

UNIVERSITY OF LISBON

FACULTY OF FARMACY



Developments in Tumor Targeting and Internalizing Peptides

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Integrated Master in Pharmaceutical Sciences

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presented to Faculty of Pharmacy of the University of Lisbon**

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Resumo

A grande limitação do uso de fármacos citotóxicos em terapias antitumorais resume-se à falta de capacidade destas moléculas em distinguir células tumorais e células saudáveis. Os Tumor Targeting Peptides (TTPs), desde a sua descoberta há 30 anos atrás, têm-se tornado numa ferramenta útil para o tratamento e diagnóstico do cancro, uma vez que reconhecem alvos moleculares tumorais específicos, tal como os anticorpos (Abs), mas sem as suas desvantagens estruturais. Assim, a sua utilização no desenvolvimento de conjugados terapêuticos peptídicos (PDCs) é bastante promissora, embora ainda nenhum destes sistemas de entrega de fármacos esteja aprovado para uso clínico.

O objetivo deste trabalho consistiu na revisão de literatura relacionada com abordagens terapêuticas alvo-dirigidas e na organização e resumo dos TTPs que contribuíram de uma forma mais valiosa para os avanços no desenvolvimento de conjugados terapêuticos.

Neste trabalho, foram apresentados 9 péptidos com capacidade de reconhecimento alvo-específico, representativos de cada classe de TTPs e promissores quanto a uma futura aplicação clínica como parte integrante de conjugados terapêuticos - RGD, NGR, F3, Octreotide, LyP-1, Bombesin, Angiopep 2, CREKA e M2pep. Desta forma, foram focados para cada um dos TTPs, pontos-chave relativos a características estruturais e funcionais, alvos celulares específicos, estudos de relação estrutura-atividade (SAR), capacidade de internalização e aplicações no desenvolvimento de conjugados terapêuticos.

Palavras-Chave: Reconhecimento alvo-específico, tumor targeting peptides, RGD, NGR, conjugados terapêuticos.

Abstract

The major inconvenient of the use of cytotoxic drugs in anti-tumor therapies is their poor ability to distinguish tumor cells from the normal cells. Since their discovery about 30 years ago, tumor targeting peptides (TTPs) have become an useful tool for the treatment and diagnosis of cancer, once they are able to recognize specific tumor targets, such as antibodies (Abs) but without their structural disadvantages. Thus, their use in the development of peptide-drug conjugates (PDCs) is promising, although none of these drug delivery systems (DDS) are approved for clinical use yet.

The aim of this work was to review the literature about tumor targeting approaches and to organize and summarize the TTPs that contributed in a significant way to the advances of this field, including the development of therapeutic conjugates.

In this work, nine peptides with targeting ability, representative of each class of TTPs and promising for a future clinical application as an integral part of drug conjugates were presented - RGD, NGR, F3, Octreotide, LyP-1, Bombesin, Angiopep 2, CREKA and M2pep. This way, structural and functional features, addresses, structure-activity relationship (SAR) studies, internalization capacity and applications in the development of therapeutic conjugates were focused as key-points for each targeting peptide.

Key-words: Targeting, tumor targeting peptides, RGD, NGR, peptide-drug conjugates.

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Abbreviations and Symbols

α - Alfa

β – Beta

γ – Gamma

Abs – Antibodies

ADCs – Antibody-drug conjugates

AMPs – Antimicrobial peptides

APN – Aminopeptidase N

BB1 – Bombesin receptor type 1

BB2 – Bombesin receptor type 2

BB3 – Bombesin receptor type 3

BB4 – Bombesin receptor type 4

BBB – Blood brain barrier

BBRs – Bombesin receptors

CDRs – Complementarity determining regions

CendR – C-end Rule

CNS – Central nervous system

CPPs – Cell penetrating peptides

CPT - Camptothecin

cRGD – Cyclic RGD

DDS – Drug delivery systems

DOTA – 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid

DOX – Doxorubicin

DTPA – Diethylenetriaminepentaacetic acid

ECM – Extracellular matrix

EPR – Enhanced permeability and retention

GnRH – Gonadotropin releasing hormone

GRPR – Gastrin realizing peptide receptor

HMG₂N – Human high mobility group protein 2

IFN- γ – Interferon- γ

Ig G – Immunoglobulin G
IL-12 – Interleukin-12
iNGR – Internalizing NGR
iRGD – Internalizing RGD
isoDGR - Isoaspartate-glycine-arginine
KLA – KLAKLAKKLAKLAK sequence
LDL - Low density lipoprotein
LHRH – Luteinizing hormone-releasing hormone
LRP-1 – Lipoprotein receptor-related protein 1
mRNA – Messenger ribonucleic acid
MTX – Methotrexate
NCL – Nucleolin
NMBR - Neuromedin B receptor
NRP-1 – Neuropilin-1
OBOC – One-bead one-compound
PDCs – Peptide-drug conjugates
PEG – Poly(ethylene glycol)
PET – Positron emission tomography
PIMT – Protein-L-isoAsp-O-methyltransferase
PTX – Paclitaxel
rRNA – Ribosomal ribonucleic acid
SAR – Structure-activity relationship
SMIs – Small molecules inhibitors
SST – Somatostatin
SSTR – Somatostatin receptors
TAMs - Tumor-associated macrophages
TNF α – Tumor necrosis factor α
TTPs – Tumor targeting peptides
VEGF – Vascular endothelial growth factor

1. Introduction

Cancer is characterized by the uncontrolled growth of abnormal cells due to mutations of genes involved in cells growth, proliferation or survival, creating a complex chain of events and modifying basic biological operations of cells, such as the ability to respond to growth signals, invade tissues and regulate cell death programs. Genetic alterations may be inherited or accrued after birth, causing activation of oncogenes or inhibition or deletion of tumor suppressor genes.¹⁻⁴ Cancer cells can evolve to benign or malignant tumors. Whereas benign tumors are encapsulated and do not invade surrounding tissue, malignant tumors have less well differentiated cells, grow more rapidly and invade and destroy adjacent normal tissue, being able to generate secondary tumors (metastases) in distant organs in the body through blood vessels and lymphatic channels.

Tumor cells are capable to grow even in the face of starvation of the host, causing morbidity or his death. The most common effects on patients are cachexia, hemorrhage and infection. In most cases, the origin of the cancer is not clear, but it is known that external (such as cigarette smoking and radiation) and internal (like immune system defects, genetic predisposition) factors can contribute to cancer development.¹ Current cancer treatments are surgery, radiation and chemotherapy.^{2,5} Whereas surgery allows the removal of solid tumors (in other words, tumors confined to the anatomical area of origin or primary tumors), radiation therapy is based on the biological effect of ionizing radiations in tumor localization. Moreover, chemotherapy, the basis for the treatment of disseminated tumor, uses chemicals with cytotoxic properties.^{3,6-8} Although classical chemotherapy assumes that cytotoxic drugs target the faster proliferating tumor cells, they still interfere with normal cells, inducing systemic toxicity and causing severe secondary effects like nausea, vomiting, hair loss, damages to liver, bone marrow and kidney.⁹

Over the past two decades, it was verified a continuous decline in number of deaths by cancer in consequence of significant advances in the development of anticancer drugs.^{10,11} Still, cancer remains one of the leading cause of death worldwide, with metastases being the major complication in patients with cancer.^{1,10-13} Besides, generalized anatomic imaging techniques has lack of sensitivity to provide early diagnosis for cancer.¹²

1.1. Tumor Microenvironment

In primary tumors, cancer cells are surrounded by a complex microenvironment with unique physicochemical properties, since the fast tumor growth demands a higher consumption of energy and oxygen, leading to secondary acidic metabolites accumulation, hypoxia and hyperthermia (Figure 1). Here, the most abundant cells type is cancer-associated fibroblasts that contribute to extracellular matrix remodeling and cellular growth. Moreover, the predominant inflammatory cells, called tumor-associated macrophages (TAMs), can differentiate into M2-like macrophages, endowed with immunosuppressive features. Other types of cells are encountered in tumor microenvironment, including endothelial cells and their precursors, granulocytes, lymphocytes, natural killer cells and antigen-presenting cells, like dendritic cells. Interactions between tumor cells and their surrounding cells stimulates tumor development by production of enzymes, cytokines, chemokines and growth factors, angiogenesis occurring, immune escape and extracellular matrix disarrangement.^{2,14,15}

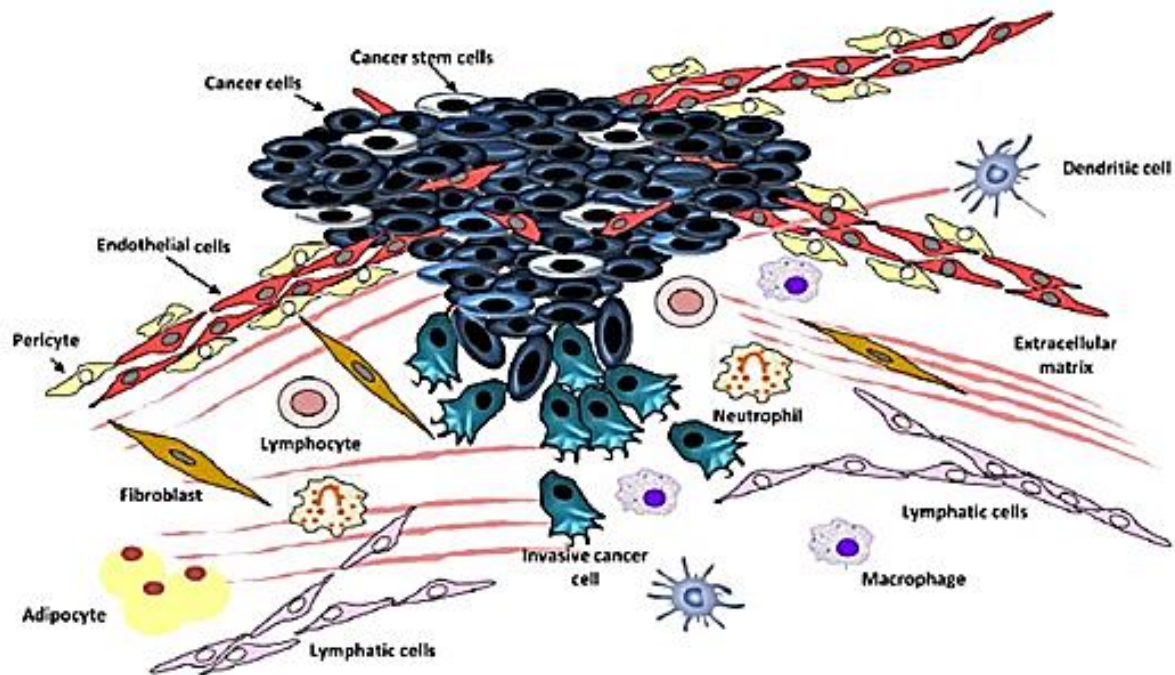


Figure 1: Tumor microenvironment constitutes. Adapted.¹⁴

1.1.1. Plasma Membrane

The plasma membrane has a crucial role in cell's behavior, namely in communication with other cells, cell movement, migration and adherence to other cells or structures, access to nutrients in the microenvironment and recognition by the body's immune system. On the plasma membrane of malignant cells, a number of

biochemical changes are verified, mainly the appearance of new surface antigens, proteoglycans, glycolipids and mucins, and alteration of cell to cell or cell to extracellular matrix communication. These changes induce loss of density-dependent inhibition of growth, decrease of adhesiveness and loss of anchorage dependence.^{1,2}

In cancer cells membrane, there is an increased negative charge due to loss of symmetry between zwitterionic phospholipids and the consequent exposure of anionic phosphatidylserine in the external surface of the plasma membrane. On the other hand, the presence of linked sialic acid and glycolipids or glycoproteins like mucins, proteoglycans, heparin sulfate and chondroitin sulfate also contributes to increase the negative charge of tumor cell membranes.¹

1.1.2. Tumor Vasculature

Angiogenesis consists in the formation of new blood vessels from pre-existing ones, that normally occurs in inflammations or tissues regeneration processes, such as tissue growth, wound healing, menstrual cycle and placental implantation.^{2,16} In solid tumors with 1-2 mm or more in diameter, pathological angiogenesis occurs, not only to increase supply nutrients and oxygen, but also to carry tumor cells to adjacent and/or distant organs.^{1,11-13,16} In this case, angiogenesis is triggered by hypoxia that activates endothelial cells through cell surface and secreted proteins, mostly integrins and matrix metalloproteinases, leading to the formation of malformed and dysfunctional new blood vessels.¹⁵

Tumor blood vessels exhibit structural and morphological differences from the normal vascular system.¹⁴ The vasculature in the majority of healthy tissues has a pore size of 2 nm and more specifically, post-capillary venules have 6 nm of pore, while in the tortuous tumor blood vessels, pores vary in size from 100 to 780 nm.¹⁷ In addition, solid tumors have poor blood flow through their vessels, but the presence of the enhanced permeability and retention (EPR) effect (an alteration in fluid dynamics due to the rapid growing and the abnormal tumor neo-vasculature) makes the tumor vasculature leaky and porous, enabling cellular metabolites accumulation at higher concentration than in normal tissues.^{2,14} Moreover, tumor endothelial cells express a different set of molecules on their surface, that can be angiogenesis-related (like the vascular endothelial growth factor (VEGF), heparin sulfate and nucleolin) or tumor type-specific.^{12,14,16,18}

Up to now, very little is known about tumor lymphatic vasculature. In normal tissues, lymphatic vessels are responsible for interstitial fluid and macromolecules

transportation from tissues to the bloodstream, in a unidirectional way. Thus, like in tumor blood vessels, tumor lymphatic vessels may connect to healthy vasculature and allow metastases spread.¹⁸

1.2. Tumor Targeting Therapy

The major inconvenient of cytotoxic drugs are their poor ability to distinguish cancer cells from the normal ones.^{2,11} Particularities of tumor cells, as well of tumor microenvironment (increased negative charge of cell membranes, lack of lymphatic drainage from the tumor and EPR effect) provides the delivery and uptake of antitumor drugs.^{2,17} In this case, the selectivity to tumor cells is based on a passive targeting mechanism, since there is a selective extravasation and accumulation of molecules in tumor tissues.¹⁷ However, it is common for antitumor drug formulations to have poor efficacy due to fail of tumor specificity, insufficient drug accumulation inside the tumor microenvironment and drug resistance (in other words, inactivation of the drug *in vivo* or modification of drug targets or drug efflux, through genetic and epigenetic modifications of tumor cells) that leads to unwanted side effects, therapeutic failure or cancer recurrence.^{2,3}

Tumor cells and tumor-associated tissues express different or overexpress surface antigens or receptors (tissue specific markers also known as vascular bed-specific zip codes or addresses) compared to normal tissue, which makes them into molecular targets to deliver antitumor molecules through an active targeting mechanism (manly known as tumor targeting approach), based on conjugation of addresses with their respective high affinity ligands (or tumor targeting molecules).^{1,2,19} Surface receptors like integrins, vascular epidermal growth factor (VEGF), folate, transferrin, luteinizing hormone-releasing hormone (LHRH), and somatostatin (SST) have shown high potential in tumor targeting therapy.¹⁴

1.2.1. Antibodies

Antibodies (Abs) are the natural targeting proteins of the organism. Consequently, Abs or their fragments are the most common targeting molecules employed in the delivery of antitumor drugs and imaging agents, since they provide antigen-specific binding affinity and can stimulate the immune system to fight or inhibit tumor growth.^{4,14,20} At clinical practice, Abs are utilized as an integral part of antibody-drug conjugates (ADCs) for cancer therapy, like Trastuzumab and Cetuximab, for breast cancer, and colorectal and head and neck cancer, respectively (Figure 2B and

C).^{12,14} Others ADCs, have been approved for radionuclides delivery, immunotoxins and antitumor antibiotics.

