

# Regression Tables with huxreg

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## Regression tables with huxreg

Huxtable includes the function `huxreg` to build a table of regressions.

You call `huxreg` with a list of models. These models can be of any class which has a `tidy` method defined in the `broom` package. The method should return a list of regression coefficients with names `term`, `estimate`, `std.error` and `p.value`. That covers most standard regression packages.

Let's start by running some regressions to predict a diamond's price.

```
data(diamonds, package = "ggplot2")
diamonds <- diamonds[1:100,]

lm1 <- lm(price ~ carat + depth, diamonds)
lm2 <- lm(price ~ depth + factor(color, ordered = FALSE), diamonds)
lm3 <- lm(log(price) ~ carat + depth, diamonds)
```

Now, we use `huxreg` to display the regression output side by side.

```
huxreg(lm1, lm2, lm3)
```

	(1)	(2)	(3)
(Intercept)	981.607 (720.175)	900.067 (2431.815)	6.269 *** (0.782)
carat	4328.324 *** (136.755)		3.531 *** (0.149)
depth	-27.785 * (11.656)	-6.804 (39.293)	-0.019 (0.013)
factor(color, ordered = FALSE)E		449.490 (239.388)	
factor(color, ordered = FALSE)F		391.705 (290.880)	
factor(color, ordered = FALSE)G		583.111 (308.513)	
factor(color, ordered = FALSE)H		126.916 (256.367)	
factor(color, ordered = FALSE)I		-47.220 (253.092)	
factor(color, ordered = FALSE)J		-123.430 (269.157)	
N	100	100	100
R2	0.912	0.123	0.854
logLik	-675.703	-790.788	6.822
AIC	1359.405	1599.576	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

The basic output includes estimates, standard errors and summary statistics.

Some of those variable names are hard to read. We can change them by providing a named vector of variables in the `coefs` argument.

```
color_names <- grep("factor", names(coef(lm2)), value = TRUE)
names(color_names) <- gsub(".*)(.)", "Color: \\1", color_names)

huxreg(lm1, lm2, lm3, coefs = c("Carat" = "carat", "Depth" = "depth", color_names))
```

	(1)	(2)	(3)
Carat	4328.324 *** (136.755)		3.531 *** (0.149)
Depth	-27.785 * (11.656)	-6.804 (39.293)	-0.019 (0.013)
Color: E		449.490 (239.388)	
Color: F		391.705 (290.880)	
Color: G		583.111 (308.513)	
Color: H		126.916 (256.367)	
Color: I		-47.220 (253.092)	
Color: J		-123.430 (269.157)	
N	100	100	100
R2	0.912	0.123	0.854
logLik	-675.703	-790.788	6.822
AIC	1359.405	1599.576	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

Or, since the output from `huxreg` is just a huxtable, we could just edit its contents directly.

```
diamond_regs <- huxreg(lm1, lm2, lm3)
diamond_regs[seq(8, 18, 2), 1] <- paste("Color:", LETTERS[5:10])

# prints the same as above
```

Of course, we aren't limited to just changing names. We can also make our table prettier. Let's put our footnote in italic, add a caption, and highlight the cell background of significant coefficients. All of these are just standard huxtable commands.

```
suppressPackageStartupMessages(library(dplyr))

diamond_regs |>
  map_background_color(-1, -1, by_regex(
    "\\*" = "yellow"
  )) |>
  set_italic(final(1), 1) |>
  set_caption("Linear regressions of diamond prices")
```

Table 1: Linear regressions of diamond prices

	(1)	(2)	(3)
(Intercept)	981.607 (720.175)	900.067 (2431.815)	6.269 *** (0.782)
carat	4328.324 *** (136.755)		3.531 *** (0.149)
depth	-27.785 * (11.656)	-6.804 (39.293)	-0.019 (0.013)
Color: E		449.490 (239.388)	
Color: F		391.705 (290.880)	
Color: G		583.111 (308.513)	
Color: H		126.916 (256.367)	
Color: I		-47.220 (253.092)	
Color: J		-123.430 (269.157)	
N	100	100	100
R2	0.912	0.123	0.854
logLik	-675.703	-790.788	6.822
AIC	1359.405	1599.576	-5.644

\*\*\*  $p < 0.001$ ; \*\*  $p < 0.01$ ; \*  $p < 0.05$ .

