# Package 'myplots.ew'

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Title This Package (	Contains Useful Plotting Functions		
Version 0.0.0.9000  Description This package will contain meaninful functions that will assit in the plotting.  License GPL (>= 3)  Encoding UTF-8  Roxygen list(markdown = TRUE)  RoxygenNote 7.3.0  R topics documented:  ggraph			
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		ggraph	Create a quick scatter plot in ggplot.
		Description	
		This will graph two given vectors in a ggplot-style scatter plot with the x-axis labeled "x" and the y-axis labeled "y".	
		Usage	
		ggraph(x, y,	point_color = "black", point_size = 1.5, point_shape = 19)
		Arguments	
x	This is the first vector to be plotted.		
У	This is the first vector to be plotted.		
point_color	This is the color of the points that will be plotted.		
<pre>point_size</pre>	This is the size of the points that will be plotted. The default is size 1.5.		
point_shape	This is the shape of the points that will be plotted. The default is 19: a filled circle		

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#### Value

This function returns a ggplot scatter plot object.

### **Examples**

```
## Create a scatter plot of y vs x.
x <- rnorm(100)
y <- x + rnorm(100, 0, 0.3)
ggraph(x, y)</pre>
```

influence\_plots

Influence Plots

### **Description**

This will compute many common residual and influence plots to check adequacy of a given model.

## Usage

```
influence_plots(model)
```

#### **Arguments**

model

This is a lm or glm (with binomial family) object

## Value

This function returns plots for the jackknife (externally studentized) residuals vs index, jackknife residuals vs fitted values, leverage values vs index, Cook's distance vs index, DfFits vs index, and all DfBetas vs index.

## **Examples**

```
## Randomly generate correlated variables x and y and then create the ## influence plots for them x <- rnorm(100) y <- x + rnorm(100, 0, 0.3) influence_plots(lm(y \sim x))
```

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