It is presently accepted that angiogenesis is the most limiting factor to tumor growth, since the loss of blood vessels leads to tumor shrinkage. Thus, it was promoted the development of conjugates with the ability to target tumor endothelial cells, ADCs or antibody-fusion proteins, like Avastin (bevacizumab) and VEGF-Trap, respectively.¹⁴

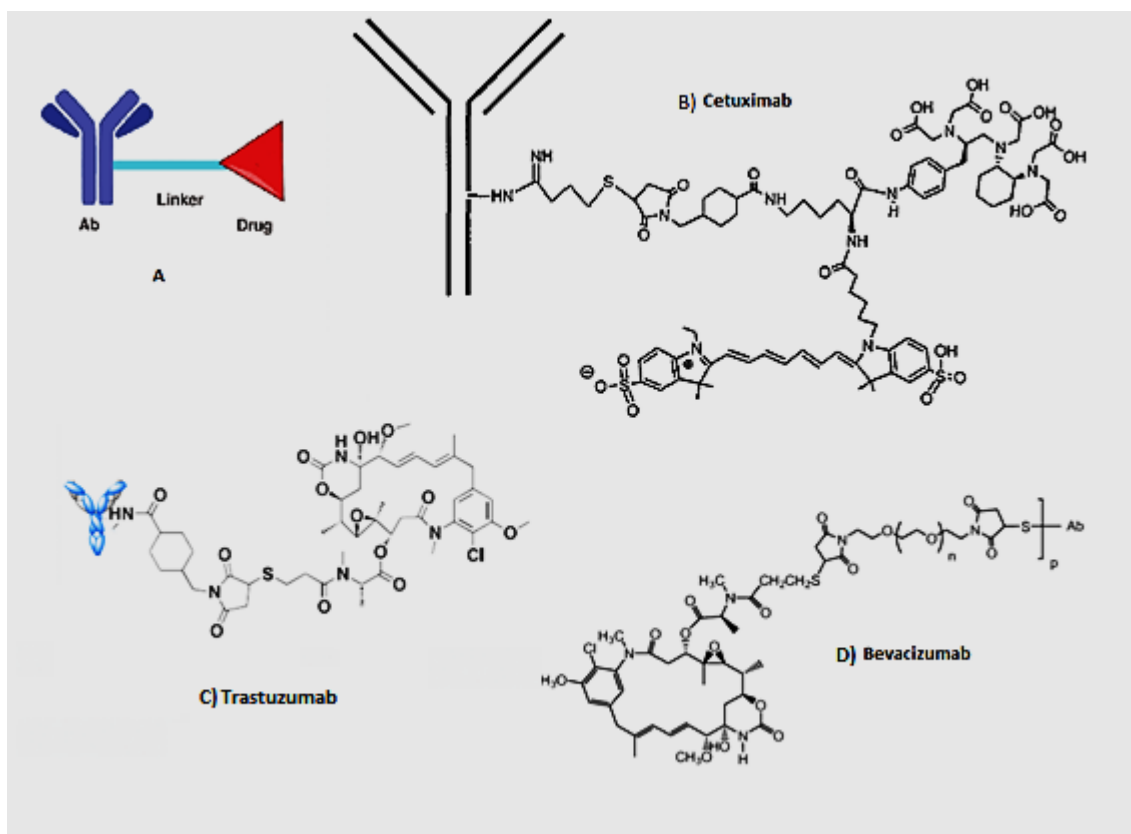


Figure 2: ADCs. A) General structure of ADCs; B) Cetuximab; C) Trastuzumab; D) Bevacizumab. Ab – Antibody. Adapted.²¹⁻²⁴

Despite many decades of research, tumor targeting therapy did not reach significant clinical success. Abs and large protein ligands have several limitations, such as the low tumor tissue penetration ability because of their large size and dose-limiting toxicity to the liver, spleen and bone marrow, due to non-specific uptake into the reticulo-endothelial system. Besides, Abs have a difficult and expensive commercial-scale production and there is the possibility of anti-idiotypic Abs generation and the forming of immune complexes.¹⁴

In an attempt to counter these disadvantages, there were developed small molecules inhibitors (SMIs). SMIs can specifically target tumor addresses in order to block cellular pathways or mutant proteins required for cancer cell growth and survival.^{3,25} Today, a lot of SMIs are under pre-clinical and clinical trials, being tyrosine kinase inhibitors group the most approved for tumor therapy, such as imatinib, sorafenib, erlotinib and lapatinib (**Figure 3**).^{3,25,26}

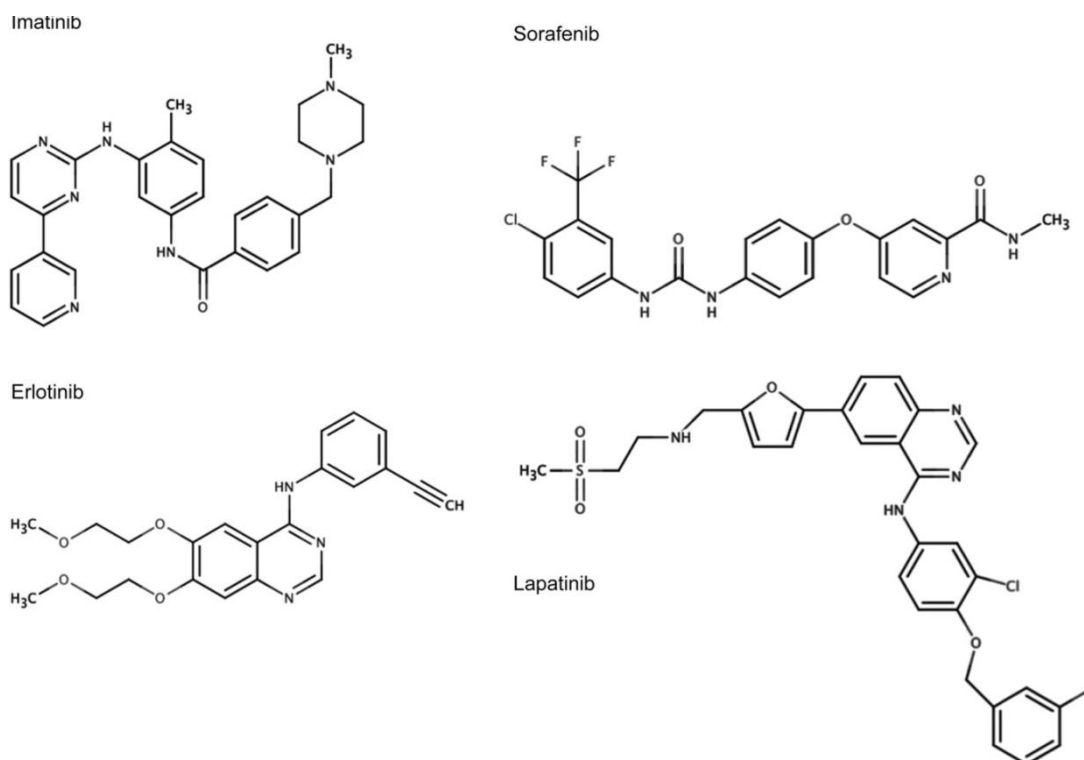


Figure 3: Chemical structures of SMIs examples. Adapted.²⁷

1.3. Peptides in Cancer Therapy

Peptides consist in small proteins, constituted by 50-100 amino acids (**Table 1**). Like proteins, peptides can organize structurally in a primary structure composed by an amino acids sequence and in a secondary structure with a special arrangement, like α helixes and β sheets.²⁸

Table 1: Amino acids and their abbreviations. Taken from ²⁹

Amino acid	Three-letter abbreviation	One-letter abbreviation	Amino acid	Three-letter abbreviation	One-letter abbreviation
Alanine	Ala	A	Methionine	Met	M
Arginine	Arg	R	Phenylalanine	Phe	F
Asparagine	Asn	N	Proline	Pro	P
Aspartic acid	Asp	D	Serine	Ser	S
Cysteine	Cys	C	Threonine	Thr	T
Glutamine	Gln	Q	Tryptophan	Trp	W
Glutamic acid	Glu	E	Tyrosine	Tyr	Y
Glycine	Gly	G	Valine	Val	V
Histidine	His	H	Asparagine or aspartic acid	Asx	B
Isoleucine	Ile	I	Glutamine or glutamic acid	Glx	Z
Leucine	Leu	L			
Lysine	Lys	K			

In cancer therapy, peptides are functional domains of proteins with specific bioactivities, including binding to receptors, structural sensitivity to chemical conditions (like acidity and temperature), penetration of the plasma membrane and activation or inhibition of cellular pathways.^{2,5} Taking into account anti-tumor therapy, there are three main groups of peptides with increased value: antimicrobial peptides or pore-forming peptides (AMPs), cell penetrating peptides (CCPs) and tumor targeting peptides (TTPs) (**Figure 4**).¹⁵

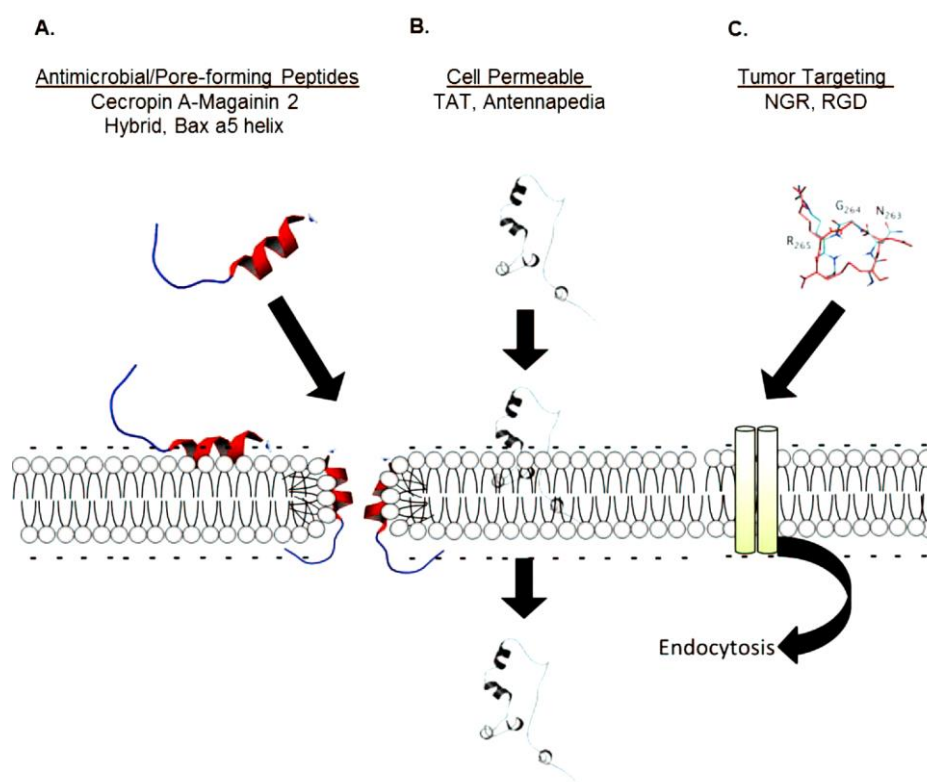


Figure 4: Classes of peptides with advantageous features in cancer therapy and respective examples. A) AMPs B) CPPs; C) TTPs. TAT - trans-activating transcriptional activator. Taken from ²

AMPs (also called host defense peptides or cationic antimicrobial peptides) are cytotoxic agents based in naturally occurring peptides of protective innate immune response to microbes in many species. This peptides has a cationic amphipathic structure that allows them to interact with anionic lipid membranes and induce pores formation or membranes disruption, causing cellular necrosis or apoptosis. Whereas necrosis is the result of AMPs targeting lipids membrane, leading to cells lysis, apoptosis is triggered by mitochondrial membrane disruption. AMPs constitute alternative anti-tumor drugs since their cytotoxicity occurs within minutes, decreasing drug resistance.²

