

CEAMEC shiny App v1.0 User Manual

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1. Overview

CEAMEC (Cost-Effective Animal Management via Environmental Capacity), a *shiny* app in the HTML user interface (UI) programmed with *R* language, is a tool to provide managers with cost estimation of ecological-based management strategies in the population control of over-abundant nuisance species in the anthropogenic environments. Integrated with hierarchical modelling functions in *R* package *unmarked* (Fiske & Chandler, 2011) to identify the association between population density and the environmental resources, *CEAMEC* computes the change between pre-management (observed and may be subject to extant management) and post-management (user-defined management target) environmental carrying capacity. Based on the change, *CEAMEC* optimizes the quantity of different resources to be manipulated at the lowest cost. In this version, *CEAMEC* works for population survey data of distance sampling, repeated counts, removal sampling and double observer sampling (corresponding to *unmarked*'s hierarchical modelling functions of *distsamp*, *pcount* and *multinomPois*).

2. Access CEAMEC

Users can run *CEAMEC* online in the Shiny Cloud:

<https://qt37t247.shinyapps.io/CEAMEC/>

Otherwise, users can run *CEAMEC* through *R* on a local device after installing from GitHub:

<https://github.com/qt37t247/CEAMEC>

Source code, example data files and *R* scripts for the demonstration of non-interactive *CEAMEC* run (without using UI) are available from GitHub.

3. The UI

UI of *CEAMEC* (see Figure 1) comprises three tabs: the “Field data input” tab for survey data input and hierarchical modelling, the “CEAMEC” tab for density visualization, cost-effectiveness analysis and results output, and “Help” tab for the access to useful links. Under the “field data input” tab, there are three sub-tabs (namely “Distance sampling”, “Repeated count” and “Removal sampling or double observer sampling”) corresponding different types of population surveys and different hierarchical

modelling methods. Under each sub-tab, there are three sections (titles of sections highlighted with brown bold font): “**Survey information**”, “**Modelling with covariates**” and “**Models with covariates**”. Detailed explanations of all items in the UI are listed in the [tooltips section](#) at the end of the manual.

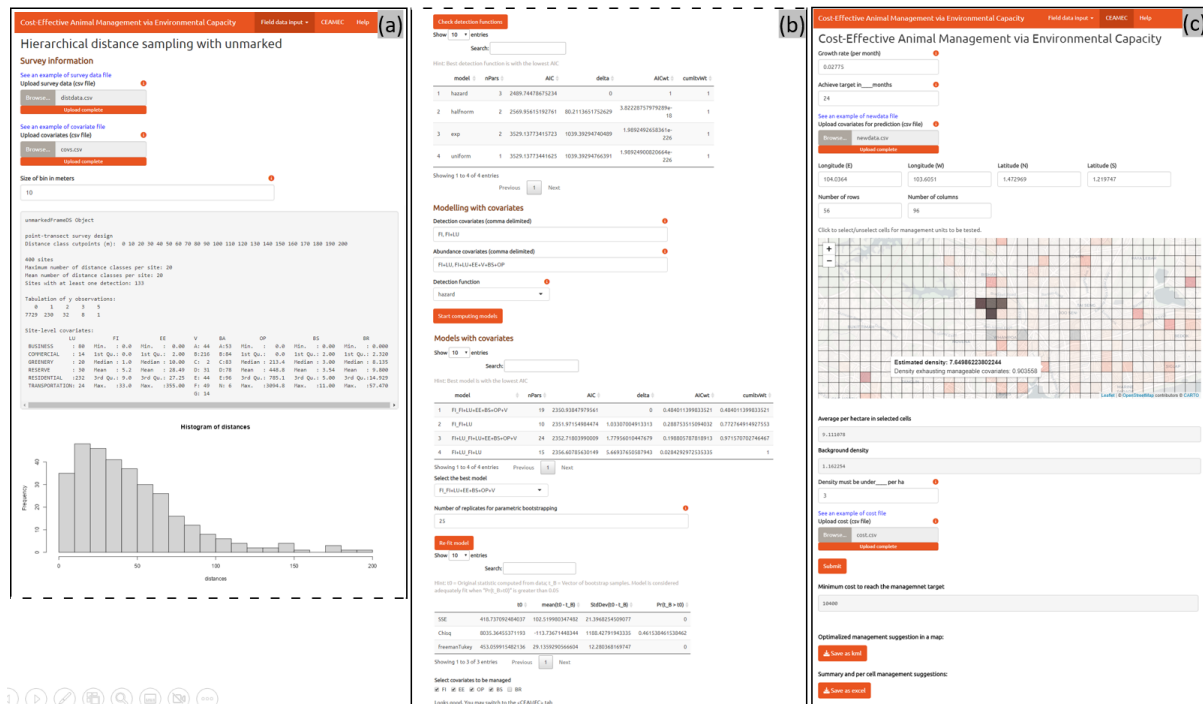


Figure 1. Screenshot of fully executed *CEAMEC* UI with the demonstration dataset of pigeons in Singapore. (a) and (b) present interface under the “Field data input” tab (Distance sampling sub-tab); (c) presents interface under the “CEAMEC” tab.

