## 11.0 Appendix A

**Appendix A - Table A1.** Summary of the <u>assumptions</u> and pros/cons of the different <u>modelling approaches</u> (adapted from Wearn & Glover-Kapfer [2017] and Clarke et al. [2022]).

<b>Objective</b>	Approach	Assumptions	Pros	Cons	References
<u>Species</u> inventory	<u>Species</u> inventory	• No formal <u>assumptions</u> <sup>1</sup>	<ul> <li>Maximum flexibility for study design (e.g., <u>camera days per</u> <u>camera location</u> or use of <u>lure</u><sup>2</sup>)<sup>1</sup></li> </ul>	<ul> <li>Not reliable estimates for inference ("considered as unfinished, working drafts")<sup>1</sup></li> </ul>	<sup>1</sup> Wearn & Glover- Kapfer, 2017
Species richness	Species richness	<ul> <li><u>Camera locations</u> are <u>randomly</u> <u>placed</u><sup>1</sup></li> <li><u>Camera locations</u> are independent<sup>1</sup></li> <li><u>Detection probability</u> of different species remains the same<sup>1</sup> ("true" species richness estimation involves attempting to correct for "<u>imperfect</u> <u>detection</u>"<sup>1</sup>)</li> <li>Sampling effort is comparable between <u>camera locations</u><sup>31</sup></li> </ul>	<ul> <li>Fundamental to ecological theory and often a key metric used in management<sup>1</sup></li> <li>Simple to analyze, interpret and communicate<sup>1</sup></li> <li>Models exist to estimate asymptotic species richness, including unseen species (simple versions of these models - "EstimateS" and the "vegan" R-packages)<sup>1</sup></li> </ul>	<ul> <li>Dependent on the scale (as captured in the species-area relationship)<sup>1</sup></li> <li>All species have equal weight in calculations, and community evenness is disregarded<sup>1</sup></li> <li>Insensitive to changes in abundance, community structure and community composition<sup>1</sup></li> </ul>	<ul> <li><sup>2</sup> Rovero et al., 2013</li> <li><sup>3</sup> MacKenzie et al., 2002</li> <li><sup>4</sup> MacKenzie et al., 2006</li> <li><sup>5</sup> Rowcliffe &amp; Carbone, 2008</li> <li><sup>6</sup> Lambert, 1992</li> </ul>
Species diversity	Species diversity	<ul> <li><u>Camera locations</u> are <u>randomly</u> <u>placed</u><sup>1</sup></li> <li><u>Camera locations</u> are independent<sup>1</sup></li> <li><u>Detection probability</u> of different species remains the same<sup>1</sup></li> </ul>	<ul> <li>Captures evenness and richness (although some indices only reflect evenness)<sup>1</sup></li> <li>Most indices are easy to calculate and widely implemented in software packages (e.g., "EstimateS" and "vegan" in R)<sup>1</sup></li> </ul>	<ul> <li>Many indices exist, and it can be difficult to choose the most appropriate<sup>1</sup></li> <li>Comparing measures across space, time and studies can be very difficult<sup>1</sup></li> <li>Insensitive to changes in community composition<sup>1</sup> (however, this may be conditional on study design)</li> </ul>	<ul> <li><sup>7</sup> Mullahy, 1986</li> <li><sup>8</sup> McCullagh &amp; Nelder, 1989</li> <li><sup>9</sup> Heilbron 1994</li> <li><sup>10</sup> Karanth &amp; Nichols, 1998</li> <li><sup>11</sup> Karanth, 1995</li> </ul>
Species diversity	β-diversity	<u>Camera locations</u> are <u>randomly</u> <u>placed</u> <sup>1</sup>	<ul> <li>Can be used to track changes in community composition<sup>1</sup></li> </ul>	<ul> <li>No single best measure for all purposes<sup>1</sup></li> </ul>	<sup>12</sup> Clarke et al., 2023

<u>Objective</u>	Approach	Assumptions	Pros	Cons	References
		<ul> <li>Randomness and independence<sup>1</sup></li> <li>Samples are assumed to have been taken at random from the broader population of sites<sup>1</sup></li> </ul>	<ul> <li>Plays a critical role in effective conservation prioritization (e.g., designing reserve networks)<sup>1</sup></li> <li>Important for detecting changes in the fundamental processes<sup>1</sup></li> </ul>	<ul> <li>Interpretation/communication not always straightforward<sup>1</sup></li> <li>Scale-dependent (i.e., influenced by the size of the communities that are being included)<sup>1</sup></li> </ul>	<ul> <li><sup>13</sup> Noss et al., 2003</li> <li><sup>14</sup> Kelly et al., 2008</li> <li><sup>15</sup> Moeller et al., 2018</li> <li><sup>16</sup> Chandler &amp;</li> </ul>