By default, standard errors are shown below coefficient estimates. To display them in a column to the right, use `error_pos = "right"`:

```
huxreg(lm1, lm3, error_pos = "right")
```

	(1)		(2)	
(Intercept)	981.607	(720.175)	6.269 ***	(0.782)
carat	4328.324 ***	(136.755)	3.531 ***	(0.149)
depth	-27.785 *	(11.656)	-0.019	(0.013)
N	100		100	
R2	0.912		0.854	
logLik	-675.703		6.822	
AIC	1359.405		-5.644	

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

This will give column headings a column span of 2.

To display standard errors in the same cell as estimates, use `error_pos = "same"`:

```
huxreg(lm1, lm3, error_pos = "same")
```

	(1)		(2)	
(Intercept)	981.607	(720.175)	6.269 ***	(0.782)
carat	4328.324 ***	(136.755)	3.531 ***	(0.149)
depth	-27.785 *	(11.656)	-0.019	(0.013)
N	100		100	
R2	0.912		0.854	
logLik	-675.703		6.822	
AIC	1359.405		-5.644	

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

You can change the default column headings by naming the model arguments:

```
huxreg("Price" = lm1, "Log price" = lm3)
```

	Price	Log price
(Intercept)	981.607 (720.175)	6.269 *** (0.782)
carat	4328.324 *** (136.755)	3.531 *** (0.149)
depth	-27.785 * (11.656)	-0.019 (0.013)
N	100	100
R2	0.912	0.854
logLik	-675.703	6.822
AIC	1359.405	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

To display a particular row of summary statistics, use the `statistics` parameter. This should be a character vector. Valid values are anything returned from your models by `broom::glance`:

```
gl <- as_hux(broom::glance(lm1))
gl |>
  restack_down(cols = 3, on_remainder = "fill") |>
  set_bold(odds, everywhere)
```

<b>r.squared</b>	<b>adj.r.squared</b>	<b>sigma</b>
0.912	0.91	211
<b>statistic</b>	<b>p.value</b>	<b>df</b>
504	5.65e-52	2
<b>logLik</b>	<b>AIC</b>	<b>BIC</b>
-676	1.36e+03	1.37e+03
<b>deviance</b>	<b>df.residual</b>	<b>nobs</b>
4.33e+06	97	100

Another value you can use is `"nobs"`, which returns the number of observations from the regression. If the `statistics` vector has names, these will be used for row headings:

```
huxreg(lm1, lm3, statistics = c("N. obs." = "nobs",
  "R squared" = "r.squared", "F statistic" = "statistic",
  "P value" = "p.value"))
```

	(1)	(2)
(Intercept)	981.607 (720.175)	6.269 *** (0.782)
carat	4328.324 *** (136.755)	3.531 *** (0.149)
depth	-27.785 * (11.656)	-0.019 (0.013)
N. obs.	100	100
R squared	0.912	0.854
F statistic	504.082	283.881
P value	0.000	0.000

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

By default, `huxreg` displays significance stars. You can alter the symbols used and significance levels with the `stars` parameter, or set `stars = NULL` to turn off significance stars completely.

```
huxreg(lm1, lm3, stars = c(`*` = 0.1, `**` = 0.05, `***` = 0.01)) # a little boastful?
```



	(1)	(2)
(Intercept)	981.607	6.269 ***
	(720.175)	(0.782)
carat	4328.324 ***	3.531 ***
	(136.755)	(0.149)
depth	-27.785 **	-0.019
	(11.656)	(0.013)
N	100	100
R2	0.912	0.854
logLik	-675.703	6.822
AIC	1359.405	-5.644

\*\*\* p < 0.01; \*\* p < 0.05; \* p < 0.1.

You aren't limited to displaying standard errors of the estimates. If you prefer, you can display t statistics or p values, using the `error_format` option. Any column from `tidy` can be used by putting it in curly brackets:

```
# Another useful column: p.value
huxreg(
  lm1, lm3,
  error_format = "[{statistic}]",
  note         = "{stars}. T statistics in brackets."
)
```

	(1)	(2)
(Intercept)	981.607 [1.363]	6.269 *** [8.016]
carat	4328.324 *** [31.650]	3.531 *** [23.773]
depth	-27.785 * [-2.384]	-0.019 [-1.499]
N	100	100
R2	0.912	0.854
logLik	-675.703	6.822
AIC	1359.405	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05. T statistics in brackets.

Here we also changed the footnote, using **note**. If **note** contains the string "{stars}" it will be replaced by a description of the significance stars used. If you don't want a footnote, just set **note** = NULL.

