially directed to the most salient points in this map. Inhibiting this location automatically allows the system to attend to the next most salient location. The main conclusions are: with a similar average number of keypoints, our 3D keypoint detector outperforms the other eight 3D keypoint detectors evaluated by achiving the best result in 32 of the evaluated metrics in the category and object recognition experiments, when the second best detector only obtained the best result in 8 of these metrics. The unique drawback is the computational time, since BIK-BUS is slower than the other detectors. Given that differences are big in terms of recognition performance, size and time requirements, the selection of the keypoint detector and descriptor has to be matched to the desired task and we give some directions to facilitate this choice.

After proposing the 3D keypoint detector, the research focused on a robust detection and tracking method for 3D objects by using keypoint information in a particle filter. This method consists of three distinct steps: Segmentation, Tracking Initialization and Tracking. The segmentation is made to remove all the background information, reducing the number of points for further processing. In the initialization, we use a keypoint detector with biological inspiration. The information of the object that we want to follow is given by the extracted keypoints. The particle filter does the tracking of the keypoints, so with that we can predict where the keypoints will be in the next frame. In a recognition system, one of the problems is the computational cost of keypoint detectors with this we intend to solve this problem. The experiments with PFBIK-Tracking method are done indoors in an office/home environment, where personal robots are expected to operate. The Tracking Error evaluates the stability of the general tracking method. We also quantitatively evaluate this method using a "Tracking Error". Our evaluation is done by the computation of the keypoint and particle centroid. Comparing our system that the tracking method which exists in the Point Cloud Library, we archive better results, with a much smaller number of points and computational time. Our method is faster and more robust to occlusion when compared to the OpenniTracker.

Keywords

Visual Attention; Human Visual System; Saliency; Regions of Interest; Biologically Motivated Computer Vision; Keypoints Detectors; Interest Points; 3D Object Recognition; Feature Extraction; Descriptors; Performance Evaluation; Tracking; Particle Filter; Machine Learning.

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- 3DSC 3D Shape Context
- AIT Anterior Inferotemporal
- AUC Area Under the ROC Curve
- BIK-BUS Biologically Inspired 3D Keypoint based on Bottom-Up Saliency
- BIMP Biologically Inspired keyPoints
- BMMSKD Biological Motivated Multi-Scale Keypoint Detector
- CE Cluster Extraction
- CM Cognitive Mapping
- CRF Conditional Random Field
- CVFH Clustered Viewpoint Feature Histogram
- CVS Computer Vision Systems
- DEC Decidability
- DoG Difference of Gaussians
- **ESF** Ensemble of Shape Functions
- FDN Frequency Domain Divisive Normalization
- FEF Frontal Eye Field
- fMRI functional Magnetic Resonance Imaging
- FOA Focus of Attention
- FPFH Fast Point Feature Histograms
- GPGPU General-Purpose Computation on Graphics Processing Unit
- I-POMDP Infomax Partially Observable Markov Decision Processes
- ICA Independent Component Analysis
- ICL Incremental Coding Length
- Infomax Information Maximization
- IOR Inhibition-Of-Return
- ISS3D Intrinsic Shape Signatures 3D
- IT Inferotemporal
- HC Histogram-based Contrast
- HVS Human Visual System
- KL Kullback-Leibler
- KLT Kanade-Lucas-Tomasi

- LGN Lateral Geniculate Nucleus
- LIP Lateral Intraparietal
- LOOCV Leave-One-Out Cross-Validation
- LTM Long-Term Memory
- MM Motor Mapping
- MST Medial Superior Temporal
- MT Middle Temporal
- OMM Observable Markov Model
- **OpenCV** Open Source Computer Vision
- **ORM** Object-Recognition Memory
- OUR-CVFH Oriented, Unique and Repeatable Clustered Viewpoint Feature Histogram
- PCA Principal Component Analysis
- PCE Principal Curvatures Estimation
- PCL Point Cloud Library
- PFBIK-Tracking Particle Filter with Bio-Inspired Keypoints Tracking
- PFDN Piecewise Frequency Domain Divisive Normalization
- PFH Point Feature Histograms
- PIT Posterior Inferotemporal
- **PNN** Probabilistic Neural Network
- PO Parieto-Occipital
- POMDP Partially Observable Markov Decision Processes
- PP Posterior Parietal
- PPF Point Pair Feature
- PQFT Phase spectrum of Quaternion Fourier Transform
- PS Planar Segmentation
- PTF Pass Through Filter
- **RANSAC** Random Sample Consensus
- RAS Reticular Activating System
- RC Region-based Contrast
- RLVC Reinforcement Learning of Visual Classes
- ROC Receiver Operator Characteristic

- ROI Region of Interest
- RPI Residual Perceptual Information
- SC Superior Colliculus
- **SDC** Shape Distribution Component
- SEF Supplementary Eye Field
- SGURF Semi-Global Unique Reference Frames
- **SHOT** Signature of Histograms of OrienTations
- SIFT Scale Invariant Feature Transform
- SM Sensory Mapping
- SPL Superior Parietal Lobe
- SUN Saliency Using Natural statistics
- SURF Speeded-Up Robust Features
- SUSAN Smallest Univalue Segment Assimilating Nucleus
- SVM Support Vector Machine
- USAN Univalue Segment Assimilating Nucleus
- USC Unique Shape Context
- VFH Viewpoint Feature Histogram
- VOCUS Visual Object Detection with a Computational Attention System

Chapter 1

Introduction

This thesis addresses the subject of keypoint detection, proposing new methods with biological inspiration and evaluating them against the state-of-art methods in an object recognition framework. The context and focus of the thesis are further described in this chapter, with the problem definition, motivation and objectives, the thesis statement, the main contributions, and the thesis organization.

1.1 Thesis Motivation and Objectives

We live in a world full of visual data. The continuous flow of visual data flowing towards our retinas needs to be processed to extract the information important for our actions. To select the important information from the large amount of data received, the brain must filter its inputs. The same problem is faced by many modern technical systems. Computer Vision Systems (CVS) have to deal with a very large number of pixels in each frame, as well as with the high computational complexity of many approaches related to the interpretation of image data [1], making the task specially difficult if the system has to function in real time.

Selective visual attention provides a mechanism for the brain to focus computational resources on an object at a time, either guided by low-level image properties (*Bottom-Up* attention) or based on a specific task (*Top-Down* attention). Recognizing objects on different locations is achieved by focusing the attention on one location at a time. For many years, research in this area has been mainly of theoretical interest, given the computational requirements of the models presented. For example, Koch and Ullman [2] presented the first theoretical model of selective attention in monkeys but only Itti et al. [3] that could reproduce this model on a computer. Since then, the computing power has increased substantially, allowing the appearance of more implementations of computational attention systems that are useful in practical applications.

First, this thesis intends to present both faces of visual attention systems, from neuroscience to computational systems. For researchers interested in computer attention systems, the necessary neuroscience knowledge on human visual attention is given (in chapter 2). While for neuroscientists, the various types of available computational approaches for the simulation of human visual attention based on *Bottom-Up* attention are presented (in chapter 3). This work presents not only the biologically plausible approaches, but discusses also the computational and hybrid approaches (a mixture of biological and computational concepts). Heinke and Humphreys [4] conducted a review of the computational attention models with a psychological purpose. On the other hand, a study on computational models of attention inspired by neurobiology and psychophysics is presented by Rothenstein and Tsotsos [1]. Finally, Bundesen and Habekost [5] presents a comprehensive review of psychological attention models in general.

An area that has attracted a lot of attention in the computer vision community is the area of keypoint detection, with the development of a series of methods which are stable under a wide range of transformations [7]. The keypoints are points of interest which can be considered to help humans on the recognition of objects in a computational way. Some of them are developed based on general features [8], specific [7,9,10] or a mixture of them [11]. Given the number of existing keypoint detectors, it is surprising that many of top recognition systems do not use these detectors. Instead, they process the entire image, either by pre-processing it to obtain feature vectors [12], by sampling descriptors on a dense grid [13] or by processing entire images hierarchically and detecting salient features in the process [14]. These approaches provide a lot of data that helps classification, but also introduce much redundancy [15] or high computacional cost [13]. Typically, the largest computational cost of these systems is in the stage of feature computation (or descriptors in 3D). Therefor, it makes sense to use a non-redundant subset of points from the input image or point cloud: as the computational cost of descriptors is generally high, it does not make sense to extract descriptors from all points. Thus, keypoint detectors are used to select interesting points which descriptors are then computed in these locations. The purpose of the keypoint detectors is to determine the points that are different in order to allow an efficient object description and correspondence with respect to point-of-view variations [16].

Motivated by the need to quantitatively compare different keypoint detector approache, in a common and well established experimentally framework inspired by the work on 2D [17, 18] and 3D features [19] a comparison of several 3D keypoint detectors is made. In relation to the work of Schmid et al. [17] and Salti et al. [19], the novelties are: it uses a real database instead of an artificial one; the large number of 3D point clouds; and different keypoint detectors. The benefit of using real 3D point clouds is that it reflects what happens in real life (e.g. robot vision). Robots never "see" a perfect or complete object, like the ones simulated by artificial objects. To evaluate the invariance of keypoint detection methods, the keypoints are extracted directly from the original cloud. Moreover, a transformation to the original 3D point cloud before extracting a second set of keypoints is applied. Once we have those keypoints from the transformed cloud, it is possible to apply an inverse transformation, so that they can be compared with the keypoints extracted from the original cloud. If a particular method is invariant to the applied transformation, the keypoints extracted directly from the original cloud should correspond to the keypoints extracted from the cloud where the transformation was applied.

The interest on using depth information in computer vision applications has been growing recently due to the decreasing prices of 3D cameras such as *Kinect* and *Asus Xtion*. This type of cameras renders, it is possible to make a 2D and 3D analysis of the captured objects, as depth information improves object perception, allowing the determination of its shape or geometry. The cameras can return directly the 2D image and the corresponding cloud point, which is composed by the RGB and depth information. Depth information improves object perception, as it allows the determination of its shape or geometry. A useful resource for users of this type of sensors is the Point Cloud Library (PCL) [20] which contains many algorithms that deal with point cloud data, from segmentation to recognition, from search to input/output. This library is used in this work to deal with real 3D data and also to evaluate the robustness of the detectors with variations of the point-of-view in real 3D data.

In this thesis, a new 2D keypoint detector is also presented. The method is a biologically motivated multi-scale keypoint detector, which uses color and intensity channels of an image. our approach is based on the Biologically Inspired keyPoints (BIMP) [7], which is a fast keypoint detector inspired by the biology of the human visual cortex, extended by introducing color analysis, similar to what is done in the human retina. A comparative evaluation is conducted on a large public RGB-D Object Dataset [21], which is composed by 300 real objects from 51 categories. The evaluation of the proposed method and the state-of-art keypoint detectors

is based on category and object recognition using 3D descriptors. This dataset contains the location of each point in the 2D space, which allows us to use 2D keypoint detector methods on the point clouds.

Furthermore, a 3D keypoint detector is also proposed, which consists on a saliency model based on spatial attention derived from the biologically plausible architecture proposed by Koch and Ullman [2] and Itty et al. [3]. It uses three feature channels: color, intensity and orientation. The computational algorithm of this saliency model has been presented in [3] and it remains the basis of later models and the standard saliency benchmark in 2D images. We present the 3D version of this saliency detector and demonstrate how keypoints can be extracted from a saliency map. The 3D keypoint detectors and descriptors that are compared can be found in version 1.7 of the PCL [20]. It is then possible to find what is the best pair of keypoint detector/descriptor for 3D point cloud objects. This is done to overcome the difficulty that arises when choosing the most suitable pair of keypoint detector and descriptor for use in a particular task, which is archived with the public RGB-D Object Dataset.

In his work [22], Alexandre focuses on the descriptors available in PCL, explaining how they work and making a comparative evaluation on the same dataset. It compares descriptors based on two methods for keypoint extraction: the first one is a keypoint detector; and the second approach consists on sub-sampling the input cloud with two different sizes, using a voxelgrid with 1 and 2 centimeter leaf size, being the sub-sampled points considered keypoints. One conclusion in his work is that the increased number of keypoints improves recognition results at the expense of size and computational time. In this study, we further explore that approach demonstrating that the results also depend on the keypoint location. The same author studies the accuracy of the distances both for objects and category recognition and finds that simple distances give competitive results [23].

This thesis ends by proposing a system for tracking keypoints, where the tracking is the process of following moving objects over time using a camera. There is a vast range of applications for tracking, such as, vehicle collision warning and avoidance, mobile robotics, speaker localization, people and animal tracking, tracking a military target and medical imaging. To perform tracking an algorithm analyzes sequential video frames and outputs the location of targets on each frame. There are two major components of a visual tracking system: target representation and filtering. Target representation is mostly a bottom-up process, whereas filtering is mostly a top-down process. These methods give a variety of tools for identifying the moving object. We identified the following as the most common target representation algorithms: Blob tracking, Kernel-based or mean-shift tracking and contour tracking. Filtering involves incorporating prior information about the scene or object, dealing with object dynamics, and evaluating different hypotheses. These methods allow the tracking of complex objects along with more complex object interaction (e.g. tracking objects moving behind obstructions [24]).

In this thesis, we use the information given by a Kinect camera directly. With this camera, it is unnecessary to spend time producing the depth map, since it is given by the camera. In traditional stereo vision systems, two cameras are placed horizontally from one another and used to obtain two differing views of the scene, in a similar manner to the human binocular vision.

1.2 Main Contributions

This section briefly describes the four main scientific contributions resulting from the research work in this thesis.

The first contribution is an in-depth analysis of 3D keypoint detectors that are publicly available in the PCL library, with their description and an evaluation of the invariance. The invariance of the 3D keypoint detectors is evaluated according to rotations, scale changes and translations. The evaluation criteria used are the absolute and the relative repeatability rate. Using these criteria, the robustness of the detectors is evaluated with for changes of point-of-view. This study is part of chapter 4, which consists of an article published in the 9th Conference on Telecommunications (Conftele'13) [25] and extended to the 9th International Joint Conference on Computer Vision, Imaging and Computer Graphics Theory and Applications (VIS-APP'14) [26].

The second contribution of this thesis is the proposal of a 2D keypoint detector containing biological inspiration. The method can be regarded as a color extension of the BIMP keypoint detector [7], where the color information is included in a biological plausible way and reproduces the color information in the retina. Multi-scale image features are combined into a single keypoints map. The detector is compared against state-of-art detectors and is particularly well-suited for tasks such as category and object recognition. The evaluation gave the best pair 2D keypoint detector/descriptor on a RGB-D object dataset. This 2D keypoint detector is presented in chapter 5 and published in the 10th International Symposium on Visual Computing (ISVC'14) [27].

The third contribution of this thesis consists of a 3D keypoint detector based on saliency and inspired by the behavior and neural architecture of the primate visual system. The keypoints are extracted based on a bottom-up 3D saliency map, which is a map that encodes the saliency of objects in the visual environment. The saliency map is determined by computing conspicuity maps (a combination across different modalities) of the orientation, intensity and color information in a bottom-up and in a purely stimulus-driven manner. These three conspicuity maps are fused into a 3D saliency map and, finally, the focus of attention (or "keypoint location") is sequentially directed to the most salient points in this map. Inhibiting this location automatically allows the system to attend to the next most salient location. A benchmarking between each pair of 3D keypoint detector and 3D descriptor is performed, to evaluate their performance on object and category recognition. These evaluations are done in a public database of real 3D objects. This 3D keypoint detector is described in chapter 6, which consists of an article published in the 20th Portuguese Conference on Pattern Recognition (RecPad'14) [28] and extended to the IEEE Transactions on Image Processing (IEEE TIP) [29].

The last contribution of this thesis is the proposal of a robust detection and tracking method for 3D objects by using keypoint information in a particle filter. The method is composed by three distinct steps: Segmentation, Tracking Initialization and Tracking. The segmentation step is performed to remove the background information, thus reducing the number of points for further processing. The initial information of the tracked object is given by the extracted keypoints. The particle filter does the tracking of the keypoints, so with that we can predict where the keypoints location will be in the next frame. This tracker is presented in chapter 7 and published in the 10th IEEE Symposium Series on Computational Intelligence (IEEE SSCI'14) [30].

1.3 Thesis Outline

This thesis is organized in eight chapters. The present chapter describes the context, focus and the problems addressed in the research work, as well as the thesis motivation, objectives, statement and the adopted approach for solving the problem. A summary of the main contributions of this doctoral program is also included, followed by a description of its organization and structure. The remainder chapters of this thesis can are summarized as follows.

Chapter 2 provides an overview on the Human Visual System (HVS), describing how it process the visual signals captured by the eyes. That description is based on the opinion of neuroscientists and psychologists, being focused on the area of human visual attention. This chapter is added in this thesis to give support to the analysis of the differences between applications with biological inspiration and computational ones, presented in chapter 3.

Chapter 3 presents the state-of-art of bottom-up saliency methods. The methods are categorized based on whether they are biologically plausible, purely computational, or hybrid. When a method is classified as biologically plausible it means that it follows the knowledge of the HVS. Other methods are purely computational and not based on any of the biological principles of vision. The methods classified as hybrid are those that incorporate ideas that are partially based on biological models.

Chapter 4 is composed by three parts: 1) description of the 2D and 3D keypoint detectors that will be used in later chapters; 2) description of 3D descriptors that will be used to evaluate the keypoint detectors and get the best pair of keypoint detector/descriptor in object recognition; 3) a repeatability evaluation of the 3D keypoint detectors, to measure the invariance of the methods relatively to the rotation, scale change and translation.

Chapter 5 presents a new keypoint detection method with biological inspiration and compares it against state-of-the art methods in object recognition. A retinal color extension was also developed for an existing keypoint detector in the literature, inspired by the HVS.

Chapter 6 proposes a 3D keypoint detector based on a biological bottom-up saliency method, which is evaluated in the same way as presented earlier. The conspicuity maps obtained from the intensity and orientation are fused in order to produce the saliency map. With that, the attention can be directed to the most salient point, thus considering a keypoint.

Chapter 7 presents a particle filter framework for 3D keypoint tracking and it is composed by three main steps: segmentation, tracking initialization and tracking. This method is compared against the one presented in the used library. The experiments are done indoors in an office/home environment, where personal robots are expected to operate.

Chapter 8 presents the conclusions and contributions of this thesis and discusses directions for future research work.

Biologically Motivated Keypoint Detection for RGB-D Data

Chapter 2

The Human Visual Attention: Neuroscientists and Psychologists Perspectives

This chapter introduces the topic of human visual attention as seen by neuroscientists and psychologists, in order to facilitate the understanding of how information processing in the HVS is done. Most of the information comes from an area commonly referred to as *Computational Neuroscience*, defined by Trappenberg as: "the theoretical study of the brain used to discover the principles and mechanisms that guide the development, organization, information processing and mental abilities of the nervous system" [31].

2.1 Visual System

In this section, an introduction on the anatomy and physiology of the visual system is presented. More detailed information can be found in, for example, Hubel [32] and Kolb et al. [33].

2.1.1 Retina

The retina is part of the brain, having been sequestered from it early in development but having kept its connections with the brain proper through a bundle of the optic nerve. It is responsible for the formation of images, i.e., the sense of sight [32].

In each retina there are about 120 million photoreceptors (rods and cones) that release neurotransmitter molecules at a rate that is maximal in darkness and decreases, logarithmically, with increasing light intensity. This signal is then transmitted to a local chain of bipolar cells and ganglion cells.

There are about 1 million ganglion cells in the retina and their axons form the optic nerve (see figure 2.1). There are, therefore, about 100 photoreceptors per ganglion cell; however, each ganglion cell receives signals from a *receptive field* on the retina, a roughly circular area that covers thousands of photoreceptors.

Between the photoreceptors and bipolar cells, there is a horizontal layer of cells (called *horizontal cells*) linked together so that the potential of each is a weighted average of its neighbors' potential. Each bipolar cell receives input from a photoreceptor and a horizontal cell, producing a signal that is proportional to the logarithm of the difference between the signals produced by the other two cells. As a result, large areas with uniform illumination produce very weak signals, and areas with illumination changes, such as object contours, result in strong signals. That is, the retina detects essentially brightness variations.

As horizontal cells have a relatively slow response, when a photoreceptor detects a moving object, they still have information about the previous object position. This way, the output signal of the bipolar cells (after passing through the layer of amacrine cell to the ganglion cells) contains useful information for motion detection.

Biologically Motivated Keypoint Detection for RGB-D Data



Figure 2.1: The optic nervous system. The visual system includes the eyes, the connecting pathways through to the visual cortex and other parts of the brain in the mammalian system (figure adapted from [142]).

An image is produced by the excitation of the rods and cones in the retina. The excitation is processed by various parts of the brain that work in parallel to form a representation of the external environment in the brain.

Rods, which are far more numerous than cones, are responsible for our vision in dim light and saturated at daylight levels and don't contribute to the image formation [34, 35]. Cones do not respond to dim light but are responsible for our ability to see fine detail and for our color vision [32]. The light in most office settings falls between these two levels. At these light levels, both rods and cones are actively contributing to the information patterns coming out of the eye.

In humans, there are three types of cones sensitive to three different spectra, resulting in *trichromatic color vision* [144]. The cones are conventionally labeled according to the ordering of the peak wavelengths of their spectral sensitivities: short, medium, and long cone types [145]. These three types do not correspond well to particular colors as we know them, but the short, medium and long wavelengths are considered as a representation of the blue, green and red colors, respectively [144, 146]. It is the lack of one or more of the sub-types of cones that causes individuals to have deficiencies in color vision or other types of color blindness [147]. These individuals are not blind to the objects of a certain color, but experience the inability to distinguish between two groups of colors that can be distinguished by people with normal vision.

Retinal ganglion cells have two types of response, depending on the cell's receptive field: *ON cells* and *OFF cells* (see figure 2.2). These receptive fields comprise a central region approximately circular, where the light has an effect on the firing of the cell, and a ring around it. In ON cells, an increment in light intensity in the center of the receptive field causes the firing rate to increase. In OFF cells, it makes it decrease [32]. In a linear model, this response profile is well described by a difference of Gaussians and is the basis for many edge detection algorithms. In addition to these simple differences in ganglion cells, they are also differentiated by chromatic sensitivity and the type of spatial sum they employ [32].



Figure 2.2: Receptive field with center-surround organization. The upper photoreceptors are composed by *OFF-Center* and *ON-Surround*, and the bottom one is the inverse (figure adapted from [143]).

Regarding the visual signal transfer to the brain via the visual system, the retina is vertically divided in two: the temporal half (closer to the temple) and the nasal half (closer to the nose), as shown in figure 2.1. The axons from the nasal half cross the brain at the optic chiasm (see figure 2.1) to join with axons from the temporal half of the other eye before reaching the Lateral Geniculate Nucleus (LGN).

Although there are over 120 million receptive cells in the retina, there are only about 1.2 million fibers (axons) in the optic nerve, thus a large amount of pre-processing is performed inside of the retina. The *fovea* produces the most accurate information. Although it occupies about 0.01% of the visual field (less than 2° of visual angle), about 10% of axons in the optic nerve are devoted to it. The resolution limit of the fovea was determined at around 10,000 points. The information capacity is estimated at 500,000 bits per second, discolored, or about 600,000 bits per second, with color.

The retina, unlike a camera, doesn't just send an image to the brain. It spatially encodes (compress) the image to fit the limited capacity of the optic nerve. Compression is necessary because there are 100 times more photoreceptor cells as ganglion cells, as mentioned above. In the retina, the spatial coding is performed by the *center-surround* structures as implemented by bipolar and ganglion cells. There are two types of center-surround structures in the retina (see figure 2.2): *ON-Center* and *OFF-Center*. The ON-Center use a positive weighed center and negatively weighed neighbors. The OFF-Center use exactly the opposite. The positive weighing is better known as *excitatory* and the negative as *inhibitory* [32].

These center-surround structures are not physical in the sense that they can be seen by staining tissue samples and examining the anatomy of the retina. The center-surround structures are logical (i.e., mathematically abstract) in the sense that they depend on the strength of connection between bipolar and ganglion cells. It is believed that the connection strength between cells depends on the number and types of ion channels embedded in the synapses between bipolar and ganglion cells. Kuffler, in the 1950s, was the first to begin to understand these center-surround structures in the retina of cats [148].