On the other hand, CPPs have the ability to deliver cargos into cells, from small molecules to large microparticles, without having tumor cell specificity. CPPs are able to penetrate cell membranes directly through the plasma membrane or through several internalization mechanisms, such as clathrin- or caveolin-mediated endocytic pathway, micropinocytosis or an endocytosis-independent mechanism, including the carpet, inverted micelle, barrel stave pore and toroidal models (**Figure 5**).^{2,15} CPPs can be organized into cationic, hydrophobic and amphipathic groups.¹⁵

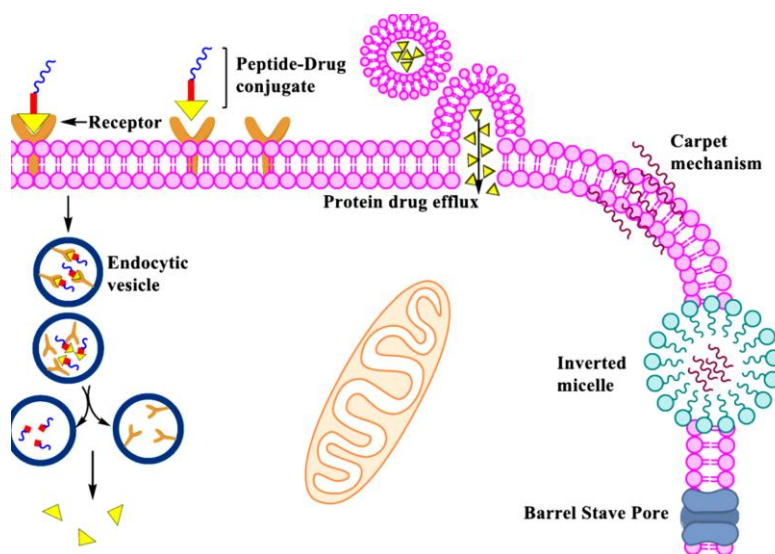


Figure 5: Different mechanisms used by CCPs to cellular internalization. Adapted.¹¹

1.3.1. Peptides in Tumor Targeting Therapy

Since their discovery about 30 years ago, TTPs, also known as tumor homing peptides, have become an useful tool for tumor therapy and diagnosis, due to their ability to mimic targeting properties of antibodies without macromolecular disadvantages.¹⁴ Thus, TTPs bind to specific addresses of tumor tissues with low to no affinity to normal cells. In comparison to antibodies, TTPs have a much smaller size

libraries (natural peptides or antibodies), one-bead one-compound (OBOC) libraries and phage display peptide libraries.¹⁴ In focused Abs libraries, peptides displaying specific antigen affinity are identified by amino acid sequences of complementarity determining regions (CDRs) of Abs, variable domains of Abs responsible for antigen recognition. However, recent findings show that not all CDRs regions are important for antigen binding and CDRs outside regions can contribute to its coupling.

OBOC library screen methods are based on a chemical technique composed of 90 μm -sized beads, each containing a different peptide ligand.¹⁴ In this technique, peptides are subject to a split mix strategy and screened against cell surface targets previously labeled. The advantage of this method is the possibility of incorporating non-natural components in peptides ligands, like D-amino acids, cyclization and branches.^{14,35}

Phage display peptide libraries, mostly utilize to identify TTPs, are focused in a biological approach that creates a random combinatorial library, through genetic modifications of the DNA of phages, allowing the expression of surface ligands.^{14,36} These type of libraries are exposed to protein targets (for instance, through whole cells, tissue samples and live animals) and then phages with binding capacity are analyzed by DNA sequencing, immunohistochemistry, *in vivo* imaging and mass spectrometry to identify target-binding peptides.^{13,14,37} In this method, there are multiple repetitions of selection and amplification steps (panning) to enrich the number of surface targets with the highest binding affinity, allowing the screening of a large number of peptide sequences without the need to have previous knowledge of the molecular composition of the binding site (**Figure 7**).^{14,35}

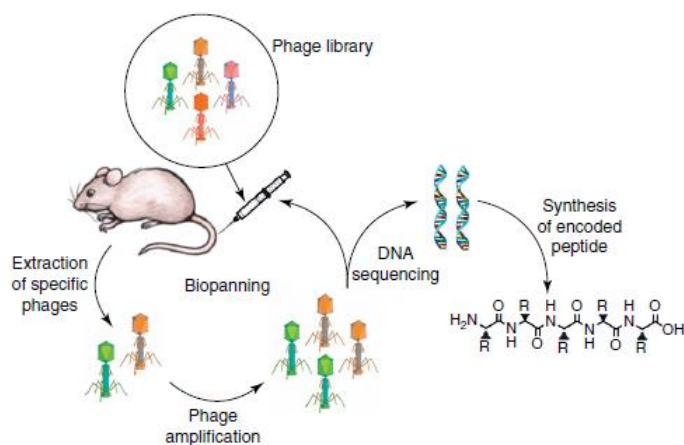


Figure 7: Steps of screening and identification of TTPs using the *in vivo* phage display technology. Taken from ¹⁹

1.5. Targeted Drug Delivery Systems

Nowadays, both in academia and industry, the development of site-specific drug delivery systems (DDS) and efficient drug targeting approaches are being investigated to surpass the systemic “off-target” effects shown by antitumor therapies and to maximize their therapeutic efficacy.^{11,13,17,38}

Generally, the starting point of a drug targeting approach is a prodrug construction by a covalent modification of the drug with a carrier unit, a small moiety that inactivates the pharmacological effect of the drug during delivery and provides adequate pharmacokinetic properties.³⁸ The carrier unit and the drug are attached by a chemical linker that controls the intracellular release of the drug (for example, acid-cleavable and reduction-sensitivity spacers, where environment changes trigger their intracellular cleavage).¹¹

Alternatively, delivery vehicles (also known as particulate drug carriers) may be used to encapsulate the drug and control its biodistribution (**Figure 8**).³⁸ Delivery vehicles are molecular assemblies with high loading capacity that confine the drug in a loading space (commonly the core of the particulate), providing protection against enzymatic inactivation. Due to the absence of covalent conjugation to entrap the drug molecule, delivery vehicles allow the use of a single carrier unit to formulate several drugs. Furthermore, the composition and size of vehicles can be adjusted to determine particulates with ideal physicochemical properties, which diversifies their stability and, per consequence, the rate of drug release. In the field of cancer therapy, the ideal size is in the 50-150 nm range (nanocarriers) to avoid extravasation into normal tissues and, at the same time, enabling the extravasation from most tumor blood vessels into tumor interstitium.¹¹

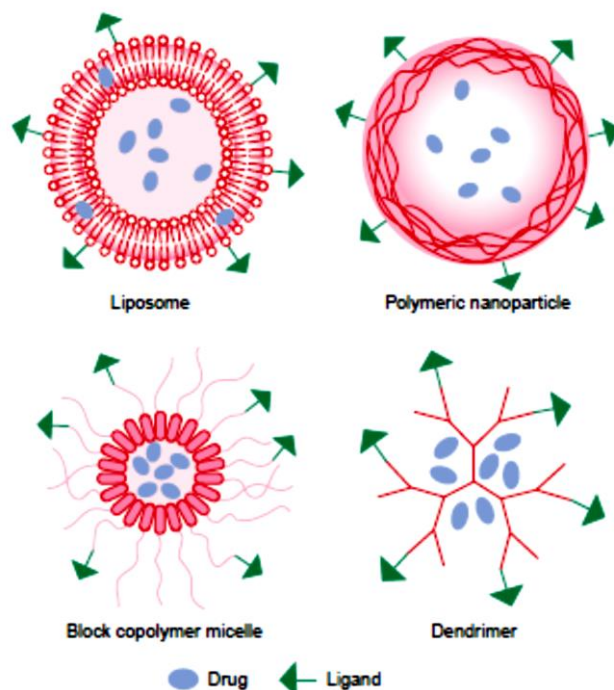


Figure 8: Delivery vehicles used in targeted DDS to cancer therapy application. Adapted.¹⁷

The most common nanocarriers are liposomes, polymeric nanoparticles and block copolymer micelles. Whereas liposomes are lipid molecules orderly in a lamellar disposition, generally used as unilamellar vesicles, polymeric nanoparticles are biocompatible polymers where the drug is entrapped to a vesicle surrounded by a polymer membrane or dispersed in a matrix, named nanocapsules or nanospheres, respectively. Moreover, block copolymer micelles are characteristically spherical amphiphilic copolymers, whose loading space accommodates hydrophobic drugs, and the hydrophilic outer layer promotes dispersal of the micelles in water.¹⁷ As drug delivery nanocarriers or imaging probes, nanoparticles have been shown more advantages in comparison with others nanocarriers, including the better tumor penetrating ability, higher ability to carry cargos and quality of image information.¹² However, the prodrug approach tends to prevent premature drug liberation and decreases inert materials quantity.³⁸

It is common to conjugate hydrophilic polymers, like poly(ethylene glycol) (PEG), onto carrier units' surface to inhibit their uptake by the reticuloendothelial system, therefore prolonging the *in vivo* half-life and, consequently, the targeting potential. PEG forms a hydrophilic barrier around the particulate that blocks proteins and other macromolecules to reach the particulate surface, including antibodies raised against the particulate. In this case, the ligand must be conjugated to the terminal of

the stabilizing component to prevent the shielding effect that leads to a severe reduction of ligand-receptor interaction.¹⁷

For the majority of targeted DDS, the cargo needs to be internalized into cells to exercise its pharmacological activity. After the peptide-receptor interaction, conjugates can be internalized through receptor-mediated endocytosis, where they may move through the early and late endosomes and finally, into lysosomes. Furthermore, to exert pharmacological activity, the cargo has to exit these organelles to reach the cytosol or the nucleus (the two sites of action of intracellularly active drugs). In lysosomes, due to enzymes and/ or pH dropping, the linker between the cargo and the carrier is broken, enabling the cargo escape to cytosol while, in the meantime, receptors are recycled (**Figure 9**).^{17,39} If the microenvironmental conditions do not allow the rapid disintegration of the particle, the diffusion of the drug out of the endosomes is necessary, through lytic peptides, pH-sensitive polymers and swellable dendritic polymers.^{17,32} In the case of nanocarriers, internalization of the drug can occur without concomitant internalization of the particulate, due to atypical conditions of the tumor microenvironment (like acidic pH, presence of lipases, enzymes and oxidizing agents), which results in the accelerated release of the drug and its diffusion into tumor cell, through passive diffusion or active transport.¹⁷

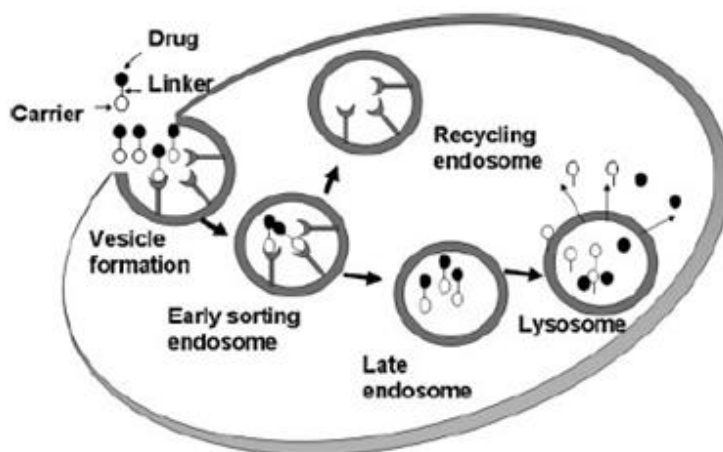


Figure 9: Schematic cell internalization process for targeted DDS. Adapted.³⁹

1.5.1. TTPs as Carrier Units in DDS

During the past two decades, more efficient functional peptides were developed.¹⁵ In particular, TTPs have been studied as carrier units of peptide-based drug conjugates (PDCs), an emerging class of prodrugs, formed through the covalent

attachment of a specific peptide sequence to a drug via a cleavable linker (**Figure 10**).^{11,38}

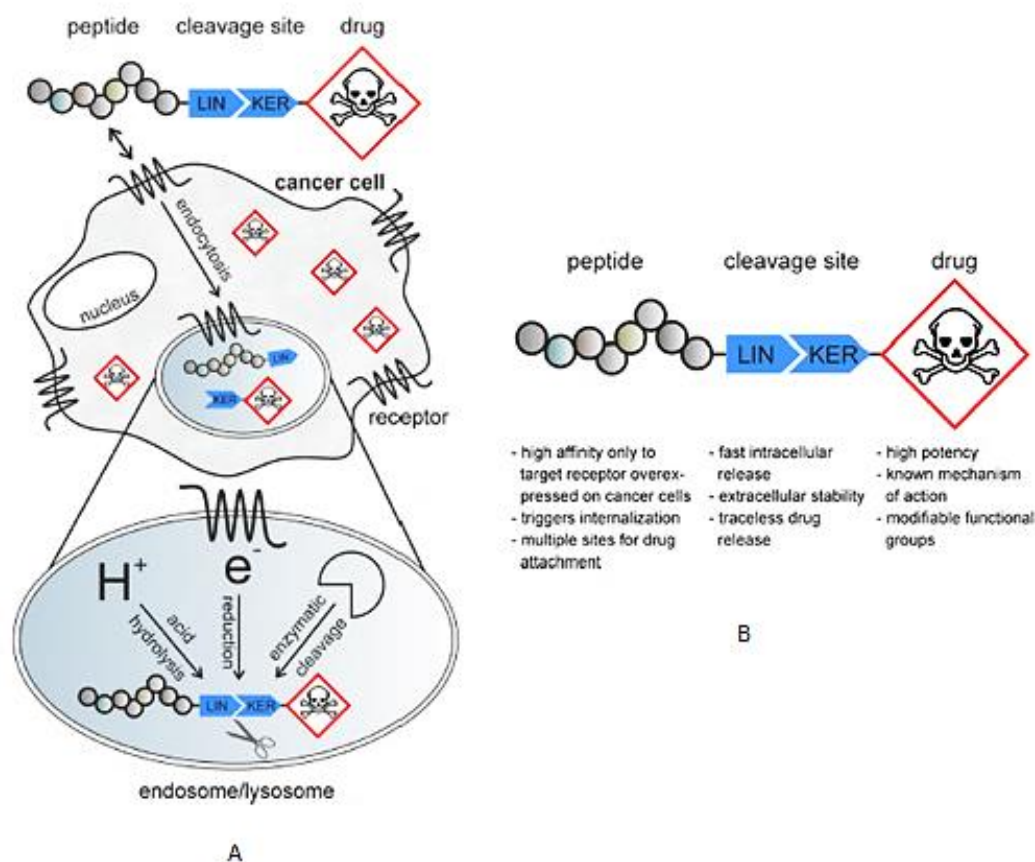


Figure 10: Schematic representation of PDCs. A) Schematic representation of tumor targeting approaches using PDCs; B) Necessary key-properties of PDCs components. Adapted.¹¹