4. Workflow

In general, *CEAMEC* intakes population survey data and the covariates, the variables collected along with the survey, to compute the correlation between abundance and covariates using hierarchical modelling implemented in *R* package *unmarked* (Fiske & Chandler, 2011). The model, which best describes the correlation between the abundance and the environment, is then used to compute the change of carrying capacity, which is calculated with user-defined management target. Based on the change of carrying capacity and the unit monetary costs provided by users for the management of different resources, *CEAMEC* applies genetic algorithm to compute cost-effectiveness for the combinations of resource reduction (resources correspond to covariates).

the units. For example, instead of using “0.43” kilometer, users may need to use “430” meters. Or, instead of using “0.652”, users may need to use “65.2” percent for the environmental variables in proper fractions.

CEAMEC intakes a variety of population survey data and computes hierarchical models with the corresponding functions of *unmarked*. As hierarchical models aim to explore the correlation between the abundance of species and the environmental context, it is always ideal to acquire descent number of environmental variables, especially the ones users think to contribute to the high density of the targeted species. Moreover, it is recommended to use multi-session surveys across seasons for species display significant seasonal behavior.

4.2. Survey data input

For [distance sampling](#), users need to upload a separate file for the site covariates. Moreover, users need to enter the size to bin distance classes for the detection modelling in the input textbox “[Size of bin in meters](#)”. After inputting the files and parameters, a summary of your survey data and a histogram of data distribution presents. If an error message displayed instead of the summary and histogram, it means that there are formatting error in the input files. Users may need to clear the issue based on what the message suggests and reload the input files to prevent the program from crashing in the subsequent steps. Once the summary of data looks correct, users may proceed to click on the “[Check detection functions](#)” button to see which detection function best describes the observation. A table appears shortly, presenting the comparisons of null models with four detection functions. In the later step, users may need to use the best detection function, with the lowest Akaike information criterion (AIC) value, to create models with the covariates.

For [repeated count](#), [removal sampling and double observer sampling](#), once survey data file is uploaded, UI displays the column names. Users need to check the checkboxes to specify which columns users store the data of “counts”, “site covariates” and “observation covariates” respectively. Users also need to define the area of the survey in the input textbox “[Area of each survey site in hectare](#)”. In addition, for [repeated count](#), users could check which function best describes the abundance distribution by click on the “[Check abundance distribution](#)” button. A table appears shortly, presenting the comparisons of null models with three abundance distributions. In the later step, user may need to use the best abundance distribution, with the lowest AIC value, to create models with the covariates. For [removal sampling and double observer sampling](#), users need to select one from the three survey

types (“Removal sampling”, “Standard double observer sampling” and “Dependent double observer sampling”) respectively.

4.3. Hierarchical modelling

In the “**Modelling with covariates**” section, users need to specify what combinations of covariates are used in modelling detection (can be both observation and site covariates, input textbox “[Detection covariates \(comma delimited\)](#)”) and abundance (site covariates only, input textbox “[Abundance covariates \(comma delimited\)](#)”) respectively. Please do not proceed if a message appears indicating inputting error. Double check if the covariates names used in the model names are consistent with the covariates names in the survey data file (*CEAMEC* is case-sensitive). For distance sampling, users need to select the detection function, which is normally the one with lowest AIC value from the four detection functions computed in the previous section. For repeated count, users need to select the latent abundance distribution, which is normally the one with lowest AIC value from the three abundance distributions computed in the previous section. Click the button “[Start computing models](#)” to start while users are unable to access the UI until computing is finished.

4.4. Model selection and manamgenet method design

Once the hierarchical modelling with all the listed combinations of covariates is finished, a table presents at the top of the “**Models with covariates**” section. According to the table, users may select the best model, normally the one with lowest AIC value. *CEAMEC* also provides an option for users to validate the model of choice by inspecting the distribution of simulated refitted dataset. Users may input a integer number (>10 is recommended but larger number leads to longer waiting time) in the input textbox “[Number of replicates for parametric bootstrapping](#)” and hit the “[Re-fit model](#)” button. After bootstrapping, a table presents three sets of statistics, sum of squared errors (“SSE”), Pearson’s Chi-squared (“Chisq”), and Freeman-Tukey Chi-squared (“freemanTukey”), comparing observed data with simulated refitted data. Ideally, we consider the model is adequately fit when the observed data (t_0) is not significantly different (“ $\Pr(t_B > t_0)$ ” is greater than 0.05) from the simulated data (t_B). The abundance model in the model of choice is used for the subsequent analyses of density estimation and cost-effectiveness computation. The covariates in the best abundance model are the determinant covariates (Figure 3). Considering that resources in some of the determinant covariates are not easily identified or easily managed, users may subset the determinant covariates for the management method designs. In *CEAMEC*, users need to check the checkboxes displaying all available site covariates, to tell the program which are chosen for the covariates to be managed (Figure 3).

