

# oSCR :: CHEAT SHEET



The oSCR package, pronounced “Oscar”, provides a set of functions for working with Spatial Capture Recapture (SCR) models.

## Getting the package

Package hosted on [GitHub](#)

```
library(devtools)
install_github("jaroy1e/oSCR")
library(oSCR)
```

## Workflow

- Every model you run on oSCR has the following 4 basic steps.
- Modeled after [unmarked](#) workflow

### 1. Format the sampling data

One file for each one:

- Spatial encounter histories
- Detector information

### 2. Define and format the State Space

- Size and resolution of the *state space*
- Spatial covariates for density

### 3. Analyze the data - model fitting

- Likelihood based: use AIC to do model selection
- No need to use other packages, oSCR has helper functions to do the model selection.

### 4. Post processing model output for inference:

- This means that now that you have your parameters all you have to do is interpret your results!

## Modelling framework

### A. Single-session models

- Repeated sample occasions on a single population of individuals using a single array of traps.

### B. Multi-session models

- Data grouped in strata or groups which are independent in space or time.

### C. Explicit sex-structured models

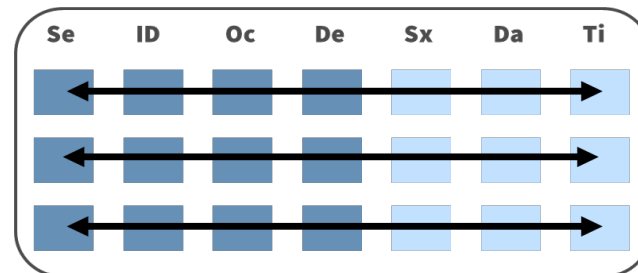
### D. Multi-session sex-structured models

## 1. Format sampling data

Before starting to use oSCR you need to format the datafiles in a scrFrame which consists of two basic spreadsheets: **edf** and **tdf**.

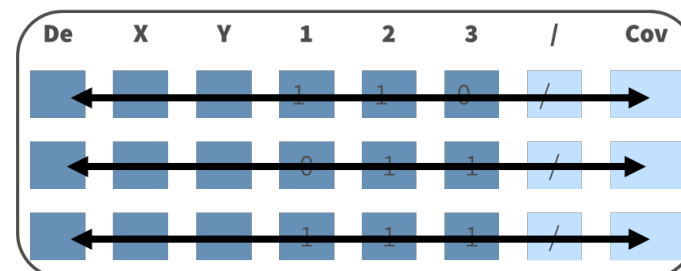
### 1.1 edf: encounter data file.

- Single **data frame**.
- Each row has individual detection events.
- Dark blue = required; light blue = optional.
- Columns contain capture information:
  - Session (Se)                      - Sex (Sx)
  - Individual ID (ID)               - Date (Da)
  - Occasion (Oc)                   - Time (Ti)
  - Detector\* (De)



### 1.2 tdf: trap deployment data file.

- A **list** with information for each session (tdf1, tdf2, ...).
- Each row is a trap.
- Columns contain trap information
  - Detector\* (De)                      1, 2, 3, ... n)
  - X (required, UTM)               - Separator (e.g., /)
  - Y (required, UTM)               - Trap level covariates
  - Binary trap operation            (different column per covariate)



\*Notice that both edf and tdf have the same **Detector (De)** column that **MUST** match (same name, class, relational database).

### 1.3 data2oscr(): is a function that links edf and tdf files via the detector\* names. Creates scrFrame.

```
data <- data2oscr(
  edf, # encounter data file
  tdf, # list containing trap deployment file
  sess.col*, # session col number or name in edf
  id.col*, # individual ID col # or name in edf
  occ.col, # occasion col number or name in edf
  trap.col*, # detector col number or name in edf
  sex.col*, # sex col number or name in edf
  sex.nacode, # character for unknown sex in edf
  K, # number of occasions
  ntraps, # number of traps
  trapcov.names, # vector of un-numbered cov names
  tdf.sep) # separator (e.g., "/")
```

\* `which(colnames(edf) %in% "name of column in edf")`

### 1.4 Summary functions for scrFrame :

- scrFrame contains information from the **edf** and **tdf** via detector names.

