# Absolute abundance estimates from shallow water baited underwater camera surveys; a stochastic modelling approach tested against field data 

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#### Abstract

Baited underwater cameras are becoming a popular tool to monitor fish and invertebrate populations within protected and inshore environments where trawl surveys are unsuitable. Modelling the arrival times of deep-sea grenadiers using an inverse square relationship has enabled abundance estimates, comparable to those from bottom trawl surveys, to be gathered from deep-sea baited camera surveys. Baited underwater camera systems in the shallow water environments are however, currently limited to relative comparisons of assemblages based on simple metrics such as $\operatorname{Max}_{\mathrm{N}}$ (maximum number of fish seen at any one time). This study describes a stochastic simulation approach used to model the behaviour of fish and invertebrates around a BUC system to enable absolute abundance estimates to be generated from arrival patterns. Species-specific models were developed for the tropical reef fishes the black tip grouper (Epinephelus fasciatus) and moray eel (Gymnothorax spp.) and the Antarctic scavengers; the asteroid (Odontaster validus) and the nemertean worm (Parbolasia corrugatus). A sensitivity analysis explored the impact of input parameters on the arrival patterns $\left(\operatorname{Max}_{\mathrm{N}}\right.$, time to the arrival of the first individual and the time to reach $\mathrm{Max}_{\mathrm{N}}$ ) for each species generated by the model. Sensitivity analysis showed a particularly strong link between $\mathrm{Max}_{\mathrm{N}}$ and abundance indicating that this model could be used to generate absolute abundances from existing or future $\mathrm{Max}_{\mathrm{N}}$ data. It in effect allows the slope of the $\mathrm{Max}_{\mathrm{N}}$ vs. abundance relationship to be estimated. Arrival patterns generated by each model were used to estimate population density for the focal species and these estimates were compared to data from underwater visual census transects. Using a BlandAltman analysis, baited underwater camera data processed using this model were shown to generate absolute abundance estimates that were comparable to underwater visual census data.


## Highlights:

- Modelling the behaviour of fish and invertebrates around a baited camera system
- Models developed for tropical fish and Antarctic invertebrates
- Abundance estimates calculated and compared to data from visual census transects
- Comparable abundance estimates generated by the model and transects

Keywords: baited underwater cameras; modeling; fish and invertebrate surveys; underwater visual census

## Abbreviations:

BUC: Baited underwater camera
Max $_{N}$ : Maximum number of individuals, of the same species, appearing on the field of view in any one frame over the whole deployment
$\mathrm{T}_{\text {arrival }}$ : Time to the arrival of the first individual from each species
$\mathrm{T}_{\operatorname{maxN}}$ : Time to the maximum number of individuals observed at one time
UVC: Underwater visual census

## 1. Introduction

Abundance estimates of marine populations, that are both accurate, close to the true abundance, and precise, repeatable under the same conditions, are important to understand changes in marine populations or communities (Farnsworth et al., 2007) and to help achieve sustainable management and effective conservation objectives (Collins et al., 2002). For marine fish and invertebrate populations the majority of this data has been collected using trawl surveys (Fitzpatrick et al., 2012; Johnson et al., 2012), which are difficult in abyssal environments and unsuitable in marine protected areas (Bailey et al., 2007). Baited underwater camera (BUC) systems have therefore been used in many studies to gather data on deep-sea scavenging fauna (Farnsworth et al., 2007) and fish assemblages in protected areas (Willis and Babcock, 2000; McLean et al., 2010). However, to use BUC data to produce absolute abundance estimates of fish and invertebrate populations requires a detailed understanding of the physical and biological parameters involved in the process of animals detecting and following the bait plume to the camera (Priede et al., 1994; Bailey et al., 2007).

Bait plume dispersal from a point source, its detection by fish or invertebrates and their arrival at the source, is influenced by a number of environmental and biological factors (Collins et al., 2002; Stoner, 2004). The odour from the bait disperses as a plume into the surrounding water on currents (Reidenbach and Koehl, 2011). The velocity and direction of currents will affect the length and lateral dispersal of the plume as well as its dispersal direction (Bailey and Priede, 2002; Dorman et al., 2012). The dispersal of odour plumes is also affected by turbulence within the aquatic environment (Meager and Batty, 2007), the topography over which it travels (Collins et al., 1999; Collins et al., 2002; Reidenbach and Koehl, 2011) and the characteristics and persistence of the bait (Bailey and Priede, 2002; Stoner, 2004). Fish and invertebrates have evolved olfactory organs with chemosensory abilities that allow them to detect odour plumes and follow them to their source (Reidenbach and Koehl, 2011). The area within the odour plume where the odour concentration is above the threshold which organisms can detect is known as the 'active space' (Sigler, 2000; Stoner, 2004). The probability of the fish entering the active space of the bait plume will be dependent on their search behaviours (Dorman et al., 2012), including their swimming speed and position in the water when foraging (Stoner, 2004), as well as the abundance and distribution of the
population (Armstrong et al., 1992). Once the plume has been detected, the fish will decide whether to follow it based on the feeding motivation that the bait provides (Dorman et al., 2012). The time that individuals remain at the bait will be determined by the availability of food within the environment (Charnov, 1976) as well as the competition and interactions with other scavengers at the bait (Armstrong et al., 1992; Bailey and Priede, 2002; Dunlop et al., 2014).