Occupancy 3	Occupancy models <sup>3</sup>	<ul> <li>Occupancy is constant<sup>[3]</sup> (abundance is constant)<sup>4</sup></li> <li>Camera locations are independent<sup>4</sup></li> <li>Detections are independent<sup>4</sup></li> <li>The probability of occupancy and detection are constant across all camera locations within a stratum or can be modelled using covariates<sup>4</sup></li> <li>Species are not misidentified<sup>4</sup></li> </ul>	<ul> <li>Does not require individual identification<sup>4</sup></li> <li>Only requires detection/non- detection data for each site<sup>1</sup></li> <li>Relatively easy-to-use software exists for fitting models (PRESENCE, MARK, and the "unmarked" R package)<sup>1</sup></li> <li>"Open" models exist that allow for the estimation of site colonization and extinction rates<sup>1,4</sup></li> <li>Multi-species <u>occupancy</u> <u>models<sup>[3]</sup> allow the inclusion of interactions among species while controlling for <u>imperfect</u> <u>detection</u><sup>1</sup></u></li> </ul>	<ul> <li>Occupancy<sup>[3]</sup> only measures distribution; it may be a misleading indicator of changes in abundance<sup>1</sup></li> <li>Interpretation/communication of results may not be straightforward (if the scale of movement is much larger than the <u>camera spacing</u> the results should be interpreted as "probability of use" rather than <u>occupancy</u>)<sup>1</sup></li> </ul>	Royle, 2013 <sup>17</sup> Royle et al., 2009 <sup>18</sup> Borchers & Efford, 2008 <sup>19</sup> Efford, 2004 <sup>20</sup> Royle & Young, 2008 <sup>21</sup> O'Brien et al., 2011 <sup>22</sup> Doran- Myers, 2018 <sup>23</sup> Morin et al., 2022 <sup>24</sup> Green et al., 2020
Relative abundance indices	Poisson Zero-inflated Poisson (ZIP) <sup>6</sup> Negative binomial (NB) <sup>7</sup>	• Many <u>assumptions</u> exist (since used for many approaches) <sup>1</sup>	<ul> <li>Simple to calculate and technically possible (even with small sample sizes when robust methods might fail)<sup>1</sup></li> <li><u>Relative abundance indices</u> often do correlate with abundance<sup>1</sup></li> </ul>	<ul> <li>Difficult to draw inferences (a large number of <u>assumptions</u>); comparisons across space, time, species, and studies are difficult<sup>1</sup></li> <li>Requires stringent <u>study design</u> (e.g., random sampling, standardized methods)<sup>1</sup></li> </ul>	<ul> <li><sup>25</sup> Parmenter et al., 2003</li> <li><sup>26</sup> Noss et al., 2012</li> <li><sup>27</sup> Sollmann et al., 2013a</li> </ul>

Objective Approac	h <u>Assumptions</u>	Pros	Cons	References
Population     Capture-	Demographic closure (i.e., no births	Calibration with independent <u>density</u> estimates is possible <sup>1</sup>	<ul> <li>Detection rates from remote cameras cannot be used as an index to compare relative abundance across species<sup>5</sup></li> <li>Requires that individuals are</li> </ul>	<ul> <li><sup>28</sup> Sollmann et al., 2013b</li> <li><sup>29</sup> Rich et al., 2014</li> <li><sup>30</sup> Whittington et al., 2018</li> <li><sup>31</sup> Royle &amp; Nichols, 2003</li> <li><sup>32</sup> Efford et al., 2009b</li> </ul>
size / recapture Absolute (CR) / abundance capture- / vital rates mark- / Density; recapture Marked (CMR) <sup>10,11</sup> population	or deaths) <sup>1</sup> <ul> <li>Geographic closure (i.e., no immigration or emigration)<sup>1</sup></li> <li>All individuals have at least some probability of being detected<sup>2</sup></li> <li>Sampled area encompasses the full extent of individuals' movements<sup>2,10</sup></li> <li>Activity centres are randomly dispersed<sup>12</sup></li> <li>Activity centres are stationary<sup>12</sup></li> </ul>	<ul> <li>abundance index that controls for imperfect detection<sup>1</sup></li> <li>Easy-to-use software exists to implement (e.g., CAPTURE); MARK Implements more complicated models with covariates (and must be used for mark-resight modelling)<sup>1</sup></li> <li>Can use the robust design with "open" models to obtain recruitment and survival rate estimates<sup>1</sup></li> </ul>	<ul> <li>distinguishable.