

	Method	Argument Name	Argument Value	Argument info	Method info
<b>g=</b>  <b>g(ind_row,ind_col)=</b>	<b>gramm(</b>	<b>'x'</b>	x variable	1D array/cellstr of length N, Matrix of size (N,M) , (N,1) cell of 1D arrays	Constructor for the class. Must be called first and result assigned to a variable Use to provide the data to be plotted
		<b>'y'</b>	y variable	1D array of length N, Matrix of size (N,M) , (N,1) cell of 1D arrays	
		<b>'color'</b>	color grouping/continuous variable	1D array/cellstr of length N	
		<b>'lightness'</b>	lightness grouping variable	1D array/cellstr of length N	
		<b>'linestyle'</b>	linestyle grouping variable	1D array/cellstr of length N	
		<b>'marker'</b>	marker grouping variable	1D array/cellstr of length N	
		<b>'size'</b>	size grouping variable	1D array/cellstr of length N	
<b>g.</b>  <b>g(ind_row,ind_col).</b>	<b>facet_grid(</b>		row grouping variable	1D array/cellstr of length N	
			column grouping variable	1D array/cellstr of length N	
		<b>'scale'</b>	<b>'fixed'</b>	Same x and y limits on all subplots	
			<b>'free_x'</b>	Same y limits on all subplots, same x limits within columns	
			<b>'free_y'</b>	Same x limits on all subplots, same y limits within rows	Use to provide data that will determine separation between subblots rows and columns. First argument provided will separate along rows, second will separate along columns
			<b>'free'</b>	Same x limits within columns, same y limits within rows	
			<b>'independent'</b>	Independent limits on each plot	
		<b>'force_ticks'</b>	<b>true/false</b>	Do we override defaults and force ticks on all subplots	
	<b>facet_wrap(</b>		column grouping variable	1D array/cellstr of length N	Use to provide data that will determine separation between subblots columns, with a wrapping: a new row of subplots is created when ncols is reached
		<b>'ncols'</b>	<b>4</b>	After how many columns do we wrap and create a new row	
		<b>'scale'</b>	<b>...</b>	Same as argument in gramm facet_grid()	
		<b>'force_ticks'</b>	<b>true/false</b>	Do we override defaults and force ticks on all subplots	
	<b>geom_point(</b>				Represent raw data as points (supports color, lightness, marker, size)
	<b>geom_jitter(</b>	<b>'width'</b>	<b>0.5</b>	How much are the points jittered in horizontal direction (in data units)	Represent raw data as jittered points, useful when lots of overlapping points, e.g. with discrete values (supports color, lightness, marker, size)
		<b>'height'</b>	<b>0.1</b>	How much are the points jittered in vertical direction (in data units)	
	<b>geom_line(</b>				Represent raw data with lines (supports color, lightness, marker, size). If x and y are 1D arrays, all points within a group will be connected !
	<b>geom_raster(</b>	<b>'geom'</b>	<b>'point'</b>	raster elements are points	Represents raw x data as a raster plot
			<b>'line'</b>	raster elements are lines	
	<b>geom_bar(</b>	<b>'width'</b>	<b>0.8</b>		
	<b>stat_summary(</b>	<b>'type'</b>	<b>'ci'</b>	mean & basic 95% CI of the mean (1.96 * sem)	
			<b>'bootci'</b>	mean & bootstrapped 95%CI of the mean	
			<b>'sem'</b>	mean and standard error of the mean	
			<b>'std'</b>	mean and standard deviation	
			<b>'quartile'</b>	median and quartiles	Represents summarized Y data per unique values of X. By default, it will group all Y values that have the same X value, compute the summary variables of interest ('type' argument), and plot it according to the 'geom' argument.
			<b>'95percentile'</b>	median and 95% percentiles	
			<b>'fitnormalci'</b>	mean and 95% CI of the mean from fitted normal distribution	
			<b>'fitpoissonci'</b>	mean and 95% CI of the mean from fitted Poisson distribution	
			<b>'fitbinomialci'</b>	mean and 95% CI of the mean from fitted binomial distribution	
			<b>'geom'</b>	means connected by a line, CI as shaded transparent area	
			<b>'area'</b>	means connected by a line, CI as thin lines	If X and Y are provided as 1D arrays but X values are not discrete enough, it is possible to compute the Y summaries over X bins with the 'bin_in' argument
			<b>'lines'</b>	means connected by a line	
			<b>'line'</b>	means connected by a line	
			<b>'solid_area'</b>	means connected by a line, CI as solid shaded area (use for vector exports in pre 2014b versions)	
			<b>'black_errorbar'</b>	CI as black errorbar	If X is provided as a matrix or a cell of arrays but every element has non-aligned X values, the argument 'interp_in' can be used to create aligned X values by interpolation over X.