CEAMEC only allow numeric covariates to be managed, if there are categorical determinant covariates that may contribute essentially to the species' density, please consider converting them into numeric. For example, instead of using vegetation categories as a site covariate, users may consider using the area or proportion of a specific vegetation category. The determinant covariates not to be managed are still included in the density estimation and subsequent analyses as contributors of background density.

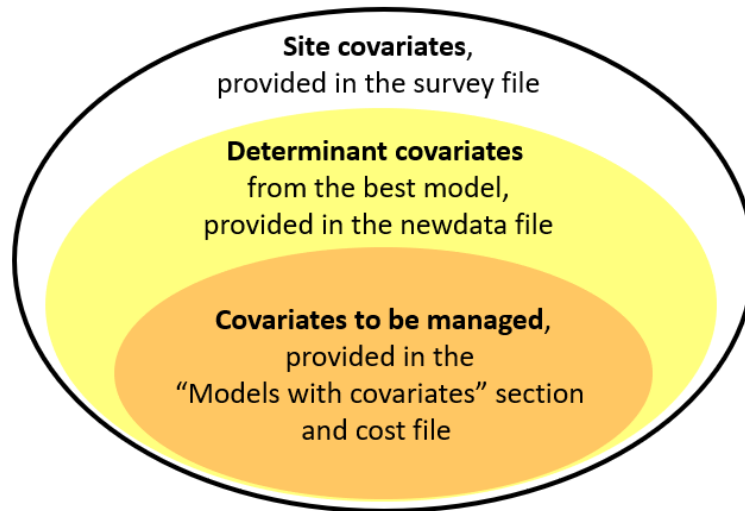


Figure 3. Graphic explanation of relations among covariates used in *CEAMEC*.

To design appropriate management methods, we have presented [a case study of pigeons in Singapore in the following section](#) as an example on how to design management methods corresponding with specific determinant covariates and calculate unit costs for each management method. In general, to calculate the cost of managing a covariate, *CEAMEC* adopt a linear equation with respect to period of management (t) and changes of covariates (Δx):

$$V = a\Delta x t + b\Delta x + ct + d$$

where a is cost per unit of the covariate per unit time, b is cost per unit of the covariate, c is cost per unit time, and d is fixed cost. Unit costs (a , b , c and d) of all covariates to be managed need to be stored in a csv file (the [cost file](#)) and to be uploaded for the cost-effectiveness compulation in the later step.

4.5. Density estimation and management unit selection

After selecting covariates to be managed, users may switch to the “*CEAMEC*” tab. First, users need to input the [growth rate](#), which is often estimated from reproductive biology studies of the species. Also, users need to specify the [period of management](#) here. To estimate the population density of the area of study, users need to provide a raster of environmental variables over the area of interest in a

[newdata file](#). In addition to uploading the newdata file, users also need to provide the extent and dimension of the raster. The resolution of the environmental variables used in the newdata file for pre-management density estimation defines the basic spatial unit, the management unit, in the *CEAMEC* analyses. *CEAMEC* considers each management unit, where an estimated density value is given, as a closed system in which dynamics of the population is subjected to the logistic population growth model with negligible exchange of members to adjacent spatial units. Subsequently, the cost-effectiveness and the management suggestions are calculated independently for each management unit.

Once the newdata file and extent and dimension of the raster are provided, UI presents a map of density estimation. Estimated density are considered as pre-management density. The pre-management density is the species density before the planning management project, which may have been already subjected to extant management effort. The [map](#) allows users to select the management units for the subsequent optimization process.

4.6. Cost-effectiveness computation

Once management units are selected, users can see the average density and the background density of the selected cells, refer to which, users may input the post-management density in the input textbox “[Density must under _____ per ha](#)”. Finally, users need to upload the [cost file](#), which is prepared during the management design. Warning message appears if there are formatting error in the [cost file](#). Users may need to clear the issues according to the message before hitting the “Submit” button. The process takes approximately one minutes per management unit selected. Once the process finished, the total cost of management for all selected management units is displayed. Users can download the map, in the kml format, and visualize the optimal management suggestions for the selected management units with Google Earth or other GIS tools. In addition, users can download an excel sheet of multiple tabs, in which the first tab records the summary of the most cost-effective managment suggestions for the selected management units whereas subsequent tabs (one management unit per tab) record the comparisons between the optimal management suggestions with other managment scenarios.