The process of bait plume detection, attraction and arrival of the deep sea grenadier Coryphaenoides armatus at a BUC was modelled using an inverse square relationship:
$n=c / t_{\text {arr }}{ }^{2}$
where $n$ is the number of fish per square kilometre and $c$ is a constant, dependent upon the current velocity and through water swimming speed of the fish towards the BUC system (Priede et al., 1990; Priede and Bagley, 2000). $\mathrm{t}_{\text {arr }}$ represents the time elapsed between the beginning of the camera deployment and the arrival of the first fish. The model was developed by Priede et al., (1990) to allow scavenger density to be estimated from their arrival rates at the BUC in conjunction with information on the odour plume spreading characteristics, current velocities and fish swimming speed. The staying time of deep-sea grenadiers at the BUC can be estimated using the relationship:

$$
N_{\beta}=\frac{\alpha_{0}}{x}\left(1-e^{-\beta x}\right)
$$

where $N_{\beta}$ is the maximum number of fish present after a certain period of time, $\alpha_{0}$ the initial rate of fish arrival at time zero, $e$ the exponential constant and $x$ a constant representing the decay of the odour plume from dilution and bait consumption (Priede et al., 1990). Arrival rates are of interest as a bait placed amongst an abundant scavenger population has a greater chance of being reached by an individual quickly (Bassett and Montgomery, 2011). The arrival times of deep-sea grenadiers at a BUC in two sites in the North Atlantic were modelled in the above manner to produce estimates of abundance which were comparable to those from bottom trawl surveys from approximately the same area and time (Armstrong et al., 1992; Priede and Merrett, 1996). However, when applied to fish arrival times on the Mid-Atlantic Ridge there was no correlation between BUC generated abundances and those estimated from trawls (Bailey et al., 2007).

The use of BUC systems in shallow waters have enabled relative comparisons of both fish and invertebrate assemblages in the tropical (McLean et al., 2010; Moore et al., 2010), temperate (Willis et al., 2003) and the Antarctic environments (Smale et al., 2007) between areas of different protection status (Willis and Babcock, 2000; Westera et al., 2003), habitat type (Moore et al., 2010) and disturbance pressure (Smale et al., 2007). The majority of studies have used the maximum number of individuals, of the same species, appearing in the field of view in any one frame over the whole deployment ( $\mathrm{Max}_{\mathrm{N}}$ ) as an index of relative abundance (Willis and Babcock, 2000; Stoner et al., 2008). Max ${ }_{N}$ avoids the repeated recording of individuals that leave and re-enter the camera field of view and usually less than the count of all animals
visiting the bait (McLean et al., 2010; Harvey et al., 2012). Some surveys have also used the time to the arrival of the first individual from each species ( $\mathrm{t}_{\text {arrival }}$ ) and time to the maximum number of individuals observed at one time $\left(\mathrm{t}_{\operatorname{maxN}}\right)$ (Willis and Babcock, 2000; Jones et al., 2003). In the shallow water environment however, the development of models of the process of fish or invertebrate arrival at BUCs has been limited (Stoner et al., 2008; Langlois et al., 2012). Heagney et al., (2007) investigated whether abyssal scavenger arrival models could be applied to shallow mid-water baited underwater video data. Existing models appropriate for deep-sea BUC studies with long soak times and where scavengers approached more slowly, were found unsuitable for shallow water BUC studies with much shorter soak times and which attract many fast moving species (Heagney et al., 2007). Rapid arrival patterns of shallow water fish result in overestimated abundance due to the inverse square law of the abyssal model (King et al., 2006; Stobart et al., 2007). Compared to the shallow water environment, currents in the abyss are relatively constant, so an assumption of a constant current speed and direction is more suitable (Heagney et al., 2007; King et al., 2008). The assumptions of deep-sea models also cannot be applied to describe the foraging behaviours of shallow water fish species, which also use sight, as well as chemoreception, to find food (Ellis and DeMartini, 1995; Stobart et al., 2007). The time related metrics used in the deep-sea such as, $\mathrm{t}_{\text {arrival }}$ and $\mathrm{t}_{\text {maxN }}$, have not correlated well with other surveys methods in some shallow water BUC surveys (Stoner et al., 2008; Willis and Babcock, 2000).

The area sampled by the active space of the odour plume is largely unknown in shallow BUC surveys. Concerns have been raised regarding the effect of localised environmental conditions, such as topography and current conditions, on plume dynamics making it difficult to make comparisons between areas (Taylor et al., 2013; Watson et al., 2009). Surveys assume that a comparable area is sampled by each deployment, however, this will often be untrue if current conditions vary (Heagney et al., 2007). The importance of the currents on the dynamics of bait plume dispersal and subsequent fish arrival patterns have been highlighted in several studies in the mid water (Heagney et al., 2007) and demersal environments (Dorman et al., 2012). The unknown sample area of shallow water BUC surveys also makes it difficult to make comparisons with abundance estimates from other survey methods. Several studies have investigated the differences in fish and invertebrate studies recorded by BUC and UVC surveys (Langlois, 2006; Watson et al., 2010), however, conclusions regarding comparisons have been difficult as the area sampled cannot be directly compared (Langlois et al., 2010).

A model to determine the absolute measures of shallow water fish or invertebrate abundance from arrival patterns at a BUC would involve developing an area based bait dispersion model using in-situ measurements of current speed and direction (Heagney et al., 2007). The mechanistic models outlined by Priede et al., (1990) to estimate the abundance of deep-sea demersal fish from first arrival times are deterministic. However, the arrival rate of fish is stochastically related to population abundance and the factors governing aspects of shallow water fish movement are often assumed to be well represented by random distribution (Farnsworth et al., 2007). This means it is important to include stochastic elements to mechanistic models. The physical factors, current distribution and velocity, observed around the camera system also have a random distribution within a particular range. Therefore it is important to introduce this
random aspect into models to describe fish attraction and arrival at a BUC system. Stochastic models that incorporate both the predictable and random aspects of a process, are increasingly being used to build our understanding of complex natural ecosystems (Brown and Kulasiri, 1996). Farnsworth et al., (2007) also modelled the arrival process of deep-sea demersal scavengers at the BUC using the addition of stochastic elements to deterministic models. Farnsworth's (2007) models unfortunately did not include a mechanism to reverse the process and calculate abundances from arrival patterns. The models also required a very large number of assumptions and parameters, making them difficult to implement for many BUC users.