`sf <- data$scrFrame`

<b>sf\$caphist</b>	Array of individual-by-trap-by-occasion (n x J x K). Binary or counts.
<b>sf\$traps</b>	Data frame containing at least trap ID and coordinates of traps. Best with UTM.
<b>sf\$indcovs</b>	Sex data (0 female, 1 male) or any bivariate covariate. NAs allowed.
<b>sf\$trapCovs</b>	List of session specific trap covariates. Row per trap, and column per covariate.
<b>sf\$sigCovs</b>	A data frame of covariates that affect space use (sigma, $\sigma$ ).
<b>sf\$trapOperation</b>	A list of session specific information on trap operational data.
<b>sf\$occasions</b>	A vector of number of occasions per session .
<b>sf\$mmdm</b>	Mean maximum distance moved pooled across sessions. $\frac{1}{2} \text{mmdm} \sim \sigma$
<b>sf\$mdm</b>	Maximum distance moved pooled across sessions.
<b>\$telemetry</b>	Telemetry object for fitting resource selection models.

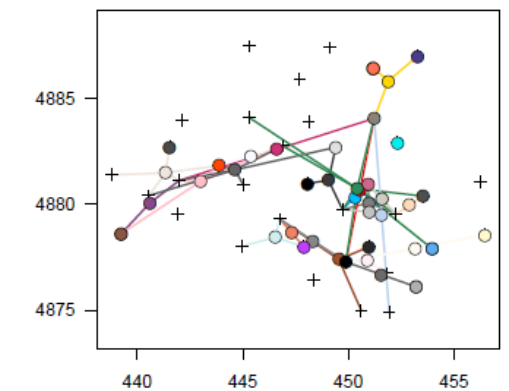
### 1.5 Summary of scrFrame sf

```
S1
n individuals 47
n traps       38
n occasions   8

S1
avg caps      3.21
avg spatial caps 2.02
mmdm         4.65
```

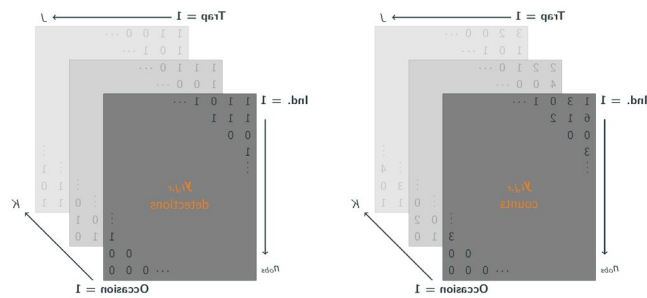
### 1.6 Spatial captures per session

`plot(sf)` #y and x are UTM





## 1.4.1 Navigating the scrFrame



### Capture history

- Session 1, all individuals, all traps, occasion 3  
`sf$caphist[[1]][ , 3]`
- Session 1, individual 4, all traps, all occasions  
`sf$caphist[[1]][4, ]`

### Traps

- Session 1 trap coordinates  
`sf$traps[[1]]`

### Trap covariates

- Trap covariate df session 1 occasion 4  
`sf$trapCovs[[1]][[4]]`

### Trap operation

- Session 1 trap trap operation matrix  
`sf$trapOperation [[1]]`

### Covariates that affect sigma ( $\sigma$ )

- These covariates are NOT session specific. This is a sessions=rows dataframe  
`sf$ sigCovs[[1]]`

### Vectors and single numbers

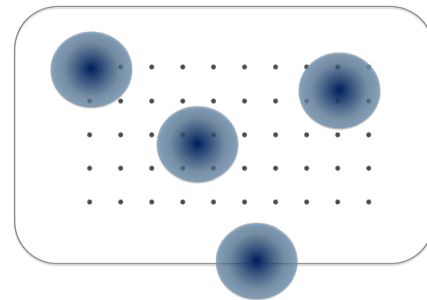
- `sf$ occasions`
- `sf$mmdm`
- `sf$mdm`

## Datasets available

- ```
> data(package = "oSCR")
> data(ocelot)
> data("beardata")
> data("nybears")
> data("peromyscus")
> data("mink")
```

## 2. Create the State Space

The **State Space (S)** is the core element of SCR models. It defines where individuals can live and should represent activity centers of all sampled individuals.