5. Example data demonstration

We have provided an example dataset for the demonstration of using *CEAMEC*. The dataset is modified from a case study of feral pigeon population modelling in Singapore based on distance sampling survey carried out in 2016 (Tang et al., 2018) with additional covariates. The timing provided were measured on a laptop of six-core, Intel i7 (2.6GHz) 12-processor with 16GB RAM.

Field data input tab (Distance sampling sub-tab)

Survey information

We uploaded the distance sampling survey data file, [distdata.csv](#) (file description see appendix 8.1.), in the fileinput “[Upload survey data \(csv file\)](#)”. We uploaded the coveriates with the file [covs.csv](#) (file description see appendix 8.1.) in the fileinput “[Upload covariates \(csv file\)](#)”. As we put “0” for the “length” in the [covs.csv](#), CEAMEC automatically identify that we used point transects for the survey. We entered “10” for the input textbox “[Size of bin in meters](#)” to generate discrete distance classes for modelling the correlation between detection and distance. We checked that the summary and the data distribution (see Figure 1a) and clicked on button “[Check detection functions](#)” to run four null models with different detection functions (the process took ~18 seconds). We confirmed that the detection of pigeons in this distance sampling case study was best described with a “hazard” model (based on the lowest AIC value across all four detection functions, see Figure 1b).

Modelling with covariates

We entered two combinations “FI, FI+LU” in the input textbox of “[Detection covariates \(comma delimited\)](#)” and two combinations “FI+LU, FI+LU+EE+BS+OP+V” in the input textbox “[Abundance covariates \(comma delimited\)](#)” (Figure 1b). CEAMEC generated below four models for best model computation (please refer to the description of [covs.csv](#) in the appendix 8.1. for the of expansion of covariates’ names):

1. FI_FI+LU: FI corresponds detection while FI and LU correspond abundance;
2. FI+LU_FI+LU: FI and LU correspond detection while FI and LU correspond abundance;
3. FI_FI+LU+EE+BS+OP+V: FI corresponds detection while FI, LU, EE, BS, OP and V correspond abundance;
4. FI+LU_FI+LU+EE+BS+OP+V: FI and LU correspond detection while FI, LU, EE, BS, OP and V correspond abundance.

We selected “hazard” in the dropdown menu “[Detection function](#)” as it best described the field observations as shown in the result of “[Check detection functions](#)”.

Models with covariates

After model computation (the process took ~16 minutes), models were sorted by ascending AIC values in the table generated (Figure 1b). We selected “FI_FI+LU+V+EE+BS+OP” in the dropdown menu “[Select the best model](#)” as it exhibited the lowest AIC value. The model consists of a detection model and an abundance model, whose covariates are connected by “_”. The name of the best model also

suggest that the detectability of pigeons is likely correlated to the number of feeding incidents (FI), whereas the abundance of pigeons is likely correlated to six covariates (FI, LU, V, EE, BS and OP). The determinant covariates identified are “FI” (number of feeding incidents), “LU” (land use types), “V” (vegetation types), “EE” (number of eating establishments), “BS” (number of bus stops) and “OP” (length of overpasses).

We input “25” in the input textbox “[Number of replicates for parametric bootstrapping](#)” to simulate data with the chosen model for 25 times and compare with the observed data to check adequacy of model fit. Then we clicked on button “[Re-fit model](#)” to proceed the simulation (the process took ~16 minutes). As results, a table lists statistics of comparisons between simulated data and observed data is generated after bootstrapping (see Figure 1b). We decided to proceed with the model (a model with $\Pr(t_B > t_0)$ is greater than 0.05 is consider adequately fit).

We checked four covariates to be managed (“FI”, “EE”, “BS”, “OP”) in the checkboxes under “[Select covariates to be managed](#)”. We chose these four from the six determinant covariates for two reasons: first, they are directly associated with resources that support the population density of pigeons in Singapore: “FI”, “EE” and “BS” are associated with food sources whereas “OP” is associated with sheltered roosts; second, they make it straightforward to demonstrate the relationship between the quantity of covariates and the quantity of resources during [management method design](#) (please read the details of manamgenet method design in response to the quantity of covariates and estimation of costs in the appendix 8.2.).