The primary objective of the present study was to develop a stochastic modelling approach to enable the estimation of the absolute abundance of fish and invertebrates using arrival data collected using a shallow water BUC system. This involved the development of species-specific models for two fish and two invertebrate species observed in tropical and Antarctic BUC surveys. A global sensitivity analysis was used to determine the impact of model parameters on the arrival pattern produced by the model. A secondary objective, following the development of an effective modelling methodology, was to demonstrate how absolute abundance estimates can be generated from BUC data using the methodology. The achievement of this objective was assessed by comparing the model absolute abundance outputs to those from corresponding underwater visual census (UVC) transects. It was hypothesised that 1 ), the sensitivity analysis would show which model variables have an effect upon the arrival pattern of fish or invertebrates at the BUC and what aspects of the arrival pattern variable are affected the most (i.e. $\operatorname{Max}_{\mathrm{N}}, \mathrm{t}_{\text {arrival }}$ and $\mathrm{t}_{\operatorname{maxN}}$ ) and 2), that the modelling methodology would generate absolute abundance estimate that were comparable to those from corresponding UVC surveys.

## 2. Materials and Methods

### 2.1. Model outline

The simulation was built in MATLAB (R2010b) using the movement of an individual fish around a BUC system within a designated area. A bait plume was plotted and the area covered ( $B_{a}, \mathrm{~m}^{2}$ ) was described as a sector of a circle, using the three equations below. The length of the plume $\left(L_{p l}, \mathrm{~m}\right)$ was calculated using a radius described as the mean current speed $\left(V_{w}, \mathrm{~ms}^{-1}\right)$ recorded throughout the deployment multiplied by the simulation time ( $T$, seconds). The plume therefore expanded with every time step of the simulation. The plume angle ( $P l_{\theta}$, radians) was calculated from the inverse tangent of the diffusional velocity $\left(B_{y}, \mathrm{~ms}^{-1}\right)$, divided by the current speed $\left(V_{w}, \mathrm{~ms}^{-1}\right)$. The relationship between these model parameters is described in the equations:

$$
\begin{gathered}
L_{p l}=V_{w} T \\
P l_{\theta}=2 \tan ^{-1}\left(\frac{B_{y}}{V_{w}}\right) \\
B_{a}=\left(\frac{\theta}{2}\right) L_{p l}^{2}
\end{gathered}
$$

Simulations depict the movement of a population of a fixed abundance within a defined area (A, $\mathrm{m}^{2}$ ). Prior to detection of the bait plume fish move at a cruising speed $\left(V_{c r}, \mathrm{~ms}^{-1}\right)$ or are stationary, and turned a random
number of times ( $T_{r}$ ) within a set time period known as the turning interval ( $I n t_{t r}$, seconds). The direction within which the fish travels after each turn ( $D_{r}$, radians) was randomly selected (independently for each individual).

$$
\begin{gathered}
D_{r}=\operatorname{rand}(0,360) \\
\operatorname{Int}_{t r}=\operatorname{rand}\left(0, T_{r}\right) * T_{r}
\end{gathered}
$$

The starting point $\left(P_{s t},(x, y)\right)$ was selected (again independently for each individual) from a random position within the simulation area $\left(\mathrm{A}, \mathrm{m}^{-2}\right)$ using the formula below:

$$
\left(P_{s t},(x, y)\right)=\operatorname{rand}\left(-\frac{A}{2}, \frac{A}{2}\right)
$$

The distance travelled per time step $\left(D_{S}, \mathrm{~m}\right)$ was calculated by dividing the cruising speed by the time resolution ( $T_{r}$, seconds). Distance travelled in the x and y axis $\left(D_{s}(x, y)\right)$ was found by multiplying the cruise speed divided by the simulation time resolution (length of the time-step used in simulations) and multiplying this by sine and cosine of the direction ( $D_{r}$, radians):

$$
\begin{gathered}
D_{s}(x)=\frac{V_{c r}}{T_{r}} \sin \left(D_{r}\right) \\
D_{s}(y)=\frac{V_{c r}}{T_{r}} \cos \left(D_{r}\right)
\end{gathered}
$$

The distance to the camera $\left(D_{c m}(x, y)\right)$ was calculated by taking the square root of the distance travelled in the x and y axis:

$$
D_{c m}(x, y)=\sqrt{D_{s}(x, y)^{2}}
$$

When the distance to the camera $\left(D_{c m}(x, y)\right)$ is less than the radius associated with the circular bait area $\left(B_{a}, \mathrm{~m}^{2}\right)$ the fish is considered to have encountered the bait plume area. On encounter the fish turns into an approach angle $\operatorname{app}(\theta)$ calculated using:

$$
\operatorname{app}(\theta)=\left(180,360,0,-180 \tan ^{-1} D_{s}\left(\frac{x}{y}\right)\right)
$$

(the angle used in this equation depends upon the position on the fish when the bait plume is encountered). This change in direction causes the fish to swim directly upstream towards the bait at a through-water approach speed up the plume towards the camera $\left(V_{f s a}, \mathrm{~ms}^{-1}\right)$. This speed is faster than the cruising swimming speed and was calculated from observation of fish max swimming speed in previous published studies. Current speed ( $V_{w}, \mathrm{~ms}^{-1}$ ) is subtracted to account for the fish swimming upstream against the current. Once in the bait plume the distance travelled towards the camera and its relation to the camera position is recalculated using the through-water approach speed $\left(V_{f s a}, \mathrm{~ms}^{-1}\right)$ :

$$
\begin{gathered}
D_{s}(x)=\frac{V_{f s a}}{T_{r}} \sin \left(D_{r}\right) \\
D_{s}(y)=\frac{V_{f s a}}{T_{r}} \cos \left(D_{r}\right)
\end{gathered}
$$

Upon reaching the bait the individual will remain there for a "staying time" ( $S_{t}$, seconds) found by taking a random time between a pre-determined interval. This was multiplied by the time resolution ( $T_{r}$, seconds) of the simulation:

$$
S_{t}=\operatorname{rand}\left(\left[1800, T_{r}\right]\right)
$$

After remaining at the camera for the staying time the fish is removed from the simulation as it is assumed to have reached satiation or decided to forage elsewhere. Simulations run for 60 or 90 minutes and record the total number of fish, or invertebrates, present at the bait every 30 seconds, the same interval is used in the in-situ BUC studies. For the invertebrates studied here staying time was set till the simulation end. The model is depicted in as a diagram in Fig. 1.