### ssDF: the State Space Data Frame

- List with spatially explicit information from each session.
- At least include the coordinates (X, Y) of the discrete state space (UTM).
- Can include spatial covariates for a continuous state space to study variation in Density.
- Non habitat can be removed by removing unwanted coordinates (e.g., parking lot).

| X | Y | Cov.1 | Cov.2 | Cov.n |
|---|---|-------|-------|-------|
| █ | █ | █     | █     | █     |
| █ | █ | █     | █     | █     |

### 2.1. make.ssDF():

- Remember that  $\frac{1}{2} \text{mmdm} \sim \sigma$
- Extracts covariates and removes non habitat

```
ss <- make.ssDF(scrFrame,
  buffer, #~3 to 4σ around traps
  res) # ≤ σ
```

### 2.2. Plot the state space

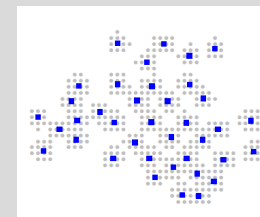
- Plot state space  
`plot(ss)`
- Plot state space & traps  
`plot(ss, sf)`



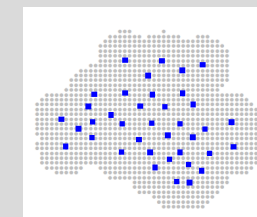
### Vary the buffer and/or resolution

Varying buffer, fixed resolution

```
make.ssDF(sf,
  buffer = 1,
  res = 0.5)
```

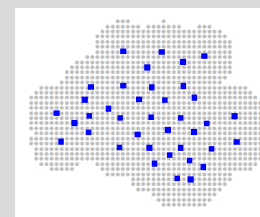


```
make.ssDF(sf,
  buffer = 3,
  res = 0.5)
```

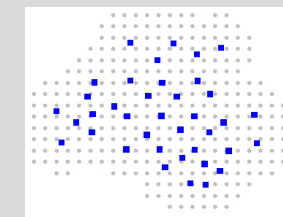


Fixed buffer, varying resolution

```
make.ssDF(sf,
  buffer = 3,
  res = 0.1)
```



```
make.ssDF(sf,
  buffer = 3,
  res = 0.5)
```



## 3. Fit the model

### 3.1. Single-session model: Fit the model with oSCR.fit():

```
sf <- data$scrFrame
mod <- oSCR.fit(model,
  scrFrame, #sf
  ssDF, ...)
```

- See pg. 3 for null model and multi-session models.

**model** is a list with 3 basic formulations:

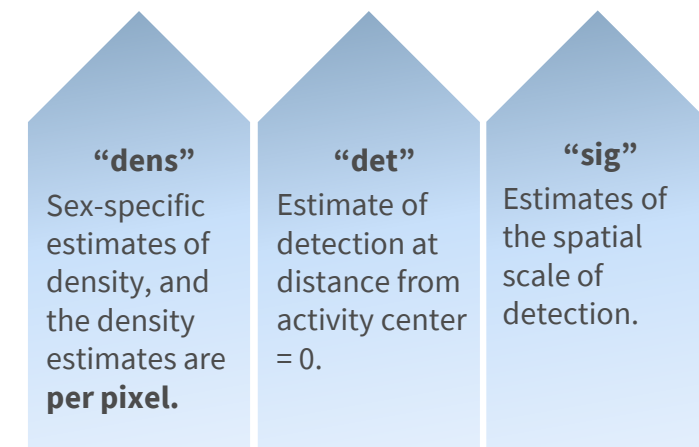
```
list(D ~ 1, p0 ~ 1, sig ~ 1)
```

| Variation in... |                              |
|-----------------|------------------------------|
| <b>D</b>        | pixel density                |
| <b>p0</b>       | baseline encounter prob/rate |
| <b>sig</b>      | sigma ( $\sigma$ )           |