To ensure drug delivery to tumor cells, it is necessary to improve stability *in vivo* and evaluate pharmacokinetic properties of PDCs, that will improve selectivity for cancer cells.² Generally, peptides have relatively large size, hydrophilicity and hydrogen-binding potential, which makes them unsuitable to be orally administered, due to the difficult passage by intestinal mucosal barriers. Thus, the parenteral route is the most chosen to deliver TTPs conjugates.³⁹ Moreover, it is possible to conjugate multiple TTPs to a single delivery vehicle to increase cargo delivery and prevent peptides degradation by proteolysis, by blocking N- and C-terminally, through incorporation of D-amino acids (unnatural amino acids), cyclization and insertion of reduced peptide bonds.^{14,31} The most commonly utilized cyclization method is disulfide bridge formation from two Cys residues.⁴⁰ Other important property of PDCs that promotes efficient delivery of drugs is the internalizing ability of some TTPs into cell membranes.¹⁷ Since some TTPs cannot internalize their cargos, CPPs-TTPs

conjugates have been developed, which contribute for a better intracellular drug delivery mechanism (**Figure 11**).¹¹

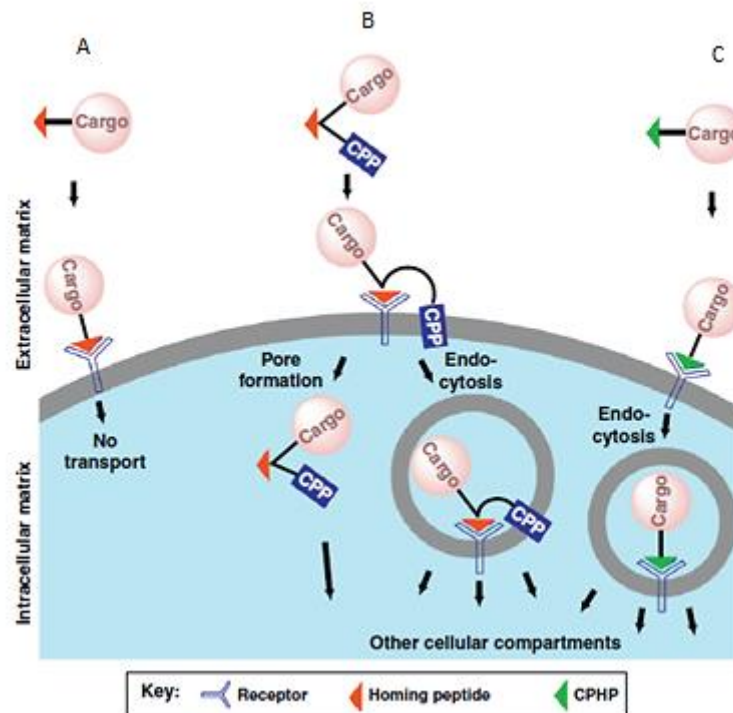


Figure 11: Principal of cell-selective peptide targeting and delivery. A) TTPs without internalization ability; B) TTPs-CPPs conjugates; C) TTPs with cell penetration ability. CPHP – Cell-penetrating homing peptides. Adapted.¹⁹

TTPs conjugates with cytotoxic drugs, in addition to being exhaustively studied in several preclinical and clinical studies, were also investigated as carriers of radioisotopes.^{12,30} In this case, there is an attachment of a radio-ligand to the peptide carrier through the aid of a chelator, such as diethylenetriaminepentaacetic acid (DTPA) and 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid (DOTA), but there is no release of the cargo, since it is an unnecessary process. Radionuclides like ^{99m}Tc , ^{111}In , ^{68}Ga , ^{123}I , ^{64}Cu and ^{18}F can be use in cancer radiotherapy, due to their longer half-life.³¹

2. Objectives

Nowadays, cytotoxic drugs used in antitumor therapies are not very effective and cause serious adverse effects due to their poor ability to distinguish tumor cells from the normal cells. To solve this problem, molecular systems based on conjugation of a cytotoxic drug and an antibody responsible for the tumor targeting ability are available on the market. However, these therapeutic conjugates have a high economic cost and structural disadvantages. On the other hand, since their discovery about 30 years ago, tumor targeting peptides (TTPs) are able to recognize specifically tumor targets, such as Abs but without their structural disadvantages. Thus, the use of these peptides in targeted therapeutic delivery systems to cure cancer patients is very promising. However, many of the features of these peptides are not yet fully understood.

The aim of this work is to review the literature regarding tumor targeting approaches and to organize and summarize the TTPs that contributed in a significant way to the advances of this field, including the development of therapeutic conjugates. This way, it is intended to focus their structural and functional characteristics, addresses, structure-activity relationship (SAR) studies, internalizing capacity and application in the development of drug conjugates.

3. Materials and Methods

3.1. Materials

In this work journal articles were used, mainly review articles published in several international scientific journals, as sources of information. These articles were collected and selected from the National Library of Medicine's search service (PubMed database), that provides access to millions of references in United States National Library of Medicine and other related databases.

3.2. Methods

Firstly, the literature review started with the collection and reading of articles (especially review articles) regarding tumor targeting-based approaches and peptides with tumor targeting abilities. Then, the most representative peptides of the scientific advances made in this field were selected based on these articles. The selection criteria were based on the number of articles that referred the respective peptides, the quantity of information available, the actuality of the articles and the tumor addresses. Finally, others articles of each selected peptide were collected to summarize information of each, in order to highlight structural and functional features, addresses, structure-activity relationship (SAR) studies, internalization capacity and applications in the development of therapeutic conjugates (**Table 2**).

Table 2: Key-words and expressions utilized to obtain the selected articles.

Key words and expressions	References
Tumor targeting internalizing peptides review	17
Tumor targeting peptides review	7,11,12,14–16,30–32,37,41
Tumor homing peptides review	2,13,18,19,42
Peptides targeting cancer review	5,35
Peptide drug conjugates review	38,39
Tumor therapy review	1,8,10,26
Cancer therapies review	3,25
Novel cancer treatment review	4,43
F3 peptide nucleolin	44–47
RGD	48–63
NGR	36,64–66
NGR structure activity relationship	67
NGR structure activity	68
NGR pharmacophore	69
Lyp	70,71
Lyp peptide binding	72
Bombesin	73
Bombesin receptor cancer review	74
Bombesin receptor interaction structure	75–77
CREKA	78–82
SSTR mediated endocytosis	83–86
Angiopep 2	87–91
M2pep	40,92–96

4. Results

Following the literature review, nine peptides were selected: RGD, NGR, F3, Octreotide, LyP-1, Bombesin, Angiopep 2, CREKA and M2pep. These peptides were organized according to the targeting localization type: tumor vasculature, tumor lymphatic vessels, tumor cells and tumor microenvironment. Relevant and recent information concerning others TTPs was collected and organized (**Annexes 1 and 2**).

4.1. Peptides Targeting Tumor Vasculature

4.1.1. RGD

The RGD tripeptide (Arg-Gly-Asp) is the most widely investigated among all TTPs (Figure 12A).¹⁴ This tumor targeting peptide was discovered by phage display techniques, as an essential cell recognition site of several blood, extracellular matrix (ECM) and cell surface proteins like fibronectin, vitronectin and fibrinogen.^{2,7,15,35,42,48,49} The RGD peptide can selectively target endothelial cells of tumor blood vessels expressing $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins receptors.^{13,14}

Integrins are cell adhesion receptors for ECM proteins, immunoglobulins, growth factors, cytokines and matrix-degrading proteases. These divalent heterodimeric membrane glycoproteins, composed by non-covalently associated α - and β -subunits are expressed at the surface of normal tissue and blood vessels.^{7,14,42,55} Despite 24 integrins subtypes being known, endothelial cells of angiogenic vessels express a different set of integrins, being $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins specifically upregulated in tumor vasculature.^{5,14,35} Furthermore, $\alpha_v\beta_3$, $\alpha_v\beta_5$, $\alpha_5\beta_1$, $\alpha_6\beta_4$, $\alpha_4\beta_1$ and $\alpha_v\beta_6$ integrins are also overexpressed in tumor cells, being the most studied integrins in cancer.⁵¹ However, RGD can be recognize by 8-12 of the 24 known integrins.³⁵ The activation of this receptors triggers cellular pathways involved in tumor angiogenesis and metastasis promotion.^{11,37} In particular, $\alpha_v\beta_3$ integrin regulates intracellular signaling that protects tumor cells from the anti-proliferative action of anti-tumor drugs.³⁷

There are several RGD recognition sites in α and β subunits of integrins. In all cases, these binding sites are localized at or near a binding site for divalent cations, being the β_3 subunit of $\alpha_v\beta_3$ integrins the primary site for RGD binding.^{48,57,58} Adjacent binding sites of divalent cations helps to keep a favorable binding conformation. By itself, a simple linear RGD ligand presents low affinity for integrins receptors. This is

related to the conformation freedom of the RGD peptide that determines its selectivity.^{5,60} For example, the RGD-TNF α conjugate is promising, although the presence of four Cys residues in the peptide structure makes it difficult to fold in a homogeneous manner, compromising its effectiveness.^{14,42} This way, to obtain a biologically active conformation, several cyclic RGD analogues (cRGD) were developed via “head-to-tail” modification, to create a more rigid structure, more stable at neutral pH and better at resisting proteolysis.^{5,52,54,60,62}

In cyclic peptides, the RGD motif is flanked by other amino acids to build a ring system, which may induce receptor affinity or selectivity and other biological properties.⁵⁵ cRGD peptides include RGD-4C (ACDCRGDCFCG) and Cilengitide (**Figure 12**).^{5,54} Cilengitide, the salt of the cyclic pentapeptide with the sequence Arg-Gly-Asp-DPhe-(NMeVal) - c(RGDf[NMe]V) - demonstrated encouraging results in patients with glioblastoma, which implies penetration of the blood brain barrier (BBB), although clinical trials have not fully proved its effectiveness.^{2,7,11,19,35,37,59} Cilengitide can act as a specific antagonist of $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins.^{9,35} Others structural modifications in Cilengitide, including the introduction of the unnatural D-conformation of Phe amino acid and N-methylation, improve its receptor affinity and pharmacokinetic properties.⁶⁰ More recently, three PDCs were developed based on this pentapeptide, in which the mutilated Val was mutated to Lys or Ser amino acids for the creation of a primary amine or hydroxyl group, potential sites for drug conjugation.³⁵

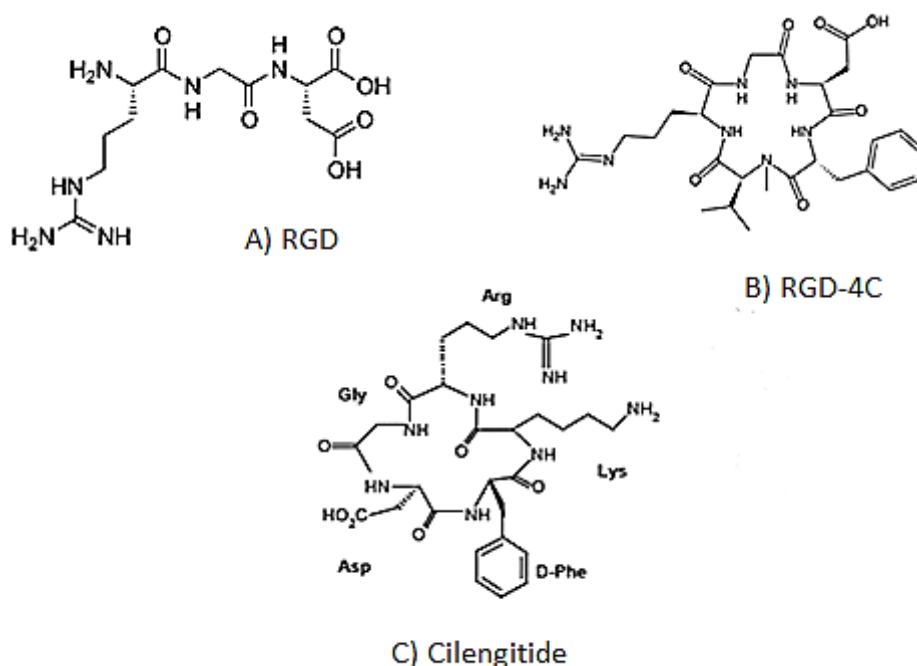


Figure 12: Chemical structure of RGD peptides. Adapted.⁵⁵

On the other hand, others RGD peptides were designed to improve tumor penetration.² Internalizing RGD (iRGD) is a 9-amino acid, disulfide-bridged cyclic peptide with the ability to facilitate the internalization into endothelial cells, in addition to target $\alpha_v\beta_3$ integrins.^{2,13,14,97} This extra capacity allows the intracellular uptake of the conjugated cargos, contributing to the internalizing process of antitumor drugs. The penetration capacity is due to a specific sequence contained in the peptide, the sequence R/KXXR/K (X: any amino acid) at C-terminal, named C-end Rule (CendR), that binds to neuropilin-1 (NRP-1), a transmembrane protein expressed on membranes of endothelial cells that, when activated, triggers downstream signal pathways to increase cell permeability.^{2,13,14,63} After iRGD connection to an integrin, a protease cleaves the peptide to produce CRGDK/R and exposes the CendR motif that can interact with NRP-1 receptors (**Figure 13**).^{55,61} The CendR penetration is receptor-mediated, energy-dependent, effective with small molecules and causes extravasation of the peptide (or their cargos) within minutes.⁶¹