### 2.2. General assumptions

Fish or invertebrates are assumed to act independently of each other at all stages of the simulation and to always react to the bait plume on encounter. The bait plume was always spread from the origin of the coordinate system used in the simulations and assumed to disperse in a single direction. The present model assumes a constant plume concentration and represents a framework that can be combined with fluid dynamics models of bait plume dispersal from a point source in the future to enable the dilution of the plume concentration and changes in current direction to be incorporated into the models.

Simulations were developed for four species; the grouper Epinephelus fasciatus and moray eels of the genus Gymnothorax spp. recorded in the tropical Gulf of Aqaba and the Antarctic scavenging invertebrates Odontaster validus and Parbolasia corrugatus. The BUC system consisted of a digital stills camera (SeaLife DC800 or DC1000) enclosed in an underwater housing. No additional light was required for work in the Gulf of Aqaba, but in Antarctica the camera was synchronized, via optical cables, with two variablepower digital slave strobe light units (Epoque ES-23DS). The camera was placed in time lapse mode (30 s intervals). The camera equipment was supported on an L-shaped frame of aluminium tubing. A u-shaped bracket holding the camera was bolted to the vertical element of the frame and angled downwards at $60^{\circ}$ to view the mesh bait bag attached to the far end of a horizontal pole. 200 g of either chopped fish (Sparus aurata and Dicentrarchus labrax ) in the Gulf of Aqaba or chopped Antarctic invertebrates (Ophionotus victoriae, O. validus, Sterechinus neumayeri and Laternula elliptica) were used as bait. The system was deployed from a boat and lowered to the seabed or placed by a SCUBA diver. A ballast weight (c10kg) held the camera system to the seabed and it was held upright in the water column by two small mid-water buoys. At the end of deployments the camera system was recovered either by hauling on a recovery line or by attachment and inflation of a lifting bag by SCUBA divers.

Data on swimming or crawling speeds, the turning frequency and aspects of the foraging behaviours for each species were determined from published studies (Fulton, 2007; D'Aout and Aerts, 1999; Clarke and Prothero-Thomas, 1997; Kidawa, 2001; Bshary et al., 2006) (Table 1). Estimations of staying time were based on observation of individuals in BUC deployments. For the tropical species it was difficult to identify individuals to calculate their staying time at the bait and estimations were taken from observation of the number of consecutive images an individual of that species was observed in. Current velocity was recorded during Antarctic deployments using a Nortek Aquadopp Acoustic Doppler current meter (Aquadopp Current Meter, Nortek, USA) while for the Gulf of Aqaba data an Acoustic Doppler Current Profiler between 10 m and 1 km from BUC deployments was used. Current meter measurements provided the current ranges within which the simulation could operate.

All BUC deployments had a matching underwater visual census (UVC) transect at the same location and depth making up on station. In the Gulf of Aqaba an area of $100 \mathrm{~m}^{2}$ was swum once ( $50 \times 2 \mathrm{~m}$ transect) and the numbers of E. fasciatus and Gymnothorax. spp. were recorded on a slate ( 32 stations total, eight at each at 5, 10, 15 and 20 m ). In Antarctica the density of $O$. validus and $P$. corrugatus was recorded from analysis of images from a $25 \times 0.5 \mathrm{~m}$ UVC transect of continuous stills images ( 18 stations total, six each at 5,10 and 25 m ).

The ranges of input parameters for each model are described in Table 1. The current speeds observed during the BUC deployments in both the Antarctic and the Gulf of Aqaba were approximately comparable to the current speeds measured in the deep-sea environment by Sainte-Marie and Hargrave, (1987).
Therefore, due to the lack of measurements of the diffusional velocities the same velocity, $10^{-3} \mathrm{~m} \mathrm{~s}^{-1}$, used to model the arrival of scavengers at a baited camera by Sainte-Marie and Hargrave, (1987) was used.

Moray eels of the genus Gymnothorax and blacktip groupers (E. fasciatus) are ambush predators highly associated with rocky reefs and crevices and will defend a small territory (Gibran, 2007). Therefore in simulations of Gymnothorax. spp. and E. fasciatus movement around the BUC system individuals were relatively slow moving prior to the detection of the bait plume. Antarctic invertebrate scavengers are slow moving compared to the tropical fish therefore BUC deployments in the shallow water Antarctic environment lasted for 1.5 h . The invertebrates also crawl along the seabed so current velocity was not subtracted from the approach velocity. Both Antarctic scavengers remained stationary prior to the detection of an odour plume and on reaching the bait scavengers remained there till the end of the simulation as observed in BUC deployments.