Alternatively, you can display confidence intervals. Use **ci\_level** to set the confidence level for the interval, then use {conf.low} and {conf.high} in **error\_format**:

```
huxreg(lm1, lm3, ci_level = .99, error_format = "{conf.low} -- {conf.high}")
```

	(1)	(2)
(Intercept)	981.607	6.269 ***
	(-910.629 – 2873.844)	(4.214 – 8.324)
carat	4328.324 ***	3.531 ***
	(3969.004 – 4687.643)	(3.140 – 3.921)
depth	-27.785 *	-0.019
	(-58.411 – 2.842)	(-0.052 – 0.014)
N	100	100
R2	0.912	0.854
logLik	-675.703	6.822
AIC	1359.405	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

To change number formatting, set the `number_format` parameter. This works the same as the `number_format` property for a huxtable - if it is numeric, numbers will be rounded to that many decimal places; if it is character, it will be taken as a format to the base R `sprintf` function. `huxreg` tries to be smart and to format summary statistics like `nobs` as integers.

```
huxreg(lm1, lm3, number_format = 2)
```

	(1)	(2)
(Intercept)	981.61	6.27 ***
	(720.17)	(0.78)
carat	4328.32 ***	3.53 ***
	(136.75)	(0.15)
depth	-27.78 *	-0.02
	(11.66)	(0.01)
N	100	100
R2	0.91	0.85
logLik	-675.70	6.82
AIC	1359.41	-5.64

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

Lastly, if you want to bold all significant coefficients, set the parameter `bold_signif` to a maximum significance level:

```
huxreg(lm1, lm3, bold_signif = 0.05)
```

	(1)	(2)
(Intercept)	981.607	<b>6.269 ***</b>
	(720.175)	<b>(0.782)</b>
carat	<b>4328.324 ***</b>	<b>3.531 ***</b>
	<b>(136.755)</b>	<b>(0.149)</b>
depth	<b>-27.785 *</b>	-0.019
	<b>(11.656)</b>	(0.013)
N	100	100
R2	0.912	0.854
logLik	-675.703	6.822
AIC	1359.405	-5.644

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

## Altering data

Sometimes, you want to report different statistics for a model. For example, you might want to use robust standard errors.

One way to do this is to pass a tidy-able test object into `huxreg`. The function `coeftest` in the “lmtest” package has tidy methods defined:

```
library(lmtest)
library(sandwich)
lm_robust <- coeftest(lm1, vcov = vcovHC, save = TRUE)
huxreg("Normal SEs" = lm1, "Robust SEs" = lm_robust)
```

	Normal SEs	Robust SEs
(Intercept)	981.607 (720.175)	981.607 (1117.654)
carat	4328.324 *** (136.755)	4328.324 *** (293.929)
depth	-27.785 * (11.656)	-27.785 (17.995)
N	100	100
R2	0.912	0.912
logLik	-675.703	-675.703
AIC	1359.405	1359.405

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

If that is not possible, you can compute statistics yourself and add them to your model using the `tidy_override` function:

```
lm_fixed <- tidy_override(lm1, p.value = c(0.5, 0.2, 0.06))
huxreg("Normal p values" = lm1, "Supplied p values" = lm_fixed)
```

	Normal p values	Supplied p values
(Intercept)	981.607	981.607
	(720.175)	(720.175)
carat	4328.324 ***	4328.324
	(136.755)	(136.755)
depth	-27.785 *	-27.785
	(11.656)	(11.656)
N	100	100
R2	0.912	0.912
logLik	-675.703	-675.703
AIC	1359.405	1359.405

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.

You can override any statistics returned by `tidy` or `glance`.

If you want to completely replace the output of `tidy`, use the `tidy_replace()` function. For example, here's how to print different coefficients for a multinomial model.

```
mnl <- nnet::multinom(gear ~ mpg, mtcars)
```

```
## # weights:  9 (4 variable)
## initial  value 35.155593
## iter  10 value 23.131901
## final   value 23.129234
## converged
```

```
tidied <- broom::tidy(mnl)
models <- list()
models[["4 gears"]] <- tidy_replace(mnl, tidied[tidied$y.level == 4, ])
models[["5 gears"]] <- tidy_replace(mnl, tidied[tidied$y.level == 5, ])
huxreg(models, statistics = "AIC")
```

	4 gears	5 gears
(Intercept)	-9.502 ** (3.262)	-7.691 * (3.232)
mpg	0.475 ** (0.168)	0.358 * (0.168)
AIC	54.258	54.258

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05.