<sup>1</sup> However, CR<sup>[10,11]</sup> has also been used to estimate abundance of species that lack natural markers but that have phenotypic and/or environment-induced characteristics<sup>2,13,14</sup></li> <li>When the sample size is large enough to reliably estimate density with CR, <sup>[10,11]</sup> individuals are unlikely to have a unique marker<sup>2,13,14</sup></li> <li>Dependent on the surveyed area, which is difficult to track and calculate<sup>1</sup></li> <li>Requires a minimum number of captures and recaptures<sup>1</sup></li> <li>Relatively stringent requirements for study design (e.g., no "holes" in the trapping grid)<sup>1</sup></li> <li>Geographic closure at the plot level, which is often unrealistic<sup>1</sup></li> </ul>	2009b <sup>33</sup> Royle et al., 2014 <sup>34</sup> Augustine et al., 2019 <sup>35</sup> Burgar et al., 2018 <sup>36</sup> Sun et al., 2022 <sup>37</sup> Sollmann, 2018 <sup>38</sup> Augustine et al., 2018 <sup>39</sup> Davis et al., 2021 <sup>40</sup> Rowcliffe et al., 2013

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
Density /	<u>Spatially</u> explicit	• Demographic closure (i.e., no births or deaths) <sup>1</sup>	<ul> <li>Produces direct estimates of density or population size for</li> </ul>	<ul> <li>Assumes a specific relationship between abundance and detection<sup>1</sup></li> <li>Density cannot be explicitly estimated because the true area animals occupy is never measured (only approximated)<sup>16</sup></li> <li>Requires that individuals are identifiable<sup>1</sup></li> </ul>	<ul> <li><sup>42</sup> Rowcliffe et al., 2014</li> <li><sup>43</sup> Rowcliffe et al., 2016</li> <li><sup>44</sup> Rowcliffe et al., 2011</li> <li><sup>45</sup> Cusack et al., 2015</li> <li><sup>46</sup> Nakashima</li> </ul>
size; <u>Marked</u> population	capture recapture (SECR) <sup>17–20</sup> (also referred to as <u>Spatial</u> <u>capture-</u> <u>recapture</u> [SCR])	<ul> <li>Detection probability of different individuals is equal<sup>1</sup> <ul> <li>or, for SECR, individuals have equal detection probability at a given distance from the centre of their home range<sup>1</sup></li> </ul> </li> <li>Detections of different individuals are independent<sup>1</sup></li> <li>Detections of different individuals are independent<sup>1</sup></li> <li>Behaviour is unaffected by cameras and marking<sup>1</sup></li> <li>Individuals do not lose marks<sup>1</sup></li> <li>Individuals are not misidentified<sup>1</sup></li> <li>Surveys are independent<sup>1</sup></li> <li>For conventional models, geographic closure (i.e., no immigration or emigration)<sup>1</sup></li> <li>Spatially explicit models have further assumptions about animal movement<sup>1,17,21</sup>; these include: <ul> <li>Home ranges are stable<sup>1</sup></li> <li>Movement is unaffected by cameras<sup>1</sup></li> <li>Camera locations are randomly placed with respect to the</li> </ul> </li> </ul>	<ul> <li>explicit spatial regions<sup>16</sup></li> <li>Allows researchers to mark a subset of the population/to take advantage of natural markings<sup>1</sup></li> <li>Estimates are fully comparable across space, time, species and studies<sup>1</sup></li> <li><u>Density</u> estimates obtained in a single model, fully incorporate spatial information of locations and individuals<sup>1</sup></li> <li>Both likelihood-based and Bayesian versions of the model have been implemented in relatively easy-to-use software (DENSITY and SPACECAP, respectively, as well as associated R packages)<sup>1</sup></li> <li>Flexibility in study design (e.g., "holes" in the trapping grid)<sup>11</sup></li> <li>"Open" <u>SECR</u><sup>[17–20]</sup> models exist that allow for estimation of recruitment and survival rates<sup>1</sup></li> </ul>	<ul> <li>Requires that a minimum number of individuals are trapped (each recaptured multiple times ideally)<sup>1</sup></li> <li>Requires that each individual is captured at a number of <u>camera locations</u><sup>1</sup></li> <li>Multiple cameras per station may be required to identify individuals; difficult to implement at large spatial scales as it requires a high density of cameras<sup>12,23</sup></li> <li>May not be precise enough for long-term monitoring<sup>24</sup></li> <li>Cameras must be close enough that animals are detected at multiple <u>camera locations</u><sup>1</sup> (may be challenging to implement at large scales