The center-surround structures are mathematically equivalent to edge detection algo-

rithms used by computer programmers to extract or enhance the edges in an image. Thus, the retina performs operations on the object edges within the visual field. After the image is spatially encoded by the center-surround structures, the signal is sent through the optic nerve (via the axons of ganglion cells) across the optic chiasm to the LGN, as shown in figure 2.1.

2.1.2 Lateral Geniculate Nucleus (LGN)

The LGN is the primary broadcasting center for visual information received from the retina and is found inside the thalamus. The LGN receives information directly from retinal ganglion cells via the optic tract and from the Reticular Activating System (RAS). RAS is an area of the brain responsible for regulating arousal (physiological and psychological state of being awake or reactive to stimuli) and sleep-wake transitions. The neurons in the LGN send their axons through the optic radiation, a direct pathway to the primary visual cortex, as shown in figure 2.3. In mammals, the two strongest paths that connect the eye to the brain are those that are designed for LGNd (dorsal part of the LGN in the thalamus), and for the Superior Colliculus (SC) [36].

In humans and monkeys, the LGN is normally described as having six distinct layers. The two inner layers, 1-2, are called the *magnocellular* layers, while the four outer layers, 3-6, are called *parvocellular* layers [151].

Both the LGN of the right and left hemispheres receive inputs from each eye. However, each LGN receives information from only one half of the visual field. This is due to the axons of ganglion cells of the inner half of the retina (nasal side), crossing to the other side of the brain through the chiasm, as shown in figure 2.1. The axons of ganglion cells of the outer half of the retina (temporal sides) remain on the same side of the brain. Therefore, the right hemisphere receives visual information from the left visual field, and the left hemisphere receives visual information from the right visual field [37].

The LGN receives input from some sources, including the cortex and sends its output to



Figure 2.3: Block diagram of the connections between the visual reception and the visual specialized brain lobes of the HVS. The dorsal pathway comprises several cortical areas, including the Middle Temporal (MT) or V5, the Medial Superior Temporal (MST) area, and the ventral and lateral intraparietal areas (VIP and LIP). The Parieto-Occipital (PO) sulcus separates the parietal and occipital lobes. Visual areas TE and TEO (for which Anterior Inferotemporal (AIT) and Posterior Inferotemporal (PIT) are alternative names [149,150], respectively) have a significant reciprocal connection with the perirhinal cortex (cortical region in the Inferotemporal (IT) cortex), figure adapted from [142].

the cortex. The LGN receives some input from the optic tract, as shown in figure 2.1. The axons that leave the LGN go to the V1 visual cortex (see figure 2.3). Both magnocellular layers 1 - 2 and the parvocellular layers 3 - 6 send their axons to layer 4 of V1. Studies involving blindsight people have suggested that the projections of the LGN not only travel to the primary visual cortex, but also to higher cortical areas V2 and V3 [152].

The precise function of the LGN is unknown. It has been shown that while the retina performs spatial decorrelation through center-surround inhibition, the LGN performs temporal decorrelation [153]. However, there is certainly much more going on. Recent experiments in humans with functional Magnetic Resonance Imaging (fMRI) found that both spatial attention and saccadic eye movements can modulate the activity in the LGN [154].

2.1.3 Visual Cortex

The brain's visual cortex is the part of the cortex responsible for processing visual information. It is located in the occipital lobe, at the back of the brain (see figure 2.4). The term visual cortex refers to the primary visual cortex (also known as striate cortex or V1) and areas of extrastriate visual cortex such as V2, V3, V4 and V5. The primary visual cortex is anatomically equivalent to the area *Brodmann 17* [155]. The areas of the extrastriate cortex consist of *Brodmann 18* and *Brodmann 19* [155].

The dichotomy between the way you enter dorsal/ventral (also called flow `where/what' or `action/perception' [58]) was defined by [156] and is still a controversial topic among vision scientists and psychologists. It is probably an excessive simplification of the true state of affairs



Figure 2.4: Specialized brain lobes (left hemisphere) and the ventral and dorsal streams (figure adapted from [142]).

in the visual cortex. It is based on the study of visual illusions, such as the Ebbinghaus illusion, that may distort judgments of nature perception, but when the subject responds with an action, such as grasping, no distortion occurs. However, recent work, like in [157], suggests that both systems of action and perception are equally fooled by such illusions.

The neurons in the visual cortex allow the development of an action when visual stimuli appear within their receptive field. By definition, the receptive field is the region within the entire visual field that causes an *action potential* (in physiology, an action potential is a short-lasting event in which the electrical membrane potential of a cell rapidly rises and falls, following a consistent trajectory). But any given neuron can better respond to a subset of stimuli within its receptive field. For example, a neuron in V1 may fire to any vertical stimulus in its receptive field and ignore other types of stimulus. In the earlier visual areas, like in the IT cortex (see figure 2.3), a neuron can only fire when a certain face appears in its receptive field.

2.1.3.1 Primary Visual Cortex (V1)

The primary visual cortex is the best-studied area of the visual system. In all studied mammals, it is located in the posterior pole of the occipital cortex (the occipital cortex is responsible for processing visual stimuli), as shown in figures 2.3 and 2.4. It is the simplest and oldest part of the visual cortical area. It is strongly specialized in processing information about static and moving objects, and is excellent in pattern recognition. The primary visual cortex is divided into six distinct functional layers, labeled 1 to 6. The 4th layer is the one that receives more visual input from the LGN. The average number of neurons in the primary visual cortex of an adult human being was estimated to be 280 million [158].

V1 has a well defined map of visual spatial information. For example, in humans the whole top of the calcarine sulcus responds strongly to the lower half of the visual field, and the bottom of the calcarine to the upper half of the visual field. Conceptually, this retinotopic mapping is a transformation of the visual image of the retina to V1. The correspondence between a given location in V1 in the subjective field of vision is very precise: even the blind spots are mapped in V1. In humans and animals with a fovea on the retina the proportion of the central visual field, a phenomenon known as *cortical magnification*. Perhaps for the purpose of precise spatial encoding, neurons in V1 have the smallest receptive field size in all regions of the visual cortex.

The tuning properties of V1 neurons differ greatly. That is, the responses can discriminate small changes in visual orientations, spatial frequencies and colors. In addition, individual V1 neurons in human and animals with binocular vision have ocular dominance (namely, there is a preference for one of the eyes). In V1, and in the primary sensory cortex in general, neurons with similar properties tend to cortical columns. Hubel and Wiesel [159] proposed a new organization of classic ice cube cortical columns for two tuning properties: ocular dominance and orientation. However, this model can not accommodate color, spatial frequency and many other features to which neurons are tuned. The exact arrangement of all these cortical columns within V1 remains a research topic.

The current consensus seems to be that the initial responses of V1 neurons are composed of sets of tiled selective spatio-temporal filters. In space, the operation of the V1 can be thought of as similar to many local spatial functions, complex Fourier transforms, or more precisely Gabor transforms. Theoretically, these filters together can carry out the neural processing of spatial frequencies, orientations, movements, directions, speeds (thus temporal frequency), and many other spatio-temporal features.

V1 neurons are also sensitive to the global scene organization [38]. These response prop-

erties probably result from recurrent processing and lateral connections in the neuron pyramids [39]. *Feedforward* connections are mostly driving, and *feedback* connections are mostly modulatory in their effects [40, 41]. Evidence shows that the *feedback* connections originating from higher-level areas such as V4, IT, MT, with larger and more complex receptive fields, can change and shape responses in V1, representing contextual or extra-classical receptive field effects [160--163].

Computational theories of spatial attention in the visual system propose that the attention modulation increases the responses of neurons in many areas of the visual cortex [42--44]. The natural place where it is possible predict an early increase of this type is V1 and recent fMRI evidence shows that the striate cortex can be modulated by attention in a manner consistent with this theory [45].

Studies in macaque disagree as to the presence and amount of attention modulation in V1 [164]. Experiments performed with human neuroimaging are less ambiguous and invariably find robust changes of attention in primary visual cortex [45, 165] and in subcortical structures [166, 167]. Electrophysiological studies in humans often report an attention modulation in the visual cortex [136, 137], but the spatial origins of these signals are normally only estimated by indirect means (for example, the response time of the signal or the relation of waveforms on stimulus location). Direct estimates of neural modulation from electrophysiological measurements with higher spatial precision differ considerably [168]. Another recent study of attention modulation in human V1 using intracranial electrodes in a single subject could not find attention effects [169]. One possible explanation for this result is that this task was not demanding enough to generate strong modulation of the neural response. Generally, it appears that attention can act at early places in the visual stream and modulate neural responses, but the effects may be weaker than those seen in higher areas or confined to a subset of the neural population [164].

2.1.3.2 Visual Area V2

The visual area V2, also called the prestriate cortex [46], is the second largest area of the visual cortex, and the first region within the visual association area. It receives strong *feedforward* connections from V1 and sends strong connections to V3, V4 and V5. It also sends strong *feedback* connections to V1.

Anatomically, V2 is divided into four quadrants, a dorsal and ventral representation in each hemisphere (left and right). Together, these four regions provide a complete map of the visual world. Functionally, V2 has many properties in common with V1. Recent research has shown that cells in V2 show a small amount of attention modulation (more than in V1, less than in V4), are set to moderately complex patterns, and can be driven by multiple orientations in different sub-regions within a single receptive field [47, 48].

It is argued that the entire ventral stream (see figure 2.4) is important for visual memory [49]. This theory predicts that the Object-Recognition Memory (ORM) undergoes changes that may result in the manipulation of V2. In a recent study it was revealed that certain V2 cells play a very important role in storage in the ORM, and the conversion of short-term memories into long-term memories [50]. This area is highly interconnected within the ventral stream of the visual cortex. In the monkey brain, this area receives strong *feedforward* connections from the primary visual cortex and sends strong projections to the other secondary visual cortex areas (V3, V4 and V5) [170, 171] (see figure 2.3). Most neurons in this area respond to simple visual features such as orientation, spatial frequency, size, color and shape [51--53]. V2 cells also respond to various characteristics of complex shapes, such as the orientation of illusory

contours [51] and if the stimulus is part of the foreground or the background [54, 55].

2.1.3.3 Visual Area V3

The V3 cortex region is located immediately in front of V2. There is still some controversy over the exact extent of area V3, with some researchers proposing that the cortex located in front of V2 may include two or three functional subdivisions. For example, Felleman et al. [56] proposed the existence of a dorsal V3 in the upper hemisphere, which is distinct from ventral V3 located at the bottom of the brain. The dorsal and ventral V3 have distinct connections with other parts of the brain, appear in different sections stained with a variety of methods, and contain neurons that respond to different combinations of visual stimuli.

The dorsal V3 is usually considered part of the dorsal stream (shown in figure 2.4), receiving inputs from V2 and the primary visual area and projecting to the Posterior Parietal (PP) cortex. Some studies with fMRI suggested that the V3 area may play a role in the processing of global motion [172]. Other studies considered the dorsal V3 as part of a larger area called the dorsomedial area, which contains a representation of the entire visual field. The neurons in the dorsomedial area respond to the coherent motion of large patterns covering large portions of the visual field [173].

The ventral V3 has considerably weaker connections to the primary visual area, and stronger connections to the IT cortex. While previous studies suggested that the ventral V3 had only a representation of the upper visual field, a more recent work indicates that this area is more extensive than previously appreciated, and like the other visual areas, may contain a complete visual representation [174].

2.1.3.4 Visual Area V4

The visual area V4 is one of the visual areas of the extrastriate visual cortex. It is located before V2 and after the PIT area, as shown in figure 2.3. V4 is the third cortical area in the ventral stream, receiving strong *feedforward* input from V2 and sending strong connections to the PIT.

V4 is the first area in the ventral stream that has a strong attention modulation. Most studies indicate that selective attention can change firing rates in V4 by about 20%. Moran and Desimone [57] characterize these effects, and this was the first study to find effects of attention anywhere in the visual cortex [58].

Like the V1, V4 is tuned at the level of orientation, spatial frequency, and color. But unlike V1, V4 is set to feature extraction of objects of intermediate complexity, like simple geometric shapes, although no one develop a complete description of the V4 parameter space. The visual area V4 is not tuned for complex objects like faces, as the areas of the IT cortex.

The firing properties of V4 were first described, at the end of 1970, by Zeki [66], that also named the area. Originally, Zeki argued that the purpose of V4 was to be responsible for processing color information. At the beginning of 1980, it was proved that V4 was directly involved in the shape recognition, as previous areas of the cortex. This research supported the hypothesis of the two streams first presented by Ungerleider and Mishkin [156].

2.1.3.5 Visual Area V5 or MT

The visual area V5, also known as Middle Temporal (MT) visual area, is a region of the extrastriate visual cortex, which is thought to play an important role in the perception of motion,
the integration of local motion signals into global perceptions and orientations of some eye movements [59]. The MT is connected to a large variety of cerebral cortical and subcortical areas. Its inputs include visual cortical areas V1, V2 and dorsal V3 [60,61], koniocellulare regions of the LGN [62] and inferior pulvinar. The pattern of projections to MT changes somewhat between the representations of the foveal and peripheral visual fields [63]. The MT was shown to be organized in steering columns [175]. DeAmgelis and Newsome [64] argued that the neurons in MT have also been organized based on its fit to the binocular disparity.

The standard view is that V1 provides the most important input to the MT [59] (see figure 2.3). However, several studies have shown that neurons in MT are able to respond to visual information, often in a selective manner, even after V1 is destroyed or disabled [65]. In addition, research carried out by Zeki [66] suggests that certain types of visual information may reach MT before it even reaches V1.

The MT sends its outputs to areas located in the immediately surrounding cortex. Early studies of the electro-physiological properties of neurons in MT showed that a large proportion of cells were in tune with the speed and direction of visual stimuli on the move [176, 177]. These results suggest that MT plays a significant role in the processing of visual motion.

The study of lesions also supports the role of MT in motion perception and eye movement. Neuropsychological studies of a patient who could not see motion, seeing the world as a series of static frames instead, suggested that MT in the primates is homologous to V5 in humans [178, 179]. There is still much controversy about the exact form of the calculations performed in MT [180] and some research suggests that movement perception is already available at lower levels of the visual system (such as on the V1 [181, 182]). These results left open the question of precisely what MT could do that V1 could not. Much work has been performed in this region, since it seems to integrate the local signals of visual motion in the global movement of complex objects [183]. For example, an injury in V5 can cause a deficit in motion perception and processing of complex stimuli. It contains many neurons tuned to the motion of complex visual features. The micro-stimulation of a neuron located in the V5 affects the perception of motion. For example, if someone finds a neuron with a preference for upward motion, and then uses an electrode to stimulate it, the monkey becomes more sensitive to upward motion [184].

2.2 Visual Attention

In this section, several concepts regarding visual attention are discussed. More detailed information can be found in, for example, Pashler [67, 68], Style [69], and Johnson and Proctor [70].

Generally, we seem to keep a rich representation of our visual world and major changes to our environment will attract our attention. Only a small region of the scene is analyzed in detail, in each moment: the attention focus region. This is usually, but not always, the same region which is fixed by the eyes [71, 72]. The order in which a scene is investigated is determined by the mechanisms of selective attention. Corbetta proposed the following definition of attention: "defines the mental ability to select stimuli, responses, memories, or thoughts that are behaviorally relevant among many others that are behaviorally irrelevant" [134].

There are two categories of factors that motivate attention: the *bottom-up* factors and *top-down* factors [73]. Corbetta and Shulman [74] review the evidence on partially segregated networks of brain areas that perform different attention functions. The preparation and application of goal directed (top-down) selection of stimuli is performed by a system that includes

parts of the intraparietal cortex and superior frontal cortex, which is also modulated by the detection of stimuli. Another system, which the top-down selection is not involved with, is largely lateralized to the right hemisphere, where it includes the temporoparietal cortex and inferior frontal cortex. This system is specialized on the detection of behaviorally relevant stimuli, particularly when they are *salient* or unexpected. Thus, it is possible to indicate that there are two separate areas of the brain that are involved in attention. According to Theeuwes [75], the bottom-up influence is not voluntarily suppressible: a highly salient region captures the Focus of Attention (FOA), regardless of the task.

The bottom-up factors are derived solely from the visual scene [76]. The regions of interest that attract our attention in a bottom-up way are called *salient* and the feature responsible for this reaction must be sufficiently discriminating in relation to surrounding features. Beyond the bottom-up attention, this mechanism is also called exogenous attention, automatic, reflexive, or peripherally directed attention [77].

In contrast, top-down attention is spurred by cognitive factors like knowledge expectations and current goals [74]. For example, car drivers are more likely to see gas stations on a street and cyclists to notice the existence of bicycle paths [78].

In psychophysics, top-down influences are often investigated, through so-called signaling experiments. In these experiments, a signal directs the attention to the target. The signals may have different characteristics: they can indicate where the target is, for example, a central arrow that points to the direction of the target [69, 135], or what destiny will be, for example, the signal is a target image or a word that describes the target [185, 186].

The mechanisms of bottom-up attention have been more thoroughly investigated than the mechanisms of top-down attention. One reason is that the data driven stimuli are easier to control than cognitive factors, such as knowledge and expectations, although little is known about the interaction between the two processes.

The mechanisms of selective attention in the human brain still remain open in the field of perception research. The nonexistence of a brain area solely oriented for visual attention [79--81] is one of the most important discoveries in neurophysiology, but the visual selection appears to be present in almost all areas of the brain associated with visual processing [82]. Additionally, new discoveries indicate that many areas of the brain share the processing of information from different senses and there is growing evidence that large parts of the cortex are multi-sensory [83]. A network of anatomical areas performs the mechanisms of attention [74]. The major areas of this network are the PP cortex, the SC, the Lateral Intraparietal (LIP) area, the Frontal Eye Field (FEF) and the pulvinar, as shown in figures 2.1 and 2.3. Opinions differ on the question of what areas perform certain tasks.

Posner and Peterson [187] describe three important functions related to attention: the orientation of attention, target detection and alertness. According to them, the first function, the orientation of attention to a salient stimulus, is accomplished by the interaction of three areas: PP, SC, and the pulvinar. The PP is responsible for the release of the focus of attention from its current location, Inhibition-Of-Return (IOR), the SC shifts attention to a new location and the pulvinar is specialized in reading data from the indexed location and is called by the posterior attention system. The second function of attention, target detection, is performed by what the authors call *anterior attention system*. Finally, they state that the alertness to high-priority signals is dependent on activity in the norepinephrine system arising in the locus coeruleus [188].

The FEF and the SC are the brain areas involved in eye movements. Recently, Bichot [189] points out that the FEF is the place where a kind of map projections is located, which derives

information from bottom-up, as well as top-down influences. The saliency maps, for other researchers, are located in different areas, for instance at LIP [190], at SC [191], at V1 [192] or at V4 [193].

The source of top-down signals may derive from a network of areas in the parietal and frontal cortex. Based on [194], these areas include the Superior Parietal Lobe (SPL), the FEF and the Supplementary Eye Field (SEF). The transient response of the signal in the occipital lobe and more sustained responses in the dorsal PP cortex along the intraparietal sulcus (IPs) and frontal cortex in or near the putative human homologue of the FEFs were found by Corbetta and Shulman [74]. The interaction of bottom-up and top-down signals occurs in V4, in the view of Ogawa and Komatsu [195].

2.3 Summary

With this chapter, an overview of the HVS was provided. Visual attention is a highly interdisciplinary field and researchers in this area come from different backgrounds [131]. For psychologists, research conducted in the area of human behavior is performed by isolating certain specific tasks, in order to understand the internal processes of the brain, often resulting in psychophysical theories or models [134]. While neurobiologists observe the brain's response to certain stimuli [135], using techniques such as fMRI, having therefore a direct view of the brain areas that are active under certain conditions [45, 136, 137].

In recent years, these different areas have profited considerably from each other. Psychologists use research conducted by neurobiologists, in order to improve their attention models, while neurobiologists consider psychological experiments to interpret their data [134]. In addition, psychologists began to implement computer models or use computer models previously developed, to verify that they have a similar behavior to that of human perception. Thus, psychologists tend to improve the understanding of the mechanisms and help the development of better computational models.

Chapter 3

Saliency, the Computational Models of Visual Attention

Selective visual attention, initially proposed by Koch and Ullman [2], is used by many computational models of visual attention. *Saliency* maps is a term which was introduced by Itti et al. [3] in their work on the rapid scene analysis, and by Tsotsos et al. [84] and Olshausen et al. [85] in their work on visual attention. In some studies, for example in [84, 86], the term saliency appears referred to as *visual attention* or in [87, 88] as unpredictability, rarity or surprise. The saliency maps are used as a two-dimensional scalar map of values representing the visual saliency of the corresponding location, independently of the particular stimulus that makes the location salient [1].

With the emerging interest in *active vision*, the computer vision researchers have been increasingly concerned with the mechanisms of attention and have proposed a number of computational models of attention. An active vision system is one that can manipulate the point of view of the camera(s) in order to analyze the surrounding environment and to obtain better information from it.

The methods that will be presented can be categorized based on whether they are biologically plausible, purely computational, or hybrid [89]. Other types of categories are described in [6]. In general, all methods employ a low-level approach by determining the contrast of the image regions relative to their surroundings, using one or more features of intensity, color and orientation. When a method is said to be biologically plausible it means that it follows the knowledge of the HVS. There is usually an attempt to combine known features, extracted by the retina, LGN, primary visual cortex (V1), or by other visual fields (such as V2, V3, V4 and V5). Itti et al. [3], for example, base their method on a biologically plausible architecture proposed in [2]. They determine the center-surround contrast with the Difference of Gaussians (DoG) approach. Frintrop et al. [90] present a method inspired by Itti's method, but the center-surround differences are obtained using square filters and integral images to reduce the processing time.

Other methods are purely computational and not based on any of the biological principles of vision. Ma and Zhang [86] and Achanta et al. [91] estimate the saliency using the distances from the center-surround. While Hu et al. [92] estimate the saliency through the application of heuristic measures on the initial saliency measures obtained by histogram thresholding of feature maps. The maximization of mutual information between the distributions of features in the center and surround of an image is made [93]. Hou and Zhang [94] perform the processing in the frequency domain.

The methods classified as hybrid are those that incorporate ideas that are partially based on biological models. Here, the method of Itti et al. [3] is used by Harrel et al. [95] to generate the characteristics maps; the normalization is performed by an approach based on graphs. Other methods use computational approaches like maximization of information [96] that represent biological plausible models of saliency detection.

In the rest of this chapter, we focus only on those models which can process arbitrary digital images and return corresponding saliency maps. In each of the following sections, the models are introduced in chronological order.

3.1 Biological Plausible Methods

Many plausible biological models that have strong links with the psychological or neurophysiological findings are described in this section. Almost all these attention models are directly or indirectly inspired by cognitive concepts.

A visual attention model is proposed by Ahmad [196]. It consists of a network propagation corresponding to the pulvinar, whose outputs correspond to the areas V4, IT and MT. The main network corresponds to the SC, FEF and PP areas, while the control network corresponds to the PP areas and the working memory corresponds to the prefrontal cortex.

The attention model proposed by Niebur and Koch [197] scans the scene both in the form of a rapid, bottom-up, saliency-driven and task-independent manner and in a slower, top-down, volition-controlled and task-dependent manner, based on the work in [2, 198, 199].

Itti et al. [3, 200] built a saliency model based on spatial attention derived from the biologically plausible architecture proposed by Koch and Ullman [2], using three feature channels: color, intensity, and orientation. This model has been the basis of later models and the standard benchmark for comparison. It has been shown to correlate with human eye movements in free-viewing tasks [201, 202]. The features are calculated by a linear set of center-surround operations, similar to visual receptive fields. Typically visual neurons are more sensitive to a small region of the visual space (the center), while stimuli presented in a broader concentric region with the center (the surround) inhibit the neuronal response. This type of architecture, sensitive to local spatial discontinuities, is particularly well-suited to detect locations which stand out from their surroundings and is a general computational principle in the retina, LGN, and primary visual cortex [122]. Based on retinal input, given the intensity, color, and orientations, they create Gaussian pyramids. The center-surround contrast is determined using a DoG approach from the Gaussian pyramids, creating "feature maps". An input image is subsampled into a Gaussian pyramid and each pyramid level σ is decomposed into channels for Red (R), Green (G), Blue (B), Yellow (Y), Intensity (I), and local orientations (O_{σ}) . From these channels, center-surround feature maps f_l for different features l are constructed and normalized ($\mathcal{N}(\cdot)$). In each channel, maps are summed across scale and normalized again:

$$f_l = \mathcal{N}\left(\sum_{c=2}^{4}\sum_{s=c+3}^{c+4}f_{l,c,s}\right), \forall \in L_l \cup L_C \cup L_O \quad (3.1)$$

$$L_I = \{I\}, L_C = \{RG, BY\}, L_O = \{0^o, 45^o, 90^o, 135^o\}$$
 (3.2)

The feature maps are combined into a single "conspicuity map" for each feature type:

$$C_{l} = f_{l}, C_{C} = \mathcal{N}\left(\sum_{l \in L_{C}} f_{l}\right), C_{O} = \mathcal{N}\left(\sum_{l \in L_{O}} f_{l}\right) \quad (3.3)$$

The three conspicuity maps then are summed into the unique bottom-up saliency map:

$$S = \frac{1}{3} \sum_{k \in \{I, C, O\}} C_k.$$
 (3.4)

In [203], this model was extended by adding motion and flicker contrasts to video domain.