### 2.3. Data analysis

Models generated an arrival patterns for fish or invertebrates at the bait based on a predicted number present every 30 s , to produce a dataset in the same form as that from in-situ BUC deployments. $\mathrm{Max}_{\mathrm{N}}$, $t_{\text {arrival }}$ and $t_{\text {maxN }}$ were used to describe the arrival pattern of fish or invertebrates at the BUC. This sensitivity analysis enabled the dependence of the fish or invertebrate arrival pattern output by the model on input parameters to be determined and was used to test hypothesis one. A global sensitivity analysis was
performed on each species-specific model to determine the impact of the input parameters; population abundance, current speed, diffusional velocity, swimming speed before contact with the odour plume, approach speed and staying time (Table. 1). Each input parameter was set to be randomly selected from the full range of potential values and each of the four models was run 300 times to ensure that the full range of potential input parameters was considered. This was checked by plotting a histogram of the distribution of the input parameters and was also used to ensure that the range of input values had a random distribution. Both the marginal and bivariate simulated factor distributions were explored to ensure that coverage of the factor space was extensive (Saltelli, 2000).

A stepwise regression was performed in R (version 3.0.2, The R Development Core Team, 2013) to examine the relationship between the input parameters and the model output abundance indices; $\mathrm{Max}_{\mathrm{N}}$, $t_{\text {arrival }}$ and $t_{\text {maxN }}$. The relationship between any input parameter identified as having a significant effect on $\operatorname{Max}_{\mathrm{N}}, \mathrm{t}_{\text {arrival }}$ and $\mathrm{t}_{\text {maxN }}$ was plotted in a scatter plot. The relationship between the model parameters and the BUC abundance indices were unknown as this early stage of model development and the stepwise regression was used as a tool to explore these relationships. The analysis of the influence of model input parameters on the resultant fish or invertebrate arrival pattern highlighted which parameters were important to calibrate with in-situ measurements.

### 2.4. Producing absolute abundance estimates from BUC data

Any parameters with a significant effect were parameterised using an in-situ measurement of this variable where available. For example, if current speed had a significant impact on the $\operatorname{Max}_{N}$ then the current speed from the in-situ BUC deployment providing the camera data was used to produce an abundance estimate was used as a model input. Those identified as having no significant impact on the model output were set to be selected randomly from a range of suitable values for that measure. However, for some parameters an in-situ measurement was not available and values within the models had to remain as the estimates ranges. These parameters were highlighted as those requiring future measurement to improve the accuracy of the model outputs.

To produce absolute abundance estimates using the modelling methodology a suitable range of estimated population abundances must be first input into the model. In practice these estimates could be derived from previous surveys using other methods, literature for similar areas or be best guesses. In the case of this validation exercise corresponding UVC surveys from the same position and approximately the same time as the BUC deployments were used to find a suitable abundance range for the tropical and Antarctic models. Each single population abundance input into the model produced a BUC arrival pattern. For example, if an abundance range of 1-100 individuals was used 99 arrival patterns would be produced. The arrival patterns produced by the model were compared to the arrival patterns produced by the corresponding BUC survey. The R-squared value of the slope fitted to the arrival curve of individuals at the camera with time was used to find a match between model and BUC arrival patterns. Once a match was found the population abundance input into the model to produce that arrival pattern is recorded as the model's best estimate of the
absolute abundance of the fish or invertebrate population surveyed by the BUC system. This process is illustrated in Fig. 2 where the arrival pattern from five model runs of the model of E. fasciatus movement around the BUC can be compared to that of the in-situ BUC arrival pattern.

The absolute abundance estimate produced using the model methodology and field BUC data were compared to those generated by corresponding UVC surveys to validate the ability of the model to produce accurate abundance estimates. Models describing the movement of the two tropical fish species and the two Antarctic invertebrate scavengers in relation to the BUC system were validated using transect data. BUC absolute abundance estimates were compared to those from the corresponding UVC surveys using a BlandAltman analysis (Bland and Altman, 1986). A Bland-Altman analysis is used to compare two methods of measurement, usually a new method with an established one (Bland and Altman, 1986). In this study the UVC represents the established method for measuring fish and invertebrate absolute abundance and the BUC the new method. The Bland-Altman plots show the mean difference between the two corresponding measurements from both methods, known as 'the bias', and the $95 \%$ limits of agreement as $+/-1.95$ SD of the mean difference. The plot enables visual judgement of the agreement between the measurements and the smaller the range between the measurements the better the match (Bland and Altman, 1986; Bland and Altman, 1995). An analysis showing no significant systematic bias between the two methods would show the majority of the data points within the confidence limits and that points would have a symmetrical around zero. A Bland and Altman analysis was performed in the R package 'MethComp' and a Bland-Altman plot and measures of the test bias test were produced to compare the measurements of absolute abundance using the UVC and tropical and Antarctic BUC models (Fig. 3).

## 3. Results

### 3.1. Sensitivity analysis

The input parameters (abundance, current speed, approach speed, cruising speed, diffusional velocity and staying time) produced by 300 runs of the 4 models were plotted in frequency histograms and their distribution was random and encompassed the full range of potential input parameters. Sensitivity analysis revealed that the model input parameters explained a large proportion of the variability in the $\mathrm{Max}_{\mathrm{N}}$ output of the 4 models. Input parameters explained less of the variability in the time-based metrics ( $\mathrm{t}_{\text {arrival }}$ and $\left.t_{\operatorname{maxN}}\right)$. Abundance was the model input parameter that had the greatest impact on the $\mathrm{Max}_{\mathrm{N}}, \mathrm{t}_{\text {arrival }}$ and $\mathrm{t}_{\operatorname{maxN}}$ outputs from the model for all 4 species.