CEAMEC tab

After switching to the “CEAMEC” tab, we set “0.02775” in the input textbox “[Growth rate \(per month\)](#)” for pigeons as suggested by previous studies (Johnston and Janiga, 1995). This growth rate is based on the assumption that around one third of pigeons in the population breeds every year; that each pair produces an average of five fledged offspring per year; and that around half of the population dies every year. We set “24” in the input textbox “[Achieve target in months](#)” for period of management. We uploaded [newdata.csv](#) (file description see appendix 8.1.) in the fileinput “[Upload covariates for prediction \(csv file\)](#)”. Finally, we set the study area to encompass the entire Singapore with geographic limits at 104.0364°E to 103.6051°E (east-west) and 1.472969°N to 1.219747°N (north-south). We rasterized the study area into 56 (rows) × 96 (columns) raster cells (500m × 500m for each raster cell). A [map](#) presented displaying estimated densities across all management units (Figure 1c). Redder colour hues imply a higher pigeon density in each management unit and vice versa. Hovering over a

management unit with the cursor triggers a pop-up that displays the pre-management density and the background density of the management unit. As we only declared four of the six determinant covariates as being subject to management and manipulation, the remaining two determinant covariates (“LU” and “V”) contributed to the background density, which is the minimum pigeon density the management unit can reach when all the four covariates available for manipulation are being made exhaustive use of. In this example application, we selected four management units with a relatively high pigeon density (average of ~9 pigeons per hectare, see output textbox **“Average per hectare in selected cell”** in Figure 1c) and background density of ~1 pigeons per hectare (see output textbox **“Background density”** in Figure 1c). We input “3” in the input textbox **“Density must under per ha”** to set out reducing the density below three pigeons per hectare for all four selected management units. We uploaded the [cost.csv](#) (file description see appendix 8.1.) and hit the **“Submit”** button to initiate the cost-effectiveness computation (the process took ~6 minutes).

In the end, *CEAMEC* produced an optimal management plan that entails a cost of 10,400 Singapore dollar (see output textbox **“Minimum cost to reach the management target”** in Figure 1c) to reduce pigeons to fewer than three per hectare within 24 months for the four management units.

We clicked **“Save as kml”** and downloaded the kml file to view the detailed management suggestions for each management unit. The kml file can be opened in Google Earth and visualized as polygons (a management unit per polygon) over the map (Figure 4a). In the pop-up table above the clicked management unit, *CEAMEC* lists the management unit ID (“layer”), management costs, pre- and post-management densities of pigeons and the quantity of covariates to be managed. The example result can be interpreted as follows: in the management unit (cell number: 2838), 3625 Singapore dollar must be spent for averting 15 feeding incidents and installing warning signs at five bus stops in order to reduce pigeons from more than 12 per hectare to less than three per hectare in two years. We clicked **“Save as excel”** and downloaded the Excel file for the summary of optimal management suggestions across all selected management units in the first tab (Figure 4b) and per management unit comparison between the best management suggestion and other, financially less optimal combinations of management methods in the subsequent tabs (Figure 4c).

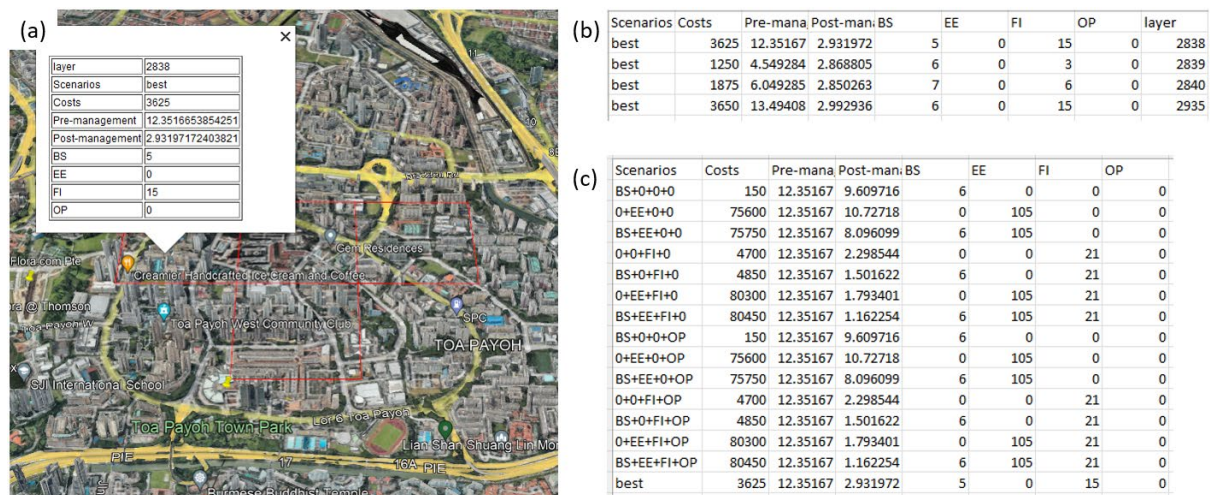


Figure 4. Results from the demonstration run of *CEAMEC*. (a) screenshot of the kml file opened in Google Earth, (b) summary table of optimal methods, and (c) per management unit table of methods comparisons.