as many cameras are needed)"<sup>16</sup></li> <li>½ MMDM (Mean Maximum Distance Moved) will usually lead to an under -estimation of home range size and thus overestimation of <u>density</u><sup>1,25,26</sup></li> </ul>	<ul> <li><sup>40</sup> Nakasnima et al., 2018</li> <li><sup>47</sup> Meek et al., 2016</li> <li><sup>48</sup> Anile &amp; Devillard, 2016</li> <li><sup>49</sup> Huggard, 2018</li> <li><sup>50</sup> Becker et al., 2022</li> <li><sup>51</sup> Warbington &amp; Boyce, 2020</li> <li><sup>52</sup> Howe et al., 2017</li> <li><sup>53</sup> Borchers &amp; Marques, 2017</li> <li><sup>54</sup> Palencia et al., 2021</li> <li><sup>55</sup> Gilbert et al., 2021</li> </ul>

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
Objective	Approach Spatial mark- resight (SMR) (type of SCR model) <sup>16,27,28</sup>	Assumptions         distribution and orientation of home ranges1         o Distribution of home range centres follows a defined distribution (Poisson, or other, e.g., negative binomial)1         • Demographic closure (i.e., no births or deaths) <sup>12,16</sup> • Geographic closure (i.e., no immigration or emigration) <sup>12,16</sup> • Individuals do not lose marks1 (for maximum precision), but SMR <sup>[16,27,28]</sup> does allow for inclusion of marked but unidentified resighting detections <sup>27,29</sup> • Individuals are not misidentified1         • Failure to identify marked individuals is random <sup>12,30</sup> • Marked animals are a random	<ul> <li>Pros</li> <li><u>SECR</u><sup>[17–20]</sup> accounts for variation in individual <u>detection</u> <u>probability</u>; can produce spatial variation in <u>density</u>; <u>SECR</u><sup>[17–20]</sup> more sensitive "to detect moderate-to-major populations changes" (+/-20-80%)<sup>12,23</sup></li> <li>Estimates are fully comparable to <u>SECR</u><sup>[17–20]</sup> of <u>marked</u> species<sup>1</sup></li> <li>Can be applied to a broader range of species than <u>SECR</u><sup>[17–20]1</sup></li> <li>Allows researcher to take advantage of natural markings<sup>1</sup></li> <li>Allows researcher to mark a subset of the population (note - precision is dependent on number of <u>marked</u> individuals in a population)<sup>1</sup></li> </ul>	<ul> <li>Cons</li> <li>Animals may have to be physically captured and <u>marked</u> if natural marks do not exist on enough individuals<sup>1</sup></li> <li>All individuals must be identifiable<sup>1</sup></li> <li>Allows for <u>density</u> estimation for a <u>unmarked population</u>, but the precision of the <u>density</u> estimates are likely to be very low value<sup>1</sup></li> <li>Remains poorly tested with camera data, although it offers promise<sup>1</sup></li> </ul>	References <sup>56</sup> Twining et al., 2022 <sup>57</sup> Bessone et al., 2020 <sup>58</sup> Loonam et al., 2021 <sup>59</sup> Bridges & Noss, 2011 <sup>60</sup> Rovero & Zimmermann , 2016
		<ul> <li>sample of the population with home ranges located inside the state space<sup>28,29</sup></li> <li>Detections are <u>independent</u><sup>12,16</sup></li> <li>Individuals have equal <u>detection</u> <u>probability</u> at a given distance from the centre of their home range<sup>1</sup></li> <li>Detections of different individuals are <u>independent</u><sup>1</sup></li> <li>Movement is unaffected by cameras<sup>1</sup></li> <li>Behaviour is unaffected by cameras and marking<sup>1</sup></li> <li><u>Camera locations</u> are <u>randomly</u> <u>placed</u> relative to the distribution and orientation of home range<sup>1</sup></li> </ul>		<ul> <li>Density estimates are likely less precise than with <u>SECR</u><sup>[17–20]</sup> or <u>REM</u>, unless a large proportion of the population have marks<sup>1</sup></li> <li>Requires sampling points to be close enough that individuals encounter multiple cameras<sup>1</sup></li> </ul>	

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
<u>Objective</u>	Approach	Assumptions         • Camera locations are close enough together that animals are detected at multiple cameras <sup>12,16</sup> • Surveys are independent <sup>1</sup> • Home ranges are stable <sup>1</sup> • Distribution of home range centres follows a defined distribution (Poisson, or other, e.g., negative binomial) <sup>1</sup>	Pros	Cons	References