The saliency maps produced by this approach have been used by other researchers for applications that apply image processing in small devices [204] and unsupervised object segmentation [205, 206]. For example, a Markov random field model is used to integrate the seed values from the saliency map along the low-level color features, texture, and edges to grow the salient object regions [205]. The Winner-Take-All detects the most salient location and directs attention towards it, such that only features from this location reach a more central representation for further analysis. An IOR mechanism transiently suppresses this location in the saliency map, such that attention is autonomously directed to the next most salient image location.

Itti's framework [3] and Ahmad's model [196] build up an elegant mapping from computational attention model to biological theories. However, the high computational complexity in these systems requires a massively parallel method to obtain fast responses, which is a common character of biological structure based attention models. For high-level applications as robot vision and video content analysis, it is necessary that these models produce fast responses.

Rosenholtz et al. [207, 208] proposed a visual search model. This model could also be used to predict the saliency over an image, in this case the participants were free to rotate the eyes in order to detect the target, if necessary. Initially, features of each point are derived in an appropriate uniform feature space. Then, they computed the distractor features based on the features distribution, mean, μ , and covariance, \sum . With that, the model defines saliency target as the Mahalanobis distance, Δ , between the target feature vector, T, and the mean of the distractor distribution, where

$$\Delta^2 = (T - \mu)' \sum_{n=1}^{-1} (T - \mu). \quad (3.5)$$

This model is similar to [209--211] in the sense that it estimates 1/P(x) (rarity of a feature or self-information) for each image location x. This model also underlies a clutter measure of natural scenes [212].

Li [213] introduced a neural implementation for the saliency map of the V1 area that can also account for search difficulty in pop-out and conjunction search tasks, as in [203].

In [214], a saliency map was computed by extracting primary visual features using the method of Itti et al. [3]. Besides the primary visual features, this method also detected pop-out objects based on their social relevance, in particular, human faces by the method proposed in [215]. The final activation map was obtained by combining the scalar saliency map and the detected faces. Given the computed activation map, the fixation points were defined as the peak locations of the activation map while fixation field sizes were estimated by an adaptable retinal filter centered on the fixation points. The FOA was moved serially over the detected Region of Interests (ROIs) by a decision theoretic mechanism. The generated sequence of eye fixations was determined from a perceptual benefit function based on perceptual costs and rewards, while the time distribution of different ROIs was estimated by memory learning and decaying.

Le Meur et al. [216] proposed a method for bottom-up saliency based on the HVS. Initial, they conducted several eye-tracking experiments in order to infer the mechanisms in HVS. With that, they implemented features like center-surround interactions, visual masking, perceptual decomposition and contrast sensitivity functions to build the saliency model. They extended this model in [217] to the spatio-temporal domain by fusing temporal, chromatic and achromatic information. Here, visual features are extracted from the visual input into several separate

parallel channels. A feature map is obtained for each channel, then a unique saliency map is built from the combination of those channels. The major novelty lies in the inclusion of the temporal dimension as well as the addition of a coherent normalization scheme.

Navalpakkam and Itti [218, 219] proposed a computational model for the task-specific guidance of visual attention in real-world scenes. Their model emphasizes four aspects that are important in biological vision: determining task-relevance of an entity, biasing attention for the low-level visual features of desired targets, recognizing these targets using the same low-level features, and incrementally building a visual map of task-relevance at every scene location. Given a task definition in the form of keywords, the model first determines and stores the task-relevant entities in working memory, using prior knowledge stored in Long-Term Memory (LTM). It attempts to detect the most relevant entity by biasing its visual attention system with the entity's learned low-level features. It attends to the most salient location in the scene, and attempts to recognize the attended object through hierarchical matching against object representations stored in LTM. It updates its working memory with the task-relevance of the recognized entity and updates a topographic task-relevance map with the location and relevance of the recognized entity. The model's performance on search for single features and feature conjunctions is consistent with existing psychophysical data. These results of their biologicallymotivated architecture suggest that the model may provide a reasonable approximation to many brain processes involved in complex task-driven visual behaviors.

Frintrop [131] presented a biologically motivated computer model called Visual Object Detection with a Computational Attention System (VOCUS). The system is based on a mechanism of human perception called selective attention. To imitate the biological process the VOCUS system first analyses the provided audio and video data, considering different features like contrast, color or intensity in parallel. Then, the saliency maps that indicate interest regions, are generated based on the analysis. If additional information is provided the maps are then processed in search for matching attributes. In other words, the model can use both bottom-up attention cues, characteristics of the image and top-down cues and emotion related aspects of the image [220]. In the end, all the information is fused and a focus region is established for more detailed analysis.

VOCUS is based on psychological models like the feature integration theory presented in [221] and on computational models as the Neuromorphic Vision Toolkit of Itti et al. [3]. It is also one of the few systems able to perform goal-directed search, which means that it is able to recognize previously defined objects in provided data. The system is also able to process data in real time and thus it may be used in robotics or monitoring systems [222, 223]. The model represents a major step forward on integration of data and model-driven mechanisms for studies of visual attention [131] and has been referenced in many works on computer vision [224]. Frintrop et al. [90,131] used images integrals [215,225] to accelerate the computation of centersurround differences to find regions salient using maps of separate features of color, intensity and orientation. Their proposal obtains a higher resolution in the saliency maps as compared to the method of [3]. For this, they resize the filter on each scale, instead of the image and thus maintain the same resolution as the original image on all scales.

Kootstra et al. [226] proposed a multi-scale extension for three symmetry-saliency operators and compared them with human eye-tracking data. This extension was applied on the isotropic symmetry and radial symmetry operators presented in [227] and the color symmetry of Heidemann [228]. The authors compared their model against the Itti et al. [3] method and showed that their performance is significantly better on symmetric stimuli.

Marat et al. [229] developed a bottom-up model for spatio-temporal saliency prediction in

video. Based on the magnocellular and parvocellular cells of the retina, their approach extracts from the video. With this, they produce a static saliency map and a dynamic one, which is fused in a single spatio-temporal map. Prediction results of this model were better for the first few frames of each clip snippet.

Bian et al. [230, 231] propose a biologically plausible frequency domain saliency detection method called Frequency Domain Divisive Normalization (FDN), which has the topology of a biologically based spatial domain model, but is conducted in the frequency domain. This reduces computational complexity because unlike all spatial domain methods, they don't need to decompose the input image into numerous feature maps separated in orientation and scale (like [3,95]), and then compute saliency at every spatial location of every feature map. Saliency is normally defined as some measure of difference between center and surround. In order to overcome this constraint, they propose a Piecewise Frequency Domain Divisive Normalization (PFDN) using Laplacian pyramids and overlapping local patches. While PFDN is slower than FDN, it is more biologically plausible and performs better in eye fixation prediction.

The PFDN algorithm from input image to final saliency map is given by:

- 1. Convert the input image to CIELab color space;
- 2. Decompose the image into a number of scales using a Laplacian pyramid;
- 3. For each scale and every color channel, separate into overlapping local patches with a shift between patches;
- 4. Perform a Fourier transform for each patch using $R[k] = F_k\{I\}$, where F denotes a Fourier transform and R[k] is the *k*-th Fourier coefficient;
- 5. Calculate normalization terms $E_i = \sqrt{\frac{w \sum_{k \in i} |R[k]|^2}{N + \sigma^2}}$ and to simplify set the constants $w = \sigma = 1$;
- 6. Obtain the divisive normalized Fourier coefficients;
- 7. Obtain the saliency maps of each patch $S = W|F^{-1}\{\hat{R}\}|^2$, where F^{-1} denotes the inverse Fourier transform, W is a windowing function to remove edge effects, and S is the corresponding spatial domain saliency map;
- 8. For each scale and color channel, recombine the saliency maps of all patches by taking the maximum value at each pixel location;
- 9. Resize all scales to be equal in size and take the spatial maximum across all scales and color channels to obtain the final saliency map;
- 10. Smooth the saliency map with a Gaussian filter.

Chikkerur et al. [232] proposed a model similar to the model of Rao [130]. The goal of their work is to infer the identity and the position of objects in visual scenes: spatial attention emerges as a strategy to reduce the uncertainty in shape information while feature-based attention reduces the uncertainty in spatial information. In this model, attention emerges as the inference in a Bayesian graphical model which implements interactions between ventral and dorsal areas. This model is able to explain some physiological data (neural responses in ventral stream (V4 and PIT) and dorsal stream (LIP and FEF)) as well as psychophysical data (human fixations in free viewing and search tasks).

Murray et al. [233] proposed a model based on a low-level vision system. This model contains a principled selection of parameters as well as an innate spatial pooling mechanism, can be generalized to obtain a saliency model. They generalize a particular low-level model developed to predict color appearance [234] and has three main levels:

- 1. Visual stimuli are processed based on what is known about the early human visual pathway (color-opponent and luminance channels, followed by a multi-scale decomposition);
- 2. A simulation of the inhibition mechanisms present in cells of the visual cortex;
- 3. Information is integrated at multiple scales by performing an inverse wavelet transform directly on weights computed from the non-linearization of the cortical outputs.

Biologically plausible models have the advantage of expanding our view of biological underpinnings of visual attention. This further helps understanding computational principles or neural mechanisms of this process as well as other complex processes such as object recognition.

3.2 Computational Methods

A serial model for visual pattern recognition based on the primate selective attention mechanism and applied it to handwritten digit and face recognition was proposed by Salah et al. [138]. In an attentive level, they constructed a bottom-up saliency map using simple features in order to decrease the computational cost. In the intermediate level, the information is extracted by dividing the image space into uniform regions and training a single-layer perceptron at each region of the image. Finally, they used an associative level with a discrete Observable Markov Model (OMM) in order to combine the information. Regions visited by a fovea are treated as states of the OMM. An IOR allows the fovea to focus on the other positions in the image.

The work of Ramström and Christensen [129] is focused on a discussion of saliency measure using multiple cues based on game theory concepts. Feature maps are integrated using a scale pyramid, inspired by the attention mechanism of Tsotsos et al. [84]. The nodes of the pyramid are subject to trading on a market and the outcome of the trading represents the saliency. They use the spotlight mechanism for finding the ROI.

In [130, 235], they proposed a template matching model by sliding a template of the desired target to every location in the image. In each location, they compute saliency as a similarity measure between the template and the local image patch.

Ma et al. [86,236] propose a method based on local contrast for generating saliency maps which operates on a scale only and is not based on any biological model. The input to this map is resized and the image color in the space *CIELuv* is subdivided into blocks of pixels. The saliency map is obtained by summing the differences of blocks of pixels in small neighborhoods. This framework extracts the regions and points of attention. For object segmentation, a fuzzygrowing method is applied in the saliency map regions. In contrast with Itti's framework [3] and Ahmad's model [196], they employ the theories on human attention mechanisms as high-level guidance for computer algorithm design. For example, they proposed a motion attention model for video skimming [237], a static attention model for image content analysis [86], and a pure computational algorithm for salient region extraction from video [238]. They have also proposed a user attention model for video summarization in [239], which integrated Itti's model as static attention model. The model of visual control presented in [240] is built around the concept of visual behaviors. They borrow the usage of behavior from the robotics community to refer to a sensory-action control module that is responsible for handling a single narrowly defined goal [241]. The key advantage of the behavior based approach is compositionality: complex control problems can be solved by sequencing and combining simple behaviors. For the purpose of modeling human performance it is assumed that each behavior has the ability to direct the eye, perform appropriate visual processing to retrieve the information necessary for performance of the behavior's task, and choose an appropriate course of action.

Hu et al. [92,242] estimate the saliency applying heuristic measures to the initial saliency obtained by histogram thresholding the feature maps. The threshold was applied to the color, intensity and orientation maps through the analysis of histogram thresholding entropy instead of an approach in different scales. Then, they use a spatial compactness measure, calculated as the area of the convex hull covering the saliency region, and saliency density, which is a function of the magnitude of saliency values in the saliency feature maps, to weigh the individual saliency maps before their combination.

Gao and Vasconcelos [93] maximize the mutual information between feature distributions of the center and surround regions of an image by proposing a specific goal for the saliency: classification. An object in the visual stimulus is classified as belonging to each class of interest (or not), and the saliency must be assigned to locations that are useful for this task. This was initially used for object detection [93], where a set of features are selected to improve the discrimination of the class of interest from all other stimuli, and the saliency is defined as the weighted sum of the responses in the set of features which are salient to this class.

In [243, 244], Gao et al. extended their static image saliency to dynamic scenes. The saliency is measured as the Kullback-Leibler (KL) divergence between the histogram of features in a location and the surrounding region, with the features implemented as optic flow. They use DoG filters and Gabor filters, to measure the saliency of a point as the KL divergence between the histogram of filter responses on the point and the histogram of filter responses in the surrounding region. Thus, they solve the complexity problem commonly faced by this type of models, as well as some models of ridge filters using linear responses as features. These models always assign high salience scores for highly textured areas. In [245], these authors used discriminant saliency model in recognition applications, which shows good performance.

Ko and Nam [206] used a Support Vector Machine (SVM) trained on the region features of the image to select the salient regions of interest from the input image, which are then clustered to extract the salient objects.

Jodogne and Piater [246] proposed reinforcement learning algorithm that can be used when the perceptual space contains images, called Reinforcement Learning of Visual Classes (RLVC). RLVC is an iterative algorithm that is suitable for learning direct image-to-action mappings by taking advantage of the local-appearance paradigm. It consists of two interleaved learning processes: Reinforcement learning of a mapping from visual classes to actions, and incremental building of a feature-based image classifier.

Hou and Zhang [94] presented a model that is independent of features, categories, or other forms of prior knowledge of the objects. Through analysis of the log-spectrum of an input image, they extract the spectral residual of an image. In this model, the spectral residual is the innovation and serves as a compressed representation of a scene. Given an input image I(x), amplitude A(f) and phase P(f) are derived. Then, the log spectrum $\mathfrak{L}(f)$ is computed from the down-sampled image. From $\mathfrak{L}(f)$, the spectral residual R(f) can be obtained by multiplying $\mathfrak{L}(f)$ with $h_n(f)$ which is an $n \times n$ local average filter. Using Inverse Fast Fourier Transform, they build, in the spatial domain, the output image called saliency map. The saliency map contains mainly the non-trivial portion of the scene. To improve the result, they smoothed the saliency map with a Gaussian filter. This process can be summarized as:

$$\mathcal{S}(x) = q(x) * \mathfrak{F}^{-1} \left(\exp \left(\mathcal{R}(f) + \mathcal{P}(f) \right) \right)^2, \quad (3.6)$$

$$\mathcal{P}(f) = \mathfrak{J}(\mathfrak{F}[\mathcal{I}(x)]), \quad (3.7)$$

 $\mathcal{R}(f) = \mathcal{L}(f) - h_n(f) * \mathcal{L}(f), \quad (3.8)$

 $\mathcal{L}(f) = \log \left(\mathcal{A}(f) \right), \quad (3.9)$

$$\mathcal{A}(f) = \mathcal{R}\left(\mathfrak{F}[\mathcal{I}(x)]\right), \quad (3.10)$$

where \mathfrak{F} and \mathfrak{F}^{-1} denote the Fourier and Inverse Fourier Transforms, respectively. *P* denotes the phase spectrum of the image, and is preserved during the process. By thresholding they find salient regions, called proto-objects, for fixation prediction.

In [247], they proposed a novel dynamic visual attention model based on the rarity of features. They introduce the Incremental Coding Length (ICL) which is a principle of how it is possible to distribute energy in the attention system. This principle aims to optimize the immediate energy distribution in the system in order to achieve an energy-economic representation of its environment. With the salient visual cues corresponding to unexpected features (according to the definition of ICL), the extracted features may elicit entropy gain in the perception state. The basis functions, described below, are used as features in the attention analysis. Specifically, they use 8×8 RGB image patches from natural scenes for training. A set of $8 \times 8 \times 3 = 192$ basis functions is then obtained. To validate this theoretical framework, they examined experimentally several aspects of visual attention. Comparing with static saliency maps, their model predicted saccades more accurately than did other mainstream models. Because the model updates its state in an online manner, they can consider the statistics of a temporal sequence and the model achieved strong results in video saliency generation. Finally, when features, based on ICL, are combined using weighted sampling, the model provides a coherent mechanism for dynamic visual search with IOR.

Boccignone [248] addressed the issue of joint segmentation and saliency computation in dynamic scenes. They used a mixture of Dirichlet processes, as a basis for computational modeling of object-based visual attention. The idea of using mixture modeling for low-level saliency was first proposed in [249], but limited to classic finite mixtures, in the context of static images and without addressing the issue of segmentation. He also proposed an approach for partitioning a video into shots based on a foveated representation of a video.

[250] propose a simplified version of the model proposed in [210], projecting it to run

in real time. They empirically evaluate the saliency model in saccades control of a camera in social robotics situations.

More recently, in [251], Zhang et al. developed an algorithm in which the saliency is updated online on each new frame. The performance of the model for predicting human fixation while watching videos is comparable to previous methods, with the advantage that it is substantially simpler.

Butko and Movellan [252] extend Najemik and Geisler framework [253] by applying longterm Partially Observable Markov Decision Processes (POMDP) [254] planning methods, in which the primary goal is to gather information. Najemik and Geisler [253] developed an Information Maximization (Infomax) model of eye movement and applied it to explain the visual search of simple objects. They modeled the visual search as a control strategy designed to detect the location of a visual target. According to these authors, the model successfully captured some aspects of human saccades, but has two important limitations: (1) Its fixation policy is glutton, maximizing the instantaneous information gain rather than the long-term gathering of information; (2) It applies only to artificially constructed images.

Butko and Movellan [252] refer to their extension of the approach presented in [253] as Infomax - Partially Observable Markov Decision Processes (I-POMDP), where they showed that long-term Infomax reduces search time. The new formulation allows them to give answers to questions about the temporal dynamics of optimal eye movement. Furthermore, the strategy search varies depending on the characteristics of the optical device that is used to search [252]. While this model is intended to solve the first limitation of the model in [253], the second limitation remained unsolved. The model was only suitable for a limited class of psychophysical stimuli, in particular images that could be described as containing a field of Gaussian noise. In [255], they present a first attempt to extend the I-POMDP model to be useful for computer vision applications. And in [132], they have a computational analysis of eye movement from the point of view of the theory of stochastic optimal control.

A model of attention with visual memory and online learning is proposed in [161], and it is composed of three parts: Sensory Mapping (SM), Cognitive Mapping (CM) and Motor Mapping (MM). The novelty of this model lies in the CM, which incorporates into the visual memory and online learning. To mimic visual memory, they present an *Amnesic Incremental Hierachical Discriminant Regression tree* which is used to guide the amnesic elimination of redundant information from the tree. The *Self-Supervised Competition Neural Network* in CM has the characteristics of online learning since its connection weights can be updated in real time according to environment changes.

Guo et al. [256] present an approach named Phase spectrum of Quaternion Fourier Transform (PQFT) that produces a spatial-temporal saliency, thereby extending the *Quaternion Fourier Transform* method [257]. More recently, in [258], these authors have applied the saliency detection method PQFT to perform the compression of images and video, taking advantage of the multi-resolution representation of wavelets.

Achanta et al. [91] present a method to determine saliency regions in images using the *CIELab* color space [259]. They use low level color and luminance features. An advantage of this method is that the saliency maps have the same size and resolution as the input image. They use the difference-of-mean filter to estimate the center-surround contrast. The lower frequencies are retained depending of the size of the largest surround filter and higher frequencies depend on the size of the smallest central filter. In this way, this approach retains all of the frequency range $]0, \pi]$ with a notch in the mean. They demonstrate the use of this algorithm in the segmentation of semantically meaningful whole objects from digital images.

More recently, in [89], Achanta et al. changed some concepts of their previous method [91] so that the objects' salient regions have more well defined limits. These limits are preserved through a substantial retention of the most frequent content from the original image. The saliency map is obtained by calculating the Euclidean distance between the mean vector *Lab* (obtained by converting the image to the *CIELab* color space) of an input image and each pixel value of a version of the image with Gaussian blur, before being converted to *CIELab* color space. They uniformly attribute saliency values to the entire salient regions, instead of only to the edges or texture regions. This is achieved by relying on the overall contrast of the pixel instead of the local contrast measured in terms of both color and intensity features.

Unlike previous models presented by these authors, Achanta and Süsstraunck [260] present a model where they are concerned with the computational efficiency, noise robust re-targeting scheme based on seam carving [261] by using saliency maps that assign higher importance to visually prominent whole regions (and not just edges). The values of the stronger saliency are not just assigned to the edge image, but to the whole region. This is accomplished by calculating global pixel saliency using intensity as well as color features. The saliency maps easily avoid the artifacts that conventional seam carving generates and are more robust in the presence of noise. In this method, the saliency maps are processed only once, independently of the number of seams added or removed.

The most recent algorithm for saliency detection presented by Achanta and Süsstraunck [262] is based on the premise that you can make assumptions about the scale of the object to be detected based on their position in the image. A pixel belonging to a salient object near the boundary will be less central than one inside the object. So, assuming that the salient object is completely within the image, and not cut-off by the image borders, it is possible to vary the bandwidth of the center-surround filter by increasing the low frequency cut-off when they approach the image borders. Indeed, as they approach the image borders they use a more local surround region. They choose to do this by making the surround symmetric around the center with respect to the edges of the image. When choosing a surrounding symmetrical to each pixel, they implicitly address each pixel to be placed in the center of its own sub-image. This method is different from the one in [89], where the entire image is used as the common global surround (abstracted as the average image *CIELab* color vector) to any given pixel, resulting in an asymmetric environment for pixels that are not in the center of the image.

Rosin [263] proposed an edge-based scheme for saliency detection over grayscale images, being composed in four steps:

A simple method for detecting salient regions in images was proposed by Rosin [263]. This method only requires: a Sobel edge detection; a threshold decomposition at multiple levels to produce a set of binary edge images; a the distance transformation on each of the binary edge images to propagate the edge information; and sum it to obtain the overall saliency map. Moreover, it avoids the need for setting any parameter values.

Seo and Milanfar [264] proposed a framework for both static and space-time saliency detection. Initially they used a local image structure at each pixel represented by a matrix of local regression kernels (equation 3.11), which are robust in the presence of noise and image distortions.

$$\mathcal{K}(x_l - x_i) = \frac{\sqrt{\det(C_i)}}{h^2} \exp\left\{\frac{(x_l - x_i)^T C_l(x_l - x_i)}{-2h^2}\right\}, \ C_l \in \mathbb{R}^{2 \times 2}, \quad (3.11)$$

where l = 1, ..., P, P is the number of the pixels in a local window, h is a global smoothing

parameter, and the matrix C_l is a covariance matrix estimated from a collection of spatial gradient vectors within the local analysis window around a sampling position $x_l = [x_1, x_2]_l^T$. Then, they use a nonparametric kernel density estimation for such features, which results in a saliency map constructed from a local "self-resemblance" measure, indicating the likelihood of saliency. A "matrix cosine similarity" (a generalization of cosine similarity) is employed to measure the resemblance of each pixel to its surroundings. For each pixel, the resulting saliency map represents the statistical likelihood of its feature matrix F_i given the feature matrices F_j of the surrounding pixels:

$$s_i = \frac{1}{\sum_{j=1}^{N} \exp\left(\frac{-1+\rho(F_i, F_j)}{\sigma^2}\right)},$$
 (3.12)

where $\rho(F_i, F_j)$ is the matrix cosine similarity between two feature maps F_i and F_j , and σ is a local weighting parameter.