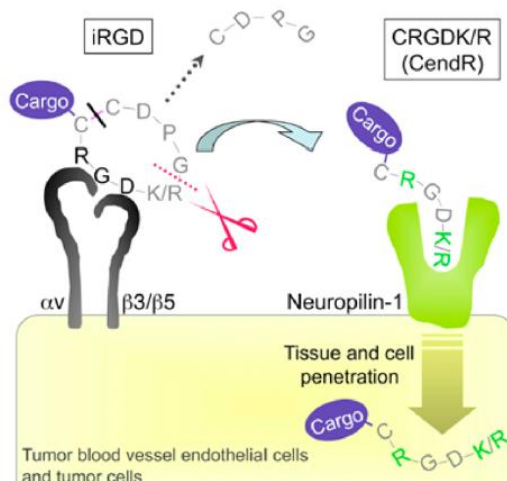


Figure 13: Penetration mechanism of iRGD. Taken from ⁵⁵

The current clinical trials involving RGD motif (Table 7, Annex 3) are testing its application not only in tumor therapy, but also in diagnostic imaging, through conjugation with cytotoxic drugs, peptides or proteins, nucleic acids, radionuclides and contrast agents. For instance, the first RGD-modified positron emission tomography (PET) tracer in clinical trials was Galacto-RGD, an RGD analog obtain by cyclization of a RGD pentapeptide and modification of Phe to the unnatural configuration – c(RGDfv).^{5,52}

More specifically, in clinical trials, the improvement of the delivery and therapy efficacy of tumor therapy agents when conjugated with RGD peptide or analogues,

namely cytotoxic drugs like paclitaxel (PTX) and doxorubicin (DOX), therapeutic peptides such as KLAKLAKKLAKLAK (KLA), cytokines like tumor necrosis factor α (TNF α), interferon- γ (IFN- γ) and interleukin-12 (IL-12), Abs and their fragments such as the Fc fragment of immunoglobulin G (Ig G) was proven.^{12,13} Furthermore, RGD-targeted nanocarriers benefit of the possibility to enhance the internalization process via integrin-mediated endocytosis, instead of standard uptake mechanisms after drug release, such as the RGD-modified PEGylated liposome-encapsulates DOX.^{5,55}

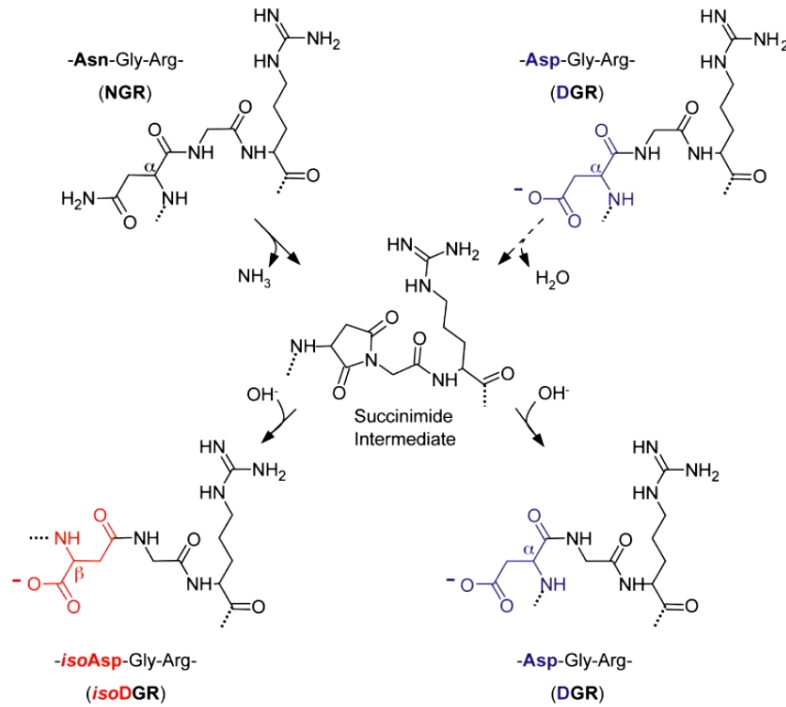
4.1.2. NGR

The NGR tripeptide (Asn-Gly-Arg), discovered by the *in vivo* phage display method, specifically binds to tumor endothelial cells expressing the aminopeptidase N (APN) receptor, also called CD13.^{13,37,64} Like RGD, this sequence are encountered in natural proteins, such as fibronectine.³⁶

CD13 is a membrane-bound and highly glycosylated metalloproteinase, with a significantly role in protein degradation and regulation of cytokines, antigen presentation, cell proliferation and migration and angiogenesis.^{13,42,66} Although it is overexpressed in tumor blood vessels, CD13 can be find in tumor cells, fibroblasts and pericytes, and in normal tissue like mast cells, keratinocytes, proximal renal tubules, myeloid cells and epithelial cells.^{13,36,42} There are several CD13 isoforms, which could explain the binding of NGR peptides to tumor vasculature but not to other CD13-rich tissues.^{14,42,68} Furthermore, in endothelial cells, CD13 interacts with galectin-3 (a proangiogenic protein) in a carbohydrate recognition-dependent manner, forming a complex that, together with the abnormal architecture of tumor blood vessels, may be related to differential glycosylation or conformational changes and, therefore, with the selectivity of NGR motif to endothelial CD13.³⁶

Recent studies suggest that NGR can interact with integrins by a spontaneous and unusual mechanism that consists in the rapid deamidation of the Asn residue, through the formation of a succinimide ring, followed by hydrolysis, forming isoaspartate and transforming NGR into the isoaspartate-glycine-arginine (isoDGR) derivate, a cell adhesion motif with high affinity for $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins (**Figure 14A**).^{36,65,68,69,98} Thus, isoDGR can promote endothelial cell adhesion, having a binding site located within the RGD binding pocket, being as well an antagonist of $\alpha_v\beta_3$ integrin.^{36,65} However, mostly in injured tissues and wound healing, cells produce protein-L-isoAsp-O-methyltransferase (PIMT), an enzyme that converts isoaspartate to aspartate, causing the deactivation of isoDGR function (**Figure 14B**).³⁶

A



B

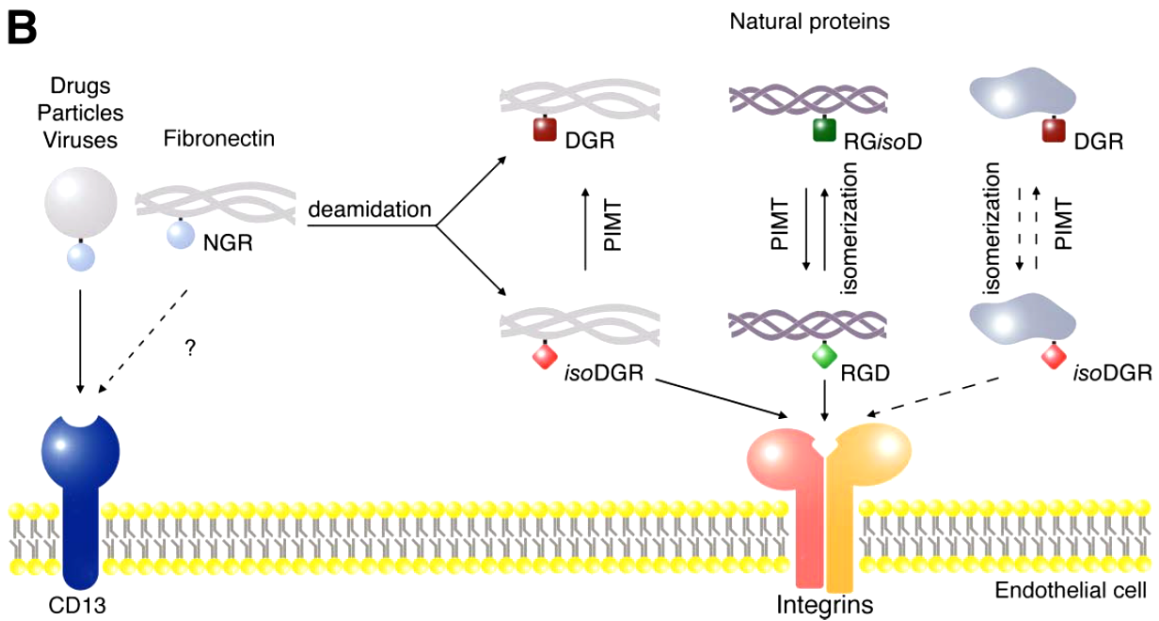


Figure 14: Formation of isoDGR (A) and schematic representation of effects on receptor interactions (B).

Taken from ³⁶

Several therapeutic molecules have been conjugated to NGR peptides, namely cytotoxic drugs, therapeutic proteins, proapoptotic peptides, viral particles, imaging agents and DNA complexes. Specific examples include conjugation of NGR with DOX, the proapoptotic peptide D and $\text{TNF}\alpha$.^{13,64} This conjugates may penetrate cell membrane via receptor mediated endocytosis.⁶⁸

Generally, small cyclic NGR peptides are more selective than linear forms, due to conformational constraining that improves binding affinity. The most effective cyclic NGR peptides developed are c(CNGRC) and (KNGRE)-NH₂ (**Figure 15A and B**). The c(CNGRC) was designed with the disulfide bonding of the two Cys residues, increasing the targeting efficiency due to stabilization of the bent conformation.^{14,64,65,68} Particularly, cCNGRC-TNF α (currently tested in Phase III clinical trials) (Table 8, Annex 3) have shown a promising potent anticancer activity due to its capacity to facilitate drug penetration and infiltration of immune cells, since TNF α increases intracellular adhesion molecules on endothelial cells, expression of pro-inflammatory cytokines and recruitment of tumor-specific cytotoxic T cells.^{13,64} Moreover, to surpasses the disulfide bond disadvantages (susceptibility to biodegradation and chemical modification), the “head-to-side-chain” amide bond cyclized peptide - c(KNGRE)-NH₂ - was developed.^{65,68} Other examples of NGR peptides are GNGRG, NGRAHA and CVLNGRMEC.⁶⁴ SAR studies of GNGRG (**Figure 15C**) suggest that the linear NGR peptide without disulfide constraints is more thermodynamically favorable when its configuration is based in a β -turn in Gly3 and Arg4 and hydrogen bonding interactions between Asn2 and Cys5. Thus, like the RGD receptor binding, there is necessary a folded structure for receptor binding, with intramolecular stabilizing interactions (like hydrogen bonds) that stabilize the folded state.⁶⁷

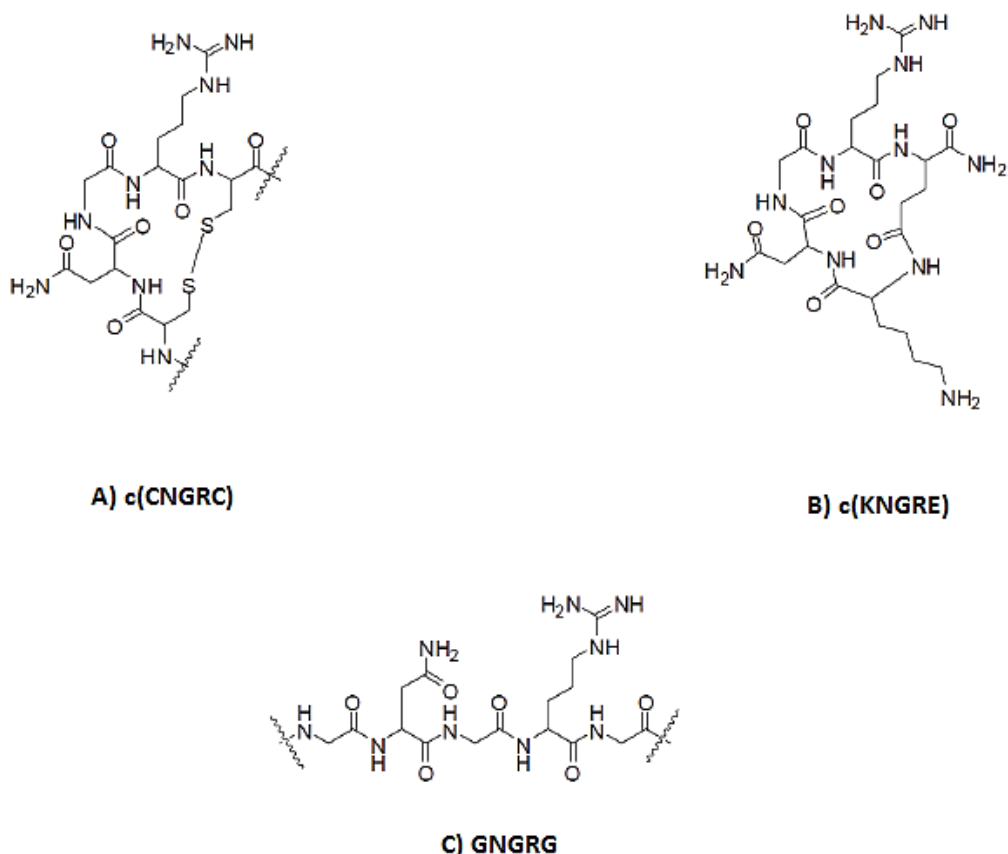


Figure 15: Structure of common NGR peptides. Adapted.⁶⁵

Similar to the iRGD, internalizing NGR (iNGR) possesses the CendR motif and, therefore, the ability to bind to NRP-1 and penetrate tumor endothelial cells.¹³

4.1.3. F3

F3 peptide (KDEPQRARSARLSAKPAPPKPEPKPKKAPAKK) is a 34–amino acid fragment of the human high mobility group protein 2 (HMG₂N), a nucleosomal protein that participates in unfolding of chromatin structure, facilitating the transcriptional activation of genes.^{44,46,47,99} This tumor targeting peptide, discovered by the phage display libraries technique, binds to the nucleolin (NCL) receptor overexpressed in tumor cells, tumor endothelial cells and in a bone marrow subpopulation, precursor of endothelial cells.^{44,46,99}

