

# Package: navmix (via r-universe)

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**Type** Package

**Title** A package for performing directional clustering by fitting a noise-augmented von Mises-Fisher mixture model

**Version** 0.2.1

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**Description** A package for performing directional clustering by fitting a noise-augmented von Mises-Fisher mixture model

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**URL** <https://github.com/aj-grant/navmix>

**Imports** expm, ggplot2, plotrix, RColorBrewer, skmeans, tidyr

**Repository** <https://mrcieu.r-universe.dev>

**RemoteUrl** <https://github.com/aj-grant/navmix>

**RemoteRef** HEAD

**RemoteSha** c3feac409f14f1bd622a4363dbe9403c558f3b79

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navmix

*Noise-augmented directional clustering***Description**

Performs directional clustering by fitting a noise-augmented von Mises-Fisher mixture model

**Usage**

```
navmix(
  x,
  K = 10,
  select_K = TRUE,
  common_kappa = FALSE,
  pj_ini = 0.05,
  no_ini = 5,
  tol = 1e-04,
  max_iter = 100,
  plot = FALSE,
  plot_heat = TRUE,
  plot_heat_mu = FALSE,
  plot_parallel = TRUE,
  plot_radial = FALSE,
  plot_radial_options = list(plot_radial_separate = FALSE, radial_legend_pos = c(-2.5,
    2.7), radial_separate_col = 2)
)
```

**Arguments**

<code>x</code>	Matrix of values where rows represent observations and columns represent features.
<code>K</code>	The number of clusters to fit.
<code>select_K</code>	If TRUE (the default setting), the number of clusters will be chosen by BIC, with K the maximum number of clusters considered. If FALSE, then a model with K clusters will be fit.
<code>pj_ini</code>	The initial proportion of observations which belong in the noise cluster. Must be a number greater or equal to 0 and strictly less than 1. The default value is 0.05. If set to 0, no observations will be placed in the noise cluster.
<code>no_ini</code>	The number of time the algorithm is run with different initialisations. Must be a number greater than zero. The default value is 5.
<code>tol</code>	The tolerance threshold for convergence of the EM algorithm. Must be a number greater than 0. The default value is 1.0e-4.
<code>max_iter</code>	The maximum number of iterations of the EM algorithm. Must be a number greater than 0. The default value is 100.
<code>plot</code>	Plots of the results will be produced if set to TRUE. Default is FALSE.

<code>plot_heat</code>	Produces a heatmap of the results if <code>plot</code> is set to <code>TRUE</code> . The heatmap will also be returned as a <code>ggplot</code> object.
<code>plot_radial</code>	Produces (a) radial plot(s) of the results if <code>plot</code> is set to <code>TRUE</code> .
<code>common_kapp</code>	If <code>TRUE</code> , then model will force the <code>kappa</code> parameter to be equal for all clusters, except the noise cluster.
<code>plot_radial_separate</code>	If set to <code>FALSE</code> (the default value), the fitted means of each cluster are plotted on the same radial plot. If set to <code>TRUE</code> , they are plotted on separate radial plots.
<code>radial_legend_pos</code>	Adjusts the position of the legend for a radial plots with all fitted means plotted together.
<code>radial_separate_col</code>	Adjusts the format of the output of radial plots on separate plots.

### Value

Returned are the BIC values for each model fitted (`$BIC`), the final fitted model (`$fit`) and, if produced, the heatmap as a `ggplot` object (`$heatmap_plot`). The fitted model has the following.

<code>mu</code>	A matrix where each column represents the mean of the fitted von Mises-Fisher distribution for each cluster.
<code>kappa</code>	A row vector where each element represents the <code>kappa</code> parameter of the fitted von Mises-Fisher distribution for each cluster.
<code>g</code>	A matrix of probabilities for each observation belonging to each cluster. The value in the <code>j</code> th row and <code>k</code> th column represents the probability that the <code>j</code> th observation belongs to the <code>k</code> th cluster.
<code>z</code>	A vector of the cluster membership of each observation when allocated according to the cluster for which it has the highest probability of membership (hard clustering).
<code>bic</code>	The BIC for the fitted model.
<code>l</code>	The value of the likelihood function at the estimated parameters.

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