Yu et al. [265] attempt to simulate top-down influences. Five components of top-down influences are modeled: structure of object representation for LTM, learning of object representations, deduction of task-relevant features, estimation of top-down biases, mediation between bottom-up and top-down models, and perceptual completion. This model builds a dual-coding object representation for LTM. It consists of local and global codings, characterizing internal properties and global attributes of an object. Probabilistic Neural Networks (PNNs) are used for object representation in that they can model probabilistic distribution of an object through combination of confident values. A dynamically constructive learning algorithm was developed to train PNNs when an object is attended. Given a task-specific object, this proposed model recalls the corresponding object representation from PNNs, deduces the task-relevant feature dimensions and evaluates top-down biases. Bottom-up and top-down biases are mediated to yield a primitive grouping based saliency map. The most salient primitive grouping is finally put into the perceptual completion processing module to yield an accurate and complete object representation for attention.

Mahadevan and Vasconcelos [266] introduced a spatio-temporal saliency algorithm based on a center-surround, which extends a discriminant formulation of center-surround saliency previously proposed for static imagery in [93]. This method is inspired by the biological mechanisms of motion-based perceptual grouping. The combination of discriminant center-surround saliency with dynamic textures produced spatio-temporal saliency algorithm, applicable to scenes with highly dynamic backgrounds and moving cameras.

Li et al. [267] proposed a visual saliency model based on conditional entropy for both image and video. Saliency was defined as the minimum uncertainty of a local region given its surrounding area (namely the minimum conditional entropy), when perceptional distortion is considered. They approximated the conditional entropy by the lossy coding length of multivariate Gaussian data. The final saliency map was accumulated pixel-by-pixel and further segmented to detect the proto-objects. In [268], they proposed a newer version of this model by adding a multi-resolution scheme to it.

Avraham and Lindenbaum [269] presented a bottom-up attention mechanism, based on a validated stochastic model to estimate the probability that an image part is of interest. They refer to the probability as saliency, and therefore specify the saliency mathematically. The model quantifies the various intuitive observations, such as increased correspondence likelihood between visually similar regions of the image. The algorithm starts with a pre-attentive

segmentation and then uses a rough approximation to a graphical model to efficiently reveal which segments are more likely to be of interest. They prefer the objects belonging to smaller groups than those which are relatively very different from the rest of the image. Their approach to the design of the saliency algorithm is to quantify the labels that are target and non-target candidates in a probabilistic model, which would eventually identify the salience of a candidate with their likelihood of being a target.

Borji et al. [133] proposed a three-layered approach for interactive object-based attention inspired by RLVC and U-Tree algorithms [270]. Each time the object that is most important to disambiguate appears, a partially unknown state is attended by the biased bottom-up saliency model and recognized. Then the appropriate action for the scene is performed.

Based on the principle of the Infomax, Wang et al. [271] introduced a computational model to simulate human saccadic scanpaths on natural images. The model integrates some factors that drives human attention reflected by eye movements: reference sensory responses, fovea-periphery resolution discrepancy, and visual working memory. They compute three multiband filter response maps for each eye movement which are then combined into multi-band residual filter response maps. Finally, they compute Residual Perceptual Information (RPI) at each location, which is a dynamic saliency map varying along with eye movements. The next fixation is selected as the location with the maximal RPI value.

3.3 Hybrid Methods

Lee and Yu [272] proposed that mutual information among the cortical representations of the retinal image, the priors constructed from our long-term visual experience, and a dynamic short-term internal representation constructed from recent saccades, all provide a map for guiding eye navigations. By directing the eyes to locations of maximum complexity in neuronal ensemble responses at each step, the automatic saccadic eye movement system greedily collects information about the external world while modifying the neural representations in the process. This model is close to the work presented in [253].

Renninger et al. [273] built a model based on the idea that humans fixate at those informative points in an image which reduce our overall uncertainty about the visual stimulus, and it is similar to the approach presented in [272]. This model is a sequential information maximization approach whereby each fixation is aimed at the most informative image location given the knowledge acquired at each point. A foveated representation is incorporated by reducing resolution as distance increases from the center. Shape histogram edges are used as features.

Peters et al. [274--277] presented a model of spatial attention that can be applied to arbitrary static and dynamic image sequences with interactive tasks. The claimed novelty lies in the combination of these elements and in the fully computational nature of the model. The bottom-up component computes a saliency map from 12 low-level multi-scale visual features. The top-down component computes a low-level signature of the entire image, and learns to associate different classes of signatures with the different gaze patterns recorded from human subjects performing a task of interest. They measured the ability of this model to predict the eye movements of people playing contemporary video games. They showed that a point-wise multiplication of bottom-up saliency with the top-down map learned in this way results in high prediction performance.

Harelet al. [95] proposed a new visual saliency model, called Graph-Based Visual Saliency. It consists of two steps: first forming activation maps on certain feature channels, and then normalizing them in a way which highlights conspicuity and admits combination with other map.

Oliva et al. [278] and Torralba et al. [249, 279] proposed a Bayesian framework for visual search tasks, where they use biologically inspired linear filters for different orientations and scales. The filter responses are known to correlate with each other. For example, a vertical bar in the image will activate an adjusted filter to the vertical bars but will also activate (with less intensity) an adjusted filter for bars with an inclination of 45 degrees. The joint probability of a feature vector is estimated using multivariate Gaussian distributions [278] and posteriorly by multivariate generalized Gaussian distributions [279]. Bottom-up saliency is derived from their formulation as $\frac{1}{p(f|f_C)}$ where f_G represents a global feature that summarizes the probability density of presence of the target object in the scene, based on analysis of the scene gist.

Torralba et al. [279] present their model of Contextual Guidance, which is a Bayesian formulation of visual salience. The Contextual Guidance model makes use of global features, forming a holistic representation of the scene to guide attention to locations in a scene that might contain the target. Global features are calculated by forming a low-dimensional representation of a scene, combining the low-level features in large parts of the image and using Principal Component Analysis (PCA) to further reduce the dimensionality.

Bruce and Tsotsos [96] define bottom-up saliency based on maximum information sampling. The information in this model is computed based on Shannon's self-information. The distribution of the features is estimated from a neighborhood of a point, which can be as large as the entire image. When the vicinity of each point is defined as the entire image of interest, as implemented in [96], the definition of projection becomes identical to the term of the bottom-up saliency from the work presented in [278, 279]. It is noteworthy, however, that the feature spaces used in the two models are different. They present a model for calculating the visual saliency constructed on the basis of a theoretical formulation of information. The model employs features that were learned from natural images using Independent Component Analysis (ICA). These were shown to resemble the receptive fields of neurons in the primary visual cortex (V1), and their answers have the desirable property of dispersion. Moreover, the features learned are approximately independent, simplifying the likelihood estimation without making independence assumptions.

The Bayesian Surprise theory of Itti et al. [88, 280, 281] define saliency as a deviation from what is expected based on a set of internal models of the local visual world. According to this theory, the organisms form models of their environment and assign probability distributions over the possible models. With the arrival of new data, the distributions of the possible models are updated with the Bayes rule and the KL divergence between the prior distributions, where the posterior distributions is measured. The new data forces the distribution to be altered, the greater the divergence. These KL scores of different distributions with respect to the models are combined to produce a saliency score. Their implementation of this theory leads to an algorithm that determines the saliency as a kind of deviation of the features present in the closest neighbors, but extends the concept of neighborhood to the spatio-temporal realm.

Kienzle et al. [282, 283] addressed the bottom-up influence of local image information on human eye movements. The model consists of a non-parametric bottom-up approach for learning attention directly from human eye-tracking data. The saliency function is determined by its maximization of prediction performance on the observed data. A SVM was trained to determine the saliency using the local intensities. Also this method produces center-surround operators analogous to receptive fields of neurons in early visual areas (LGN and V1).

Liu et al. [284--286] formulate salient object detection as a problem of image segmentation, where they separate the salient object from the background. In [284], they propose a set of features including multi-scale contrast, center-surround histogram and color spatial distribution to describe a salient object locally, regionally and globally. The Conditional Random Field (CRF) is learned to effectively combine these features to detect salient objects.

In [285, 286], they extended the local, regional and global salient features to the field of motion, so as to be applied not only to images, but also to videos. They designed a dynamic programming algorithm to solve a global optimization problem, with a rectangle to represent each salient object. The salient object sequence detection is defined as an energy minimization problem (like a binary labeling) using a CRF framework, while static and dynamic salience, spatial and temporal coherence, and the global topic model are well defined and integrated to identify a salient object sequence.

Zhang et al. [15, 210] proposed a model called Saliency Using Natural statistics (SUN), by considering what the visual system is trying to optimize when directing attention. The resulting model is a Bayesian framework in which bottom-up saliency emerges naturally as the self-information of visual features, and overall saliency (incorporating top-down information with bottom-up saliency) emerges as the point-wise mutual information between local image features and the search target's features when searching for a target. Self-information in this context, learned from natural images statistics, corresponds to the findings of the new items that attract attention in visual search [287]. The SUN formula for bottom-up saliency is similar to the ones in [96, 249, 278, 279], which are all based on the concept of self-information or a Bayesian formulation. The statistical differences between the current image statistics and natural ones leads to radically different types of self-information. The motivation for using self-information with the statistics of the current image is that a foreground object is likely to have features that are distinct from the features of the background. Since targets are observed less frequently than background during an organism's lifetime, rare features are more likely to indicate targets.

The idea that the salience of an item depends on its deviation from the average statistics of the image can find its roots in the model of visual search proposed in [207], which represented a number of motion pop-out phenomena, and can be seen as a generalization of the saliency of the center-surround-base found in [2].

In [210], the saliency is calculated locally, which means that the model is consistent with the early visual system neuroanatomy and results in an efficient algorithm, with very few free parameters. They extend the model in [15] to temporally dynamic scenes, and characterize the video statistics around each pixel using a bank of spatio-temporal filters with separable space-time components. The joint spatio-temporal impulse response of these filters is the product of a spatial and a temporal impulse response. In [210], the spatial impulse responses are DoG, which model the properties of neurons in the LGN.

To predict the likelihood of where humans typically focus on a video scene, Pang et al. [288] proposed a stochastic model of visual attention by introducing a dynamic Bayesian network to predict where humans typically focus in a video scene.. Their model simulates and combines the visual saliency response and the cognitive state of a person to estimate the most probable attended regions [289]. They reported that the HVS are not deterministic and people may attend to different locations on the same visual input on different occasions.

Garcia-Diaz et al. [211, 290, 291] introduced an approach to visual saliency that relies on a contextually adapted representation produced through adaptive whitening of color and scale features. The proposed approach is based on the classic hierarchical decomposition of images which are initially separate chromatic components (*CIELab* color space). The luminance channel is decomposed into various orientations and scale representation by means of Gabor bank of filters. This approach is inspired by the image representation described in the early stages of the visual pathway. Then, whitening is imposed to groups of oriented scales for each whitened chromatic component. This strategy keeps the number of components involved in whitening limited, overcoming the problems of computational complexity. The decorrelation is achieved by applying PCA on the multi-scale responses, extracting from them a local measure of variability. Furthermore, a local mean is performed to obtain a unified and efficient measure of saliency. As a result, a specifically adapted image representation arises. The resulting image components have zero mean and unit variance. To obtain a saliency map, they simply compute point distinctiveness by taking, for each pixel, the squared vector norm in this representation divided by the sum of the representation across all pixels (Hotelling's T^2 statistic [292]).

we generalize our saliency framework to dynamic scenes and develop a simple, efficient, and online bottom-up saliency algorithm.

Zhang et al. [251] extended the SUN model to dynamic scenes and develop a simple, efficient, and online bottom-up saliency algorithm. The first step starts by applying a bank of spatio-temporal filters to each video. These filters are designed to be both efficient and in line with the HVS. The probability distributions of these spatio-temporal features were learned from a set of videos from natural environments. Finally, the model calculates features and estimates the bottom-up saliency for each point as $-\log p(F = f_z)$.

Following the same direction as the one presented in [249, 278, 279, 293], they linearly integrated three components (bottom-up saliency, gist, and object features) for explaining eye movements in looking for people in a database of about 900 natural scenes.

Judd et al. [97] use a learning approach and train a linear SVM classifier, similarly to one in [283], directly from human eye tracking data and they also present a new public database. The classifier trained by these authors uses low, mid and high level features extracted directly from the images with a resolution of 200×200 . The low-level biologically plausible features used were: local energy of the steerable pyramid filters [294], the simple saliency model described in [207, 209], based on subband pyramids, and intensity, orientation and color contrast features corresponding to the image features calculated in [3]. The mid-level features are due to the horizon, where human beings naturally look more to the objects. At this feature level, they introduced the horizontal line detector from mid-level gist features [209]. Finally, the high-level features used are the Viola and Jones face detector [215, 225] and the Felzenszwalb et al. person detector [295].

Using the database presented in [97, 296] measured the amount of visual information that is available from blurred images. Moreover, they separated the natural images in easy, medium and hard based on their complexity using the following informal criterion: each image was displayed at various resolutions and the authors estimated the lowest resolution in which the image could be understood. The images perceived as having low resolution were classified initially and images understood as containing higher resolutions were classified at the end. Easy images are those which tend to contain a large object or a simple landscape and can be understood using squared images with only 16 or 32 pixels resolution. The medium images have multiple objects or are more complex and can be understood using squared images have many small details or are often abstract and need 64 - 128 pixels of resolution. To reduce the resolution of each image, they used the same method as in [297]. They applied a binomial low-pass filter to each color channel, and then the filtered image was downsampled by a factor of two. As color is an important feature, they preserved the color range of the blurred version of the images. To do this, they scaled the range of each downsampled image as large as possible within the range of 0 - 1, maintaining the same mean

values of pixel luminance.

Li et al. [298] presented a probabilistic multi-task learning approach for visual saliency estimation in video. Here, the problem of visual saliency estimation was modeled by simultaneously considering the stimulus-driven and task-related factors in a probabilistic framework. The stimulus-driven component simulates the low-level processes in HVS using multi-scale wavelet decomposition and unbiased feature competition, while a task-related component simulates the high-level processes to bias the competition of the input features. The algorithm learns various fusion strategies, which are used to integrate the stimulus-driven and task-related components to obtain the visual saliency, similar to the one presented in [276]. Experiments on two eye-fixation datasets [299] and one regional saliency dataset [300] show that this model outperforms seven existing bottom-up approaches presented in [3, 94, 95, 256, 280, 301, 302].

Goferman et al. [303, 304] propose a context-aware saliency that detects regions of the image representing the scene. The goal is to identify both fixation points and detect the dominant object. In conformity with this setting, they present a detection algorithm that is based on four observations realized in the psychological literature [221, 305--307]:

- 1. Local low-level considerations, such as contrast and color;
- 2. Global considerations, suppressing features which occur frequently, maintaining the features that deviate from the norm;
- 3. Visual organization rules, indicating that the visual forms may have one or more centers of gravity depending on how they are organized;
- 4. High-level factors, such as human faces.

In conformity with the first observation, the areas that have different colors or patterns must obtain higher saliency values. Moreover, the homogeneous or blurred areas must obtain lower saliency values. According to the second observation, the features that occur more often should be removed. For the third observation, the salient pixels are grouped and not spread throughout the image. In this algorithm, they first define the saliency using multiple scales. In the next step, they modify the saliency to further accommodate the third principle. Finally, observation four is implemented as post-processing.

Cheng et al. [308] focus on data-driven bottom-up saliency using the detection of the image contrast, based on the work presented in [309], where they believe that cortical cells can be hard wired to respond preferentially to stimuli of high contrast in their receptive fields. They propose a contrast analysis for extracting high-resolution, full-field saliency maps based on the following observations:

- 1. A method based on global contrast, that separates a large-scale object from its surroundings, is preferred to obtain the local contrast;
- 2. Global considerations enable the assignment of comparable saliency values to similar image regions, and can uniformly highlight entire objects;
- 3. The saliency of a region depends mainly on its contrast with neighboring regions;
- 4. Saliency maps should be rapid and easy to generate, in order to allow the processing of large collections of files, and facilitate efficient classification of the images.

They propose the Histogram-based Contrast (HC) method to measure the saliency. The HC assigns pixel-wise saliency values based simply on the color separation from all other image pixels to produce complete resolution saliency maps. They use a histogram-based approach for efficient processing, while employing a smoothing procedure to control the quantization artifacts. As an improvement over HC maps, they incorporate spatial relations to produce Region-based Contrast (RC) maps, where they first segment the input image into regions and then assign saliency values to the segmented regions. The saliency value of a region is calculated using a global contrast score, measured by contrast and spatial distances from a region to other regions in the image.

Klein and Frintrop [310] based their model on the structure of cognitive visual attention, and the saliency calculation is performed in each feature dimension. The method allows a consistent computation of all the feature channels and a well-founded fusion (based on theoretical information) of these channels into a saliency map.

Riche et al. [311] proposed to extract multi-scale rarities from YCbCr color space using multiple Gabor filters.

3.4 Examples of Saliency Detection

In this section, we will present some results obtained by some of the methods described in this chapter. The assessment was carried out in two databases, which we will describe.

The first database, called Toronto, was presented by [96]. It contains 120 images captured indoors and outdoors with a resolution of 681×511 pixels. For eye tracking, images were presented randomly to each of 20 persons and between each image a gray mask was presented during 2s on a 21-inch CRT monitor. The persons were at a distance of 0.75m from the monitor. Stimuli were color images and the task was free viewing.

The second database is called MIT and was introduced by [97]. Images were collected from Flicker creative commons and LabelMe datasets. In this database there are 779 landscape images and 228 portrait images. Images were freely viewed with 1s gray screen between each and the eye tracking camera was re-calibrated after every 50 images.



Table 3.1: Comparing saliency maps and average execution time of some models in images from Toronto and MIT databases.

Table 3.1: Comparing saliency maps and average execution time of some models in images from Toronto and MIT databases.

	Databases			
	Toronto	міт		
Original				
Harel et al. [95]	Mean Time 0.61s	Mean Time 0.60s		
Hou and Zhang [94]	Mean Time 0.0029s	Mean Time 0.0031s		
Liu et al. [284]	Mean Time 13.61s	Mean Time 15.36s		
Achanta et al. [91]	Mean Time 10.98s	Mean Time 69.74s		
Zhang et al. [210]	Mean Time 1.79s	Mean Time 4.39s		
Achanta et al. [89]	Mean Time 0.07s	Mean Time 0.12s		
Bruce and Tsotsos [312]	Mean Time 5.08s	Mean Time 12.73s		
Judd et al. [97]	Mean Time 10.32s	Mean Time 13.82s		
Achanta and Süsstrunk [262]	Mean Time 4.19s	Mean Time 23.81s		

	Databases			
	Toronto	MIT		
Original				
Goferman et al. [303]	Mean Time 57.28s	Mean Time 49.43s		
Hou et al. [313]	Mean Time 0.02s	Mean Time 0.01s		

Table 3.1: Comparing saliency maps and average execution time of some models in images from Toronto and MIT databases.

Table 3.1 presents the saliency maps produced by 13 methods on 3 images from each of the databases, and the average computational time needed to obtain them.

3.5 Applications

Until now, the attention has concentrated on the concepts of the human visual attention and present psychological and neurological theories on what is known about the HVS which has influenced the computational attention models. There was also a description of the general structure and features of computational models of attention, giving an overview of the stateof-the art in this area. There are, however, many technological applications of these models which have been developed over the years and which have further increased interest in attention modeling. The applications of attention modeling are organized into four categories: image, object, robotic and video, as shown in table 3.2.

The images category was divided into five sub-categories: assembling, compression, quality evaluation, resolution and target. In the assembling sub-category, we consider methods that use saliency maps to perform image summarization through collage creation. Collage is a difficult and time consuming task, since the pieces should be nicely cut and matched [303]. Saliency maps can also be used in image compression. Here, the compression rate in a region of the image will depend on whether this is a salient region or not. If a region is not salient the applied compression rate is higher. The use of saliency maps to evaluate the quality of images consists in finding an automatic algorithm that evaluates the quality of pictures or video as a human observer would do [318]. The image resolution task aims to arbitrarily change image aspect ratios while preserving visually prominent features. The target sub-category consists in identifying a target, no matter how many distractors are present.

There are also some applications adapted to work with videos. The difference between a method that can only work with static images and another that can work with videos is linked to its computational complexity, because if we wish to analyze all the video frames the method has to be extremely fast. Besides, the operations that can be performed in a video using saliency

Applications Category		References		
Image	Assembling Compression Quality Evaluation Resolution Target	[303, 314, 315] [258, 316] [317319] [216, 260, 320324] [216, 303, 320, 321]		
Object	Detection Recognition Segmentation Tracking	[89, 91, 218, 250, 255, 260, 262, 284, 285, 293] [131, 138, 139, 325, 326] [205, 206, 327329] [97, 266, 330]		
Robotic	Active Vision Localization Navigation	[133] [131, 316, 326] [90, 199, 326]		
Video	Compression Detection Summarization	[203, 258, 331, 332] [248] [229, 236]		

Table 3.2: Applications that use computational models of attention.

maps are very similar to the ones used in the images category.

The division made for the object category is as follows: detection, recognition, segmentation and tracking. The object detection is a very important step in computer vision and it can be done using saliency maps, as demonstrated by several authors. When referring to the recognition sub-category, the saliency maps are used as a basis for a recognition system, as in [326] where landmarks are recognized so that the robot may move to a given location.

The methods presented that focus on segmentation using saliency maps are methods that give more importance to the edges of objects, making it easier to define the object. The tracking sub-category, considers problems where there is the need to track the eyes.

The presented applications that involved robots use the saliency maps for robot navigation and localization.

3.6 Summary

Computational attention has gained a significant popularity in the last decade. Engineers use the discoveries made by psychologists and neurobiologists, explained in chapter 2, and attempt to reproduce them in computational models, so that they can reduce the processing time in some applications [42--44]. One of the contributors to the increase in popularity was the improvement in computational resources. Another contribution was the performance gains obtained from the inclusion of visual attention (or saliency detection) modules in object recognition systems [131, 138, 139].

Most of the research presented, has been focused on the bottom-up component of visual attention. While previous efforts are appreciated, the field of visual attention still lacks computational principles for task-driven attention. A promising direction for future research is the development of models that take into account time varying task demands, especially in interactive, complex, and dynamic environments. In addition, there is not yet a principled computational understanding of visual attention. The solution is beyond the scope of a single area. In order to obtain a solution it is necessary to have the cooperation of the several areas, from the machine learning community, computer vision and also the biological areas as well as neurology and psychology.

Chapter 4

Keypoint Detectors, Descriptors and Evaluation

When processing image or 3D point cloud data, features must be extracted from a small set of points, usually called keypoints. This is done to avoid the computational complexity required to extract features from all points. There are many keypoint detectors and this suggests the need of a comparative evaluation. When the keypoint detectors are applied to objects, the aim is to detect a few salient structures which can be used, instead of the whole object, for applications like object registration, retrieval and data simplification. In this chapter, the description of some 2D and 3D keypoint detectors (focusing more on 3D), and also 3D descriptors is made. Finally, an evaluation of 3D keypoint detectors, available in PCL, is made with real objects on 3D point clouds. The invariance of the 3D keypoint detectors is evaluated according to rotations, scale changes and translations. The evaluation criteria used are the absolute and the relative repeatability rate. Using these criteria, the robustness of the detectors is evaluated with respect to changes of point-of-view.

4.1 Keypoint Detectors

4.1.1 Harris 3D

The Harris method [333] is a corner and edge based method and these types of methods are characterized by their high-intensity changes. These features can be used in shape and motion analysis and they can be detected directly from the grayscale images. For the 3D case, the adjustment made in PCL for the Harris3D detector replaces the image gradients by surface normals, where the covariance matrix *Cov* will be calculated. The *keypoints response* measured at each pixel coordinate (x, y, z) is then defined by

$$r(x, y, z) = det(Cov(x, y, z)) - k(trace(Cov(x, y, z)))^{2},$$
 (4.1)

where k is a positive real valued parameter and a thresholding process is used to suppress weak keypoints around the stronger ones. The keypoint responses are positive in the corner region, negative in the edge regions, and small in flat regions [333]. If the contrast of the point cloud increases, the magnitude of the keypoint responses also increase. The flat region is specified by the *trace* falling below some selected threshold.