6. Tooltips

6.1. Field data input tab

6.1.1. Distance sampling sub-tab

CEAMEC adopts hierarchical distance sampling functions from the *R* package *unmarked*, in-depth explanations of distance sampling survey input file structure and parameters can be found in the *unmarked* documents in the links below:

<https://rdrr.io/cran/unmarked/f/inst/doc/distsamp.pdf>

<https://rdrr.io/cran/unmarked/man/unmarkedFrameDS.html>

Survey information section

Upload survey data (csv file): A csv format data frame where each row is a detected individual. Must have two columns. One (named “distance”) for distances to the detected individuals and the other (named “transect”) for transect names.

Example file in GitHub: [distdata.csv](#)

Upload covariates (csv file): A csv format data frame of environmental variables (covariates) that vary at the site level. Number of rows must match number of transects. Number of columns should equal to number of covariates with one column per covariate. An additional column (named “length”) should be attached as the last column for the length of transects (in the unit of meter, put 0 if using point transects).

Example file in GitHub: [covs.csv](#)

Size of bin in meters: Distance data is binned into discrete distance classes with the size provided.

Check detection functions: Run null models (without covariates) across four detection functions and identify which function best describes the correlation between distance with probability of detection. The detection process is modeled as multinomial: $y_{ij} \sim \text{Multinomial}(N_i, p_{i1}, p_{i2}, \dots, p_{iJ})$, where p_{ij} is the multinomial cell probability for transect i in distance class j . These are computed based upon a detection function $g(x / \sigma)$, such as the half-normal (“halfnorm”), negative exponential (“exp”), uniform (“uniform”) or hazard rate (“hazard”). Output a table comparing the four null models.

Modelling with covariates section

Detection covariates (comma delimited): List all covariates or combinations of covariates to model detection (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Abundance covariates (comma delimited): List all covariates or combinations of covariates to model abundance (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Detection function: Select from one of the four detection functions: “halfnorm”, “hazard”, “exp”, or “uniform”. You could choose based on the result table if you ran check detection functions (normally the detection function with lowest AIC).

Start computing models: Compute models with all combinations of detection covariates and abundance covariates. If you input x combinations of covariates in detection covariates textbox and y combinations of covariates in abundance textbox, this step will produce $x \times y$ models. Each model includes two formulas for the covariates’ correlation with detection probability and abundance respectively. At the end of modelling, a table will be generated to compare the statistics of all the models.

Models with covariates section

Select the best model: Select one of the models computed with covariates, normally the model with lowest AIC.

Number of replicates for parametric bootstrapping: Number of bootstrap replicates (must be integer) to check adequacy of model fit. Can be time consuming (>1 hour) if a large number (>100) is chosen.

Re-fit model: This step simulates datasets based upon the chosen best model, refits the model, and evaluates a user-specified fit-statistic for each simulation. Comparing this sampling distribution to the observed statistic provides a means of evaluating goodness-of-fit or assessing uncertainty in a quantity of interest.

Select covariates to be managed: Check the checkboxes for the corresponding covariates as covariates to be managed.

6.1.2 Repeated count sub-tab

CEAMEC adopts hierarchical modelling functions for repeated count survey from the *R* package *unmarked*, in-depth explanations of repeated count survey input file structure and parameters can be found in the *unmarked* documents in the links below:

<https://rdr.io/cran/unmarked/man/unmarkedFramePCount.html>

<https://studylib.net/doc/6696451/fitting-royle-s-n-mixture-model-with-package-unmarked-in-...>

Survey information section

Upload repeated count data (csv file): A csv format data frame of the repeated count data with observation and site covariates appended. A transect per row. Columns contains counts (one session per column), observation covariates (one session per column) and site covariates (one covariate per column). Different sessions can be identified with '#' (e.g. '.1', '.2', '.3') in the column names.

Example file in GitHub: `mld_pcount.csv`

Column names for counts: Check the names of columns contain counts of all sessions. The checkboxes appear on UI after uploading the data file.

Column names for site covariates: Check the names of columns contain site covariates. The checkboxes appear on UI after uploading the data file.

Column names for observation covariates: Check the names of columns contain observation covariates of all sessions. The checkboxes appear on UI after uploading the data file.

Area of each survey site in hectares: Normally to be the size of the transect. But if using traps, you may need to estimate the area that the trap may cover.

Check abundance distribution: Run null models (without covariates) across three abundance distributions and identify which best describe the abundance distribution. The latent abundance distribution, $f(N \mid \theta)$ can be set as a Poisson, negative binomial, or zero-inflated Poisson random variable, depending on the setting of the mixture argument, mixture = "P", mixture = "NB", mixture = "ZIP" respectively. For the first two distributions, the mean of N_i is λ_i . If $N_i \sim NB$, then an additional parameter, α , describes dispersion (lower α implies higher variance). For the ZIP distribution, the mean is $\lambda_i(1-\psi)$, where ψ is the zero-inflation parameter. Output a table comparing the three null models.