			<b>'errorbar'</b>	CI as colored errorbar	
			<b>'bar'</b>	means as colored bars	
			<b>'point'</b>	means as points	
		<b>'dodge'</b>	<b>true/false</b>	Do we dodge on x when using multiple colors (useful for bar and errobar geoms)	
		<b>'setylim'</b>	<b>true/false</b>	Do we set the YLim for the subplot according to the summary or the data?	
		<b>'interp'</b>	<b>'linear'</b>	Provide to interpolate the output (corresponds to the methods argument of interp1)	
		<b>'interp_in'</b>	<b>100</b>	Provide to linearly interpolate the input over x (corresponds to number of x points)	
		<b>'bin_in'</b>	<b>10</b>	Provide to bin inputs over x values (corresponds to number of bins)	
		<b>'width'</b>	<b>0.6</b>	Provide to set the width of bars and errorbars	
		<b>'dodge'</b>	<b>0.7</b>	When using multiple colors, use to dodge graphical elements between colors with the same x value	
	<b>stat_smooth(</b>	<b>'lambda'</b>	<b>1000</b>	Smoothing parameter (low values smooth less)	Represents fast spline smoothed Y data with confidence interval. This is not proper to use when X/Y are matrices or cells of arrays
		<b>'geom'</b>	<b>...</b>	Same geom as in gramm stat_summary()	
	<b>stat_glm(</b>	<b>'distribution'</b>	<b>'normal'</b>	Same argument as fitglm()	Fits and displays generalized linear models to the data.
			<b>...</b>		
		<b>'geom'</b>	<b>...</b>	Same geom as in gramm stat_summary()	
		<b>'fullrange'</b>	<b>true/false</b>	Do we display the fit over the whole x axis, or just on the range of the value used for the fit	
		<b>'disp_fit'</b>	<b>true/false</b>	Do we display the fitted equations (with pvals stars)	
	<b>stat_fit(</b>	<b>'fun'</b>	<b>@(param1,param2,x)x.^param1+param2</b>	Anonymous function with parameters to fit as first arguments and x as last argument	Fits and displays a provided custom function to the data
		<b>'StartPoint'</b>	<b>[param1_start param2_start]</b>	Array with starting values of parameters	
		<b>'intopt'</b>	<b>'observation'</b>	95% bounds on a new observation (see option of predint())	
			<b>'functional'</b>	95% bounds for the fitted function	
		<b>'fullrange'</b>	<b>true/false</b>	Do we display the fit over the whole x axis, or just on the range of the value used for the fit	
		<b>'disp_fit'</b>	<b>true/false</b>	Do we display the fitted equations	
	<b>stat_bin(</b>	<b>'geom'</b>	<b>...</b>	Same geom as in gramm stat_summary()	
		<b>'nbins'</b>	<b>30</b>	Number of bins	
		<b>'edges'</b>	<b>-20 : 0.5 : 20</b>	Edges ovf bins (overrides 'nbins')	
		<b>'geom'</b>	<b>'bar'</b>	Results as dodged bars	
			<b>'line'</b>	Results connected by a line	
			<b>'overlaid_bar'</b>	Results as overlaid bars (use transparency)	
			<b>'stackedBars'</b>	Results as stacked bars	
			<b>'stairs'</b>	Results as stair line	
			<b>'point'</b>	Results as points	
		<b>'normalization'</b>	<b>'count'</b>		
		<b>'fill'</b>	<b>...</b>	Same as 'Normalization' argument of histcounts()	
			<b>'face'</b>		
			<b>'edge'</b>		
			<b>'all'</b>		
			<b>'transparent'</b>		
		<b>'width'</b>	<b>0.6</b>	Provide to specify width of bars	
		<b>'dodge'</b>	<b>0.7</b>	Provide to specify dodging between elements	

	Method	Argument Name	Argument Value	Argument info	Method info
	stat_density()	'bandwidth'		Same argument as ksdensity()	
		'function'	'pdf'		
			...	Same argument as ksdensity()	
		'kernel'	'normal'		
			...	Same argument as ksdensity()	
	stat_bin2d()	'npoints'	100	How many points are used to plot the density	
		'extra_x'	10	Extend the x value range over which the density is evaluated	
	stat_bin2d()	'nbins'	[n_xbins n_ybins]		