		<ul> <li>Animals' activity centres are randomly dispersed<sup>12,16</sup></li> <li>Animals' activity centres are stationary<sup>12,16</sup></li> <li>All animals have stable activity centres within home ranges where detection probability is greatest<sup>27,31,32</sup></li> </ul>			
Density; Unmarked population	Spatial count (SC) / Unmarked spatial capture- recapture (type of SCR model) <sup>16,33</sup>	<ul> <li><u>Camera locations</u> are close enough together that animals are detected at multiple cameras<sup>12,16</sup></li> <li>Demographic closure (i.e., no births or deaths)<sup>12,16</sup></li> <li>Geographic closure (i.e., no immigration or emigration)<sup>12,16</sup></li> <li>Detections are <u>independent</u><sup>12,16</sup></li> <li>Animals' activity centres are randomly dispersed<sup>12,16</sup></li> <li>Animals' activity centres are stationary<sup>12,16</sup></li> </ul>	• Does not require individual identification <sup>12</sup>	<ul> <li>Produces imprecise estimates even under ideal circumstances unless supplemented with auxiliary data (e.g., telemetry)<sup>16,22,27,28</sup></li> <li>Precision decreases with an increasing number of individuals detected at a camera"<sup>23</sup> (as overlap of individuals' home ranges increases) <sup>12,34</sup></li> <li>Not appropriate for low <u>density</u> or elusive species when recaptures too few to confidently infer the number and location of activity centres"<sup>12,35</sup></li> <li>Not appropriate for high-<u>density</u> populations with evenly spaced activity centres (camera[- specific] counts will be too similar and impair activity centre inference)"<sup>12</sup></li> </ul>	

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
				<ul> <li>Ill-suited to populations that exhibit group-travelling behaviour"<sup>12,36</sup></li> </ul>	
				<ul> <li>Study design (camera arrangement) can dramatically affect the accuracy and precision of <u>density</u> estimates"<sup>12,37</sup></li> </ul>	
				<ul> <li>Cameras must be close enough that animals are detected at multiple <u>camera locations</u> (may be challenging at large scales as many cameras are needed)"<sup>12,16</sup></li> </ul>	
Density / population size; Partially Marked population	Spatial Partial Identity Model (Categorical SPIM; catSPIM) <sup>34,36</sup> (Extension of SC model using animal traits (e.g., Sex Class, antler points) and model parameters)	<ul> <li>Same as <u>SC</u><sup>12,34,36</sup> <ul> <li>Camera must be close enough together that animals are detected at multiple cameras<sup>12,16</sup></li> <li>Demographic closure (i.e., no births or deaths)<sup>12,16</sup></li> <li>Geographic closure (i.e., no immigration or emigration)<sup>12,16</sup></li> <li>Detections are independent<sup>12,16</sup></li> </ul> </li> <li>Activity centres are randomly dispersed<sup>12,16</sup></li> <li>Activity centres are stationary<sup>12,16</sup></li> <li>Each categorical identifier (e.g., male/female, collared/not collared, etc) has fixed number of possibilities<sup>36</sup></li> <li>All possible values of categorical identifiers occur in the population with probabilities that can be estimated<sup>12,34,36</sup></li> <li>Every individual is assigned "full categorical identify" (i.e., "set of</li> </ul>	• May produce more precise and less biased <u>density</u> estimates than <u>SC</u> with less information <sup>12,36</sup>	<ul> <li>Sensitive to non-independent movement (e.g., group-travel; can cause over-dispersion and bias estimates<sup>12,36</sup>); may limit application to solitary species only<sup>12,36</sup></li> <li>May produce be less reliable/accurate estimates for high-<u>density</u> populations<sup>12,36</sup></li> <li>Too few categorical identifiers/ possibilities can result in mis- assignments and overestimating <u>density</u><sup>12,25,34</sup></li> </ul>	

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
		<ul> <li>traits given all categorical identifiers and possibilities")<sup>12,34</sup></li> <li>Individuals' identifying traits do not change during the survey (e.g., antlers present/absent)<sup>34</sup></li> </ul>			