Modelling with covariates section

Detection covariates (comma delimited): List all covariates or combinations of covariates (can be either/both observation and site covariates) to model detection (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Abundance covariates (comma delimited): List all covariates or combinations of covariates (site covariates only) to model abundance (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Latent abundance distribution: Select one of the three mixture: 'P' (Poisson), 'NB' (negative binomial) or 'ZIP' (zero-inflated Poisson). You could choose based on the table produced by “**Check abundance distribution**” (normally the mixture with lowest AIC).

Start computing models: Compute models with all combinations of detection covariates and abundance covariates. If you input x combinations of detection covariates and y abundance covariates, this step will produce $x \times y$ models. Each model includes two formulas for the covariates' correlation with detection probability and abundance respectively. At the end of modelling, a table will be generated to compare the statistics of all the models.

Models with covariates section

Select the best model: Select one of the models computed with covariates, normally the model with lowest AIC.

Number of replicates for parametric bootstrapping: Number of bootstrap replicates (must be integer) to check adequacy of model fit. Can be time consuming (>1 hour) if a large number (>100) is chosen.

Re-fit model: This step simulates datasets based upon the chosen best model, refits the model, and evaluates a user-specified fit-statistic for each simulation. Comparing this sampling distribution to the observed statistic provides a means of evaluating goodness-of-fit or assessing uncertainty in a quantity of interest.

Select covariates to be managed: Check the checkboxes for the corresponding covariates as covariates to be managed.

6.1.3 Removal sampling or double observer sampling sub-tab

CEAMEC adopts hierarchical modelling functions for removal sampling or double observer sampling survey from the *R* package *unmarked*, in-depth explanations of survey input file structure and parameters can be found in the *unmarked* documents in the links below:

Removal sampling:

<https://rdr.io/cran/unmarked/man/ovendata.html>

Double observer sampling:

<https://rdr.io/cran/unmarked/man/unmarkedFrameMPois.html>

Survey information section

Upload survey data (csv file): A csv format data frame of the count data with observation and site covariates appended. A transect per row. Columns contains counts (one session per column), observation covariates (one session per column) and site covariates (one covariate per column). Different sessions can be identified with '#' (e.g. '.1', '.2', '.3') in the column names.

Example removal sampling data file in GitHub: oven_removal.csv

Example double observer sampling data file in GitHub: fake_double.csv

Column names for counts: Check the names of columns contain counts of all sessions. The checkboxes appear on UI after uploading the data file.

Column names for site covariates: Check the names of columns contain site covariates. The checkboxes appear on UI after uploading the data file.

Column names for observation covariates: Check the names of columns contain observation covariates of all sessions. The checkboxes appear on UI after uploading the data file.

Survey type: Select one from three: removal sampling, standard double observer sampling, and dependent double observer sampling.

Area of each survey site in hectares: Normally to be the size of the transect. But if using traps, you may need to estimate the area that the trap may cover.

Modelling with covariates section

Detection covariates (comma delimited): List all covariates or combinations of covariates to model detection (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Abundance covariates (comma delimited): List all covariates or combinations of covariates to model detection (comma delimited). A combination of covariates is written as the names of covariates connected with '+'. Consider non-interactive run if many combinations are tested, otherwise very time-consuming.

Start computing models: Compute models with all combinations of detection covariates and abundance covariates. If you input x combinations of detection covariates and y abundance covariates, this step will produce $x \times y$ models. Each model includes two formulas for the covariates' correlation with detection probability and abundance respectively. At the end of modelling, a table will be generated to compare the statistics of all the models.

Models with covariates section

Select the best model: Select one of the models computed with covariates, normally the model with lowest AIC.

Number of replicates for parametric bootstrapping: Number of bootstrap replicates (must be integer) to check adequacy of model fit. Can be time consuming (>1 hour) if a large number (>100) is chosen.

Re-fit model: This step simulates datasets based upon the chosen best model, refits the model, and evaluates a user-specified fit-statistic for each simulation. Comparing this sampling distribution to the observed statistic provides a means of evaluating goodness-of-fit or assessing uncertainty in a quantity of interest.

Select covariates to be managed: Check the checkboxes for the corresponding covariates as covariates to be managed.

6.2 **CEAMEC** tab

Growth rate (per month): Growth rate can be estimated from reproductive experiments or field observations. If using annual growth rate please divide by 12.

Achieve target in ____ months: Length in months for the period of management.

Upload covariates for prediction (csv file): A csv format data frame containing environmental variable of rasterized area of study. Each row represents each cell in the rasterized study area with cell ID indicated at the first column. All determinant covariates should be included with one covariate per column starting from the second column.