NCL is a non-ribosomal protein important to polymerase I transcription, being found in high quantity in the cell nucleolus, where ribosome assembly and ribosomal ribonucleic acid (rRNA) transcription occurs. Other nucleoplasmic NCL receptors are responsible for the regulation of gene expression, mostly of oncogenes, and genome

stability, due to its interaction with RNA polymerase II. Moreover, in cytoplasm, nucleolin interacts with messenger ribonucleic acid (mRNAs) coding proteins related with cell proliferation and apoptosis. NCL structure can be divided in three domains: the negatively charged N-terminal domain (with acidic regions due to the abundance in aspartic acid and glutamic acid) disunited by basic stretches, the central domain containing four RNA binding domains and the C-terminal domain, predominantly with Gly, Arg and Phe residues.^{45,99}

In tumor cells and tumor endothelial cells, overexpressed NCL promotes the cell internalization of several ligands related with angiogenesis, proliferation and apoptosis.⁴⁵ F3 peptide can be internalized by NCL due to its NH₂-terminal domain (via receptor-mediated endocytosis) and carried into the nucleus after cell penetration.^{14,99} SAR studies show that the D-amino acid form of F3 has equal internalization ability, although it is not efficiently transported into the nucleus. This internalizing mechanism is not defined, but studies have shown energy dependence for the mechanism occurrence and the possibility of F3 interaction with NRP-1.^{14,46}

F3 peptide was conjugated with radiotherapeutic agents for tumor therapy, such as ²²⁵Ac, ²¹³Bi and ¹¹¹In.¹³ Nonetheless, there are no studies about the conjugation between this tumor targeting peptide and cancer drugs.¹⁹

4.1.4. Octreotide

SST is an endogenous acid polypeptide with the sequence Ala-Gly-c(Cys-Lys-Asn-Phe-Phe-Trp-Lys-Thr-Phe-Thr-Ser-Cys), with high-affinity for SSTRs. This peptide acts as an endogenous inhibitory regulator and has various biological functions, including inhibition of many hormone secretions, cell survival and cell proliferation.^{41,83} However, SST possesses an extremely short half-life (about 2-3 minute), due to its rapid inactivation by peptidases, being limited its clinical utility.^{41,86} Thus, several synthetic SST analogues were developed to treat endocrine tumors, through shortening of its sequence and introducing D-amino acids to increase the half-life time. SAR studies demonstrate that the key sequence for binding and biological activity was the β -turn fragment Phe-Trp-Lys-Thr sequence, the residues 7-10 of the SST, in addition to the existing disulfide bridge.^{83,86} After interacting with the receptor, SST analogues are rapidly internalized into tumor cells (via SSTR-mediated endocytosis), being able to translocate to the nucleus.^{83,84} Furthermore, these analogues have been used to develop conjugates with application in targeting therapy, radiotherapy and tumor imaging (**Figure 16**).⁸³

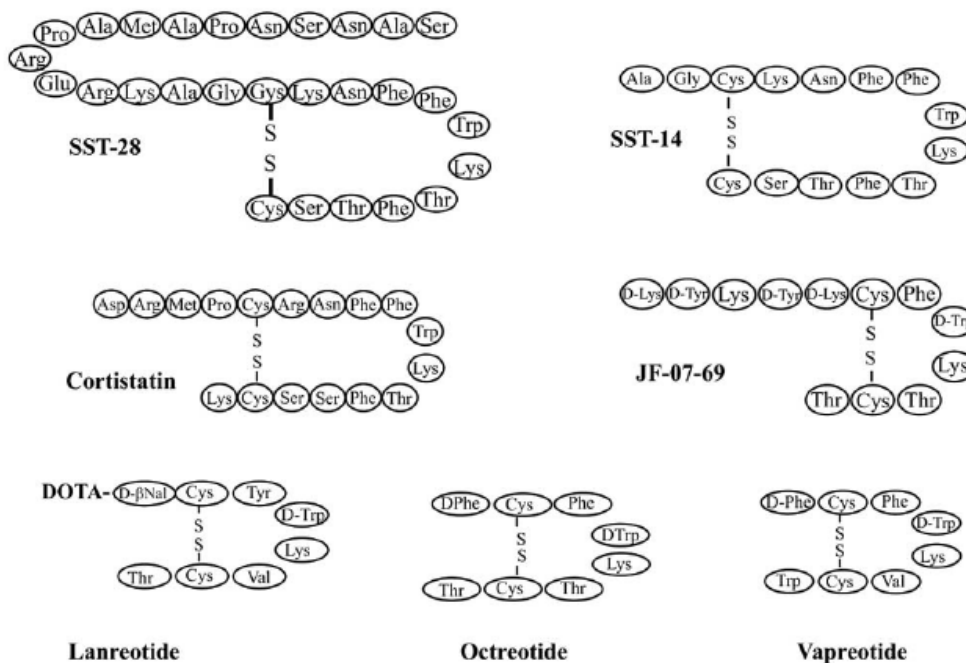


Figure 16: Schematic chemical structures of SSTs and analogues. Taken from ⁸³

SSTRs are transmembrane G-protein coupled receptors expressed in the central nervous and immune systems, endocrine tissue, gastrointestinal tract and skin, playing different physiological roles.^{35,41,85} There are five distinct SSTR subtypes, being the SST2R and SST5R receptors overexpressed in various types of tumors, not only in tumor cells, but also in tumor endothelial cells.^{32,83}

Octreotide, a cyclic octapeptide with the sequence D-Phe1-c(Cys2-Phe3-D-Trp4-Lys5-Thr6-Cys7)-Thr8-ol), is an agonist of SSTRs, namely SST2R subtype, that mimics the natural SST hormone, although it is a more potent inhibitor of growth hormone, insulin and glucagon.^{32,41,86} Nowadays, octreotide is utilized successfully against growth hormone-producing tumors in many countries.^{32,41} The remarkable feature of octreotide is its stability towards degradative enzymes.⁸⁴ The β -turn is a stable conformation due to the N- and C-terminal of the D-Phe1 and Thr(ol)8 or Trp8-NH₂ amino acids by intramolecular hydrogen bonds that stabilize the conformation of the peptide.⁸⁶ Besides, SAR studies also showed that both C- and N-terminal residues of octreotide are important for binding affinity.³⁹

Potent chemotherapeutic agents including PTX, camptothecin (CPT), DOX and methotrexate (MTX) have been used for the development of octreotide conjugates.^{32,83} For example, two molecules of PTX were coupled to the N-terminal of octreotide by an ester bond and utilizing a succinic acid spacer (**Figure 17**).³² More

recently, the redox-sensitive prodrug octreotide(Phe)-polyethylene glycol-disulfide bond-paclitaxel - OCT(Phe)-PEG-ss-PTX - was design.⁴¹ On the other hand, the first radiolabeled SST analog clinically applied was the (¹¹¹In-DTPA⁰)-octreotide.^{5,84}

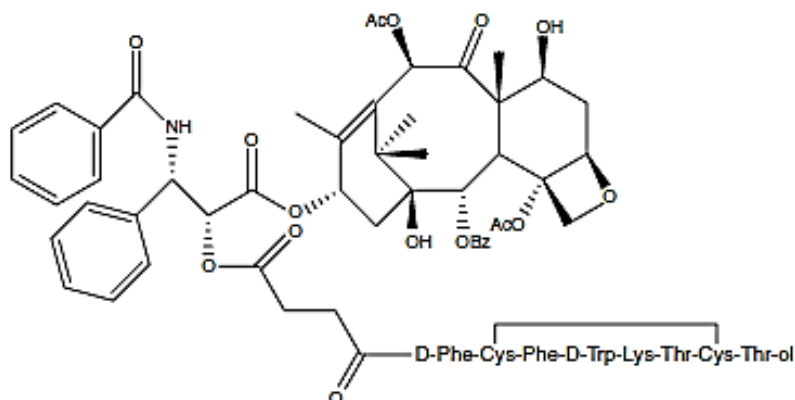


Figure 17: Chemical structure of ester bond-linked PTX-octreotide. Adapted.³¹

4.2. Peptides Targeting Tumor Lymphatic Vessels

4.2.1. LyP-1

LyP-1 is a cyclic nonapeptide (CGNKRTRGC) with the capability to target tumor lymphatic vessels and tumor cells in hypoxic areas, by interaction with p32, a mitochondrial/ cell surface protein receptor (**Figure 18**).^{14,16} LyP-1 can penetrate the plasma membrane into the cytoplasm and nucleus.⁷² By itself, the linear form of LyP-1 peptide has cytotoxic activity, since it accumulates in tumor hypoxia areas, decreasing the number of lymphatic vessels and promoting apoptosis of tumor cells undergoing stress.^{14,70,72} For instance, LyP-1 was used to improve the delivery of DOX-loaded PEGylated liposomes.¹⁵

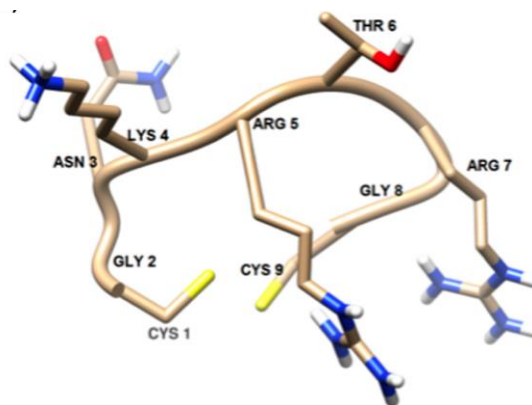


Figure 18: Cyclic structure of LyP-1. Adapted.⁷²

The p32 receptor, also called as hyaluronic acid binding protein 1, is overexpressed on tumor lymphatics, tumor cells and TAMs.^{15,16} However, the molecular mechanism of LyP-1 recognition by the receptor is poorly understood.⁷² On the other hand, LyP-1 internalization is similar to iRGD penetrating mechanism. The binding of LyP-1 to the p32 receptor triggers its proteolytic cleavage into tLyP-1 (CGNKRTR), allowing the exposure of CendR motif and the binding to NRP receptor.⁷¹

4.3. Peptides Targeting Tumor Cells

4.3.1. Bombesin

Bombesin is a 14-amino acid neuropeptide (QQRLGNQWAVGHLM), containing high affinity to the gastrin realizing peptide receptor (GRPR), a G protein-coupled receptor overexpressed in tumor cells, also named as bombesin receptor type 2 (BB2).^{11,39,41,73,75}

Mammalian bombesin receptors (BBRs) can be divided into four receptor subtypes: neuromedin B receptor (NMBR or BB1), GRPR and bombesin receptor subtypes 3 and 4 (BB3 or BRS-3 and BB4, respectively).^{35,73} These receptors can naturally occur in central nervous system (CNS) and in peripheral tissues, having a wide spectrum of actions in physiological processes.⁷⁴ The activation of BBRs promotes the release of others peptide hormones like insulin, glucagon, prolactin and gastrin and cytokine activity, being able to act as a growth factor.⁷⁵ Although there are four subtypes of BBRs, the natural ligand of BB3 is unknown and it has low affinity for bombesin peptides.⁷⁴ As tumors can secrete bombesin to stimulate tumor growth, tumor targeting approaches based on this peptide may interrupt tumor autocrine-growth.^{35,74}

BBRs antagonists can block the tumor growth stimulated by bombesin peptides. While BBRs agonists are internalized via receptor-mediated endocytosis, their antagonists have low to no internalization ability.⁷⁴ To enhance bombesin antagonist activity, it is possible to cleave the peptide bond in its active site (positions one to five or Tyr4-Lys3, and Tyr4-D-Phe12), introduce a nonpeptide between C-terminal and adjacent amino acid residues or replace them, incorporate D-amino acids, ester modifications, disulfide bridges and non-peptide bonds.⁷⁵ The selectivity of GRPR antagonists depends mostly on interactions with amino acids Thr297, Phe302, and Ser305 in the fourth extracellular domain of the receptor. All of these amino acid faces forward in the binding pocket, demonstrating that cation and hydrogen bonding interactions are essential to occur between antagonists and the three respective amino acids.^{76,77} Examples of GRPR antagonist are JMV594 [(D-Phe6, Stat13)Bn(6–14)], and JMV641 [D-Phe-Gln-Trp-Ala-Val-GlyHis-Leu(CHOH-CH₂)-(CH₂)₂-CH₃].⁷⁶

The bombesin sequence D-Tyr6-β-Ala11-Phe13-Nle14 is responsible for the rapid cell internalization in all three types of BBRs, being able to be used to develop targeting conjugates based in bombesin.^{32,41} For instance, bombesin conjugates through the coupling with MTX via a Lys spacer were developed, as well as DOX via a glutaric acid spacer and taxol using a PEG linker.^{11,39} Others conjugates of bombesin were prepared by loading them with CPT, PTX, mitochondria-disruptive peptides, marine toxins and siRNA.³⁵ Moreover, a large number of bombesin analogues were conjugated with radioisotopes like ^{99m}Tc, ¹¹¹In and ¹²⁵I and other contrast agents, and bivalent probes with ⁶⁴Cu as the cargo of the conjugate were prepared.⁷⁴

4.3.2. Angiopep 2

Angiopep 2 is a 19-amino acid peptide with the ability to cross the BBB, targeting specifically the low-density lipoprotein receptor-related protein 1 (LRP-1).¹⁴

The BBB is responsible for brain protection from potentially harmful substances circulating in the bloodstream and for maintaining the homeostatic environment of the central nervous system. This neurovascular unit is constituted by endothelial cells with extensive tight junctions, neurons, astrocytes and a contractile apparatus of smooth muscle cells and pericytes.^{89,91}