In the PCL are available two variants of the Harris3D keypoint detector: these are called Lowe [99] and Noble [100]. The differences between them are the functions that define the keypoints response (equation 4.1). Thus, for the Lowe method the keypoints response is given by:

$$r(x, y, z) = \frac{det(Cov(x, y, z))}{trace(Cov(x, y, z))^2}.$$
 (4.2)

The keypoints response for Noble method is given by:

$$r(x, y, z) = \frac{det(Cov(x, y, z))}{trace(Cov(x, y, z))}.$$
 (4.3)

In the case of the Lowe detector, the differences between the values of the keypoint responses in the corner regions, edge regions and planar regions tend to be closer to zero compared to those of the Noble detector. This means that there are more regions considered flat.

4.1.2 Kanade-Lucas-Tomasi

The Kanade-Lucas-Tomasi (KLT) detector [98] was proposed a few years after the Harris detector. In the 3D version presented in the PCL, this keypoint detector has the same basis as the Harris3D detector. The main differences are: the covariance matrix is calculated using the intensity value instead of the surface normals; and for the keypoints response they used the first eigenvalue of the covariance matrix. Finally, the suppression process is similar to the one used in the Harris3D method.

4.1.3 Curvature

The curvature method in the PCL calculates the principal surface curvatures on each point using the surface normals. The keypoints response used to suppress weak keypoints, around the stronger ones is the same as in the Harris3D.

4.1.4 Smallest Univalue Segment Assimilating Nucleus

The Smallest Univalue Segment Assimilating Nucleus (SUSAN) corner detector was introduced in [101]. SUSAN is a generic low-level image processing technique which, apart from corner detection, has also been used for edge detection and noise suppression. A geometric threshold is applied, which is simply a precise restatement of the SUSAN principle: if the nucleus (center pixel of a circular region) lies on a corner then the Univalue Segment Assimilating Nucleus (USAN) area will be less than half of its possible value.

USAN is a measure of how similar a center pixel's intensity is to those in its neighborhood. A gray value similarity function $s(g_1, g_2)$ measures the similarity between the gray values g_1 and g_2 . Summing over this kind of function for a set of pixels is equivalent to counting the number of similar pixels. It can be used to adjust the detector's sensitivity to the image's global contrast level. The smoothness plays of $s(g_1, g_2)$ an important role in noise suppression [101], since it only depends on the difference between g_1 and g_2 . To make the method more robust, points closer in value to the nucleus receive a higher weighting. Moreover, a set of rules presented in [334] are used to suppress qualitatively "bad" keypoints. Local minima of the SUSANs are then selected from the remaining candidates.

4.1.5 Scale Invariant Feature Transform

The Scale Invariant Feature Transform (SIFT) keypoint detector was proposed in [9]. This method shares similar properties with neurons in inferior temporal cortex that are used for object recognition in primate vision. In [102], the original algorithm for 3D data is presented, which uses a 3D version of the Hessian to select the interest points, which will be called SIFT3D.

The image *I* is convolved with a number of Gaussian filters whose standard deviations differ by a fixed scale factor. That is, $\sigma_{j+1} = k\sigma_j$ where *k* is a constant scalar that should be set to $\sqrt{2}$. The convolutions yield smoothed images, denoted by

$$G(x, y, \sigma_i), i = 1, ..., n.$$
 (4.4)

The adjacent smoothed images are then subtracted by

$$D(x, y, \sigma_i) = G(x, y, \sigma_{i+1}) - G(x, y, \sigma_i).$$
 (4.5)

These two steps are repeated, yielding a number of DoGs over the scale space. Once DoGs have been obtained, keypoints are identified as local minima/maxima of the DoGs across scales. This is done by comparing each point in the DoGs to its eight neighbors at the same scale and nine corresponding neighborhood points in each of the neighborhood scales. The dominant orientations are assigned to localized keypoints.

4.1.6 Speeded-Up Robust Features

Speeded-Up Robust Features (SURF) [10] is partly inspired by the SIFT descriptor. SURF is based on sums of 2D Haar wavelet responses and makes an efficient use of integral images. It uses an integer approximation to the determinant of Hessian blob detector, which can be computed extremely quickly with an integral image. For features, it uses the sum of the Haar wavelet response around the point of interest.

4.1.7 Intrinsic Shape Signatures 3D

Intrinsic Shape Signatures 3D (ISS3D) [103] is a method relying on region-wise quality measurements. This method uses the magnitude of the smallest eigenvalue (to include only points with large variations along each principal direction) and the ratio between two successive eigenvalues (to exclude points having similar spread along principal directions).

The ISS3D $S_i = \{F_i, f_i\}$ at a point p_i consists of two components: 1 -- The intrinsic reference frame $F_i = \{p_i, \{e_i^x, e_i^y, e_i^z\}\}$ where p_i is the origin, and $\{e_i^x, e_i^y, e_i^z\}$ is the set of basis vectors. The intrinsic frame is a characteristic of the local object shape and independent of viewpoint. Therefore, the view independent shape features can be computed using the frame as a reference. However, its basis $\{e_i^x, e_i^y, e_i^z\}$, which specifies the vectors of its axes in the sensor coordinate system, are view dependent and directly encode the pose transform between the sensor coordinate system and the local object-oriented intrinsic frame, thus enabling fast pose calculation and view registration. 2 -- The 3D shape feature vector $f_i = (f_{i0}, f_{i1}, \ldots, f_{iK-1})$, which is a view independent representation of the local/semi-local 3D shape. These features can be compared directly to facilitate the matching of surface patches or local shapes from different objects.

4.1.8 Biologically Inspired keyPoints

Biologically Inspired keyPoints (BIMP) [7,335] is a cortical keypoint detector for extracting meaningful points from images, solving the computational problem of [104]. The keypoints are

extracted by a series of filtering operations: simple cells, complex cells, end-stopped cells and inhibition cells. Simple cells are modeled using complex Gabor filters with phases in quadrature are given by:

$$g_{\lambda,\sigma,\theta,\phi}(x,y) = \exp\left(-\frac{\tilde{x}^2 + \gamma \tilde{y}^2}{2\sigma^2}\right) \exp\left(i\frac{2\pi \tilde{x}}{\lambda}\right), \quad (4.6)$$

where $\tilde{x} = x \cos(\theta) + y \sin(\theta)$, $\tilde{y} = y \cos(\theta) - x \sin(\theta)$, with σ the receptive field size, θ the filter orientation, λ is the wavelength and $\gamma = 0.5$. Simple cell responses are obtained by convolving the image with the complex Gabor filter: $R_{\lambda,\theta} = I * g_{\lambda,\theta}$. Complex cells are the modulus of simple cell responses $C_{\lambda,\theta} = |R_{\lambda,\theta}|$. Remaining kernels are sums of Dirac functions (δ). If $ds = 0.6\lambda \sin(\theta)$ and $dc = 0.6\lambda \cos(\theta)$, double-stopped cell kernels are defined by

$$k_{\lambda,\theta}^{D} = \delta(x,y) - \frac{\delta(x-2ds,y+2dc) + \delta(x+2ds,y-2dc)}{2} \quad (4.7)$$

and the final keypoints is given by

$$K_{\lambda}^{D} = \sum_{\theta=0}^{\pi} |C_{\lambda,\theta} k_{\lambda,\theta}^{D}|^{+} - \sum_{\theta=0}^{2\pi} |C_{\lambda,\theta} k_{\lambda,\theta}^{TI} + C_{\lambda,\theta^{\perp}} k_{\lambda,\theta}^{RI} - C_{\lambda,\theta}|^{+}, \quad (4.8)$$

where θ^{\perp} is orthogonal to θ , $|.|^+$ represents the suppression of negatives values. $k_{\lambda,\theta}^{TI}$ is the tangential inhibition kernel and $k_{\lambda,\theta}^{RI}$ the radial.

$$k_{\lambda,\theta}^{TI} = -2\delta(x,y) + \delta(x+dc,y+ds) + \delta(x-dc,y-ds)$$
(4.9)

$$k_{\lambda,\theta}^{RI} = \delta(x + dc/2, y + ds/2) + \delta(x - dc/2, y - ds/2).$$
(4.10)

4.2 3D Descriptors

4.2.1 3D Shape Context

The 3D Shape Context (3DSC) descriptor [105] is the 3D version of the Shape Context descriptor [106]. It is based on a spherical grid centered on each keypoint. The surface normal estimation is used to orient the grid to the north pole. The grid is defined by bins along the azimuth, elevation and radial dimensions. The bins along the azimuth and elevation dimensions are equally spaced, on the other hand, the radial dimension is logarithmically spaced. The final representation of the descriptor is a 3D histogram, where in each bin contains a weighted sum of the number of points falling on the grid region. These weights are inversely proportional to the bin volume and the local point density.

4.2.2 Point Feature Histograms

Descriptors such as Point Feature Histograms (PFH) [107], Fast Point Feature Histograms (FPFH) [108, 109], Viewpoint Feature Histogram (VFH) [110], Clustered Viewpoint Feature Histogram (CVFH) [111] and Oriented, Unique and Repeatable Clustered Viewpoint Feature His-

togram (OUR-CVFH) [112] can be categorized as geometry-based descriptors [336]. These type of descriptors are represented by the surface normals, curvature estimates and distances, between point pairs. The point pairs are generated by the point p and the points in its local neighborhood q. And they are represented with the angles α , ϕ and θ , which are computed based on a reference frame (u, v, w). The vector u is the surface normal at p, (n_p) , v is equal to $u \times \frac{p-q}{||p-q||_2}$ and w is the cross product of these two vectors. With this reference frame, the angles can be computed using: $\alpha = v^T \cdot n_p$, $\phi = u^T \cdot \frac{p-q}{||p-q||_2}$ and $\theta = \arctan(w^t \cdot n_p, u^T \cdot n_p)$.

PFHRGB is an version of PFH in which is included information regarding the color of the object. This variant includes three more histograms, one for the ratio between each color channel of p and the same channel of q.

4.2.3 Fast Point Feature Histograms

The FPFH descriptor [108, 109] is a simplification of the PFH. In this case, the normal orientation angles are not computed for all point pairs of p and its neighborhood. The angles are computed only from its k-nearest neighbors. The estimated values are stored into a histogram, since this represents the divisions of the feature space.

4.2.4 Viewpoint Feature Histogram

In [110], they proposed an extension of FPFH descriptor, called VFH. The main differences between this and the other two descriptors above are: the surface normal is centered on the centroid c and not in the point p (n_p); instead of computing the angles using all (PFH) or k-nearest neighbors (FPFH), it uses only the centroid of the input cloud; VFH adds a viewpoint variance using the angle $\beta = \arccos(\frac{n_p \cdot c}{||c||})$, wich represents the central viewpoint vector direction translated to each normal; and it only produces one descriptor for the input cloud.

4.2.5 Clustered Viewpoint Feature Histogram

The CVFH [111] is an extension to VFH. The idea behind this descriptor is that objects which contains stable regions S. That enable them to be divided into in a certain number of disjoint regions. Stable regions are obtained by first removing the points with high curvature and then applying a smooth region growing algorithm. For each stable regions k, they find the centroid c_k and its normal (n_{c_k}) to compute a local reference frame. It is similar to the VFH descriptor, but instead of using the centroid ant its normal of the input cloud, it is only from the stable region. The final descriptor is given by the concatenated local reference frames (u, v, w, SDC, β) , which is a histogram. The Shape Distribution Component (SDC) is equal to

$$SDC = \frac{(c - p_k)^2}{max\{(c - p_k)^2\}}, k = 1, \cdots, |S|.$$
 (4.11)

4.2.6 Oriented, Unique and Repeatable Clustered Viewpoint Feature Histogram

The OUR-CVFH [112] is a semi-global descriptor based on Semi-Global Unique Reference Frames (SGURF) and CVFH [111], which exploits the orientation provided by the reference frame to encode the geometrical properties of an object surface. For a specific surface S, it computes N triplets (c_i , n_i , RF_i) obtained from the smooth clustering and the SGURF computation. SGURF aims to solve some limitations of CVFH by defining multiple repeatable coordinate systems on S. This allows to increase the spatial descriptiveness of the descriptor and obtain the 6DoF from the alignment of the reference frames.

For the surface description, it uses an extension of CVFH in the following way: first, c_i and n_i are used to compute the first three components of CVFH and the viewpoint component as presented in [111]. The fourth component of CVFH is completely removed and instead the surface *S* is spatially described by means of the computed RF_i . To perform this, *S* is rotated and translated, so that RF_i is aligned with the x, y, z axes of the original coordinate system of *S* and centered in c_i . To take in account the perturbations on RF_i , an interpolation is performed by associating to each point p_k eight weights. The weights are computed by placing three 1-dimensional Gaussian functions over each axis centered at c_i , which are combined by means of weight multiplication. Finally, the weights associated with p_k are added to 8 histograms, its index in each histogram being selected as $\frac{c}{R_i}$, where *R* is the maximum distance between any point in *S* and c_i .

4.2.7 Point Pair Feature

The Point Pair Feature (PPF) descriptor [113] assumes that both the scene and the model are represented as a finite set of oriented points, where a normal is associated with each point. It describes the relative position and orientation of two oriented points which is similar to the surflet-pair feature from [108, 337]. If you have two points p_1 and p_2 and their normals n_1 and n_2 , the *PPF* is given by

$$PPF(p_1, p_2) = (d_2, \angle (n_1, d), \angle (n_2, d), \angle (n_1, n_2)),$$
 (4.12)

where $\angle(a, b) \in [0, \pi]$ represents the angle between a and b and $d = p_2 - p_1$.

The model is represented by a set of *PPF*'s, where similar feature vectors being grouped together. This is computed for all the pair points. The distances are sampled in d_{dist} steps and the angles in $d_{angle} = 2\pi/n_{angle}$ steps and the vectors with the same discrete representation are grouped.

An object model descriptor M can be mapped from the sampled space to the model space S. The four dimensional *PPF* defined at equation 4.12 are mapped to set A of all pairs $(m_i, m_i) \in M^2$ that define an equal feature vector.

The final local coordinates use a voting scheme, this is done in order to maximize the number of scene points that lie on the model, allowing the recovery of the global object pose. The similarities between their rotations and translations are used to obtain the pose through the voting system.

In PCL, there is also a color version, called PPFRGB. In this version, three new ratios are added, one for each color channel.

4.2.8 Signature of Histograms of OrienTations

The Signature of Histograms of OrienTations (SHOT) descriptor [114] is based on a signature histograms representing topological features, that make it invariant to translation and rotation. For a given keypoint, it computes a repeatable local reference frame using the eigenvalue decomposition around it. In order to incorporate geometric information of point locations in a spherical grid. For each spherical grid bin, a a one-dimensional histogram is obtained. This histogram is constructed by summing point counts of the angle between the normal of the keypoint and the normal of each point belonging to the spherical grid. Finally, the descriptor override all these histograms according to the local reference frame. It uses 9 values to encode the reference frame, and 11 shape bins and 32 divisions of the spherical grid, which gives an additional 352 values.

In [115], they propose two variants: one is a color version (SHOTCOLOR), where use the CIELab color space as color information; the second one (SHOTLRF), they encode only the local reference frame information, discarding the shape bins and spherical information (resulting in a 9 values to describe the local reference frame).

4.2.9 Unique Shape Context

An upgrade of the 3DSC descriptor [105] is proposed in [116], called Unique Shape Context (USC). The authors reported that one of the problems found in 3DSC is to avoid multiple descriptions for the same keypoint, based on the need to obtain as many versions of the descriptor as the number of azimuth bins. It can cause a possible ambiguity during the successive matching and classification process. To resolve that, they proposed to define only a local reference frame (as defined in [114]) for each keypoint, such that spherical grid associated to a descriptor be directed exclusively by the two main directions in relation to the normal plane. The remaining process for obtaining USC descriptor still the same as the 3DSC.

4.2.10 Ensemble of Shape Functions

In [117], they introduced the Ensemble of Shape Functions (ESF) which is a shape function describing feature properties. This is done using the three shape functions presented in [118], that are the angle, the point distance, and the area. To compute this, they use three points randomly selected, where: two of them are used to calculate the distance; the angle is defined by two lines created from all of them; and area of the triangle formed between them. An approximation (voxel grid) of the real surface is used to separate the shape functions into more descriptive histograms. These histograms will represent the point distances, angles, areas and (on, off or both) surface

4.2.11 Point Curvature Estimation

The Principal Curvatures Estimation (PCE) descriptor calculates the directions and magnitudes of principal surface curvatures (obtained using the cloud normals.) on each keypoint, eigenvectors and eigenvalues respectively. For each keypoint, it will produce a descriptor with 5 values. Three values are the principal curvature, which is the eigenvector with the largest eigenvalue and the other two values are the largest and smallest eigenvalues.

4.2.12 Descriptors Characteristics

Table 4.1 presents some features of the descriptors and is based on the one presented in [22]. The second column contains the number of points generated by each descriptor given an input point cloud with n points In this work the input cloud will be only the keypoints points. The third column shows the length of each point. The fourth column indicates if the descriptor requires the calculation of the surface normals at each point. The column 5 shows if the method is a global or a local descriptor. Global descriptors require the notion of the complete

Descriptor	N. Points	Point Size	Normals	Local/Global	Category
3DSC	п×р	1980 + 9	Y	Local	Spherical + Shape
CVFH	$m \leqslant n$	308	Y	Global	Geometry + Shape
ESF	1	640	Ν	Global	Shape
FPFH	п	33	Y	Local	Geometry
OUR-CVFH	1	308	Y	Global	Geometry + Shape
PCE	п	5	Y	Local	Shape
PFH	п	125	Y	Local	Geometry
PFHRGB	п	250	Y	Global	Geometry
PPF	п	5	Y	Global	Geometry
PPFRGB	п	8	Y	Global	Geometry
SHOT	п	352 + 9	Y	Local	Geometric + Spherical
SHOTCOLOR	п	1344 + 9	Y	Local	Geometric + Spherical
SHOTLRF	п	9	Ν	Local	Geometric + Spherical
USC	n	1980 + 9	Ν	Local	Spherical + Shape
VFH	1	308	Y	Global	Geometry

Table 4.1: Features and statistics of the evaluated descriptors in this work. n = number of points in input cloud; p = Number of Azimuth bins; m = Number of stable regions; Y = Yes; N = No.

object while local descriptors are computed locally around each keypoint and work without that assumption. The sixth column indicates if the descriptor is based on the geometry or shape of the object, and if the analysis of a point is done using a sphere.

4.3 Dataset

The large RGB-D Object Dataset¹ [21] will be used to evaluate the 2D and 3D keypoint detectors and 3D descriptors. This dataset is a hierarchical multi-view object dataset collected

¹The dataset is publicly available at http://www.cs.washington.edu/rgbd-dataset.



Figure 4.1: Examples of some objects of the RGB-D Object Dataset.

using an RGB-D camera and contains a total of 207621 segmented clouds. The dataset contains 300 physically distinct objects taken on a turntable from 4 different camera poses and the objects are organized into 51 categories. Examples of some objects are shown in figure 4.1. It's possible to see that there are some errors in the point clouds, due to segmentation errors and sometimes depth sensor noise (some materials do not reflect the infrared pattern used to obtain depth information as well). The chosen objects are commonly found in home and office environments, where personal robots are expected to operate.

4.4 Evaluation of 3D keypoint Detectors

This section is motivated by the need to quantitatively compare different keypoint detector approaches, in a common and well established experimentally framework, given the large number of available keypoint detectors. Inspired by the work on 2D features [17,18] and 3D [19], and by a similar work on descriptor evaluation [22], a comparison of several 3D keypoint detectors is made in this work. In relation to the work of [17, 19], the novelty is that it used a real database instead of an artificial, the large number of 3D point clouds and different keypoint detectors. The benefit of using real 3D point clouds is that it reflects what happens in real life, such as, with robot vision. These never "see" a perfect or complete object, like the ones present by artificial objects.

The keypoint detectors evaluation pipeline used in this section is presented in figure 4.2. To evaluate the invariance of keypoint detection methods, the keypoints are extracted directly from the original cloud. Moreover, the transformation is applied in the original 3D point cloud before extracting another set of keypoints. Getting these keypoints from the transformed cloud, the inverse transformation is applied, so that it is possible to compare these with the keypoints extracted from the original cloud. If a particular method is invariant to the applied transformation, the keypoints extracted directly from the original cloud should correspond to the keypoints extracted from the cloud where the transformation was applied.



Figure 4.2: Keypoint detectors evaluation pipeline used in this section.

4.4.1 Keypoints Correspondence

The correspondence between the keypoints extracted directly from the original cloud and the ones extracted from the transformed cloud is done using the 3D point-line distance [338]. A line in three dimensions can be specified by two points $p_1 = (x_1, y_1, z_1)$ and $p_2 = (x_2, y_2, z_2)$ lying on it, then a vector line is produced. The squared distance between a point on the line with parameter t and a point $p_0 = (x_0, y_0, z_0)$ is therefore

$$d^{2} = [(x_{1} - x_{0}) + (x_{2} - x_{1})t]^{2} + [(y_{1} - y_{0}) + (y_{2} - y_{1})t]^{2} + [(z_{1} - z_{0}) + (z_{2} - z_{1})t]^{2}.$$
 (4.13)

To minimize the distance, set $\partial(d^2)/\partial t = 0$ and solve for t to obtain

$$t = -\frac{(p_1 - p_0) \cdot (p_2 - p_1)}{|p_2 - p_1|^2}, \quad (4.14)$$

where \cdot denotes the dot product. The minimum distance can then be found by plugging *t* back into equation 4.13. Using the vector quadruple product $((A \times B)^2 = A^2 B^2 - (A \cdot B)^2)$ and taking the square root results, is obtained:

$$d = \frac{|(p_0 - p_1) \times (p_0 - p_2)|}{|p_2 - p_1|}, \quad (4.15)$$

where \times denotes the cross product. Here, the numerator is simply twice the area of the triangle formed by points p_0 , p_1 , and p_2 , and the denominator is the length of one of the bases of the triangle.

4.4.2 Repeatability Measures

The most important feature of a keypoint detector is its *repeatability*. This feature takes into account the capacity of the detector to find the same set of keypoints in different instances of a particular model. The differences may be due to noise, view-point change, occlusion or by a combination of the above.

The repeatability measure used in this section is based on the measure used in [17] for 2D keypoints and in [19] for 3D keypoints. A keypoint extracted from the model M_h , k_h^i transformed according to the rotation, translation or scale change, (R_{hl}, t_{hl}) , is said to be repeatable if the distance d (given by the equation 4.15) from its nearest neighbor, k_l^j , in the set of keypoints extracted from the scene S_l is less than a threshold ε , $d < \varepsilon$.

The overall repeatability of a detector both in relative and absolute terms is evaluated. Given the set RK_{hl} of repeatable keypoints for an experiment involving the model-scene pair (M_h, S_l) , the absolute repeatability is defined as

$$r_{abs} = |RK_{hl}| \quad \textbf{(4.16)}$$

and the relative repeatability is given by

$$r = \frac{|RK_{hl}|}{|K_{hl}|}.$$
 (4.17)

The set K_{hl} is the set of all the keypoints extracted on the model M_h that are not occluded in the scene S_l (see figure 4.3). This set is estimated by aligning the keypoints extracted on M_h according to the rotation, translation and scale and then checking for the presence of keypoints in S_l in a small neighborhood of the transformed keypoints. If at least a keypoint is present in the scene in such a neighborhood, the keypoint is added to K_{hl} .


Figure 4.3: Graphical representation of sets of keypoints.

4.4.3 Results and Discussion

In this section, the invariance of the these methods are evaluated with respect to rotation, translation and scale changes. For this, the rotation is varied according to the three axes (X, Y and Z). The rotations applied ranged from 5° to 45° , with 10° step. The translation is performed simultaneously in the three axes and the image displacement applied on each axis is obtained randomly. Finally, the scale changes are applied in a random way (between $]1 \times .5 \times]$).

In table 4.2 presents some results about each keypoint detector applied to the original clouds. The percentage of clouds where the keypoint detectors successfully extracted (more than one keypoint) is presented in column 2. In the column 3, it appears the mean number of keypoints extracted by cloud. And finally, the mean computation time (in seconds) spent by each method to extract the keypoints is presented. These times were obtained on a computer with Intel®Core™i7-980X Extreme Edition 3.33GHz with 24 GB of RAM memory.

To make a fair comparison between the detectors, all steps in the pipeline (see figure 4.2) are equal. Figures 4.4, 4.5 and 4.6 show the results of the evaluation of the different methods with various applied transformations. The threshold distances (ε) analyzed vary between $[0, 2] \ cm$, with small jumps in a total of 33 equally spaced distances calculated. As presented in section 4.1, the methods have a relatively large set of parameters to be adjusted: the values used were the ones set by default in PCL.

