

# Common statistical tests are linear models

Last updated: 29 June, 2019. Also check out the [R version](#)!

See worked examples and more details at the accompanying notebook: <https://github.com/eigenfoo/tests-as-linear>

Common name	Function in <code>scipy.stats</code>	Equivalent linear model in <code>smf.ols</code>	Exact?	The linear model in words	Icon	
<b>Simple Regression: <math>y \sim 1 + x</math></b>	<b>y is independent of x</b> P: One-sample t-test N: Wilcoxon signed-rank	<code>scipy.stats.ttest_1samp(y)</code> <code>scipy.stats.wilcoxon(y)</code>	<code>smf.ols("y ~ 1", data)</code> <code>smf.ols("y ~ 1", signed_rank(data))</code>	✓ <a href="#">for N &gt; 14</a>	One number (intercept, i.e., the mean) predicts <b>y</b> . - (Same, but it predicts the <i>signed rank</i> of <b>y</b> .)	
	P: Paired-sample t-test N: Wilcoxon matched pairs	<code>scipy.stats.ttest_rel(y1, y2)</code> <code>scipy.stats.wilcoxon(y1, y2)</code>	<code>smf.ols("y2_sub_y1 ~ 1", data)</code> <code>smf.ols("y2_sub_y1 ~ 1", signed_rank(data))</code>	✓ <a href="#">for N &gt; 14</a>	One intercept predicts the pairwise $y_2 - y_1$ differences. - (Same, but it predicts the <i>signed rank</i> of $y_2 - y_1$ .)	
<b>y ~ continuous x</b> P: Pearson correlation N: Spearman correlation	<code>scipy.stats.pearsonr(x, y)</code> <code>scipy.stats.spearmanr(x, y)</code>	<code>smf.ols("y ~ 1 + x", data)</code> <code>smf.ols("y ~ 1 + x", rank(data))</code>	✓ <a href="#">for N &gt; 10</a>	One intercept plus <b>x</b> multiplied by a number (slope) predicts <b>y</b> . - (Same, but with <i>ranked x</i> and <b>y</b> )		
<b>y ~ discrete x</b> P: Two-sample t-test P: Welch's t-test N: Mann-Whitney U	<code>scipy.stats.ttest_ind(y1, y2)</code> N/A in Python, but <a href="#">see R version</a> <code>scipy.stats.mannwhitneyu(y1, y1)</code>	<code>smf.ols("y ~ 1 + group", data)<sup>A</sup></code> N/A in Python, but <a href="#">see R version</a> <code>smf.ols("y ~ 1 + group", signed_rank(data))<sup>A</sup></code>	✓ ✓ <a href="#">for N &gt; 11</a>	An intercept for <b>group 1</b> (plus a difference if <b>group 2</b> ) predicts <b>y</b> . - (Same, but with one variance <i>per group</i> instead of one common.) - (Same, but it predicts the <i>signed rank</i> of <b>y</b> .)		
<b>Multiple regression: <math>y \sim 1 + x_1 + x_2 + \dots</math></b>	P: One-way ANOVA N: Kruskal-Wallis	<code>scipy.stats.f_oneway(a, b, c)</code> <code>scipy.stats.kruskal(a, b, c)</code>	<code>smf.ols(y ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub>)<sup>A</sup></code> <code>smf.ols(rank(y) ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub>)<sup>A</sup></code>	✓ <a href="#">for N &gt; 11</a>	An intercept for <b>group 1</b> (plus a difference if $group \neq 1$ ) predicts <b>y</b> . - (Same, but it predicts the <i>rank</i> of <b>y</b> .)	
	P: One-way ANCOVA	N/A in Python, but <a href="#">see R version</a>	<code>smf.ols("y ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub> + x", data)<sup>A</sup></code>	✓	- (Same, but plus a slope on <b>x</b> .) <i>Note: this is discrete AND continuous. ANCOVAs are ANOVAs with a continuous x.</i>	
	P: Two-way ANOVA	N/A in Python, but <a href="#">see R version</a>	<code>smf.ols("y ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub> + S<sub>2</sub> + S<sub>3</sub> + ... + S<sub>K</sub> + G<sub>2</sub>*S<sub>2</sub>+G<sub>3</sub>*S<sub>3</sub>...+G<sub>N</sub>*S<sub>K</sub>", data)</code>	✓	Interaction term: changing <b>sex</b> changes the <b>y ~ group</b> parameters. <i>Note: <math>G_{2 \dots N}</math> is an indicator (0 or 1) for each non-intercept levels of the <b>group</b> variable. Similarly for <math>S_{2 \dots K}</math> for sex. The first line (with <math>G_i</math>) is main effect of group, the second (with <math>S_i</math>) for sex and the third is the <b>group</b> <math>\times</math> <b>sex</b> interaction. For two levels (e.g. male/female), line 2 would just be "<math>S_2</math>" and line 3 would be <math>S_2</math> multiplied with each <math>G_i</math>.</i>	[Coming]
<b>Counts ~ discrete x</b> N: Chi-square test	<code>scipy.stats.chisquare(data)</code>	<b>Equivalent log-linear model</b> <code>sm.GLM(y ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub> + S<sub>2</sub> + S<sub>3</sub> + ... + S<sub>K</sub> + G<sub>2</sub>*S<sub>2</sub>+G<sub>3</sub>*S<sub>3</sub>...+G<sub>N</sub>*S<sub>K</sub>, family=...)<sup>A</sup></code>	✓	Interaction term: (Same as Two-way ANOVA.) <i>Note: Run <code>glm</code> using the following arguments: <code>glm(model, family=poisson())</code>. As linear-model, the Chi-square test is <math>\log(y_i) = \log(N) + \log(\alpha_i) + \log(\beta_i) + \log(\alpha_i\beta_i)</math> where <math>\alpha_i</math> and <math>\beta_i</math> are proportions. See more info in <a href="#">the accompanying notebook</a>.</i>	Same as Two-way ANOVA	
N: Goodness of fit	<code>scipy.stats.chi2_contingency(data)</code>	<code>sm.GLM(y ~ 1 + G<sub>2</sub> + G<sub>3</sub> + ... + G<sub>N</sub>, family=...)<sup>A</sup></code>	✓	(Same as One-way ANOVA and see Chi-Square note.)	1W-ANOVA	

List of common parametric (P) non-parametric (N) tests and equivalent linear models. The notation  $y \sim 1 + x$  is R shorthand for  $y = 1 \cdot b + a \cdot x$  which most of us learned in school. Models in similar colors are highly similar, but really, notice how similar they *all* are across colors! For non-parametric models, the linear models are reasonable approximations for non-small sample sizes (see "Exact" column and click links to see simulations). Other less accurate approximations exist, e.g., Wilcoxon for the sign test and Goodness-of-fit for the binomial test. The signed rank function is `signed_rank(df) = np.sign(df) * df.rank()`. The variables  $G_i$  and  $S_i$  are "[dummy coded" indicator variables](#) (either 0 or 1) exploiting the fact that when  $\Delta x = 1$  between categories the difference equals the slope. Subscripts (e.g.,  $G_2$  or  $y_1$ ) indicate different columns in data. lm requires long-format data for all non-continuous models. All of this is exposed in greater detail and worked examples at <https://eigenfoo.xyz/tests-as-linear/>.

<sup>A</sup> See the note to the two-way ANOVA for explanation of the notation.