LRP-1 is a member of the large receptor family of low density lipoprotein (LDL). Like other receptors of this family, the modular structure of LRP-1 includes Cys-rich complement-type repeats, a cytoplasmic domain and a transmembrane domain. This receptor is highly expressed in astrocytes, smooth muscle cells, neurons and pericytes,

but is not as expressed in the endothelium.^{89,91} LRP-1 mediates the BBB transcytosis process for several ligands such as lactoferrin, thyroglobulin, α 2 macroglobulin and tissue-type plasminogen activator. In addition to being expressed on the surface of the BBB and controlling its permeability, LRP-1 is also present on a variety of tumors, mainly in high malignant glioma cells, participating in the cytoskeleton organization and in the adhesive complex turnover by modulating integrins functions in malignant cells.^{88,89,91} Thus, although BBB has low permeability for small-molecule drugs, angiopep 2 is capable to penetrate the BBB and target brain tumor cells, using the same receptor-mediated transporter.^{14,32,87,90}

The most promising drug conjugate based on angiopep 2 consists in a 19-amino-acid linear angiopeptin-2-PTX conjugate (GRN1005). This peptide drug conjugate is composed by three PTX molecules linked to the two Lys residues (positions 5 and 9) of angiopep 2 and to the N-terminal Thr, by cleavable ester linkers.^{32,35} GRN1005 crosses the BBB by LRP-1 receptor-mediated transcytosis and is distributed broadly throughout brain parenchyma (**Figure 19**).^{32,88} However, the exact molecular mechanism of angiopep transcytosis remains unknown.⁹⁰ Further studies showed that GRN1005 is not a substrate for P-glycoprotein-mediated drug efflux, improving anti-tumor efficacy.⁸⁸

Other angiopep 2 drug conjugates were created, including angiopep-2–doxorubicin, angiopep-2–dimethylglycine etoposide and angipep 2–trastuzumab (efficient against HER2-positive intracranial tumors).³⁵

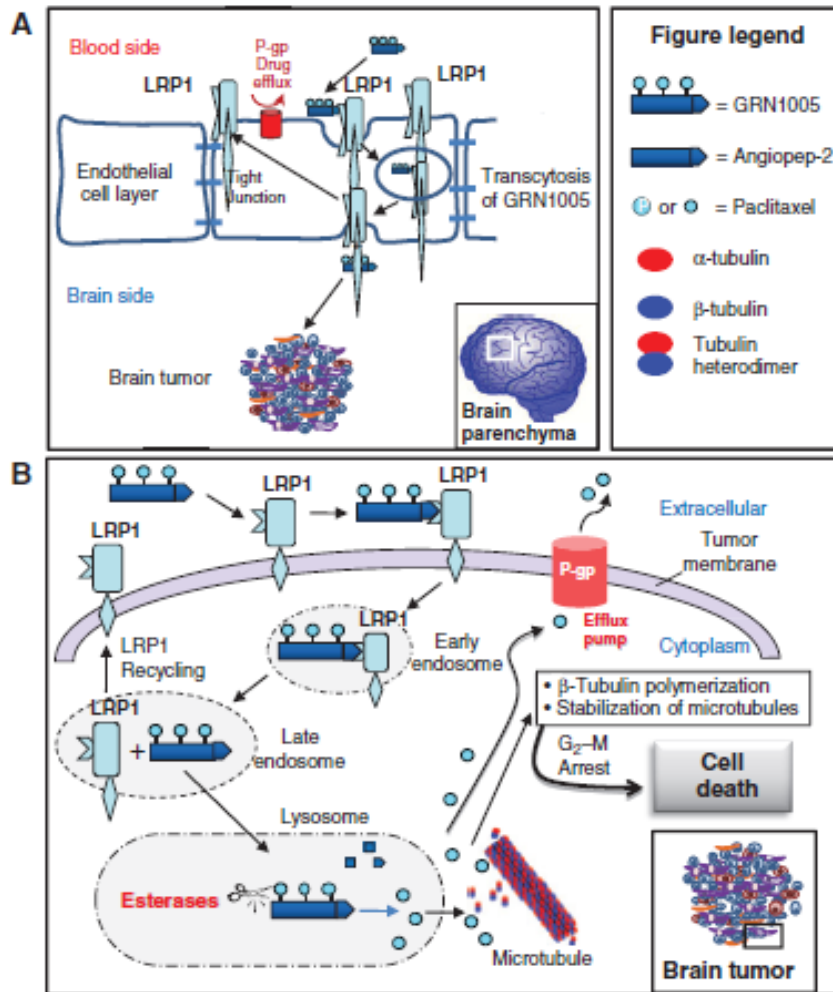


Figure 19: Internalization of GNR1005 through BBB (A) and tumor cell surface (B). Taken from ⁸⁸

4.4. Peptides Targeting the Tumor Microenvironment

4.4.1. CREKA

CREKA is a linear pentapeptide (Cys-Arg-Glu-Lys-Ala) identified by phage display libraries that targets fibrin-fibronectin complexes deposited in tumor vessel walls and tumor interstitial spaces, forming a meshwork of clotted plasma proteins detectable only in tumor tissues.^{14,15,78,81} Moreover, CREKA can induce tumor clotting, creating new binding sites in a self-amplifying effect. The exact binding site of the CREKA peptide is still unknown.

The bioactive conformation of CREKA is based on a turn-like structure where the charged groups of Glu, Lys and Arg form stable intermolecular interactions.⁸² Moreover, the sulfhydryl group of the Cys residue of CREKA is not required for binding interactions with its receptor, being the preferred site to conjugate the peptide with

other moieties.^{11,78} This knowledge promoted the design of CREKA-based peptides with enhanced features. For example, the CR(NMe)EKA has a N-metil derivate that increase tumor-homing response by protecting the peptide against proteolytic degradation (**Figure 20**).⁸² Moreover, in order to block the action of tumor-associated platelets on tumor metastatic progression, created liposomal nanoparticles bearing CREKA and ticagrelor (a platelet inhibitor) were created.⁸⁰

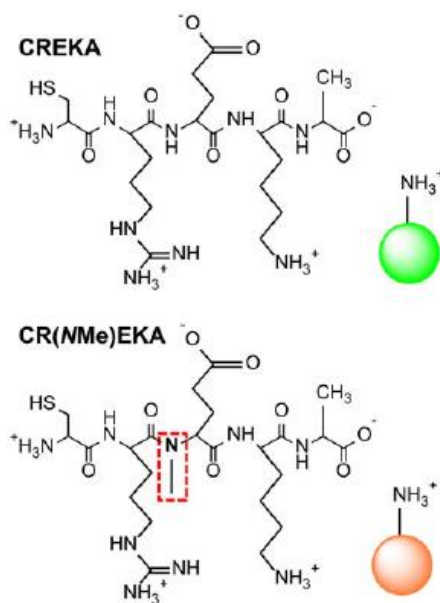


Figure 20: Chemical structures of CREKA and CR(N-Me)EKA. Taken from ⁸²

4.4.2. M2pep

The sequence YEQDPWGVKWWY, termed M2pep, identified using a phage display strategy, is a unique M2-selective peptide that specifically recognizes and internalizes M2 cells, including TAMs, having low affinity to others leukocytes.^{14,15,94}

Macrophages are phagocytic cells originated from circulating blood monocytes that extravasate into tissues and differentiate into macrophages with several functional phenotypes. Whereas M1 macrophages phenotype is stimulated by mediators, like interferon γ or lipopolysaccharide, resulting in a pro-inflammatory and microbial functional phenotype, M2 macrophages are stimulated by IL-4 and IL-13 in tissue remodeling and inflammation resolution cases.⁹⁴ TAMs are originated from circulating monocytes, differentiating within the tumor microenvironment, in M1 or M2 activated macrophages.^{93,95} Most of TAMs are M2-like macrophages and have the ability to produce immunosuppressive cytokines and have low antigen-presenting and co-stimulating capacity, facilitating tumor progression by suppressing the adaptive immune

response.^{93,94} Moreover, TAMs are responsible for immune evasion, metastasis, angiogenesis and matrix remodeling.^{14,92,93} While the increased density of TAMs is related with drug resistance, radiotherapy induces macrophage aggregation, creating radio-protective effects.⁹³

To enhance M2 macrophage-binding affinity of M2pep, a cyclic analog was build, the Cys-decafluorobiphenyl-Cys disulfide-cyclized M2pep c[DFBP-M2pep(RY)].⁴⁰ More recently, studies have shown that the divalent display of M2pep improves M2 macrophage-binding activity, while tetravalent display leads to the loss of selectivity of macrophages phenotype. Moreover, divalent and tetravalent displays of M2pep ([M2pep]₂-Biotin and [M2pep]₄-Biotin) exhibit M2 macrophage cytotoxicity. Further studies with M2pepKLA analogues ([M2pep]₂-[KLA] and [M2pep]₂-[KLA]₂) showed improvement of the cytotoxic activity (**Figure 21**).^{14,94,95} On the other hand, the optimization of the serum stability was achieved through “head-to-tail” cyclization, modifying linear M2pep(RY)Biotin with two flanking Cys, enabling disulfide cyclization, and by the replacement of the triLys spacer to D-Lys. Furthermore, N-terminal acetylation of M2pepBiotin gives protection from exolytic degradation.⁹⁶

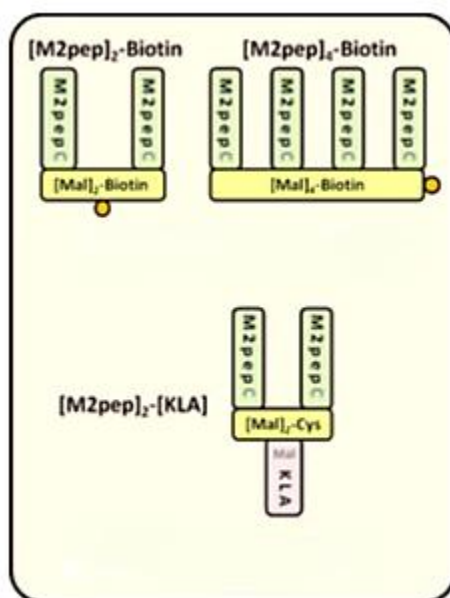


Figure 21: Schematic representation of divalent and tetravalent display of M2pep and the [M2pep]₂-[KLA] conjugate. Adapted.⁹⁵

5. Discussion

Through the analysis of the nine selected peptides, they can be divided in peptides that directly target tumor cells and peptides that target adjacent cells localized in their surrounding environment, including blood and lymphatic vessels. Targeting cells of the tumor medium prevents them of helping tumor growth, promoting an indirect death of tumor cells. Although this approach seems less effective for not directly attacking malignant cells, the access to the addresses is simpler since the peptide conjugated to the anti-tumor drug only has to reach the highly porous blood vessels of the tumor. In this regard, peptides with the ability to target tumor blood vessels show more targeting effects, reflecting a large number of studies. Furthermore, RGD and NGR peptides can be found in a wide number of clinical trials, which makes them the most suitable to integrate the first peptide-drug conjugate approved for clinical use.

Among peptides targeting tumor vasculature, RGD and NGR are those that present all the key-points of information clear and developed. Today, researchers are guiding their research to the development of new iRGD analogues and NGR double-targeting applications when converted to isoDGR. Regarding the others analyzed peptides in this targeting subtype, F3 contains exceptional internalizing properties and octreotide already has an optimized structure with excellent antagonistic capacity. However, both peptides require applicability studies in therapeutic conjugates.

Both lymphatic vessels and other elements of tumor microenvironment are still poorly understood due to the lack of available information about this subject, which has limited the development of PDCs with these peptides. Studies have not yet revealed that targeting the tumor microenvironment is enough to effectively eliminate cancer cells, whereby they are only considered as adjuvant therapy. This way, LyP-1 is a good candidate as adjuvant treatment of metastases. On the other hand, CREKA and M2pep are known to bind to specific elements of the tumor microenvironment, but their addresses are unknown, which makes difficult the increment of new binding optimization studies.

Although peptides homing tumor cells have more obstacles until reach their target, they may give additional features to PDCs. Angiopep 2 is able to cross the BBB, in addition to targeting brain tumors, which can offer significant therapeutic improvements over currently available treatments. Moreover, bombesin presents

antagonistic properties capable of duplicating the cytotoxic effect when integrating drug conjugates.

The potential utilization of tumor targeting peptides can be extended to the treatment of others diseases, since angiogenesis is not exclusive of cancer, occurring in others medical conditions like arthritis and retinopathy. Furthermore, there are others pathological clotting activity regions, where CREKA can act, and several chronic inflammation states where M2 macrophages are present, such as fibrosis, asthma and atherosclerosis.

6. Conclusion

In this work, nine peptides showed a significant contribution for the scientific advances in the development of new anti-tumor conjugates with targeting ability that may become a common reality in clinical practice: RGD, NGR, F3, Octreotide, LyP-1, Bombesin, Angiopep 2, CREKA and M2pep (**Table 3**). However, it is necessary to invest in continuous research, especially *in vivo* studies, to improve the specificity, safety, efficiency and to adequate the structural modifications of these peptides and the respective drug conjugates.

It is predictable that TTPs will reach significant clinical success and gradually evolve since, unlike Abs, the costs of production will not create a limitation to the use of this type of tumor therapy approach. In the near future, cancer patients will have access to more effective and less expensive tumor targeting therapies. Peptides-targeted drug conjugates will allow the development of more personalized tumor treatments, increasing the quality of patients' life.

Table 3: TTPs reviewed in this work.

