

# Package ‘plotmm’

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**Type** Package

**Title** Tidy Tools for Visualizing Mixture Models

**Version** 0.1.0

**BugReports** <https://github.com/pdwaggoner/plotmm/issues>

**Maintainer** Philip Waggoner <philip.waggoner@gmail.com>

**Description** The main function, `plot_mm()`, is used for plotting output from mixture models, including both densities and overlaying mixture weight component curves from the fit models. In line with the tidyverse, the package also includes the `plot_cut_point()` function to visualize the cut-point ( $\mu$ ) from the model over a histogram of the data density with several color options. Finally, the package includes the `plot_mix_comps()` helper function, which is used for both added customization as well as in the `plot_mm()` function. Supported model objects include: 'mixtools', 'EMCluster', and 'flexmix', with more from each forthcoming. Supported mixture model specifications include mixtures of univariate Gaussians, multivariate Gaussians, Gammas, logistic regressions, linear regressions, and Poisson regressions.

**Imports** methods, wesanderson, amerika, ggplot2, dplyr, patchwork

**Suggests** mixtools, EMCluster, flexmix, testthat, graphics

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**Encoding** UTF-8

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**RoxygenNote** 6.1.1

**NeedsCompilation** no

**Author** Philip Waggoner [aut, cre],  
Fong Chan [aut, ctb],  
Lu Zhang [aut, ctb]

**Repository** CRAN

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plot_cut_point	<i>Tidy Visualization of a Cut Point from a Mixture Model</i>
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### Description

Returns a plot of the data density (histogram) with an overlaid cut point generated by the fit mixture model

### Usage

```
plot_cut_point(m, plot = TRUE, color = c("grayscale", "amerika", "wesanderson"))
```

### Arguments

<code>m</code>	An object of class <code>mixEM</code> corresponding with the fit mixture model
<code>plot</code>	Logical for generating the plot. If <code>FALSE</code> , only the cut point value from the GMM is returned. If <code>TRUE</code> , histogram with the overlaid cut point is returned. Default is set to <code>TRUE</code> .
<code>color</code>	A vector of color options including "amerika" (from <code>amerika</code> package), "wesanderson" (from <code>wesanderson</code> package), and "grayscale", which is the default option.

### Details

Mixture models can be used to derive cut points separating clusters via soft assignment (See Benaglia et al. 2009 for more). `plot_cut_point()` plots data density with an overlaid cut point (the mean of the calculated  $\mu$ ) from `mixEM` objects via `mixtools`.

### References

Benaglia, T., Chauveau, D., Hunter, D. and Young, D. 2009. `mixtools`: An R package for analyzing finite mixture models. *Journal of Statistical Software*, 32(6), pp.1-29.

Ram, K., and Wickham, H. 2015. `wesanderson`: a Wes Anderson palette generator. R package version 0.3.

**Examples**

```
if(require(mixtools)){
  mixmdl <- mixtools::normalmixEM(faithful$waiting, k = 2)
}
plot_cut_point(mixmdl, plot = TRUE, color = "amerika") # returns plot, amerika
plot_cut_point(mixmdl, plot = TRUE, color = "wesanderson") # returns plot, wesanderson
plot_cut_point(mixmdl, plot = FALSE) # returns only the cut point value from the GMM
```

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plot\_gmm

*Plots Mixture Components from Gaussian Mixture Models*

---

**Description**

Generates a plot of data densities with overlaid mixture components from a Gaussian mixture model (GMM)

**Usage**

```
plot_gmm(m, k = NULL)
```

**Arguments**

m                    An object of class mixEM corresponding with the fit GMM  
k                    The number of components specified in the GMM, m

**Details**

Original function from the plotGMM package. Retained here for bridging between the packages. We recommend using instead the updated plot\_mm function.

Note: plot\_gmm requires a mixtools object to be supplied. Users must enter the same component value, k, in the plot\_gmm function, as that which was specified in the original GMM specification (also k in mixtools).

**References**

Benaglia, T., Chauveau, D., Hunter, D. and Young, D., 2009. mixtools: An R package for analyzing finite mixture models. *Journal of Statistical Software*, 32(6), pp.1-29.

Wickham, H., 2016. *ggplot2: elegant graphics for data analysis*. Springer.