		'edges'	{x_edges_array, y_edges_array}		
	stat_ellipse()	'geom'	'image'		
			'contour'		
		'type'	'95percentile'	Fit ellipse that contains 95% of the points (assuming bivariate normal)	
			'ci'	Fit ellipse that contains 95% of the bootstrapped xy means	
	stat_ellipse()	'geom'	'area'		
			'line'		
	stat_ellipse()	patch_opts			
	stat_qq()	'distribution'	makedist('Normal',0,1)	Provide a theoretical distribution to plot x against using Matlab's makedist() function. Set to 'y' to plot x against y densities.	Quantile-quantile plot
	stat_boxplot()	'width'	0.6	Width of boxes	Box and whisker plots of y data for each unique x value
		'dodge'	0.7	Dodging between boxes of different colors within unique x values	
	geom_abline()	'intercept'	0	Single value or 1D array of size P	
		'slope'	1	Single value or 1D array of size P	
	geom_abline()	'style'	'k--'	Single string or 1D cellstr of size P	
	geom_vline()	'xintercept'	1	Single value or 1D array of size P	
		'style'	'k--'	Single string or 1D cellstr of size P	
	geom_hline()	'yintercept'	1	Single value or 1D array of size P	
		'style'	'k--'	Single string or 1D cellstr of size P	
	geom_funline()	'fun'	@(x)exp(sin(x-pi))	Anonymous function or cell of anonymous functions	
		'style'	'k--'	Single string or 1D cellstr of size P	
	set_names()	'x'	'x axis legend'	Legend for the x axes	
		'y'	'y axis legend'	Legend for the y axes	
		'row'	'row legend'	Title of the row legends (actual titles will be a combination of title and value)	
		'column'	'column legend'	Title of the column legends (actual titles will be a combination of title and value)	
		'color'	'color legend'	Title of the color legend (actual legend will use the values)	
	set_title()	...		All other titles for the gramm() arguments	
	set_title()	'Title'		Desired title	
		'FontSize'	16	Any text property 'Name',value pair	
	set_polar()	'closed'	true/false	Do we connect the first and last points ?	
		'maxy'	10	Impose the max of the radial scale (default corresponds to the max of y values)	
	set_color_options()	'map'	'lch'	Default HCL-based colormap	
			'matlab'	Matlab's own post 2014b map	
			'brewer1' 'brewer2' 'brewer3'		
			'brewer_pastel' 'brewer_dark'	colorbrewer2.org colormaps	
	set_color_options()		[0.1 0 0 0 0.2 0.9]	Custom colormap as Nx3 matrix	
		'lightness_range'	[85 15]		
		'chroma_range'	[30 90]		
		'hue_range'	[25 385]		
		'lightness'	65		
	set_order_options()	'chroma'	75		
		'x'	1	Values sorted in ascending order (numeric or alphabetical)	
			0	Keep order of appearance of values in the input	
			-1	Values sorted in descending order	
	set_order_options()		[value1 value2 value3 ...] {'value1' 'value2' 'value3' ...}	Values ordered according as in the provided array/cell (all unique values have to be present in the array/cell)	
			[index1 index2 index3 ...]	Values ordered according as in the provided indices (array of indices in the sorted values array/cell)	
	set_order_options()	'color'			
		...			
	set_continuous_color()	'colormap'	'hot'		
		'LCH_colormap'	[L_start L_end; C_start C_end ; H_start H_end]		
	axe_property()	'axe_property'	axe_property_value	Pass one or multiple name,value pairs for Axes Properties (XLim,XGrid, DataAspectRatio...)	
	no_legend()				
	set_limit_extra()		0.1	How much do we extend limits of x axis (ratio wrt original limits)	
			0.1	How much do we extend limits of y axis (ratio wrt original limits)	
	set_datetick()	'x'	1	Same arguments as datetick(): tickaxis,dateformat	
		'y'	2		
g.	draw()		false	Give false as (optional) argument to allow superimposing plots	Draw the plot !
	redraw()		0.05	Redraw with custom spacing	