Target	Name	Receptor	Sequence
Tumor Vasculature	RGD	$\alpha_v\beta_3/\alpha_v\beta_5$ integrins	RGD
	NGR	Aminopeptidase N (CD13)	NGR
	F3	Nucleolin	KDEPQRRSARLSAKPA PPKPEPKPKKAPAKK
	Octreotide	Somatostatin	f-c(CFwKTCT-ol)
Tumor Lymphatic Vessels	LyP-1	P32	CGNKRTRGC
Tumor Cells	Bombesin	Bombesin	H-pEQRLGNQWAVGHLM-NH ₂
	Angiopep 2	LRP-1	TFFYGGSRGKRNNFKTEEY
Tumor Microenvironment	CREKA	Not determined	CREKA
	M2pep	Not determined	YEQDPWGVKWWY

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8 - Annexes

Annex 1: Tumor Targeting Peptides

Table 4: Peptides with the ability to target tumor blood and lymphatic vessels and tumor microenvironment. Adapted.^{11,12,14,15}

Sequence	Name	Receptor	Target cell	Examples of peptide-drug and imaging probe conjugates
CDCRGDCFC	RGD-4C	$\alpha_v\beta_3/\alpha_v\beta_5$ integrins	Angiogenic blood vasculature	Conjugated to DOX, PTX, (KLAKLAK) ₂ , TNF α , IFN γ , IL-12
CNGRCVSGCAGRC	NGR	Aminopeptidase N (CD13)		
CRGDKGPDC	iRGD	$\alpha_v\beta_3/\alpha_v\beta_5$ integrins	Blood vessels and tumor cells	Conjugated to DOX, gemcitabine, KLA, tyrosine kinase inhibitors, trastuzumab
KDEPQRRSARLSAKP	F3	Nucleolin		
PPKPEPKPKKAPAKK	Esbp	E-selectin	Activated endothelial cells	^{99m} Tc-Esbp
DITWDQLWDLMK	F56	VEGFR-1	Angiogenic blood vasculature, stromal cells, TAMs and tumor cells	DHFR-F56
WHSDMEWYLLG	K237	VEGFR -2 KDR/Flk-1	Angiogenic blood vasculature	TPC-Ahx-A7R photosensitizer
HTMYHHYQHHL	A7R			
ATWLPPR		Neuropilin-1	Angiogenic blood vessels, breast, prostate, melanoma and other cancer cells	
CCKNTDSRCK	CREKA		Fibrin-fibronectin complexes deposited as a result of subtle clotting	CREKA-coated SPIONs and liposomes
ARQLELNERTCRC	TCP-1		Blood vessels of orthotopic colorectal cancer	TCP-1-GGD(KLAKLAK) ₂
CTPSPFSHC		Chondroitin sulfate proteoglycan, NG2	Blood-vessels associated pericytes	
TAASGVRSMH	KAA		Pancreatic tumor cells	
CKAAKNK	RSR		Angiogenic islet vessels	
CRSRKG	RGR		Angiogenic and premalignant, as well as tumorigenic blood vessels	
CRGRRST	LyP-1	P32	Tumor cells, lymphatic endothelium, hypoxic areas inside tumor cells	Conjugated to fluorochromes
CGNKRTRGC	LyP-2		Lymphatic vessels	
CNRRTKAGC	IF7	Annexin 1		
IFLLWQR	CTL1		Clotted plasma proteins	
CGLLIQNEC	CTL2			
CNAGESSKNC		MMP-2/9	ECM	
CRRHWGFEC	PEGA		Tumor blood vessels	
CTTHWGFTLC			Tumor neovasculature, heparin sulfate	
CPGPEGAGC			Endothelial VCAM-1 expressing cell	
CGKRK			Proteases in breast cancer cells	
VHSPNKK				
IAGEDGDEFG				

Developments in Tumor Targeting and Internalizing Peptides

YEQDPWGVKWWY	M2pep		M2-like TAMs
FYPSYHSTPQRP	DC3		
VTLTYEFAAGPRD	P-D2	CD11c/CD18	DCs
LSLERFLRCWSDAPA	PP1	SR-A	Macrophages

Table 5: Peptides targeting tumor cells. Adapted.^{7,11,12,14,32,41}

Sequence	Name	Receptor
YHWYGYTPQNV D(CVRAC)	GE11	EGFR
WHSDMEWWYLLG CSDSWHYWC	F56 P1	VEGFR-1 VEGFR-3
CRTIGPSVC SPRPRHTLRSL	B18	Tf-R
LTVSPWY SWELYYPLRANL		Her2 E- and N-cadherins
YCAREPPTRTFAYWG WHPWSYLWTQQA	EPPT1 RP-1	uMUC-1 CD44
RLVSYNGIIFLK VLWLKNR	A5G27 FP16	CD44v3 and CD44v6 FGF3
CTVALPGGYVRVC c-dGHCitGPQ-c	pep42 OA02	GRP78 Alpha 3-integrin
TAASGVRSMH CSNRDARRC	TAA	Chondroitin sulfate proteoglycan NG2
CSSRTMHHC LLADTTHHRPWT		
VRPMLQ TSPLNIHNGQKL	HN-1	Not determined
SMSIARL QHPSFI		
EDYELMDLLAYL CGNSNPKSC	FROP-1	
SVSVGMPKSPRP LTVSPWY		erbB2
pGlu-LYENKPRRPYIL RRPYIL	Neurotensin	Neurotensin
Glp-GPWLEEEEEAYGWMDf-NH ₂ H-pEQRLGNQWAVGHLM-NH ₂	Gastrin Bombesin	Gastrin Bombesin
HSDAVFTDNYTRLRQMAVKKYLSILN- NH ₂	VIP	VIPR
CPRECESIC GTRIIYDRKFLMECRNSPVT		Aminopeptidase A GnRH
CRKRLDRN KCCYSL		IL-4 ErbB-2
EDYELMDLLAYLK FCYWKCT		Somatostatin
H-AGCKNFFWKTFTSC-OH HAIYPRH	Somatostatin	Transferrin
KAPSGRMSIVKLNQLDPSHRISDRDYMG MDF-NH ₂	CCK-33	Cholecystokinin-B/ Gastrin receptor
DYMGWMDf-NH ₂ GWMDf-NH ₂	CCK-8 Gastrin	
GNLWATGHFM-NH ₂ EWPRPQIPP- NH ₂	Neuromedin B BPP	NMBR Bradykinin Receptor B2
CYFQNCPRG-NH ₂	Vasopressin	Arginine Vasopressin Receptor

Annex 2: Tumor Targeting Peptides-based Nanomedicines

Table 6: Peptide-conjugated nanomedicines for tumor targeting. Adapted.^{5,14,35}

Tumor Targeting peptide	Nanocarrier
RGD	Doxo-liposomes (RGD-LCL)
	pHPMA-RGDfKDTX and pHPMARGD4C
	PGA-PTX-E-[c(RGDfK) ₂]
	RGD-G3-poly(Llysine)-dendrimer[DOX]/siRNA complexes
	RGD-lipopeptide 1/p53 plasmid
	cRGDfK-PEG-PCL micelles loaded with DOX and SPIO
	cRGD decorated nanoparticles or micelles (constructed from PEG-PLGA, PEG-PCL, PEGPTMC, PEG-PGA, POE-PCL, PEG-PLA)
NGR	Liposomes (TVTDOX)
	Nanoparticles (NGRPEG-PLGA(DTX))
	NGR-modified (DTX)-loaded PEG-b-PLA polymeric micelles (NGR-PMDTX) NGR-modified DSPE-PEG micelles (NGR-M-PTX)
	Polymer-drug conjugates (pHPMA-(NGR)-D(KLAKLAK) ₂)
iRGD	iRGD-abraxane nanoparticles
	iRGD-PCL-b-PVP/PTX
Lyp-1	Lyp-1-abraxane
CREKA	CREKA-abraxane
Syp-1	Syp-1-DOXliposomes
Qa-based peptide analog of sLex	Polymer conjugate (pHPMA-Qa-FITC)
Esbp	Polymer-drug conjugates (pHPMA-(Esbp)-DOX, pHPMA-(Esbp)-KLAK
F56	
A7R	F56-viral nanoparticles (VNP)
	pHPMA-(F56)-DOX
	A7RC-PTX-LIPs
	A7R- lipoplexes
GE11	(D)A7R-DOX liposomes
	pHPMA-GE11-DOX
	GE11-DOXliposomes
G3-C12	GE11-PEG-PEI polyinosine/cytosine (polyIC) polyplexes
	pHPMA-(G3-C12)- FITC
A5G27	G3-C12-HPMA- DOX
	pHPMA-(A5G27)- FITC
	mA5G27F-PEG-b-PEI/siRNA
mA5G27F-PEG-b-PEI/siRNA	Polymeric micelles (DOX-AP-pH-PM)
P-D2	P-D2-decorated PLGA
DC3	pHPMA-(DC3)-FITC DC3- PEG-b-PEI/DNA polyplexes

Annex 3: RGD and NGR Peptides under Clinic Evaluation

Table 7: RGR peptides under clinic evaluation. Taken from ¹³

Condition	Status	Intervention	Phase	Year
Glioblastoma	Ongoing	Single intratumoral injection of Delta-24-RGD; Interferon-gamma	I	2014
Glioblastoma	Ongoing	18F-Galacto-RGD PET	I	2013
Glioblastoma; Multiforme; Recurrent Tumor	Ongoing	Delta-24-RGD Temozolomide	I	2013
Prostate Cancer	Recruiting	68Ga-NOTA-BBN-RGD	I	2016
Breast Cancer	Recruiting	68Ga-NOTA-BBN-RGD	I	2016
Lung Cancer	Recruiting	68Ga-NOTA-3PTATE-RGD	I	2016
Pathological Angiogenesis	Recruiting	68Ga-NODAGA-RGD PET/CT	I	2016
Brain Cancer; Brain Neoplasm; etc	Recruiting	Delta-24-RGD	II	2016
Diffuse Large B Cell Lymphoma	Recruiting	RGD K5 PET scan	II	2015
Bronchogenic Carcinoma; Breast Carcinoma; etc	Recruiting	18F-AI-NOTA-PRGD2 PET/CT	I	2015
Advanced Head and Neck Carcinoma	Recruiting	Radiopharmaceutical (Flotegatide (18F) or RGD (68Ga)	II	2014
Advanced Non-small Cell Lung Carcinoma	Recruiting	K5-RGD PET	II	2014
Non-seminomatous Germ Cell Tumors	Recruiting	K5-RGD PET	II	2014
Metastasis				
Solid tumors	Recruiting	18F FPPRGD2 PET/CT	I&II	2013
Lung Cancer; Lung Tuberculosis	Completed	68Ga-labeled peptides of dimer RGD	I&II	2015
Breast Cancer	Completed	Breast Cancer		2010
Brain Tumor; Recurring Glioblastoma	Completed	Delta-24-RGD adenovirus	I&II	2010
Metastatic Breast Cancer; Metastatic Colon/Rectum Cancer; etc	Completed	[F-18]RGD-K5	II	2009
Ovarian Cancer	Completed	Ad5.SSTR/TK-RGD; Ganciclovir (GCV)	I	2009
Sarcoma; Melanoma; etc	Completed	F-18 RGD-K5	I	2008
Brain Cancer	Completed	Delta-24-RGD-4C	I	2008
Ovarian Cancer	Completed	Ad5-delta24RGD	I	2007
High-grade Glioma; Lung Cancer; etc	Completed	Fluciclatide Injection - (AH111585 (F18))	II	2007
Kidney Neoplasm	Terminated	18F-Fluciclatide containing the RGD	II	2013
Carcinoma, Renal Cell	Unknown	18F-RGD-PET-CT and perfusion CT scans on 3 occasions	II	2011
Head and Neck Neoplasms	Unknown	[18F]RGD-K5	II	2011
Glioma	Unknown	68Ga-BNOTA-PRGD2	I	2013
Lung Cancer	Unknown	68Ga-BNOTA-PRGD2	I	2012

Table 8: NGR peptides under clinic evaluation. Taken from ¹³

Condition	Status	Intervention	Phase	Year
Solid tumors	Ongoing	NGR-hTNF α	I	2007
Small cell lung cancer	Ongoing	NGR-hTNF: iv q3W 0.8 mcg/sqm NGR-hTNF; Doxorubicin: iv q3W 75 mg/sqm doxorubicin 60 min after NGR-hTNF infusion	II	2007
Ovarian cancer	Ongoing	NGR-hTNF α ; Pegylated liposomal-doxorubicin; Doxorubicin	II	2011
Ovarian cancer	Ongoing	NGR-hTNF: 0.8 mcg/m ² as 60-min intravenous infusion every week until confirmed evidence of disease progression or unacceptable toxicity occurs; Doxorubicin: 60 mg/m ² every 3 weeks, until cumulative dose of 550 mg/m ²	II	2011
Non-small cell lung cancer	Ongoing	NGR-hTNF α ; Cisplatin; Gemcitabine; Pemetrexed	II	2010
Soft-tissue sarcoma	Ongoing	NGR-hTNF α ; Doxorubicin	II	2007
Malignant pleural mesothelioma	Ongoing	NGR-hTNF α plus best investigator choice	III	2009
Malignant pleural mesothelioma	Recruiting	NGR-hTNF α	II	2007
Solid tumors	Completed	NGR-hTNF α : iv q3W escalating dose up 1.6 mcg/sqm	I	2007
Solid tumors	Completed	NGR-hTNF: iv q3W escalating dose NGR-hTNF up to 1.6 mcg/sqm; Cisplatin: iv q3W 80 mg/sqm 30 min after NGR-hTNF infusion for a maximum of six cycles	I	2007
Solid tumors	Completed	NGR-hTNF: 0.2, 0.4, 0.8 and 1.6 mg/m ² as 60-min intravenous infusion every 3 weeks; Doxorubicin: 75 mg/m ² intravenous infusion over 15 min (starting 1 h after the end of NGR-hTNF infusion)	I	2007
Colorectal Cancer, Head and Neck Cancer, etc	Completed	CNGRC peptide-TNF alpha conjugate	I	2006
Colorectal cancer	Completed	NGR-hTNF α : iv q3W or q1W NGR-hTNF 0.8 mg/m ²	II	2009
Colorectal cancer	Completed	NGR-hTNF α ; Oxaliplatin; Capecitabine	II	2008
Malignant pleural mesothelioma	Completed	NGR-hTNF α : iv q3W or q1W 0.8 mcg/sqm NGR-hTNF	II	2004
Hepatocellular carcinoma	Completed	NGR-hTNF α : iv q3W or q1W 0.8 mcg/sqm NGR-hTNF	II	2007