Density / population size; Partially Marked population	Spatial Partial Identity Model (2- flank SPIM) <sup>38</sup> (extension of SCR model augmented with data from partially- identifying images)	<ul> <li>Same as SCR<sup>12,38</sup></li> <li>Capture processes for left-side, right-side and both-side images are independent<sup>12,38</sup></li> </ul>	<ul> <li>Same as SCR<sup>12,38</sup></li> <li>Improved precision of <u>density</u> estimates relative to SCR<sup>12,38,39</sup></li> <li>Many study designs can be used (<u>paired sample stations</u>, single <u>camera locations</u>, and hybrids of both paired- and single <u>camera locations</u><sup>12,38,39</sup></li> <li>Can be used with single-camera and hybrid sampling designs, and therefore requires fewer cameras (or sample more area) than SCR<sup>12,38</sup></li> <li>May be more robust to non-independence than SC<sup>12,38</sup></li> </ul>	<ul> <li>Computationally intensive<sup>12,38</sup></li> <li>Increased precision is less pronounced in high-<u>density</u> populations<sup>12,38</sup></li> </ul>	
<u>Density;</u> <u>Unmarked</u>	Random encounter models (REM) <sup>40,41</sup>	<ul> <li>Demographic closure<sup>22,40</sup> (i.e., no births or deaths)</li> <li>Geographic closure<sup>22,40</sup> (i.e., no immigration or emigration)</li> <li><u>Camera locations</u> are randomly placed relative to animal movement<sup>1,40</sup></li> <li>Animal movement is unaffected by the cameras<sup>1,40</sup></li> <li>Accurate counts of independent "contacts" <u>camera locations</u><sup>1,40</sup></li> <li>Unbiased estimates of animal activity levels and speed<sup>1,42,43</sup></li> <li>Camera's detection zone can be approximated well using a 2D cone</li> </ul>	<ul> <li>Flexible study design (e.g., "holes" in grids allowed, camera spacing less important)<sup>1</sup></li> <li>Can be applied to unmarked species<sup>1</sup></li> <li>Allows community-wide <u>density</u> estimation<sup>1</sup></li> <li>Outputs also include informative parameter estimates (i.e., animal speed and activity levels, and detection zone parameters)<sup>1</sup></li> <li>Comparable estimates to SECR[<sup>17–20</sup>]<sup>1</sup></li> </ul>	<ul> <li>Requires relatively stringent study design, particularly (e.g., random sampling and use of bait or lure)<sup>1</sup></li> <li>Requires independent estimates of animal speed or measurement of animal speed within videos<sup>1</sup></li> <li>No dedicated, simple software<sup>1</sup></li> <li>Random relative to animal movement, grid preferred, avoid multiple captures of same individual, area coverage important for abundance estimation<sup>2</sup></li> <li>Possible sources of error include inaccurate measurement of</li> </ul>	

<b>Objective</b>	<u>Approach</u>	Assumptions	Pros	Cons	References
		<ul> <li>shape, defined by the radius and angle parameters<sup>44</sup></li> <li>If activity and speed are to be estimated from camera data, two additional assumptions:</li> <li>All animals are active during the peak daily activity<sup>42</sup></li> <li>Animals moving quickly past a camera are not missed<sup>43</sup></li> </ul>	<ul> <li>Does not require marked animals or identification of individuals<sup>22,40</sup></li> <li>Can use camera spacing without regard to population home range size<sup>22,40</sup></li> <li>Direct estimation of <u>density</u>; avoids ad-hoc definitions of study area<sup>40</sup></li> </ul>	detection zone and movement rate <sup>41,45</sup>	
<u>Density;</u> <u>Unmarked</u>	Random encounter and staying time (REST) <sup>46</sup>	<ul> <li>Demographic closure (i.e., no births or deaths) and geographic closure (i.e., no immigration or emigration) (animal density is constant during the survey)<sup>40</sup></li> <li>Detection is perfect<sup>1</sup> (detection probability "p" = 1) unless otherwise modelled<sup>46</sup></li> <li><u>Camera locations</u> are representative of the available habitat<sup>46</sup></li> <li><u>Camera locations</u> are randomly placed relative to the spatial distribution of animals<sup>46</sup></li> <li>Animal movement and behaviour are not affected by cameras<sup>46</sup></li> <li>Detections are independent<sup>46</sup></li> <li>The observed distribution of staying time in the focal area fits the distribution of movement<sup>46</sup></li> <li>The observed staying time must follow a given parametric distribution<sup>46</sup></li> </ul>	• Provides unbiased estimates of animal <u>density</u> , even when animal movement speed varies, and animals travel in pairs <sup>46</sup>	<ul> <li>Attraction or aversion to cameras is exhibited in some species<sup>47</sup> and could affect the time within the detection zone and subsequently affect estimates of <u>density</u><sup>22</sup></li> <li>Requires accurate measurements of the area of the camera detection zone, which has been a challenge in previous studies<sup>22,44–46,48</sup></li> <li>Mathematically challenging<sup>45</sup></li> </ul>	