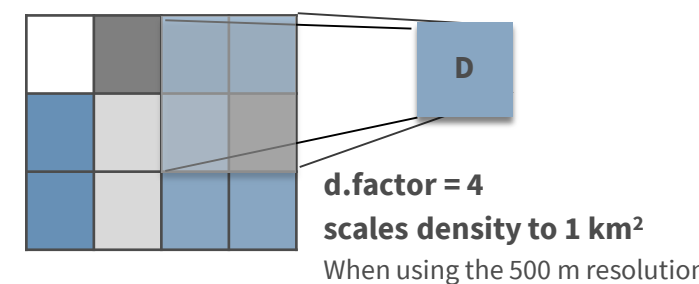
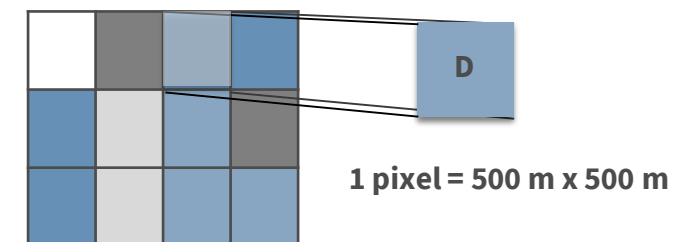
### 3.2. Backtransform to the real scale

```
get.real(model,
  newdata,
  d.factor,
  type)
```

|                 |                                                                |
|-----------------|----------------------------------------------------------------|
| <b>model</b>    | fitted model                                                   |
| <b>newdata</b>  | Optional new data object for predictions                       |
| <b>d.factor</b> | optional scale the estimates to a different resolution         |
| <b>type</b>     | density ("dens"), detection probability ("det"), sigma ("sig") |



### d.factor



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Page 3 describes the specific functions and workflow for the null model and multi-session model in the oSCR package.

## Model specifics

### Null model (SCR<sub>0</sub>)

- The null model assumes homogeneous density which means all pixels have the same expected density.
- For additional arguments see `?oSCR.fit()`

```
mod1 <- oSCR.fit(list(D ~ 1,
p0 ~ 1, sig ~ 1),
scrFrame, #scrFrame object
ssDF, #ssDF object
... ) #other arguments
mod1 #summary
```

If you included sex as a covariate in the scrFrame:

- Sex ratio  $\psi$  () will be included in the summary
- Can compare AIC with and without sex effects



### Multi-session model

Are your data organized in multi-sessions and you want to analyze all of them jointly?



**Spatial sessions:** different study areas (e.g., parks, trapping grids)



**Temporal sessions:** same areas different times (e.g. seasons, years)



Session specific **population size**  $N_g$  ( $g$ =group/session)

- Test for differences among sessions using AIC.
- Can share parameters among sessions or not.

- The **multi-session** model follows similar steps as the single session model.
- The **edf** files from multiple sessions may be merged into one data frame prior to `data2oscr`  
`edf <- rbind(edf1, edf2, ...)`
- The **tdf** files must be separate files for each session.

### 1. data2oscr for multi-session scrFrame

```
data <- data2oscr(
edf, # include session column
list(tdf1, tdf2, ...), # tdf files
sess.col*, # session col in edf
id.col*, # individual ID col in edf
occ.col, # occasion col in edf
trap.col*, # detector col in edf
sex.col*, # sex col in edf
sex.nacode, # unknown sex in edf
K, # vector with occasions per session
ntraps) # vector with traps per session
```

```
sf <- data$sf
sf # summary info per session (S1, S2..)
```