Regarding the relative repeatability (shown in figures 4.4, 4.6(a) and 4.6(c)) the methods presented have a fairly good performance in general. In relation to the rotation (see figure 4.4),

Keypoint detectors	% Keypoint clouds	Mean of extracted keypoints	Mean time (s)
Harris3D	99.99	85.63	1.05
SIFT3D	99.68	87.46	9.54
ISS3D	97.97	86.24	1.07
SUSAN	86.51	242.38	1.64
Lowe	99.99	85.12	1.02
KLT	100.00	99.16	1.03
Curvature	99.96	119.36	0.70
Noble	99.99	85.12	1.04

Table 4.2: Statistics about each keypoint detector. These values come from processing the original clouds.



Figure 4.4: Rotation results represented by the relative repeatability measure (best viewed in color). The presented neighborhood radius is in meters.

increasing the rotation angle of the methods tends to worsen the results. Ideally, the method results should not change independently of the transformations applied. Regarding the applied rotation, the method ISS3D is the one that provides the best results. In this transformation (rotation), the biggest difference that appears between the various methods is in the 5 degrees rotation. In this case, the method ISS3D achieves almost total correspondence keypoints with a distance between them of $0.25 \ cm$. Whereas for example the SIFT3D only achieves this performance for keypoints at a distance of $1 \ cm$. In both the scaling and translation (shown in figures 4.6(a) and 4.6(c)), the methods exhibit very similar results to those obtained for small rotations (5° rotation in figure 4.4(a)) with the exception of the SUSAN method, that has a relatively higher invariance to scale changes.

Figures 4.5, 4.6(b) and 4.6(d) show the absolute repeatability, that present the number of keypoints obtained by the methods. With these results, it is clear to see that the method that has higher absolute repeatability (SUSAN) is not the one that shows the best performance in terms of relative repeatability. In terms of the absolute repeatability, the ISS3D and SIFT3D have better results than the SUSAN method regarding the invariance transformations evaluated



Figure 4.5: Rotation results represented by the absolute repeatability measure (best viewed in color). The presented neighborhood radius is in meters.

in this work.

4.5 Summary

This chapter was focused on the available keypoint detectors on the PCL and Open Source Computer Vision (OpenCV) library, explaining how they work, and made a comparative evaluation on public available data with real 3D objects. The description of the 3D keypoint detectors and the repeatability evaluation of these methods was published in [25, 26].

The experimental comparison proposed in this work has outlined aspects of state-of-theart methods for 3D keypoint detectors. This work allowed us to evaluate the best performance in terms of various transformations (rotation, scaling and translation).

The novelty of this work compared with the work of [17] and [19]: a real database is used instead of an artificial, the large number of point clouds and different keypoint detectors. The benefit of using a real database is that objects have "occlusion". This type of "occlusion" is made



Figure 4.6: Relative and absolute repeatability measures for the scale change and translation clouds (best viewed in color). The relative repeatability is presented in figures (a) and (c), and the absolute repeatability in figures (b) and (d). The presented neighborhood radius is in meters.

by some kind of failure in the infrared sensor of the camera or from the segmentation method. In artificial objects this does not happen, so the keypoint methods may have better results, but the experiments reflect what can happen in real life, such as, with robot vision. Overall, SIFT3D and ISS3D yielded the best scores in terms of repeatability and ISS3D demonstrated to be the more invariant.

Chapter 5

Retinal Color Extension for a 2D Keypoint Detector

Most object recognition algorithms use a large number of descriptors extracted in a dense grid, so they have a very high computational cost, preventing real-time processing. The use of keypoint detectors allows the reduction of the processing time and the amount of redundancy in the data. Local descriptors extracted from images have been extensively reported in the computer vision literature. In this chapter, a keypoint detector inspired by the behavior of the early visual system is presented. The method is a color extension of the BIMP keypoint detector, where includes both color and intensity channels of an image. The color information is included in a biological plausible way and reproduces the color information in the retina. Multi-scale image features are combined into a single keypoints map. The detector is compared against state-of-the-art detectors and is particularly well-suited for tasks such as category and object recognition. The evaluation gave the best pair keypoint detector/descriptor on a RGB-D object dataset. Using this keypoint detector and the SHOTCOLOR descriptor a good category recognition rate is obtained and for object recognition it is with the PFHRGB descriptor that the best results are obtained.

5.1 Proposed 2D Keypoint Detector

Biological Motivated Multi-Scale Keypoint Detector (BMMSKD) is a color information extension of BIMP. The way in which the color information is added is based on a neural architecture of the primate visual system [3, 119]. Figure 5.1 presents the block diagram of the keypoint detector.

For a given color image, three images from the RGB channels are created, which are: RG, BY and grayscale image I (shown in the left column of the figure 5.2).

The *r*, *g*, and *b* channels are normalized by *I* in order to decouple hue from intensity. However, because hue variations are not perceivable at very low luminance (and hence are not salient), normalization is only applied at the locations where *I* is larger than 1/10 of its maximum



Figure 5.1: Block diagram of the proposed 2D keypoint detector method. Our method receives an image directly from the camera and generates the three new images (RG, BY and I). In each of these images the BIMP keypoint detector is applied and the result of the three detections is fused. See the text for details.





Figure 5.2: Our keypoint detection method. The first column shows the original image on the top and the keypoint fusion on the bottom. The second, third and fourth columns contain the RG, BY and gray color channels (top) and the respective keypoint detection on the bottom.

over the entire image (other locations yield zero r, g, and b). Four broadly-tuned color channels are created: R for red channel, G for green, B for blue and Y for yellow:

$$R = \frac{r - (g + b)}{2}, \quad (5.1)$$

$$G = \frac{g - (r + b)}{2}, \quad (5.2)$$

$$B = \frac{b - (r + g)}{2} \quad \text{and} \quad (5.3)$$

$$Y = \frac{r + g}{2} - \frac{|r - g|}{2 - b}. \quad (5.4)$$

Accordingly, maps

$$RG = R - G$$
 (5.5)

are created in the model to represent the red/green opponency and

$$BY = B - Y \quad (5.6)$$

for blue/yellow opponency (negative values are set to zero). For each color channel RG, BY and I, the BIMP keypoint detector is applied and the keypoint locations are fused.

Given the application of the BIMP method on each channel, three sets of keypoints k_{RG} , k_{BY} and k_I are obtained, respectively (shown in the right column of the second to fouth rows of

figure 5.2). With these three sets, a keypoint map K_m given by

$$K_m = k_{RG} \cup k_{BY} \cup k_I \quad (5.7)$$

is created. A location is considered a keypoint, if there exists another color channel, in its neighborhood, which indicates that there exists one keypoint in the region. This is:

$$k_l \in K_m : \#K_m^r(k_l) > 1$$
, (5.8)

where k_l is a keypoint location, r the neighborhood radius and $K_m^r(k_l)$ is a sub-set of K_m centered in the point k_l and with radius r. An example of the fusion result is presented in the bottom of the first column in figure 5.2.

5.2 Object Recognition Pipeline

In this section, the pipeline used in this work is presented, shown in figure 5.3. Each block will be explained in the following subsections.



Figure 5.3: Block diagram of the 3D recognition pipeline.

5.2.1 Segmented Objects and Object Database

The input camera and segmentation process is simulated by the large RGB-D Object Dataset [21]. A set of 5 images/point clouds of each physically distinct object, using a total of 1500 from each of them.

Using the 1500 images and point clouds selected, the observations are given by the Leave-One-Out Cross-Validation (LOOCV) method [339]. As the name suggests, LOOCV involves using a single observation from the original sample as the validation data, and the remaining observations as the training data. This is repeated such that each observation in the sample is used once as the validation data. This is the same as a K-fold cross-validation with K being equal to the number of observations in the original sampling. With 1500 images/point clouds and LOOCV method is possible to perform more than 2200000 comparisons for each pair keypoint detector/descriptor. A total of 60 pairs (4 keypoint detectors \times 15 descriptors) are evaluated.

5.2.2 2D Keypoint Detectors

The 2D image of the segmented object, present in the database, will feed the keypoint extraction process, which is used to reduce the computational cost of the recognition system. The keypoints implementation used is done in OpenCV library [340]. Figure 5.4 is an example of the keypoint extraction by the four methods on an image of the used dataset. In table 5.1, the average number of keypoints, mean computation time (in seconds) spent by each method to extract the keypoints and the file size (in KiloBytes) is presented. These times were obtained on a computer with Intel®Coreii7-980X Extreme Edition 3.33GHz with 24 GB of RAM memory.

5.2.3 3D Descriptors

The descriptors are extracted at the locations given by the keypoint detector obtained from the 2D images, but the processing of descriptors is done in point clouds. The point clouds have the 3D information of the segmented object, which is composed by: color (in the RGB color space) and depth information. In table 5.2 are presented some statistics about the extracted descriptors using the keypoint detectors (like in table 5.1).

5.2.4 Distance Measure and Matching

One of the stages in recognition is the correspondence between a input descriptors and a known object cloud (stored in the database). The correspondence is typically done using a distance function between the sets of descriptors. In [23], multiple distance functions are studied. In this work, the distance used is defined by

Table 5.1:	Keypoints	statistics	for 2D keypo	int detecto	rs. The r	number o	of points,	time in	seconds	(s) a	ind
size in kilo	obytes (KB)	presented	are related	to each clou	ud in the	processi	ng of the	test set	•		

Keypoint	Number of Po	Time	(s)	Size (KB)		
Detectors	Mean±Std	Median	$Mean\pm Std$	Median	Mean±Std	Median
BMMSKD	142.03±141.00	92.00	10.65±1.61	10.45	6.55±6.22	4.36
BIMP [7]	$56.05{\pm}53.07$	37.00	3.93±0.90	3.82	2.69±2.35	1.83
SIFT [9]	46.83±63.02	24.00	$0.26{\pm}0.07$	0.23	2.27±2.78	1.27
SURF [10]	47.77±60.06	24.00	0.28±0.07	0.28	2.32±2.67	1.28
Average	73.26±95.78	41.00	3.79±4.34	1.52	3.46±4.24	2.05
Original	5740.06±6851.42	3205.00			316.86±375.73	177.23



Figure 5.4: Example of the keypoints extracted by the four methods in an image.

Descriptor	Time (s	5)	Size (KB)	
Descriptor	Mean±Std	Median	Mean±Std	Median
3DSC	37.64±97.19	6.25	342.94±459.56	189.14
CVFH	0.02±0.02	0.01	0.77±0.12	0.75
ESF	0.18±0.06	0.17	7.33±0.54	7.46
FPFH	0.51±0.66	0.28	16.10±21.06	9.08
OUR-CVFH	0.01±0.01	0.01	$0.75{\pm}0.00$	0.75
PCE	0.02±0.02	0.01	$4.48{\pm}5.55$	2.63
PFH	1.12±1.97	0.47	26.81±35.95	14.97
PFHRGB	1.96±3.48	0.80	55.86±76.21	30.66
PPF	0.09±0.26	0.02	809.94±3116.22	95.29
PPFRGB	0.03±0.03	0.02	6.79±8.84	3.92
SHOT	0.05±0.06	0.03	101.37±136.74	55.68
SHOTCOLOR	0.06±0.07	0.03	310.37±419.98	169.47
SHOTLRF	0.03±0.03	0.02	$7.59{\pm}9.70$	4.34
USC	35.29±90.55	6.19	351.67±471.41	194.15
VFH	0.02±0.02	0.01	1.10±0.22	1.07
Average	6.21±40.02	0.09	141.76±875.97	9.94

Table 5.2: Descriptors statistics (for more details see caption of table 5.1).

$$D_6 = L_1(c_A, c_B) + L_1(std_A, std_B)$$
 (5.9)

that presents good results, in terms of recognition and run time, where c_A and c_B are the centroids of the sets A and B, respectively, and

$$std_A(i) = \sqrt{\frac{1}{|A| - 1} \sum_{j=1}^{|A|} (a_j(i) - c_A(i))^2}, i = 1, \dots, n,$$
 (5.10)

 $a_j(i)$ refers to the coordinate *i* of the descriptor *j*, and likewise for std_B . The L_1 distance is between descriptor (not sets) $x, y \in X$ and is given by

$$L_1(x, y) = \sum_{i=1}^{n} |x(i) - y(i)|.$$
 (5.11)

5.2.5 Recognition Measures

In order to perform the recognition evaluation will be used three measures, which are the Receiver Operator Characteristic (ROC) curve, the Area Under the ROC Curve (AUC) and the Decidability (DEC). The DEC index [341] is given by

$$DEC = |\mu_{intra} - \mu_{inter}| / \sqrt{\frac{1}{2}(\sigma_{intra}^2 + \sigma_{inter}^2)}$$
 (5.12)

is the distance between the distributions obtained for the two classical types of comparisons: between descriptors extracted from the same (*intra-class*) and different objects (*inter-class*). The μ_{intra} and μ_{inter} denote the means of the intra- and inter-class comparisons, σ_{intra}^2 and σ_{inter}^2 the respective standard deviations and the decidability can vary between $[0, \infty]$.

In statistics, a ROC is a graphical plot that illustrates the performance of a binary classifier system as its discrimination threshold is varied. The curve is created by plotting the true positive rate (TPR) against the false positive rate (FPR) at various threshold settings. It is also known as a relative operating characteristic curve, because it is a comparison of two operating characteristics (TPR and FPR) as the criterion changes [342]. The AUC is equal to the probability that a classifier will rank a randomly chosen positive instance higher than a randomly chosen negative one (assuming 'positive' ranks higher than 'negative') [343].

5.3 Results and Discussion

The obtained AUC and DEC for category and object recognition are given in tables 5.3 and 5.4. The ROCs for category and object recognition are shown in figures 5.5 and 5.6, respectively.

As shown in table 5.3 and 5.4, the method presented here increases the recognition results in both category and object recognition. Comparing this one with the original approach, it is possible to see that color information has introduced a significant improvement in both category and object recognition.

For the category recognition (table 5.3), the BMMSKD method presented here shows worst results only in three cases for the AUC measure and in six cases for the DEC. In the other pair cases, it has significant improvements compared to the other three keypoint detection methods,

	Category Recognition									
Descriptors	BMN	SKD	BIMI	P [7]	SIFT	[9]	SURF	[10]	Ave	rage
	AUC	DEC	AUC	DEC	AUC	DEC	AUC	DEC	AUC	DEC
3DSC	<u>0.667</u>	0.221	0.644	0.182	0.652	0.293	0.654	<u>0.294</u>	0.654	0.248
CVFH	<u>0.601</u>	<u>0.270</u>	0.590	0.194	0.582	0.158	0.580	0.148	0.558	0.193
ESF	0.748	0.825	0.748	0.827	<u>0.754</u>	<u>0.873</u>	0.753	0.865	0.751	0.848
FPFH	0.720	0.717	0.689	0.623	<u>0.725</u>	<u>0.796</u>	0.723	0.769	0.714	0.726
OURCVFH	<u>0.615</u>	0.338	0.588	0.291	0.585	0.275	0.595	0.284	0.596	0.297
PCE	<u>0.616</u>	0.360	0.597	0.305	0.600	0.302	0.595	0.292	0.602	0.315
PFH	<u>0.742</u>	0.843	0.716	0.746	0.739	<u>0.872</u>	0.737	0.853	0.734	0.829
PFHRGB	<u>0.773</u>	<u>1.003</u>	0.758	0.929	0.768	0.999	0.765	0.984	0.766	0.979
PPF	<u>0.626</u>	<u>0.433</u>	0.600	0.336	0.593	0.297	0.597	0.317	0.604	0.346
PPFRGB	0.551	0.011	0.553	0.013	<u>0.554</u>	<u>0.056</u>	0.537	0.028	0.549	0.027
SHOT	<u>0.631</u>	<u>0.372</u>	0.614	0.355	0.613	0.334	0.609	0.312	0.617	0.343
SHOTCOLOR	<u>0.700</u>	0.609	0.674	0.570	0.684	0.591	0.679	0.575	0.684	0.586
SHOTLRF	<u>0.681</u>	<u>0.489</u>	0.640	0.378	0.626	0.321	0.631	0.325	0.645	0.378
USC	<u>0.660</u>	0.233	0.635	0.184	0.640	0.291	0.644	0.295	0.645	0.251
VFH	<u>0.592</u>	<u>0.317</u>	0.575	0.260	0.591	0.314	0.591	0.313	0.587	0.301
Average	0.662	0.470	0.641	0.413	0.647	0.451	0.646	0.441		

Table 5.3: AUC and DEC values for the category recognition for each pair 2D keypoints/descriptor. The underline value is the best result for this descriptor and the best pair is the bold one.

	Object Recognition									
Descriptors	BMN	\SKD	BIM	BIMP [7]		SIFT [9]		⁻ [10]	Ave	rage
	AUC	DEC	AUC	DEC	AUC	DEC	AUC	DEC	AUC	DEC
3DSC	<u>0.690</u>	0.244	0.658	0.196	0.657	0.295	0.666	0.307	0.668	0.261
CVFH	<u>0.628</u>	<u>0.305</u>	0.617	0.252	0.611	0.210	0.610	0.206	0.617	0.243
ESF	0.818	1.111	0.820	1.131	0.820	1.151	<u>0.823</u>	<u>1.153</u>	0.820	1.137
FPFH	0.776	0.916	0.729	0.753	0.774	0.980	<u>0.781</u>	<u>0.997</u>	0.765	0.912
OURCVFH	<u>0.659</u>	<u>0.511</u>	0.626	0.384	0.616	0.340	0.614	0.301	0.629	0.384
PCE	<u>0.632</u>	0.407	0.606	0.329	0.603	0.311	0.614	0.352	0.614	0.350
PFH	0.794	1.062	0.759	0.910	0.792	1.105	<u>0.796</u>	<u>1.110</u>	0.785	1.047
PFHRGB	<u>0.920</u>	1.923	0.903	1.778	0.890	1.700	0.893	1.728	0.902	1.782
PPF	<u>0.655</u>	<u>0.528</u>	0.626	0.417	0.613	0.353	0.625	0.400	0.630	0.425
PPFRGB	0.568	0.031	0.542	0.055	0.579	<u>0.076</u>	<u>0.581</u>	0.034	0.568	0.049
SHOT	<u>0.661</u>	<u>0.457</u>	0.622	0.377	0.624	0.389	0.616	0.331	0.631	0.389
SHOTCOLOR	<u>0.787</u>	<u>0.911</u>	0.740	0.794	0.730	0.764	0.730	0.762	0.747	0.808
SHOTLRF	<u>0.707</u>	<u>0.564</u>	0.654	0.422	0.624	0.294	0.634	0.338	0.655	0.405
USC	<u>0.683</u>	0.260	0.655	0.214	0.656	0.330	0.659	0.321	0.663	0.281
VFH	<u>0.623</u>	<u>0.432</u>	0.595	0.334	0.609	0.372	0.604	0.367	0.608	0.376
Average	0.707	0.644	0.677	0.556	0.680	0.578	0.683	0.590		

Table 5.4: AUC and DEC values for the object recognition for each pair keypoints/descriptor. The underline value is the best result for this descriptor and the best pair is the bold one.

which are also visible in the graphs of the figure 5.5. The best result for the category recognition was obtained by the pair BMMSKD/PFHRGB, being the only one where the DEC exceeds the index 1.000. Moreover, the BMMSKD only presents a lower AUC than the average for the ESF descriptor.

Regarding the object recognition the results are presented in table 5.4 and in the graphs of the figure 5.6. Comparing the overall results of the SIFT keypoints detector in the category recognition with those of the SURF keypoints detector in the object recognition, it is found that there is an inversion. That is, while the SIFT method was the one who showed better results several times here is quite the opposite. Overall, in the object recognition there exists a improvement in the results for all the methods, because there is less variation in the data.

As can be seen, the retinal color extension, introduced in the BMMSKD keypoint detector, produces a good improvement compared to the original method. For grayscale images, the results will be the same as the BIMP, since this method generalizes it to color.

5.4 Summary

This chapter focused on keypoint detectors and presented a novel keypoint detector biologically motivated by the behavior and the neuronal architecture of the early primate visual system. This new method and part of the presented results were published in [27]. The recognition evaluation is done on a public available dataset with real 3D objects. The keypoint detectors were developed using the OpenCV library and the keypoint locations are projected to the 3D space in order to use available 3D descriptors on the PCL library.

The main conclusions of this chapter are: 1) the keypoint locations can help or degrade the recognition process; 2) a descriptor that uses color information should be used instead of

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Figure 5.5: ROCs for the category recognition experiments using 2D keypoint detectors (best viewed in color).



Figure 5.6: ROCs for the object recognition experiments using 2D keypoint detectors (best viewed in color).

a similar one that uses only shape information; 3) since there are big differences in terms of recognition performance, size and time requirements, the descriptor should be matched to the desired task; 4) to recognize the category of an object or a real-time system, it is recommended the use the SHOTCOLOR method because it presents a recognition rate of 7% below of the PFHRGB but with a much lower computational cost; and 5) to do the object recognition, the recommendation is PFHRGB because it presents a recognition rate 12.9% higher than SHOTCOLOR.

Chapter 6

Biologically Inspired 3D Keypoint Detector based on Bottom-Up Saliency

A new method for the detection of 3D keypoints on point clouds is presented. A benchmarking between each pair of 3D keypoint detector and 3D descriptor is performed, in order to evaluate their performance on object and category recognition. These evaluations are done in a public database of real 3D objects. The keypoint detector is inspired by the behavior and neural architecture of the primate visual system. The 3D keypoints are extracted based on a bottom-up 3D saliency map, that is, a map that encodes the saliency of objects in the visual environment. The saliency map is determined by computing conspicuity maps (a combination across different modalities) of the orientation, intensity and color information in a bottom-up and in a purely stimulus-driven manner. These three conspicuity maps are fused into a 3D saliency map and, finally, the focus of attention (or "keypoint location") is sequentially directed to the most salient points in this map. Inhibiting this location automatically allows the system to attend to the next most salient location.

6.1 Proposed 3D Keypoint Detector

The Biologically Inspired 3D Keypoint based on Bottom-Up Saliency (BIK-BUS) is a keypoint detector that is based on saliency maps. The saliency maps are determined by computing conspicuity maps of the features intensity, color and orientation in a bottom-up and data-driven manner. These conspicuity maps are fused into a saliency map and, finally, the focus of attention is sequentially directed to the most salient points in this map [120]. Using this theory and following the steps presented in [3, 119], a new keypoint detector is presented (shown in figure 6.1).

6.1.1 Linear Filtering

The initial part of this step is similar to the retinal color extension presented in section 5.1. Here, the four broadly-tuned color channels (R, G, B and Y) and the intensity I channel are also used.

Gaussian pyramids [121] are used in the spatial scales, which progressively low-pass and down-sample the input cloud, producing horizontal and vertical cloud-reduction factors. Five Gaussian pyramids $R(\sigma)$, $G(\sigma)$, $B(\sigma)$, $Y(\sigma)$ and $I(\sigma)$ are created from the color and intensity channels, where σ represents the standard deviation used in the Gaussian kernel.

Each Gaussian pyramid is achieved by convolving the cloud with Gaussian kernels of increasing radius, resulting in a pyramid of clouds. We apply a similar concept to search the density map D over a range of scales, where D can be {R, G, B, Y, I}. We convolve D with a set of 3D Gaussian kernels to construct a pyramid of density maps, with each layer representing the scale σ . A factor of 2 is used to down-sample the density map and the reduction of the

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Figure 6.1: General architecture of our Biologically Inspired Keypoint Detector based on Bottom-Up Saliency. Our method receives as input a point cloud similar to those shown in figures 4.1 and 6.3 and a linear filter is applied to obtain the color, intensity and orientations information. The full process is described in the text.

standard deviation of the Gaussian kernel by $\sqrt{2}$. The pyramid creation is a step similar to the DoG presented in the section 4.1.5.

Let $L(\cdot)$ (one of the five Gaussian pyramids) be a scale space for D:

$$L(x, y, z, \sigma) = D * g(x, y, z, \sigma), \quad (6.1)$$

where * is the convolution operator and $g(x, y, z, \sigma)$ is a 3D Gaussian with standard deviation σ given by:

$$g(x, y, z, \sigma) = \exp\left(\frac{-x^2 - y^2 - z^2}{2\sigma^2}\right).$$
 (6.2)

The orientation pyramids $O(\sigma, \theta)$ are obtained using the normals extracted from the intensity cloud *I*, where $\theta \in \{0^o, 45^o, 90^o, 135^o\}$ is the preferred orientation [121]. In the primary visual cortex, the impulse response of orientation-selective neurons is approximated by Gabor filters [122]. The orientation pyramids are created in a similar way to the color channels, but applying 3D Gabor filters with different orientations θ .