For both tropical models the parameter population abundance explained a large proportion of the variability in the $\mathrm{Max}_{\mathrm{N}}$ output; E. fasciatus $(\mathrm{y}=0.73 \mathrm{x}+0.71$; R-sq (adj) $=91.74 ; P<0.0001$ ) and for Gymnothorax spp. (y $=0.57 \mathrm{x}+0.49 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=97.99 ; P<0.0001)$. Input parameters explained less of the variability in the $\mathrm{t}_{\text {arrival }}$ of tropical fish at the bait. Population abundance had a small but significant effect on E. fasciatus $(\mathrm{y}=61.81 \mathrm{x}+217.89 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=18.16 ; P<0.0001)$ and Gymnothorax. spp. $\mathrm{t}_{\text {arrival }}(\mathrm{y}$ $=-54.21 \mathrm{x}+83.89 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=30.17 ; P<0.0001)$. Current speed also had a significant impact on

Gymnothorax. spp. $\mathrm{t}_{\text {arrival }}(\mathrm{y}=135.45 \mathrm{x}+42.94 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=1.47 ; P=0.02)$. Current speed explained $1.8 \%$ of the Gymnothorax. spp. $\mathrm{t}_{\operatorname{maxN}}(\mathrm{y}=315.57 \mathrm{x}+191.45 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=1.8 ; P=0.018)$ and population abundance had a significant impact on $E$. fasciatus $\mathrm{t}_{\operatorname{maxN}}(\mathrm{y}=1792.8 \mathrm{x}+6614.5 ; \mathrm{R}$-sq $(\operatorname{adj})=6.71 \% ; P<$ 0.0001 ). Staying time had no effect upon indices for both tropical models.

Only population abundance input into models of the Antarctic asteroid $O$. validus movement around the BUC explained a significant proportion of the $\mathrm{Max}_{\mathrm{N}}$ values generated $(\mathrm{y}=0.53 \mathrm{x}-0.92$; R-sq (adj) $=49.32$; $P<0.0001$ ). O. validus $\mathrm{t}_{\text {arival }}$ and $\mathrm{t}_{\text {MaxN }}$ values were also only significantly affected by input abundance ( $\mathrm{y}=$ $-234.17 \mathrm{x}+5199.6$ and $\mathrm{y}=-23.84+4915.4 ; \mathrm{R}-\mathrm{sq}(\mathrm{adj})=19.14$ and $3.37 ; P<0.0001$ and $\mathrm{P}=0.0008$ ). For $P$. corrugatus input abundance accounted for $34.48 \%$ of the variability in $\operatorname{Max}_{\mathrm{N}}(\mathrm{y}=0.2241-0.0985$; R-sq $(\mathrm{adj})=34.4 ; P<0.0001)$ and $\mathrm{t}_{\text {arrival }}$ and $\mathrm{t}_{\operatorname{maxN}} 19.29 \%$ and $1.49 \%(\mathrm{y}=-163.74+4879.5$ and $\mathrm{y}=-15.179+$ 4662.0; R-sq $(\operatorname{adj})=19.29$ and $1.49 ; P<0.0001$ and $P=0.03$. Current speed and $P$. corrugatus approach speed had no significant effect upon $\operatorname{Max}_{\mathrm{N}}, \mathrm{t}_{\text {arival }}$ and $\mathrm{t}_{\operatorname{maxN}}$ values.

### 3.2. Comparison to baited underwater camera data

The $\operatorname{Max}_{N}$ output of the models developed to describe the behaviour of the two tropical fish and Antarctic invertebrate species were all primarily affected by the input parameter population abundance. Therefore, $\mathrm{Max}_{\mathrm{N}}$ was only used to match arrival patterns from the in-situ BUC deployment and the multiple model arrival patterns. $\mathrm{T}_{\text {arrival }}$ and $\mathrm{t}_{\text {maxN }}$ were also significantly related to abundance and could also be potentially used to select model arrival patterns. There was limited evidence from the sensitivity analysis of the effect of the other model parameters on the model abundance indices therefore parameters were kept within the ranges reported in Table 1.

For 10 of the BUC deployments the corresponding UVC recorded no groupers and for three of the UVC transects that observed groupers none were observed in corresponding BUC deployments. 10 corresponding UVC and BUC pairs both recorded E. fasciatus and for 9 of these pairs the BUC model produced the same or slightly higher abundance estimates (Fig. 3a). The Bland Altman plot provides little evidence of systematic bias between the abundance estimates of the grouper E. fasciatus generated by the BUC model methodology and the UVC surveys. This is concluded as all data points are within the $+/-1.96$ SD limits of agreement in the plots and points are distributed symmetrically around the mean (Fig.4a). Only 4 corresponding UVC and BUC pairs both observed moray eels of the genus Gymnothorax and the BUC model produced higher or the same abundances. Moray eels were only observed in BUCs in 8 of the corresponding UVC and BUC pairs and only in UVC in 4 pairs. The Bland-Altman plot show that points are symmetrically distributed around the mean and that all point were within the $+/-1.96 \mathrm{SD}$ limits of agreement (Fig.4b).

In all 18 UVC and BUC pairs $O$. validus was observed and there was no clear pattern of differences between the abundance estimates recorded by each method (Fig.3c). All the data points for $O$. validus abundance estimates from the BUC model and the UVC were within or on the $+/-1.96$ SD limits of
agreement. From the plot it would however, appear that the plots were slightly asymmetrical to the zero and that average abundances from the model are slightly less than those recorded by the UVC as the abundance of $O$. validus increases (Fig.4c). For 8 of the 18 corresponding transect and BUC model pairs abundance estimates for $P$. corrugatus were only recorded by the BUC model and in a further 6 pairs the BUC model estimates were much larger than in the UVC surveys (Fig.3d). In the Bland-Altman plots two outliers were removed where abundances > 100 individuals were recorded by the BUC. All points were within the 1.96 SD limits of agreement but they were not symmetrically distributed around the mean indicating that higher abundances were measured by the BUC (Fig.4d).