**Examples**

```
if(require(mixtools)){
  mixmdl <- mixtools::normalmixEM(faithful$waiting, k = 2)
}
plot_gmm(mixmdl, 2)
```

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 plot\_mix\_comps

*Helper Function for Overlaying Mixture Components*


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### Description

Allows for plotting mixture components conditioned on a superimposed function meant for passage to ggplot's `stat_function()`

### Usage

```
plot_mix_comps(x, mu = NULL, sigma = NULL, lam = 1, beta0 = NULL,
  beta1=NULL, alpha=NULL, beta=NULL,
  normal=FALSE, logisreg=FALSE,
  gamma=FALSE, poisson=FALSE)
```

### Arguments

x	Input data
mu	Component mean
sigma	Component variance
lam	Component mixture weight
beta0	Coefficient values
beta1	Coefficient values
alpha	Initial shape parameters
beta	Initial parameter values
normal	Logical for normal distribution
logisreg	Logical for logistic regression mixtures
gamma	Logical for gamma distribution
poisson	Logical for poisson regression mixtures

### Details

Allows for component curves to be superimposed over a mixture model plot

### Examples

```
if(require(mixtools)){
  mixmdl <- mixtools::normalmixEM(faithful$waiting, k = 2)
}
x <- mixmdl$x
x <- data.frame(x)
ggplot2::ggplot(data.frame(x)) +
  ggplot2::geom_density(ggplot2::aes(x), color="black", fill="black") +
  ggplot2::stat_function(geom = "line", fun = plot_mix_comps,
    args = list(mixmdl$mu[1], mixmdl$sigma[1], lam = mixmdl$lambda[1]),
```

```

    colour = "red") +
  ggplot2::stat_function(geom = "line", fun = plot_mix_comps,
    args = list(mixmdl$mu[2], mixmdl$sigma[2], lam = mixmdl$lambda[2]),
    colour = "blue")

```

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plot\_mix\_comps\_normal *Custom Function for Overlaying Mixture Components for Normal Distributions*

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### Description

Plots a mixture component conditioned on a superimposed function

### Usage

```
plot_mix_comps_normal(x, mu, sigma, lam)
```

### Arguments

x	Input data
mu	Mean of component
sigma	Variance of component
lam	Mixture weight of component

### Details

Allows for specifying a custom function to be superimposed when plotting a mixture component assuming a normal distribution. This is the original function for the package, which is also included in the updated plot\_mix\_comps() function.

### Examples

```

if(require(mixtools)){
  mixmdl <- mixtools::normalmixEM(faithful$waiting, k = 2)
}
x <- mixmdl$x
x <- data.frame(x)
ggplot2::ggplot(data.frame(x)) +
  ggplot2::geom_density(ggplot2::aes(x), color="black", fill="black") +
  ggplot2::stat_function(geom = "line", fun = plot_mix_comps_normal,
    args = list(mixmdl$mu[1], mixmdl$sigma[1], lam = mixmdl$lambda[1]),
    colour = "red") +
  ggplot2::stat_function(geom = "line", fun = plot_mix_comps_normal,
    args = list(mixmdl$mu[2], mixmdl$sigma[2], lam = mixmdl$lambda[2]),
    colour = "blue")

```

**Description**

Generates a ggplot of data densities with overlaid mixture components from fit mixture models.

**Usage**

```
plot_mm(m, k = NULL, data = NULL)
```

**Arguments**

m	A mixture model object
k	Optional. The number of components specified in the mixture model, m
data	Name of data object required only for EMCluster objects

**Details**

This is the core function in the package, returning a ggplot object for a fit mixture model. The plot includes the data density with overlaid mixture components.

**References**

Wickham, H., 2016. ggplot2: elegant graphics for data analysis. Springer.

**Examples**

```
if(require(mixtools)){
  mixmdl1 <- mixtools::normalmixEM(faithful$waiting, k = 2)
}
plot_mm(mixmdl1, 2)

if(require(mixtools)){
  x <- c(rgamma(200, shape = 50, scale = 11), rgamma(200, shape = 28, scale = 6))
  mixmdl2 <- mixtools::gammamixEM(x, lambda = c(1, 1)/2)
}
plot_mm(mixmdl2)
```

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