6.1.2 Center-Surround Differences

In the retina, bipolar and ganglion cells encode the spatial information, using centersurround structures. The center-surround structures in the retina can be described as *on-center* and *off-center*. The *on-center* use a positive weighed center and negatively weighed neighbors. The *off-center* use exactly the opposite. The positive weighing is better known as excitatory and the negative as inhibitory [123].

Similarly to the visual receptive fields, a set of linear center-surround operations is used to compute each feature. Visual neurons are most sensitive in a small region of the visual space (the center), while stimuli in the surround inhibit neuronal response [3]. Center-surround is computed as the difference between the center pixel at scale $c \in \{2, 3, 4\}$, and the surround is the corresponding pixel at scale

$$s = c + \delta$$
 , (6.3)

with $\delta \in \{3, 4\}$. The across-scale difference between two maps (represented by \bigcirc) is obtained by interpolation to the center scale c and point-by-point subtraction.

The first set of feature maps is concerned with intensity contrast. In mammals, this is detected by neurons sensitive either to dark centers on bright surrounds (*off-center*) or to bright centers on dark surrounds (*on-center*) [3, 122]. Here, both types of sensitivities are simultaneously computed in a set of six maps I(c, s):

$$I(c, s) = |I(c) \bigoplus I(s)|.$$
 (6.4)

For the color channels, the process is similar, which, in the cortex, is called `color doubleopponent' system [3]. In the center of their receptive fields, neurons are excited by one color and inhibited by an other, while the converse is true in the surround. The existence of a spatial and chromatic opponency between color pairs in human primary visual cortex is described in [124]. Given the chromatic opponency, the maps RG(c, s) and BY(c, s) are created to take in account the red/green and green/red, and blue/yellow and yellow/blue double opponency, respectively, as:

$$RG(c, s) = |(R(c) - G(c)) \bigoplus (G(s) - R(s))|, \quad (6.5)$$

$$BY(c, s) = |(B(c) - Y(c)) \bigoplus (Y(s) - B(s))|.$$
 (6.6)

Orientation feature maps, $O(c, s, \theta)$, encode, as a group, local orientation contrast between the center and surround scales:

$$O(c, s, \theta) = |O(c, \theta) \bigoplus O(s, \theta)|.$$
 (6.7)

6.1.3 Normalization

We cannot combine directly the different feature maps because they represent different dynamic ranges and extraction mechanisms. Some salient objects appear only in a few maps, which can be masked by noise or by less salient objects present in a larger number of maps. In order to resolve that, a map normalization operator $\mathcal{N}(.)$ is used. This promotes the maps that

contain a small number of strong activity, and suppresses the peaks in the maps that have many of them [3]. $\mathcal{N}(.)$ consists of:

- 1. Large amplitude differences are eliminated by normalizing the map values to a fixed range [0..M], where M is the global maximum of the map;
- 2. Multiply the map by $(M \overline{m})^2$, where \overline{m} is the average of all its other local maxima.

The lateral cortical inhibition is the biological motivation for this normalization [344].

6.1.4 Across-Scale Combination

The conspicuity maps are the combination of the feature maps, for intensity, color and orientation. They are obtained through the reduction of each map to scale four and point-by-point addition ` \oplus ', called across-scale addition. The conspicuity maps for the intensity, \overline{I} , and color channels, \overline{C} , are given by:

$$\overline{I} = \bigoplus_{c=2}^{4} \bigoplus_{s=c+3}^{c+4} \mathcal{N}(I(c,s)) \text{ and } (6.8)$$

$$\overline{C} = \bigoplus_{c=2}^{4} \bigoplus_{s=c+3}^{c+4} \left[\mathcal{N}(RG(c,s)) + \mathcal{N}(BY(c,s)) \right].$$
 (6.9)

For orientation, we first created four intermediary maps, which are a combination of the six feature maps for a given θ . Finally, they are combined into a single orientation conspicuity map:

$$\overline{O} = \sum_{\theta \in \{0^o, 45^o, 90^o, 135^o\}} \mathcal{N} \left[\bigoplus_{c=2}^4 \bigoplus_{s=c+3}^{c+4} \mathcal{N}(O(c, s, \theta)) \right].$$
(6.10)

The three separate channels $(\overline{I}, \overline{C} \text{ and } \overline{O})$ have an independent contribution in the saliency map and where similar features will have a strong impact on the saliency.

6.1.5 Linear Combination

The final saliency map is obtained by the normalization and a linear combination between them:

$$S = \frac{1}{3} \left(\mathcal{N}(\overline{I}) + \mathcal{N}(\overline{C}) + \mathcal{N}(\overline{O}) \right). \quad (6.11)$$

6.1.6 Inhibition-Of-Return

The IOR is part of the method that is responsible for the selection of keypoints. It detects the most salient location and directs attention towards it, considering that location a keypoint. After that, the IOR mechanism transiently suppresses this location in the saliency map and its neighborhoods in a small radius, such that attention is autonomously directed to the next most salient image location. The suppression was achieved replacing saliency map values with zero. The following iteration will find the most salient point (the maximum) in different location. This iterative process stops when the maximum of the saliency map reaches a certain value (a minimum), which is defined by a threshold. Computationally, the IOR performs a similar process of selecting the global and local maximums.

6.2 3D Object Recognition Pipeline

This section presents the pipeline used in this work, shown in figure 6.2. The input clouds used are given by the RGB-D Object Dataset [21] presented in the section 4.3. These point clouds will feed the keypoint extraction process (see more details in section 6.2.1), which are used to reduce the computational cost of the recognition system.

Typically, the largest computational cost of these systems is at the stage of computing the descriptors, so, it makes sense to use only a subset of the input clouds. In figure 6.2, the cloud input also feeds the descriptors extraction, but it is only used to obtain information about the keypoints neighbors (to calculate the normals at the point). A set of object descriptors is compared to those that have been previously computed and which are in the object database. The one that presents the smallest distance is considered as the corresponding object.



Figure 6.2: Block diagram of the 3D recognition pipeline.

Portions of this pipeline, as well as the point clouds, are the same as presented in the previous chapter. The major differences between these two chapters are relative to the number of pairs of keypoint detectors/descriptors evaluated and the fact that these keypoint detectors work directly on the point clouds and not in 2D images. Here, a total of 135 pairs (9 keypoint detectors \times 15 descriptors) are evaluated.

6.2.1 Keypoint Extraction

The keypoint detection methods have many parameters to adjust, but normally the default values in the PCL are used. For all the keypoint detectors, the search radius is always the same and defined to 1cm. The Susan and SIFT3D methods were those where it was necessary to define more parameters. For the Susan method, there are two parameters: the

Keypoint	Number of Po	ints	Time	(s)	Size (KB	Size (KB)		
Detectors	Mean±Std	Median	Mean±Std	Median	Mean±Std	Median		
BIK-BUS	117.43±114.74	72.00	6.66±9.14	3.67	5.83±1.79	5.12		
Curvature	116.51 ± 125.56	68.00	0.78±1.43	0.31	5.82±1.96	5.06		
Harris3D	83.61±95.64	47.00	1.11±2.01	0.46	5.31±1.49	4.73		
ISS3D	84.54±107.34	43.00	1.13±2.01	0.46	$5.32{\pm}1.68$	4.67		
KLT	96.45±109.02	54.00	1.16±2.12	0.46	$5.51 {\pm} 1.70$	4.84		
Lowe	83.05±95.43	46.00	1.21±2.48	0.45	5.30±1.49	4.72		
Noble	83.05±95.43	46.00	1.18±2.18	0.45	5.30±1.49	4.72		
SIFT3D	85.11±103.97	43.00	2.51±3.61	1.20	5.33±1.62	4.67		
SUSAN	132.52±483.05	14.00	1.54±2.74	0.64	$6.07{\pm}7.55$	4.22		
Average	98.01±190.32	48.00	1.92±4.17	0.63	5.53±2.97	4.75		
Original	5740.06±6851.42	3205			316.86±375.73	177.23		

Table 6.1: Statistics of the 3D keypoint detectors. The parameters are the same as those presented in table 5.1.

 $distance_threshold = 0.01cm$ is used to test if the nucleus is far enough from the centroid; and the $angular_threshold = 0.01cm$ to verify if the normals are parallel. In the SIFT3D, the parameters defined are: $min_scale = 0.002$, $nr_octaves = 4$, $nr_scales_per_octave = 4$ and the $min_contrast = 1$. These parameters were adjusted with these values, such that all methods present a similar average number of the keypoints (as can be seen in table 6.1). Figure 6.3 presents a cloud of points where the several keypoint detectors were applied with these parameters.

Table 6.1 also presents some statistics about the keypoints extracted from the selected point clouds. To get an idea of the reduction between the input points clouds and the keypoints, the last row of the table contains the statistics information about the input point clouds. All the processing time was calculated based on *Intel Core I7 Extreme Edition X980* (3.3GHz), 24Gb RAM (FSB 1066) and *Fedora Core 14* operating system.

6.2.2 Descriptor Extraction

One of the goals was to evaluate the available descriptors in the current PCL version (1.7 pre-release) [20]. There are some descriptors in PCL which are not consider in this evaluation, since they are not applicable to point cloud data directly or they are not object descriptors, some of them are pose descriptors (6DoF).

It's only possible to make a fair comparison between the descriptors if they always use the same parameters in all steps of the pipeline, shown in figure 6.2. In the parametric configuration of the descriptors, the default values defined in the PCL were used. For the descriptors that use normals, a radius of 1*cm* was used for the calculus of normal and for the normal estimation radius search.

6.3 Experimental Evaluation and Discussion

The obtained AUC and DEC are given in table 6.3, while the ROCs for category and object recognition are presented in figures 6.4 and 6.5, respectively. Table 6.4 presents the infor-



Figure 6.3: Keypoint detectors applied on a "food_box" point cloud. The red points are the keypoints extracted from each detector and the number of these is presented in the legend of each sub-figure (best viewed in color).

mation about the number of times that each keypoint detector achieved the best result in the category and object recognition and the sums of these counts (Total column). When there is a tie between two methods both methods score.

Analyzing the descriptors in a generic way, the best results were obtained with the PFHRGB. It is interesting to compare it to the PFH: improvement can only be attributed to the incorporation of color information. The same is true for the SHOTCOLOR versus the SHOT descriptor. The two best results in terms of category and object recognition are presented in

Table 6.2: Statistics of the evaluated descriptors in this work. The time in seconds (s) and size in kilobytes (KB) presented are related to each cloud in the processing of the test set. To know the total time or the total size spent by a database of one of this descriptor. To obtain the total size of the database, you need to multiply the size presented by the number of clouds in the database.

Doscriptors	Time (s)		Size (KB	Size (KB)			
Descriptors	Mean±Std	Median	Mean±Std	Median			
3DSC	9181.56±21135.49	1138.86	725.28±830.02	408.02			
CVFH	0.32±0.47	0.17	$5.26{\pm}0.30$	5.20			
ESF	3.15±2.55	2.68	$6.50{\pm}0.00$	6.50			
FPFH	25.21±32.82	13.79	15.97±13.77	10.70			
OUR-CVFH	$0.32{\pm}0.59$	0.25	5.47±0.71	5.20			
PCE	1.23±2.02	0.45	5.81±2.09	5.02			
PFH	4157.63±7911.99	1129.62	49.33±52.16	29.39			
PFHRGB	8077.11±15432.91	2188.16	95.47±104.70	55.76			
PPF	1.26±2.96	0.42	398.50 ± 953.27	57.69			
PPFRGB	4.07±5.25	2.22	7.09±3.71	5.56			
SHOT	1.60±2.06	0.94	134.85±150.65	77.33			
SHOTCOLOR	1.88±3.04	1.01	494.41±564.62	278.83			
SHOTLRF	$0.72{\pm}0.80$	0.49	$7.26{\pm}3.75$	5.83			
USC	9125.88±20892.66	1135.20	728.12±831.69	408.02			
VFH	0.24±0.43	0.03	$5.20{\pm}0.00$	5.20			
Average	2362.27±10257.14	2.23	187.81±504.45	13.22			

the descriptors that use color information. The ROCs, in figures 6.4 and 6.5, also show the superiority of these two descriptors (that use color) versus the remaining. FPFH is an extension of PFH and it has a performance slightly worst than the original descriptor, but it is faster to extract and uses about half the space (shown in table 6.2), as the authors of the descriptor suggested. An interesting result is the one obtained by PPFRGB which is an color extension of PPF: in this case the none color version is better than the color version.

The USC was proposed as an upgrade to the 3DSC and our results confirm that in fact it improves the 3DSC results. Only when, the SUSAN keypoint detector was used in both recognition tasks, the 3DSC beats the USC in most of the cases.

Considering OUR-CVFH an upgrade of CVFH and this one an extension of VFH, it is not able to see where are improvements because both have lower scores and the processing times are slightly higher than the original descriptor.

In terms of computational time and space, the descriptor's requirements varies a lot. If the application needs real-time performance or when using embedded devices with limited resources there are some descriptors that cannot be considered.

Considering only the accuracy, the best combination for the category recognition is BIK-BUS /PFHRGB, closely followed by BIK-BUS/SHOTCOLOR, ISS3D/PFHRGB and ISS3D/SHOTCOLOR both in terms of AUC and DEC. The pairs BIK-BUS/PFHRGB and BIK-BUS /SHOTCOLOR have exactly the same AUC, the difference is in the DEC where it is slightly higher in the case of PFHRGB. BIK-BUS turns out again the best performer among detectors: FPFH, PPF, SHOT, SHOTCOLOR, USC and VFH. In relation to the 3DSC and SHOTLRF descriptors, our keypoint detector obtains the best DEC while the AUC is better when using Curvature keypoint detector in both descriptors.

Descriptors	Koupointo	Cate	egory	Obj	ject	Time (s)	
Descriptors	Reypoints	AUC	DEC	AUC	DEC		
	BIK-BUS	0.711	0.519	0.749	0.612	11166.25	
	Curvature	0.712	0.491	0.756	0.602	10406.55	
	Harris3D	0.706	0.472	0.740	0.539	8402.61	
	ISS3D	0.706	0.504	0.746	0.603	9581.85	
2050	KLT	0.709	0.486	0.748	0.579	9208.83	
3030	Lowe	0.705	0.468	0.746	0.560	8027.55	
	Noble	0.707	0.477	0.749	0.573	7894.47	
	SIFT3D	0.700	0.511	0.727	0.568	8745.17	
	SUSAN	0.656	0.399	0.682	0.466	12934.55	
	Average	0.701	0.483	0.738	0.567	9596.43	
	BIK-BUS	0.605	0.241	0.633	0.286	6.90	
	Curvature	0.604	0.258	0.633	0.283	1.02	
	Harris3D	0.606	0.249	0.632	0.256	1.33	
	ISS3D	0.608	0.235	0.637	0.253	1.34	
CVEL	KLT	0.606	0.252	0.633	0.270	1.38	
CVFH	Lowe	0.606	0.248	0.634	0.253	1.42	
	Noble	0.604	0.241	0.626	0.243	1.39	
	SIFT3D	0.594	0.170	0.635	0.255	2.73	
	SUSAN	0.560	0.020	0.573	0.038	2.00	
	Average	0.599	0.213	0.626	0.179	2.17	
	BIK-BUS	0.748	0.843	0.821	1.151	9.85	
	Curvature	0.746	0.817	0.817	1.109	3.83	
	Harris3D	0.747	0.821	0.822	1.133	4.35	
	ISS3D	0.747	0.827	0.818	1.130	4.30	
FSF	KLT	0.745	0.811	0.818	1.110	4.30	
	Lowe	0.746	0.815	0.818	1.110	4.32	
	Noble	0.748	0.827	0.819	1.114	4.34	
	SIFT3D	0.750	0.847	0.823	1.166	5.63	
	SUSAN	0.751	0.854	0.826	1.184	4.67	
	Average	0.748	0.829	0.820	1.257	5.07	

Descriptors	Koupointo	Cate	gory	Obj	ect	Time (s)
Descriptors	Keypoints	AUC	DEC	AUC	DEC	Time (s)
	BIK-BUS	0.844	1.434	0.900	1.833	32.67
	Curvature	0.844	1.433	0.899	1.829	25.93
	Harris3D	0.836	1.375	0.889	1.730	26.63
	ISS3D	0.843	1.429	0.900	1.841	26.39
	KLT	0.840	1.395	0.892	1.746	26.06
FPFH	Lowe	0.839	1.391	0.892	1.752	26.00
	Noble	0.840	1.397	0.893	1.753	26.14
	SIFT3D	0.837	1.377	0.897	1.806	27.60
	SUSAN	0.809	1.236	0.864	1.575	26.04
	Average	0.837	1.385	0.892	1.763	27.05
	BIK-BUS	0.600	0.222	0.629	0.274	6.91
	Curvature	0.605	0.254	0.626	0.253	1.04
	Harris3D	0.604	0.233	0.636	0.262	1.33
	ISS3D	0.606	0.224	0.635	0.241	1.36
	KLT	0.606	0.248	0.634	0.265	1.41
OUK-CVFH	Lowe	0.604	0.236	0.634	0.264	1.44
	Noble	0.602	0.225	0.635	0.271	1.39
	SIFT3D	0.593	0.159	0.626	0.218	2.73
	SUSAN	0.556	0.009	0.571	0.035	1.89
	Average	0.597	0.201	0.625	0.231	2.17
	BIK-BUS	0.614	0.393	0.639	0.470	8.01
	Curvature	0.618	0.407	0.636	0.460	2.06
	Harris3D	0.619	0.411	0.639	0.474	2.18
	ISS3D	0.623	0.427	0.645	0.495	2.28
DCF	KLT	0.625	0.432	0.646	0.503	2.28
FCL	Lowe	0.621	0.420	0.642	0.485	2.21
	Noble	0.621	0.419	0.647	0.508	2.23
	SIFT3D	0.619	0.412	0.640	0.479	3.58
	SUSAN	0.596	0.336	0.618	0.412	2.80
	Average	0.617	0.406	0.611	0.476	3.07

Descriptors	Koupointo	Cate	egory	Obj	ject	Time (s)
Descriptors	Reypoints	AUC	DEC	AUC	DEC	Time (s)
	BIK-BUS	0.848	1,488	0.893	1.832	4948.23
	Curvature	0.848	1.489	0.893	1.831	4816.54
	Harris3D	0.849	1.491	0.894	1.843	3722.78
	ISS3D	0.848	1.489	0.895	1.855	4367.38
	KLT	0.848	1.489	0.891	1.811	4202.21
PFH	Lowe	0.847	1.483	0.896	1.854	3626.09
	Noble	0.846	1.474	0.894	1.840	3651.77
	SIFT3D	0.843	1.458	0.890	1.801	3920.08
	SUSAN	0.828	1.363	0.866	1.625	6642.93
	Average	0.845	1.469	0.890	1.810	4433.11
	BIK-BUS	0.867	1.586	0.948	2.397	9567.81
	Curvature	0.859	1.535	0.938	2.267	9315.20
	Harris3D	0.859	1.533	0.941	2.303	7233.37
	ISS3D	0.866	1.585	0.948	2.394	8488.44
	KLT	0.859	1.536	0.941	2.302	8206.08
PFHRGB	Lowe	0.860	1.539	0.942	2.314	7047.42
	Noble	0.861	1.548	0.939	2.275	7116.42
	SIFT3D	0.861	1.546	0.946	2.373	7628.79
	SUSAN	0.845	1.445	0.934	2.205	12815.49
	Average	0.860	1.539	0.942	2.314	8602.11
	BIK-BUS	0.646	0.475	0.673	0.552	8.01
	Curvature	0.555	0.016	0.579	0.008	2.09
	Harris3D	0.561	0.020	0.580	0.008	1.98
	ISS3D	0.640	0.405	0.667	0.479	2.00
DDE	KLT	0.549	0.012	0.570	0.012	2.13
	Lowe	0.574	0.013	0.592	0.028	2.00
	Noble	0.576	0.021	0.592	0.007	1.92
	SIFT3D	0.641	0.434	0.666	0.510	3.51
	SUSAN	0.599	0.297	0.602	0.316	11.40
	Average	0.593	0.188	0.613	0.213	3.89

Description		Cate	gory	Obj	ect	Time (s)
Descriptors	Keypoints	AUC	DEC	AUC	DEC	Time (s)
	BIK-BUS	0 493	0.042	0 506	0.077	15 12
	Curvature	0.475	0.042	0.500	0.077	5 77
	Harris3D	0.515	0.040	0.520	0.050	5.35
		0.322	0.013	0.508	0.004	6.07
	KI T	0.400	0.004	0.500	0.024	5.72
PPFRGB		0.501	0.103	0.545	0.100	5.65
	Noblo	0.510	0.033	0.329	0.020	5.00
		0.510	0.100	0.400	0.000	0.09
		0.501	0.070	0.510	0.201	0.20
	SUSAN	0.537	0.003	0.545	0.051	17.37
	Average	0.507	0.048	0.519	0.076	8.27
	BIK-BUS	0.827	1.281	0.863	1.513	8.78
	Curvature	0.823	1.255	0.866	1.532	2.50
	Harris3D	0.817	1.224	0.858	1.490	2.58
	ISS3D	0.812	1.168	0.852	1.413	2.69
CUOT	KLT	0.820	1.235	0.855	1.448	2.77
SHUT	Lowe	0.818	1.229	0.855	1.462	2.66
	Noble	0.819	1.235	0.860	1.494	2.62
	SIFT3D	0.814	1.207	0.848	1.409	3.94
	SUSAN	0.749	0.892	0.790	1.075	3.06
	Average	0.811	1.192	0.850	1.426	3.51
	BIK-BUS	0.867	1.571	0.916	2.012	9.70
	Curvature	0.865	1.557	0.912	1.972	2.74
	Harris3D	0.858	1.519	0.906	1.918	2.72
	ISS3D	0.852	1.465	0.902	1.873	2.92
	KLT	0.861	1.542	0.908	1.935	2.95
SHUTCOLOK	Lowe	0.860	1.532	0.903	1.903	2.78
	Noble	0.859	1.530	0.907	1.930	2.80
	SIFT3D	0.839	1.394	0.896	1.792	4.18
	SUSAN	0.783	1.099	0.839	1.397	3.30
	Average	0.849	1.468	0.899	1.859	3.79

Descriptors	Keypoints	Category		Object		Time (a)
		AUC	DEC	AUC	DEC	Time (s)
SHOTLRF	BIK-BUS	0.789	1.096	0.822	1,265	7.45
	Curvature	0.790	1.062	0.814	1.188	1.54
	Harris3D	0.784	1.013	0.810	1.138	1.80
	ISS3D	0.788	1.003	0.817	1.139	1.86
	KLT	0.785	1.003	0.815	1.154	1.87
	Lowe	0.784	1.017	0.812	1.146	1.87
	Noble	0.785	1.021	0.811	1.142	1.88
	SIFT3D	0.770	0.924	0.805	1.086	3.20
	SUSAN	0.676	0.561	0.710	0.684	2.24
	Average	0.772	0.967	0.802	1.105	2.63
USC		0 720	0 (5 1	0 790	0.942	11041 92
	DIK-DUS	0.739	0.031	0.789	0.812	1041.82
	Curvature	0.730	0.631	0.780	0.778	10147.07
	HarrissD	0.728	0.599	0.778	0.743	8380.70
	1553D	0.727	0.630	0.777	0.790	9556.12
	KLI	0.731	0.609	0.784	0.774	91/3.02
	Lowe	0.729	0.604	0.781	0.765	/987.03
	Noble	0.727	0.597	0.777	0.740	7970.97
	SIFT3D	0.727	0.647	0.774	0.797	8725.92
	SUSAN	0.681	0.506	0.717	0.623	11458.16
	Average	0.725	0.597	0.774	0.758	9382.38
VFH	BIK-BUS	0.647	0.517	0.705	0.745	6.82
	Curvature	0.644	0.502	0.703	0.732	0.94
	Harris3D	0.638	0.483	0.680	0.638	1.27
	ISS3D	0.643	0.514	0.687	0.671	1.28
	KLT	0.644	0.507	0.691	0.680	1.32
	Lowe	0.638	0.481	0.687	0.670	1.37
	Noble	0.638	0.480	0.682	0.649	1.32
	SIFT3D	0.636	0.469	0.686	0.651	2.66
	SUSAN	0.584	0.295	0.615	0.404	1.70
	Average	0.635	0.472	0.835	0.649	2.37

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Kovpoint	Category		Obj	Total		
кеуропп	AUC	DEC	AUC	DEC	TULAL	
BIK-BUS	7	9	7	9	32	
Curvature	2	2	2	1	7	
Harris3D	1	1	1	0	3	
ISS3D	2	0	3	2	7	
KLT	2	1	1	0	4	
Lowe	0	0	1	0	1	
Noble	0	1	1	1	3	
SIFT3D	0	0	0	1	1	
SUSAN	2	1	2	1	6	

Table 6.4: Counting the number of times a keypoint detector has the best result in table 6.3. In case of a tie both methods score.