## 4. Discussion

Results from the sensitivity analysis indicate that for tropical and Antarctic models of fish and invertebrate movement around the BUC system the abundance of the surveyed population was the factor most strongly related to the $\mathrm{Max}_{\mathrm{N}}$. These models allow a BUC user to determine the relationship between $\mathrm{Max}_{\mathrm{N}}$ and the abundance of the focal species and allow the commonly collected $\mathrm{Max}_{\mathrm{N}}$ unit of relative abundance to be converted to absolute units. Two other commonly-recorded indices of abundance, $\mathrm{t}_{\text {arrival }}$ and $t_{\text {maxN }}$ appear to be less closely related to absolute abundance than might have been assumed, but might usefully contribute to model parameter selection where more than one abundance value results in the observed $\mathrm{Max}_{\mathrm{N}}$. Within the range of species used here, estimates of their searching speed and staying time had relatively little influence on the model $\mathrm{Max}_{\mathrm{N}}$. This is a reassuring finding as it is relatively difficult to estimate these behavioural values in wild animals.

For all species-specific, models $\mathrm{Max}_{\mathrm{N}}$ appeared to be the measure which accounted for most of the variability in the input population abundance fish or invertebrates. Measurements of $t_{\text {arrival }}$ and $t_{\text {maxN }}$ would however, reflect more about aspects of fish approach swimming speed and the current velocity observed around the BUC deployment. Stoner et al., (2008) found that a poor correlation exists between BUC time based metrics and abundance estimates of juvenile Pacific cod from corresponding seine net trawls, while $\mathrm{Max}_{\mathrm{N}}$ measures correlated well with trawl survey results. Time based metrics from BUC studies in the abyssal environment have however, been used successfully to calculate the absolute abundance of scavenging fish populations (Priede and Merrett, 1996). The current speeds observed around the BUC deployments and that were used for model ranges were relatively slow. If BUC deployments were within environments experiencing high current speeds then possibly variation in current speed would likely have a greater affect on BUC output indices and detailed current speed measurements during BUC deployments would be essential. The model framework presented here allows these different scenarios to be tested against field data. Estimates of the range of diffusional velocities experienced in the tropical and Antarctic environments were not available to investigate its potential effect upon arrival patterns, but again the framework allows easy incorporation of new field or laboratory data on diffusion to be incorporated as it becomes available. The incorporation of fluid dynamics modelling into the methodology would enable the potential effects of current speed and diffusional velocity on the arrival of fish or invertebrates at the BUC to be explored in more detail. Unlike previous models an odour plume of any shape or concentration can be
incorporated into this framework to replace the "pie segment" used here. Animals contacted by the plume or walking/swimming into the side of it would respond in the same way as those in the existing models.
Refinements such as animals resuming random movement if they leave an irregularly shaped plume would be added at this stage.

Staying time had no impact on abundance metrics even though it had been shown to affect $\mathrm{Max}_{\mathrm{N}}$ values in the deep-sea BUC studies (Priede et al., 1990). The majority of BUC studies in the abyssal northeast Atlantic found the mean staying time of the deep-sea grenadier (C. armatus) to be approximately 2 hours (Priede et al., 1994; Henriques et al., 2002). In the shallow water BUC fish arrive more rapidly and frequently, causing the staying time to likely have less of an impact on $\mathrm{Max}_{\mathrm{N}}$ values. With longer staying times the number of fish at the camera will accumulate to reach $\operatorname{Max}_{\mathrm{N}}$ and the total meaning that $\mathrm{Max}_{\mathrm{N}}$ will have more of a linear relationship with the numbers visiting the BUC. However, in the shallow water environment where more fish are coming and going from the field of view there maybe a larger difference between $\mathrm{Max}_{\mathrm{N}}$ and the total number of animals visiting the camera. These results therefore indicate that in these models accurate estimate of fish or invertebrate staying time, cruising speed or diffusional velocity are not important to the output of the model and therefore all that is necessary is the selection of a suitable range. More important factors such as fish and invertebrate approach speed and the current speed should be prioritised. The latter is certainly directly measurable at the camera, though in complex habitats the current experienced by the fauna might be quite different. Approach speed is harder to ascertain, though stereo camera systems such as BRUVS can probably provide useful information if the system lands facing downstream at the point at which animals arrive. With downward-looking cameras the field of view is often too small to get good estimates of movement speed, but not impossible, especially for slow-moving species. In our Antarctic studies we were able to directly measure invertebrate walking speed across the seabed.

The absolute abundance estimates of $E$. fasciatus and $O$. validus generated by the BUC model methodology were found to be most comparable to the abundance estimates from corresponding UVC surveys. This is because these species are visible to the UVCs as well as to the BUC. The other two species tend to be hidden in rocks (Clarke and Prothero-Thomas, 1997) or within the coral reef (Bshary et al., 2006) except when bait is present, with their occasional appearance in the open probably being caused by recent feeding or disturbance. Moray eels of the genus Gymnothorax are generally nocturnal hunters and during the day they will remain hidden within rocky refuge (Bshary et al., 2006; Bardach et al., 1959) making it difficult for daytime UVC surveys to detect them. In a number of BUC and UVC corresponding pairs the BUC survey observed moray eels when the UVC surveys recorded none causing the BUC model to estimate abundances when the UVC estimate equalled zero. The abundance estimates generated by the BUC models for the nemertean worm $P$. corrugatus were higher than those within the higher abundance estimates were produced by the BUC models for $P$. corrugatus due to the BUC recording $P$. corrgutus but none being observed in the corresponding UVC survey. This can be attributed to the species taking refuge under rocks during the day (Clarke et al., 1997) causing few to be observed in daytime transects. This will result in the model parameters being calibrated to artificially low populations densities. Little is known about the behaviour of $P$. corrugatus and it is possible that large groups of individuals congregate within refuges
(Clarke and Prothero-Thomas, 1997), violating the assumption of the model that individuals are randomly distributed and act independently of each other.