Example file in GitHub: newdata.csv (for distance sampling only)

An interactive map, covered by raster cells, visualizes density estimated with the best model. Hovering over a cell with the cursor triggers a pop-up that displays the estimated density and the background density, the minimum density can reach when all covariates to be managed are being made exhaustive use of. Single left mouse click on a management unit to select/unselect it (multiple management units can be selected for the subsequent cost-effective calculation).

Average per hectare in selected cells: average density of animals of the selected management units.

Background density: minimum density can reach when all covariates to be managed are being made exhaustive use of. Normally contributed by environmental conditions not included in the hierarchical

model or determinant covariates not manageable. If multiple management units are selected, the value is set to the management unit of highest background density.

Density must under ____ per ha: Don't set this value lower than the background density suggested in the output textbox “**Background density**”.

Upload cost (csv file): A csv format data frame with one covariate to be managed per row and one of the [four unit costs](#) per column (in the order of a, b, c, d).

Example file in GitHub: [cost.csv](#)

7. References

- Fiske, I., & Chandler, R. (2011). *Unmarked*: An R package for fitting hierarchical models of wildlife occurrence and abundance. *Journal of Statistical Software*, 43(10), 1–23.
- Johnston, R. F., & Janiga, M. (1995). *Feral pigeons* (Vol. 4). Oxford University Press on Demand.
- Tang, Q., Low, G. W., Lim, J. Y., Gwee, C. Y., & Rheindt, F. E. (2018). Human activities and landscape features interact to closely define the distribution and dispersal of an urban commensal. *Evolutionary Applications*, 11(9), 1598–1608.

8. Appendix

8.1. Input files for the demonstration run on the case study of pigeons in Singapore

CEAMEC adopts hierarchical distance sampling functions from the R package *unmarked*, in-depth explanations of distance sampling survey input file structure and parameters can be found in the *unmarked* documents in the links below:

<https://rdr.io/cran/unmarked/f/inst/doc/distsamp.pdf>

<https://rdr.io/cran/unmarked/man/unmarkedFrameDS.html>

1. **distdata.csv:** a file stores distance sampling survey data as two-column data frame, where each row is a detected individual. The first column is for distances between the observer and the detected individual; the second column is for transect names.

2. **covs.csv:** a file stores site covariates (one covariate per column) and the lengths of transects (at the last column named “length”; input “0” if using point transects) that vary at the site level (one site per row starting from the second row). The first row is for the names of the covariates (we suggest to use

abbreviation to shorten the names of models in the subsequent modelling) and “length”. In the pigeon study we demonstrated, we collected eight covariates: LU is for landuse categories; FI is for number of feeding incidences; EE is for number of eating establishments; V is for vegetation categories; BA is for building age categories; OP is for the length of over passes in meters; BS is for number of bus stops; BR is for the length of bus route in kilometers.

3. **newdata.csv**: a file stores environmental variables (one variable per column starting from the second column) across all management units of Singapore (one management unit per row starting from the second row). The first row is for the names of the covariates, which should be consistent, or subset, with the names in the covs.csv. The first column is for the cell IDs of the management units.

4. **cost.csv**: a file stores unit costs of management methods in response to covariates to be managed (one covariate per row) as five-column data frame. The first row are the names of the unit costs (a, b, c, d in response to different variables, see the [detailed explanation](#) of cost calculation in the workflow section). The first column is the the names of covariates to be managed.

8.2. Management method design and cost estimation on the case study of pigeons in Singapore

The following outlines the costs associated with each of these four management methods:

(1) To reduce feeding incidents (FI), we proposed a policy whereby persons engaging in illegal pigeon feeding (‘feeders’) are identified, approached and educated by management personnel. For each management unit, the resultant cost comprises a fixed cost (*d*) of \$500 for the investigation over the entire management unit and a cost per feeding incident (*b*) of \$200 for visiting and educating a feeder to avert one feeding incident.

(2) To reduce food sources generated by eating establishments (EE), we proposed a management plan to combine regular inspections with the disposal of exposed food waste. For each management unit, the resultant cost comprises a cost per eating establishment per month (*a*) of \$30 covering administrative fees and disposal costs.

(3) To reduce the food sources (through feeding or littering) generated at or near bus stops (BS), we proposed to install “no feeding/littering” signs and warnings that such behaviour will incur fines

when caught by surveillance cameras at bus stops. The resultant cost comprises a cost per bus stop (*b*) of \$25 for sign installations.

(4) To reduce roosts beneath overpasses (OP), we proposed to install nets to deter pigeon entry into crevices and expansion gaps. For each management unit, the resultant cost comprises a cost per meter of overpass (*b*) of \$24 for net installation and a cost per meter of overpass per month (*a*) of \$0.12 for net maintenance.

We generated the cost file with rows of selected covariates to be managed and columns of unit costs (*a*, *b*, *c* and *d*, see [equation](#) in the workflow section).