If a threshold is considered for the AUC t_{AUC} and another for the DEC t_{DEC} only two original descriptors (PFH and SHOT) and four of its variants (FPFH, PFHRGB, SHOTCOLOR and SHOTLRF). In the case SUSAN/SHOT both thresholds fail and for SHOTLRF only the threshold t_{DEC} is satisfied in seven keypoint detectors. In these descriptors, our detector only in a single case does not have the best results in both measures, and this in the case of PFH where only has a difference of 0.1%. In the other four descriptors, the recognition accuracy varies between 2.2% and 8.4%.

In terms of object recognition, the best pair is BIK-BUS/PFHRGB, but only beats the second best combination, ISS3D/PFHRGB, because it presents a better DEC. For SHOT and SHOTCOLOR descriptors if we compare our keypoint detector with the ISS3D we obtain improvements for both of 1.5% in the case of category recognition, and 1.1% and 1.4% in object recognition, respectively. The only point against our keypoint detector is relation to the processing time, since it is approximately 6 times slower than ISS3D. The processing time can be reduce by a parallel implementation or by an implementation in GPU. The architecture of the BIK-BUS, shown in figure 6.1, shows that the parallel implementation would be a good strategy to solve this problem.

6.4 Summary

In this chapter, a novel 3D keypoint detector biologically motivated by the behavior and the neuronal architecture of the early primate visual system was presented. We also made a comparative evaluation of several keypoint detectors plus descriptors on public available data with real 3D objects. This new method and the presented results were published in [28, 29].

The BIK-BUS is a keypoint detector to determine visual attention, which are also known as saliency maps. The saliency maps are determined by sets of features in a bottom-up and datadriven manner. The fusion of these sets produced the saliency map and the focus of attention is sequentially directed to the most salient points in this map, representing a keypoint location.

In the evaluation, the 3D keypoint detectors and the 3D descriptors available in the PCL library were used. The main conclusions of this chapter are: 1) a descriptor that uses color information should be used instead of a similar one that uses only shape information; 2) the descriptor should be matched to the desired task, since there are differences in terms of recognition performance, size and time requirements; 3) in terms of keypoint detectors, to obtain an



Figure 6.4: ROCs for the category recognition experiments (best viewed in color).



Figure 6.5: ROCs for the object recognition experiments (best viewed in color).

accurate recognition system is recommended the use of the BIK-BUS, since its performance was better in 32 tests, in a total of 60 tests. When the second best detector only obtained the best performance 8 times (see table 6.4); 4) for a real-time system, the ISS3D or Curvature detectors are good choices, since they have a performance that is only surpassed by BIK-BUS and are faster; 5) in terms of descriptors, if the focus is on accuracy the use of PFHRGB is recommended and for real-time a good choice is the SHOTCOLOR because it presents a good balance between recognition performance and time complexity.

Chapter 7

A 3D Keypoint Application for Tracking

In this chapter, a robust detection and tracking method for 3D objects by using keypoint information in a particle filter is proposed. Our method consists of three distinct steps: Segmentation, Tracking Initialization and Tracking. The segmentation is made in order to remove all the background information, in order to reduce the number of points for further processing. In the initialization, a keypoint detector with biological inspiration is used. The information of the followed object is given by the extracted keypoints. The particle filter does the tracking of the keypoints, so with that you can predict where the keypoints will be in the next frame. In a recognition system, one of the problems is the computational cost of keypoint detectors. This approach aims to solve this problem.

The experiments with the proposed method are done indoors in an office/home environment, where personal robots are expected to operate. The Tracking Error evaluate the stability of the general tracking method. We also evaluate quantitatively this method using a "Tracking Error". Our evaluation is done by the computation of the keypoint and particle centroid.

7.1 Particle Filter with Bio-Inspired Keypoints Tracking

This section will focus on the Particle Filter with Bio-Inspired Keypoints Tracking (PFBIK-Tracking) block presented in the figure 7.1. This method is composed by two main steps: Segmentation and Tracking, which will be described in detail below.

The Recognition block is presented in this work only in a illustrative way and it will not be discussed in this chapter. But in [22, 23, 27, 29] and the last two chapters (5 and 6) give us a good perspective on how to solve the issue of object recognition. In [22], the focus is on the descriptors available in PCL (Feature Extraction step). We briefly explain how they work and made a comparative evaluation on publicly available data.

7.1.1 Segmentation

The segmentation starts with the Pass Through Filter (PTF). This filter removes depth regions that are not contained on the desired working distances $[d_{min}, d_{max}]$, where d_{min} is the minimum distance at which the system should work and d_{max} the maximum distance. Depth regions that are not included between these distances are considered background and are discarded by the tracking system. By removing these regions (shown in figure 7.2(b)), which do not have interesting information for the object tracking system, a considerable reduction in the processing time is obtained.

The second step of the segmentation is the Planar Segmentation (PS), which is based on the Random Sample Consensus (RANSAC) algorithm [125]. It is an iterative method to estimate parameters of a mathematical model from a set of observed data which contains outliers. Basically, the data consists of "inliers" and "outliers". The distribution of the inlier's data can be explained by some set of model parameters, but may be subject to noise and outliers, which

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Figure 7.1: Setup of the recognition system. The diagram presents a complete object recognition system in order to understand better how the communication between the different stages is processed.





Figure 7.2: Representation of the segmentation steps. Figure (a) represents a cloud captured by the kinect camera. Figure (b) is the output of the pass through filter with $d_{min} = 0.0$ m and $d_{max} = 1.6$ m, and in (c) the result of the removal of planar regions. Figure (d) are the clusters of the objects, wherein each object is represented by a different color.

are data that do not fit the model. The outliers can come from extreme values of the noise or from erroneous measurements or incorrect hypotheses about the interpretation of data.

Let w be the probability of choosing an inlier each time a single point is selected, that is, $w = \frac{\#inliers}{\#points}$. Using n points, selected independently, for estimating the model, w^n is the probability that all n points are inlier's and $1 - w^n$ is the probability that at least one of the npoints is an outlier. That probability to the power of k (number of iterations) is the probability that the algorithm never selects a set of n points which all are inlier's and this must be the same as 1 - p. Where p is the probability that the RANSAC algorithm in some iteration selects only inlier's from the input cloud set when it chooses the n points from which the model parameters are estimated. Consequently,

$$1 - p = (1 - w^n)^k$$
 (7.1)

which, after taking the logarithm of both sides, leads to

$$k = \frac{\log(1-p)}{\log(1-w^n)}.$$
 (7.2)

Given the planar region estimated by RANSAC algorithm, it is possible to remove the planar regions from the cloud, keeping only the remaining objects (shown in figure 7.2(c)).

7.1.2 Tracking Initialization

In the first frame captured, to initialize the tracking, the third step of segmentation is performed, the Cluster Extraction (CE). Clustering is the process of examining a collection of "points", and grouping the points into "clusters" according to some distance measure. That is, the goal is that points in the same cluster have a small distance from one another, while points in different clusters are at a large distance from one another. This step will return a list of the clusters (shown in figure 7.2(d)), where each one contains the information of an object present in the cloud scene. In this work, Euclidean Clustering method is used. As the name implies, the distance between two points p_1, p_2 is given by the Euclidean distance:

$$D(p_1, p_2) = \sqrt{(p_{1_x} - p_{2_x})^2 + (p_{1_y} - p_{2_y})^2 + (p_{1_z} - p_{2_z})^2}.$$
 (7.3)

This PCL implementation is explained in [345].

As mentioned earlier, in [26], we presented an evaluation of the keypoint detectors available on PCL. The SIFT keypoint detector was proposed in [9]. The SIFT features are vectors that represent local cloud measurements. The 3D implementation of SIFT3D keypoint detector was presented in [102]. It uses a 3D version of the Hessian to select such interest points.

The performance of human vision is obviously far superior to that of current computer vision systems, so there is potentially much to be gained by emulating biological processes. Fortunately, there have been dramatic improvements within the past few years in understanding how object recognition is accomplished in animals and humans [99]. Some features found in IT cortex are composed by neurons that respond to various moderately complex object features, and those that cluster in a columnar region that runs perpendicular to the cortical surface respond to similar features [346]. These neurons maintain highly specific responses to shape features that appear anywhere within a large portion of the visual field and over a several octave range of scales [347]. The complexity of many of these features appears to be roughly the same

as for the SIFT. The DoG clouds are also similar to the "Place cells", which are pyramidal cells in the hippo-campus which exhibit strongly increased firing in specific spatial locations [348]. The feature responses have been shown to depend on previous visual learning from exposure to specific objects containing these features [349]. These features appear to be derived in the brain by a highly computation-intensive parallel process, which is quite different from the staged filtering given by this method. A retinotopic organization, parallel processing, feedforward, feedback and lateral connection are a complex composition of the human visual system [350]. However, the results are much the same: an image is transformed into a large set of local features that each match a small fraction of potential objects yet are largely invariant to common viewing transformations.

7.1.3 Tracking

The Tracking block presented in figure 7.1 is the Particle Filter. For this, an adaptive particle filter presented in [126, 127] is used. They presented a statistical approach to adapting the sample set size of particle filters on-the-fly. The number of the particles changes adaptively based on KL distance sampling [128], where they bind the error introduced by sample-based representation of the particle filter. The samples are generated iteratively until their number is large enough to ensure that the KL distance between the maximum likelihood estimate and the underlying posterior does not exceed a pre-specified bound. This method will choose different numbers of samples depending on the density of the 3D point cloud. If selects a small number of samples, the density is focused in a small subspace and it selects a larger number of samples, the samples have to cover most of the state space.

7.2 Results

To evaluate the performance of the method, the Euclidean distance (equation 7.3) between the centroid of the keypoints and result of the method will be used, which is the "Tracking Error". The purpose of performing this comparison is to evaluate whether a system is able to track the keypoints of an object. The tracking is done in order to remove the necessity of applying a keypoint detector in all frames. In a real-time system is not feasible to apply a keypoint detector in each frame, due to the computational time spent on their calculation.

The centroid of a finite set of p points $p_1, p_2, ..., p_k$ in \mathbb{R}^n is given by

$$C = \frac{\sum_{i=1}^{k} p_i}{k} \quad (7.4)$$

This point minimizes the sum of squared Euclidean distances between itself and each point in the set.

In order to properly evaluate the performance of the method, it will be compared with the sample-based method *OpenniTracker* available in PCL 1.7 (from the trunk). The segmentation step is applied in this tracker, where the output of this step is shown in figure 7.3. Thus, exactly the same data is given as input to both methods.

The difference between the two methods is the initialization of the particle filter. Whereas it is initialized with the results of the keypoint detector, the OpenniTracker only makes a sub-sampling. This is a very important difference in the object recognition frameworks, because the


(a) Bowl.



(b) Box_1.



(c) Box_2



(d) Mug.



Figure 7.3: Segmented point cloud sequences of the dataset. These point clouds are the inputs of the presented tracking methods, and these have already been segmented.

sub-sampling only reduces the number of points in a linear manner, while the keypoint detector is reducing the number of points based on the object characteristics.

The results presented in table 7.1, 7.2 and 7.3 are obtained using a dataset collected by us (shown in figure 7.3). This dataset contains 10 different moving objects in a total of 3300 point clouds.

In table 7.1, we can observe that method performed the tracking with a significantly lower number of points. Since the goal is to make the recognition of each object in the scene,

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Table 7.1: Mean and standard deviation number of keypoints and particles resulting from the tracker. In the OpenniTracker case, the column keypoints represents the sub-sampled cloud.

	Number of Points		
	Keypoints Particles		
PFBIK-Tracking	116.973 ± 76.251	102.107 ± 92.102	
OpenniTracker	2502.536 ± 1325.807	2132.124 ± 1987.516	

Table 7.2: Euclidean distance between the output of the tracker and the expected result.

	Distance between Centroids			
	X axis	Y axis	Z axis	All axis
PFBIK-Tracking	0.036 ± 0.032	$\textbf{0.013} \pm \textbf{0.012}$	$\textbf{0.019} \pm \textbf{0.019}$	0.045 ± 0.036
OpenniTracker	0.038 ± 0.023	0.013 ± 0.011	$\textbf{0.027} \pm \textbf{0.015}$	0.052 ± 0.022

Table 7.3: Mean and standard deviation of the Computational time (in seconds) of the evaluated methods. Here, the time of the segmentation step is discarded, because it is the same in both methods.

	Tracking Initialization	Tracking
PFBIK-Tracking OpenniTracker	$\begin{array}{c} 0.203 \pm 0.162 \\ 0.173 \pm 0.187 \end{array}$	$\begin{array}{c} 0.081 \pm 0.057 \\ 0.186 \pm 0.170 \end{array}$

the method has a stable performance with fewer points. With this number of points it is possible to think of making recognition in real time. On the other hand, with the number of points shown by the OpenniTracker this is very difficult to archive.

In table 7.2, the distance between the cloud of keypoints ('what we expect') are presented and the resulting cloud of points produced by the tracker ('what we estimate'). As already mentioned, the Euclidean distance is calculated based on the centroid of what was estimated and what we really are looking. In this table, we can see that even with a large decline in the number of points (around 50 times), this method has better performance than OpenniTracker.

In table 7.3, the mean processing time of the two evaluated methods is presented. These times were obtained on a computer with Intel @Core M2 Quad Processor Q9300 2.5GHz with 4 GB of RAM memory. Our method takes longer to initialize the tracking, but then the tracking system becomes 2.3 times faster than the other method presented. The initialization time is not a problem since this is only done once. The initialization is slower due to the fact that the keypoints are extracted, instead of the sub-sampling process used by the other method. In summary, the presented method obtains better results in terms of tracking (increased robustness to occlusion), while using less points and resulting in an improvement in terms of processing speed.

7.3 Summary

In this work, a system to perform the tracking of keypoints is presented and published in [30]. The goal is to remove the necessity of applying a keypoint detector in all frames that are analyzed. When this kind of approach is performed, the systems spend a lot of computational time and they can not operate in real time. We intend to make the tracking of keypoints because the main goal is to extract the descriptors of a particular object in the scene in order to perform the recognition. In order to do this, several segmentation steps are presented, so that the system can remove all the background and objects become isolated. When the objects are segmented, a clustering method and the SIFT3D keypoint detector are applied, which is used to initialize the particle filter. We use the SIFT3D keypoint detector because it has similar features to those in IT [99]. Once it is initialized with the intended object, it is only necessary to give as input the output of the segmentation.

With this approach, better results were obtained than using the OpenniTracker: a faster and more robust method. For future work, we will intend to do the tracking of multiple objects simultaneously and use BIK-BUS keypoint detector instead of SIFT3D.

Chapter 8

Conclusions and Further Work

This chapter presents the main conclusions that result from the research work described in this thesis. Furthermore, it discusses a few research topics related with the work developed that may be addressed in the future.

8.1 Main Conclusions

This thesis focused systems based on the human visual attention and HVS. The developed methods have characteristics that were obtained from studies in the field of neuroscience and psychology. To understand those characteristics, an overview of the HVS (chapter 2) and a review of computational methods that attempt to model visual attention (chapter 3) was provided. The focus was mostly on bottom-up attention, although some top-down models were also discussed in [129--133].

Visual attention is a highly interdisciplinary field with researchers coming from different backgrounds. For psychologists, research conducted in human behavior is performed by isolating certain specific tasks, so that the internal processes of the brain, often resulting in psychophysical theories or models [134]. Neurobiologists observe the brain's response to certain stimuli [135], using techniques such as fMRI, having therefore a direct view of the brain areas that are active under certain conditions [45, 136, 137]. Finally, engineers use the discoveries made in those areas attempting to reproduce them in computational models, so that the processing time in some applications can be reduced [42--44]. In recent years, those different areas have profited considerably from each other, psychologists use research conducted by neurobiologists, to improve their attention models, while neurobiologists consider psychological experiments to interpret their data [134]. Additionally, psychologists began to implement computer models or use computer models previously developed, to verify that they have a similar behavior to that of human perception. Thus, psychologists tend to improve the understanding of the mechanisms and help the development of better computational models.

Computational attention has gained a significant popularity in the last decade. One of the contributors to the increase in popularity was the improvement in computational resources. Another contribution was the performance gains obtained from the inclusion of visual attention (or saliency detection) modules in object recognition systems [131, 138, 139].

Most of the research presented in this thesis, was focused on the bottom-up component of visual attention. While previous efforts are appreciated, the field of visual attention still lacks computational principles for task-driven attention. A promising direction for future research is the development of models that take into account time varying task demands, especially in interactive, complex, and dynamic environments. In addition, there is not yet a principled computational understanding of the visual attention. The solution is beyond the scope of a single area. To obtain a solution it is necessary to have the cooperation of the several areas, from the machine learning community, computer vision and the biological fields as well as neurology and psychology.

Table 3.2 shows some areas where the saliency maps were applied, but with no references to whether these can be used to extract directly keypoint locations, the most nearly being the one of Rodrigues and du Buf [104]. The work of Ardizzone et al [140] compared if a particular method extracts the keypoints in salient regions. With this, an analysis was performed on the top popular keypoint detectors and presented in chapter 4, especially the ones using RGB-D information. Furthermore, a description of the 3D descriptors and an evaluation of 3D keypoint detectors, on public available data with real 3D objects, were made. The experimental comparison proposed in this work has outlined aspects of state-of-art methods for 3D keypoint detectors. This analysis allowed to evaluate the best performance in terms of multiple transformations (rotation, scaling and translation).

The novelties of this work, when compared with the work of Schmid et al. [17] and Salti et al. [19] are: we are using a real database instead of an artificial, the large number of point clouds and different keypoint detectors. The benefit of using a real database is that our objects have "occlusion", this is because some materials do not reflect the infrared or from the segmentation method. This does not happen when dealing with artificial objects, causing the keypoint methods to display better results, but our experiments reflect what can happen in real life, such as, with robot vision. Overall, SIFT3D and ISS3D yielded the best scores in terms of repeatability and ISS3D demonstrated to be the more invariant.

Another part of this research work was described in chapter 5 and encompassed the study of a 2D keypoint detector in a recognition framework. The proposal of a novel keypoint detection method is also made, called BMMSKD, that uses the retinal color information. The retinal color information was applied as an extension to the BIMP method so that color information could be used. The recognition evaluation of the proposed approach was made on public available data with real 3D objects. For this evaluation, the keypoint detectors were developed using the OpenCV library and the 2D keypoint locations were projected to the 3D space to use available 3D descriptors on the PCL library. It was possible to verify that the keypoint locations can either help or degrade the recognition process and descriptors that uses color information should be used instead of a similar one with shape information alone. Differences are big in terms of recognition performance, size and time requirements, and thus the descriptor should be matched to the desired task. If we want to recognize the category of an object or a real-time system, the recommendation is to use the SHOTCOLOR method, since it presents a recognition rate of 7% below of the PFHRGB but with a much lower computational cost. On the other hand, to perform object recognition, the recommendation is to use PFHRGB, as it presents a recognition rate 12.9% higher than SHOTCOLOR.

A novel 3D keypoint detector biologically motivated by the behavior and the neuronal architecture of the early primate visual system was presented in chapter 6. Similarly to chapter 5, a comparative evaluation was made where several keypoint detectors and descriptors were compared on public available data with real 3D objects. BIK-BUS is a keypoint detector to determine visual attention, which is also known as saliency maps. The saliency maps are determined by sets of features in a bottom-up and data-driven manner. The saliency map is then produced by the fusion of those sets, being the focus os attention sequentially directed to the most salient point of the map, that represent a keypoint location. In the evaluation, the 3D keypoint detectors and the 3D descriptors available in the PCL library were used. With a similar average number of keypoints, the proposed 3D keypoint detector outperforms all the 3D keypoint detectors evaluated, achiving the best result in 32 of the evaluated metrics in the category and object recognition experiments, while the second best detector obtained only the best result in 8 of these metrics (see table 6.4), in a total of 60 tests. The unique drawback we identified is

the computational time, since BIK-BUS is slower than the other detectors. For a real-time system, the ISS3D or Curvature detectors are more advisable choices, since theyare faster and their performance is only surpassed by BIK-BUS. Finally, in terms of descriptors the recommendation goes to either PFHRGB or SHOTCOLOR. PFHRGB should be used when one wants an accurate recognition system, while for real-time the most advisable choice is the SHOTCOLOR, since it presents a good balance between recognition performance and time complexity.

In this research work, an application for 3D keypoint detectors was also presented, called PFBIK-Tracking, consisting on a system to perform the tracking of keypoints. The goal was to remove the necessity of applying a keypoint detector in all the frames we wanted to analyze. This is caused by the keypoint detectors being applied to all the frames, and therefore the system would not be able to operate in real time. To solve this, a keypoint tracker was developed to simulate the application of keypoint detectors to all the frames, since the main propose would be to extract the descriptors of a particular object in the scene to perform the recognition. For that propose, several segmentation steps were presented, so that we can remove all the background and objects become isolated. Further to object segmentation, a clustering method and the SIFT3D keypoint detector was used because it has similar features to those in IT [99]. Once it was initialized with the intended object, it only needs to be fed with the output of the segmentation. This approach obtained better results than when the OpenniTracker was used, and thereby a faster and more robust method was presented.

The main objectives of this thesis were accomplished by the presentation of the three methods. Together, the proposed methods allow the incorporation of characteristics with biological inspiration in recognition systems. Here, the experiments were done only in an object recognition system, but it can be applied to other types, such as biometrics signal (e.g. 3D face).

8.2 Future Work

Future work that could be developed should fall into three main focuses. The first line of further research would be to reduce the computational cost of the two keypoints detectors presented. Fot that purpose, we can consider a code parallelization or an implementation on the General-Purpose Computation on Graphics Processing Unit (GPGPU) to reduce the computational time of BMMSKD and BIK-BUS. The architecture of the methods makes the code parallelization possible, as shown in figures 5.1 and 6.1.

Secondly, it would be a good idea to seek further insights on why one keypoint detector or a combination of a descriptor type and keypoint detector works better than others for a specific test. This can be accomplished by selecting a small number of top keypoint detectors and descriptors (based on the results presented in this research work) and analyze which are the best pairs on the recognition of a particular type of category or object. In this work, an analysis to cover the whole dataset and not focus on specific cases was made. A more complex analysis was not performed for two reasons: 1) the dataset used in this work is very large, it consists of 300 objects and these are divided into 51 categories; 2) were also evaluate 135 pairs of keypoint detector/descriptor and this analysis would render unfeasible using all these methods.

Finally, future work will focus on the proposed keypoint tracking system, where several possibilities are still open and can be exploited. The first possibility is the replacement of the keypoint detector to use the BIK-BUS instead of SIFT3D. The reasoning behind this proposal is

that the BIK-BUS showed better results than SIFT3D in the object recognition framework.

Another point to explore that could improve this research work would be the dataset: adding more objects and the way in which they move in the scene. This new dataset has already been captured, containing 46 different objects, organized into 24 categories and with long periods of occlusion (such as out of the camera range and behind boxes). The acquisition of objects was made using a Kinect camera placed in several locations of a room, with the objects over a remote controlled car, so that they could move around the room. To use this dataset is would be necessary to segment the objects that are moving in the scene, so that several experiments and comparisons could be performed. The first experiments should take into account the evaluation made in chapters 5 and 6. This will allow to consolidate the results presented in those chapters. With the segmented objects, it is also possible to propose a 3D CLEAR MOT Metric [141] to the 3D tracking approaches. This measure is only available for 2D methods and does not consider the depth of the object. The difference to the measure proposed in chapter 7 is that it also makes the evaluation of the overlap between the position of the real object and the approximation obtained by the tracking method. Finally, category and object recognition using particle information and descriptors can be compared to the recognition process of using keypoint detectors and descriptors.

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