Models also assume that all fish react and follow the bait plume once encountered, however factors such as satiation state, olfactory capabilities and the availability of other food sources in the environment will impact upon their decision. Due to the comparability of absolute abundance estimates from the BUC model and the UVC, it would appear that a large proportion of the nearby animals from these species reacted to the bait plume. Model assumptions include that individuals react independently of each other however, competitive behavioural interactions have been observed to occur between fish at the bait of BUC systems (Armstrong et al., 1992; Stoner et al., 2008; Dunlop et al., 2014). It has been suggested that these interactions discourage some fish from approaching the bait due to the increased chance of competition (Jones et al., 2013; Willis et al., 2003; Cappo et al., 2004) or predation (Lampitt et al., 1983; Harvey et al., 2007) presented by the other fish. It is therefore evident that in both the fish species studied competitive interactions around the BUC could potentially impact upon the arrival patterns of individuals at the bait. The effect of other species interactions on the arrival patterns of fish and invertebrates at the BUC should also be considered. Effects may include particular species posing a higher predation risk at the bait reducing the number of the other species observed. Further studies of the impact of these interactions would allow this information to be added to modelling approaches. Unlike previous models our framework would allow multiple species models to be combined using information on the species composition and potentially the effects of interactions on bait approach and staying times. Also when foraging individuals become close to the bait they are potentially attracted by the movement and sounds of others feeding (Bailey and Priede, 2002). For shallow water fish species that reply heavily upon sight for foraging and hunting (Stoner et al., 2008) this has the potential to impact on their behaviour in relation to the BUC system and thus arrival patterns. Further valuable research would be the investigation of the application of this modelling approach to other marine species, which have been found to be attracted to BUC systems. This would primarily include the large, predatory mobile species that BUC surveys have been found to effectively survey (Malcolm et al., 2007; Watson et al., 2010).

Preliminary results show that this stochastic modelling approach can generate absolute abundance estimates of some shallow water fish and invertebrate populations from BUC deployments and that these estimates are comparable to an established survey method. Discrepancies were apparently due to cryptic behaviour in some species resulting in underestimates of abundance during underwater visual census surveys. The generation of absolute abundance estimates from shallow BUC surveys improves the application of the method substantially and makes the results comparable to those of other survey methods, such as trawl surveys and transects commonly used in stock assessments and monitoring programmes. This also enables previously-collected BUC data to be reanalysed and diversity indices for these deployments to be recalculated based on the abundances of the animals present rather than combinations of $\mathrm{Max}_{N}$ values.

In conclusion, the spatial, stochastic modelling approach described and tested in this study represents one of the first attempts to model the arrival process of shallow water marine species at a BUC system.

Initial results for a small set of tropical and Antarctic species-specific models show that this method has the potential to generate absolute abundance estimates from BUC data that are comparable to UVC data. The model could be used retrospectively to re-analyse existing $\operatorname{Max}_{N}$ data. This development combined with the existing ability of BUCs to generate data in a time-and-cost efficient and non-destructive manner can significantly improve the value of this method to monitor inshore marine populations.

## Figure and Table Legends

Fig. 1. Diagram illustrating the general input and output parameters of the model simulation describing the behaviour of fish and invertebrate populations in relation to a baited underwater camera system.

Fig. 2. Example plot of the arrival pattern of the black tip grouper (Epinephelus fasciatus) at the baited underwater camera system (BUC) produced by 5 model runs and the arrival pattern from an in-situ BUC deployment.

Fig. 3. Histograms and scatter plots comparing the absolute abundance estimates generate from UVC transects (open bars) and BUC models (closed bars) for a) the grouper (Epinephelus fasciatsus), b) the moray eel species (Gymnothorax spp.), c) the Antarctic asteroid (Odontaster validus) and d) the Antarctic nemertean worm (Parbolasia corrugatus).

Fig. 4. Bland Altman plots illustrating the agreement between the abundance estimates generated by the baited underwater camera model (BUC) and the underwater visual census survey (UVC) for a) Epinephelus fasciatus, b) Gymnothorax spp., c) Odontaster validus and d) Parbolasia corrugatus.

Table 1 Input parameters ranges for Epinephelus fasciatus, Gymnothorax spp., Pollachius virens, Scyliorhinus canicula, Odontaster validus and Parbolasia corrugatus.

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Figure 1


Figure 2


Figure 3
a) Epinephelus fasciatus


b) Gymnothorax spp.


c) Odontaster validus


d) Parabolasia corrugatus




| Parameters | $\begin{aligned} & \text { Area } \\ & \left(\mathbf{m}^{2}\right) \end{aligned}$ | Current speed (m si$)$ | Abundance (individuals) | Cruising speed ( $\mathrm{m} \mathrm{s}^{-1}$ ) | Turning interval <br> (s) | Approach speed ( $\mathrm{m} \mathrm{s}^{-1}$ ) | Staying time <br> (s) | References |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Species |  |  |  |  |  |  |  |  |
| Epinephelus fasciatus | 1000 | 0.02-0.2 | 1-100 | $0-0.2$ | 0-120 | 0.294-0.365 | 0-240 | Fulton, 2007; Bshary et al., 2006 |
| Gymnothorax spp. | 1000 | 0.02-0.2 | 1-100 | 0 | 0-120 | 0.0935-0.318 | 0-180 | D'Aout and Aerts, 1999; Gibran, 2007 |
| Odontaster validus | 6.25 | 0.01-0.1 | 1-100 | 0 | $\mathrm{n} / \mathrm{a}$ | 0.0001-0.001 | To simulation end | Kidawa, 2001 |
| Parbolasia corrugatus | 6.25 | 0.01-0.1 | 1-100 | 0 | $\mathrm{n} / \mathrm{a}$ | $0.0001-0.0003$ | To simulation end | Clarke and Prothero- <br> Thomas, 1997 |