### 1.2. Summary of multi-session scrFrame

```

          S1 S2 S3 S4
n individuals 77 60 108 54
n traps       50 50  50 50
n occasions   7  5  6  4

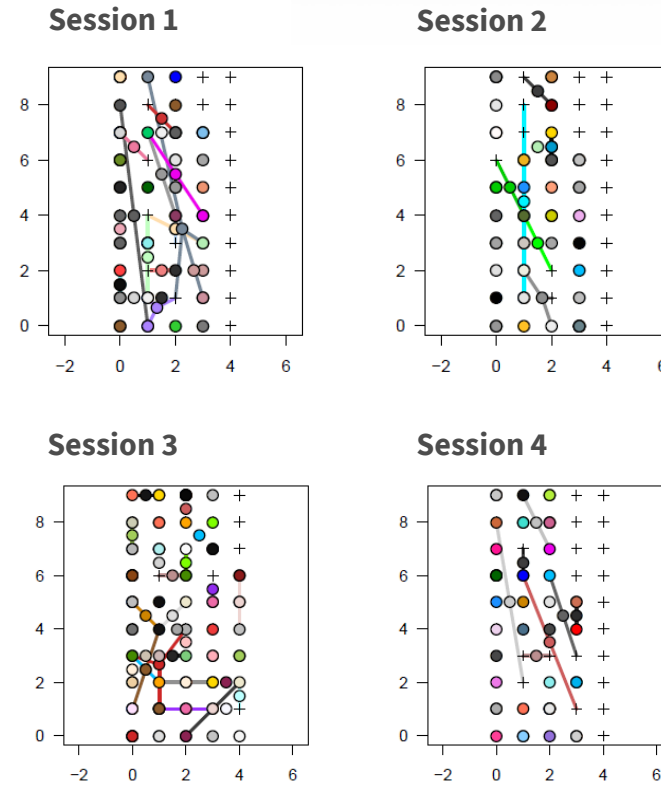
          S1 S2 S3 S4
avg caps    1.91 1.47 1.71 1.37
avg spatial caps 1.30 1.15 1.27 1.13
mmdm        2.57 2.32 1.76 2.84

Pooled MMDM: 2.21
```

### 1.3. Plot spatial captures in a multi-session scrFrame

- Use `plot(sf)` to plot a spatial capture per session

```
par(mfrow=c(1,n)) # n = sessions
plot(sf) # plot all sessions
```



### 2. Make the State Space Data Frame of a multi-session scrFrame

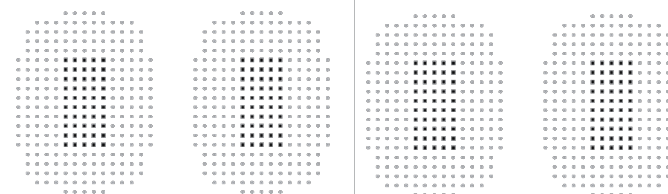
```
ss <- make.ssDF(
scrFrame, # multi-session
buffer, #buffer width
res) #state space resolution
```

- You can vary the buffer and resolution as in the single-session model.

```
?make.ssDF() # Look at the help file for other arguments
```

- Visualize the state space

```
par(mfrow=c(1,n)) # n = sessions
plot.ssDF(ss, # state space
sf) # traps
```



### 3. Model fitting

- Specify models that consider or not variation among sessions.
  - fixed vs. session specific **D**
  - fixed vs. session specific **p0**
  - fixed vs. session specific **space use ( $\sigma$ )**

| Model     | Algebra                                                         | In oSCR.fit                |
|-----------|-----------------------------------------------------------------|----------------------------|
| Density   | $\log(D(s_i)) = \beta_0$                                        | <code>D ~ 1</code>         |
| Density   | $\log(D(s_i)) = \beta_0 + \beta_{1(g)} \text{Session}_g$        | <code>D ~ session</code>   |
| Detection | $\text{logit}(p_0) = \alpha_0$                                  | <code>p0 ~ 1</code>        |
| Detection | $\text{logit}(p_0) = \alpha_0 + \alpha_{1(g)} \text{Session}_g$ | <code>p0 ~ session</code>  |
| Space use | $\log(\sigma) = \gamma_0$                                       | <code>sig ~ 1</code>       |
| Space use | $\log(\sigma) = \gamma_0 + \gamma_{1(g)} \text{Session}_g$      | <code>sig ~ session</code> |

- Include all models into a list using `fitList.oSCR()`:

```
f1 <- fitList.oSCR(
mods, # list of fitted models
rename) # if TRUE models are renamed with sensible names
```

- Compare multiple models  
`ms <- modSel.oSCR(f1)`
- Generate an AIC table to compare multiple models  
`ms$aic`

- Generate a coefficient table  
`ms$coef.tab`

- Generate a model averaged coefficients  
`ma <- ma.coef(ms)` # include a `modSel.oSCR` object

### 3.1. Back transform to the real scale

```
top.model <- m3
```

```
pred.df <- data.frame(session =
factor(c(1, 2, 3, 4, ...)))
```

```
pred.det <- get.real(
model = top.model, type = "det",
newdata = pred.df)
```