<u>Density;</u> <u>Unmarked</u>	<u>Time in front</u> of the <u>camera</u> (TIFC) <sup>49–51</sup>	<ul> <li><u>Camera locations</u> are <u>randomly</u> <u>placed</u> or representative relative to animal movement<sup>50</sup></li> <li>Movement is unaffected by the cameras<sup>50</sup></li> </ul>	<ul> <li>Does not require individual identification<sup>51</sup></li> <li>Makes no <u>assumption</u> about home range<sup>51</sup></li> </ul>	<ul> <li>Requires careful calculation of the effective area of detection<sup>51</sup></li> <li>A high level of measurement error<sup>50</sup></li> </ul>	

<b>Objective</b>	Approach	Assumptions	Pros	Cons	References
		<ul> <li>Reliable detection of animals in part of the camera's <u>FOV</u> (at least)<sup>50</sup></li> </ul>	Comparable to estimates from <u>SECR</u> <sup>[17-20]51</sup>		
Density; Unmarked	Distance sampling (DS) <sup>52,53</sup>	<ul> <li>Random or systematic random placements (consistent with the assumption that points are placed independently of animal locations)<sup>52</sup></li> <li><u>Camera locations</u> are randomly placed relative to animal movement<sup>54</sup></li> <li>Detection is perfect (detection probability "p" = 1) at focal area / distance 0<sup>54</sup></li> <li>Demographic closure (i.e., no births or deaths) and geographic closure (i.e., no immigration or emigration) (animal density is constant during the survey)<sup>54</sup></li> <li>Animal movement and behaviour are unaffected by the cameras<sup>54</sup></li> <li>Animals are detected at initial locations (e.g., they do not change course in response to the camera prior to detection)<sup>54</sup></li> <li>Distances are measured exactly (however if the data from different distances will be grouped ("binned") for analysis later, an accuracy of +/-1m may suffice)<sup>54</sup></li> <li>Snapshot moments selected independent<sup>54</sup></li> </ul>	<ul> <li>A shortcut to controlling for variation in detection distances by only counting individuals within a short distance with an unobstructed view, and well sampled across cameras and species<sup>1</sup></li> <li>Density estimates are unbiased by animal movement "since camera-animal distance is measured at a certain instant in time (intervals of duration <i>t</i> apart)"<sup>12,52</sup></li> <li>Can be applied to low-density populations<sup>12,52</sup></li> <li>Does not require individual identification<sup>52</sup></li> </ul>	<ul> <li>May require discarding a portion of the dataset (when the best fitting model truncates the dataset)<sup>1</sup></li> <li>Biased by movement speed<sup>54</sup></li> <li>Best suited to larger animals; the smaller the focal species, the lower remote cameras must be set, which reduces the depth of the viewshed, and thus sampling size and the flexibility of the model"<sup>12,52</sup></li> <li>Does not permit inference about spatial variation in abundance (unless using hierarchical distance which can model spatial variation as a function of covariates)<sup>12,55</sup></li> <li>"Calculating camera-animal distances can be labour-intensive and time-consuming (However, recently developed techniques (e.g., Johanns et al., 2022) show promise for simplifying and automating the process)"<sup>12</sup></li> <li>Requires a good understanding of the focal populations' activity patterns; density estimates can be biased (e.g., underestimated) when regular periods of inactivity are not accounted for (using detection times to infer periods of activity may help overcome this limitation)"<sup>12,52,54</sup></li> </ul>	

<u>Objective</u>	Approach	Assumptions	Pros	Cons	References
				<ul> <li>Low population density and reactivity to cameras may be major sources of bias<sup>"12,57</sup></li> </ul>	
Density; Unmarked	<u>Time-to-</u> event (TTE) model <sup>15</sup>	<ul> <li>Demographic closure (i.e., no births or deaths)<sup>15,58</sup></li> <li>Geographic closure (i.e., no immigration or emigration) at the level of the sampling frame (area of interest); this assumption does not apply at the plot-level (area sampled by the camera)<sup>15,58</sup></li> <li>Animal movement and behaviour are unaffected by the cameras<sup>54</sup></li> <li><u>Camera locations</u> placement is random, systematic, or systematic random<sup>15</sup></li> <li>Detections are independent<sup>15</sup></li> <li>Spatial counts of animals (or counts in equal subsets of the landscape) are Poisson-distributed<sup>58</sup></li> <li>Accurate estimate of movement speed<sup>58</sup></li> <li>Detection is perfect (detection probability "p" = 1)<sup>15</sup></li> </ul>	• Can be efficient for estimating abundance of common species (with a lot of images) <sup>15</sup>	<ul> <li>Requires independent estimates of movement rate (difficult to obtain without telemetry data)<sup>15</sup></li> <li>Assumes that <u>detection</u> <u>probability</u> is 1 (or apply extension to account for <u>imperfect detection</u>)<sup>15</sup></li> </ul>	
<u>Density;</u> <u>Unmarked</u>	Space-to- event (STE) models <sup>15</sup>	<ul> <li>Demographic closure (i.e., no births or deaths)<sup>15</sup></li> <li>Geographic closure (i.e., no immigration or emigration)<sup>15</sup></li> <li><u>Camera locations</u> are <u>randomly placed<sup>15</sup></u></li> <li>Detections are <u>independent<sup>15</sup></u></li> <li>Spatial counts of animals in a small area (or counts in equal subsets of the landscape) are Poisson-distributed<sup>58</sup></li> </ul>	<ul> <li>Can be efficient for estimating abundance of common species (with a lot of images)<sup>15</sup></li> <li>Does not require estimate of movement rate<sup>15</sup></li> </ul>	• Assumes that <u>detection</u> <u>probability</u> is 1 <sup>15</sup>	

<u>Objective</u>	<u>Approach</u>	Assumptions	Pros	Cons	References
		<ul> <li>Detection is perfect (<u>detection</u> <u>probability</u> "p" = 1)<sup>15</sup></li> </ul>			
<u>Density;</u> <u>Unmarked</u>	Instantaneou s sampling (IS) <sup>15</sup>	<ul> <li>Demographic closure (i.e., no births or deaths)<sup>15</sup></li> <li>Geographic closure (i.e., no immigration or emigration)<sup>15</sup></li> <li><u>Camera locations</u> are <u>randomly placed<sup>15</sup></u></li> <li>Detections are <u>independent<sup>15</sup></u></li> <li>Detection is perfect (<u>detection probability</u> "p" = 1)<sup>15</sup></li> </ul>	<ul> <li>Can be efficient for estimating abundance of common species (with a lot of images)<sup>15</sup></li> <li>Flexible <u>assumption</u> of animals' distribution<sup>15</sup></li> </ul>	<ul> <li>Requires accurate counts of animals<sup>15</sup></li> <li>Assumes that perfect (<u>detection probability</u> "<i>p</i>" = 1)<sup>15</sup></li> <li>Reduced precision<sup>15</sup></li> </ul>	
Behaviour (diel activity mating, bold	-	<ul> <li><u>Assumptions</u> vary depending on the behavioural metric<sup>1</sup></li> <li>For studies of activity patterns and temporal interactions of species: activity level is the only factor determining <u>detection rates</u>; animals are active when camera <u>detection rate</u> reaches its maximum in daily cycle<sup>33,60</sup></li> </ul>	<ul> <li>Can detect difficult to observe behaviours (i.e., boldness, or mating)<sup>59</sup></li> <li>Long-term data on behavioural changes that would be difficult to obtain otherwise (i.e., time-limited human observers, or costly GPS collars)<sup>59</sup></li> <li>Can monitor behaviour in response to specific locations (i.e., compost sites, which might be more difficult using GPS collars for example)<sup>60</sup></li> <li>Can evaluate interactions between species<sup>60</sup></li> </ul>	<ul> <li>Behavioural metrics may not reflect the behavioural state (inferred)<sup>60</sup></li> <li>Biases associated with equipment (i.e., presence of the camera itself may change behaviour studied)<sup>60</sup></li> <li>Difficult to consider individual variation<sup>60</sup></li> </ul>	