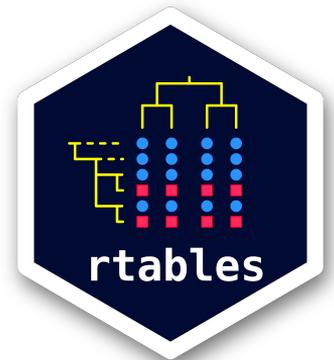


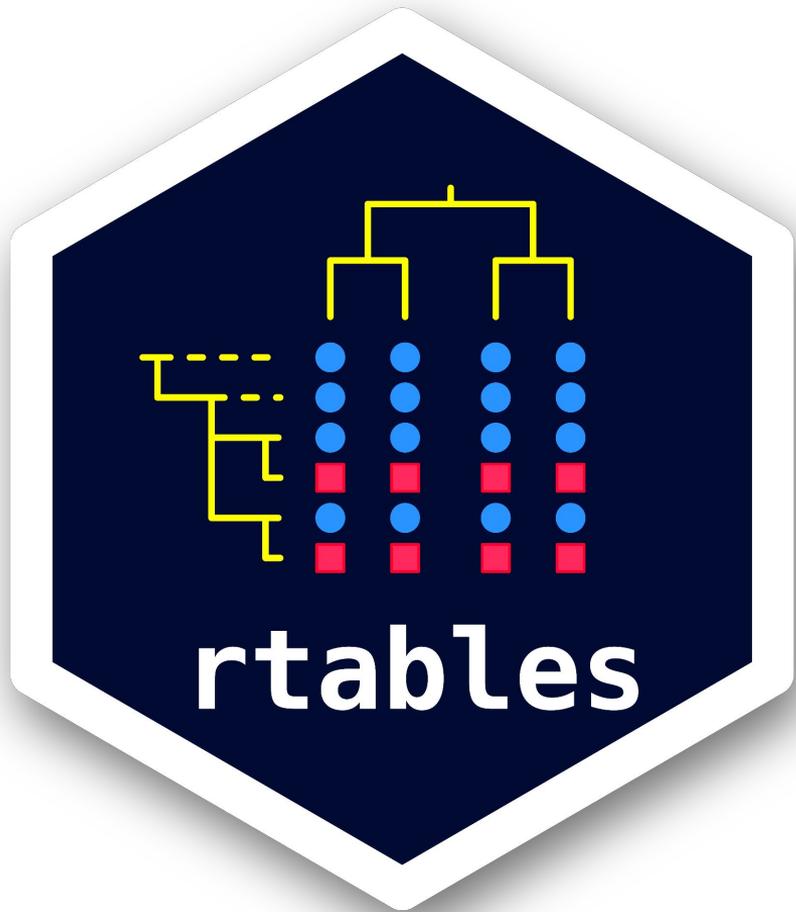
rtables For Power-Users Part 1

Nest Team Training

Jan 31, 2023

Gabriel Becker, Adrian Waddell





rtables

What you will learn

- **Tables fundamentals**
 - Conceptual model and motivations
 - Basic layout construction
- **Table Structure**
 - Pathing
 - Subsetting
 - Post-processing and mitigation
- **(Semi-) Advanced Features**
 - Specialized splitting
 - Appearance customization
 - Pagination

What Will Wait For Part 2

- (Semi-) Advanced Topics
 - Whatever we don't have time to get to today
- Advanced customization
 - Custom-written split functions
 - Advanced features of analysis function writing
- Tying it all together
 - “Thinking in `rtables`”
 - Reasoning about workarounds

What about listings?

Prototype that uses formatters (rtables rendering machinery backend) can be found here: <https://github.com/insightsengineering/rlistings>



rtables Conceptual Model

0th Law of Computing (Statistical or Otherwise)

Let the computer do tasks that are

- Tedious
- Repetitive
- Human-error prone

That's what it's there for!

The First Step In Creating a Table

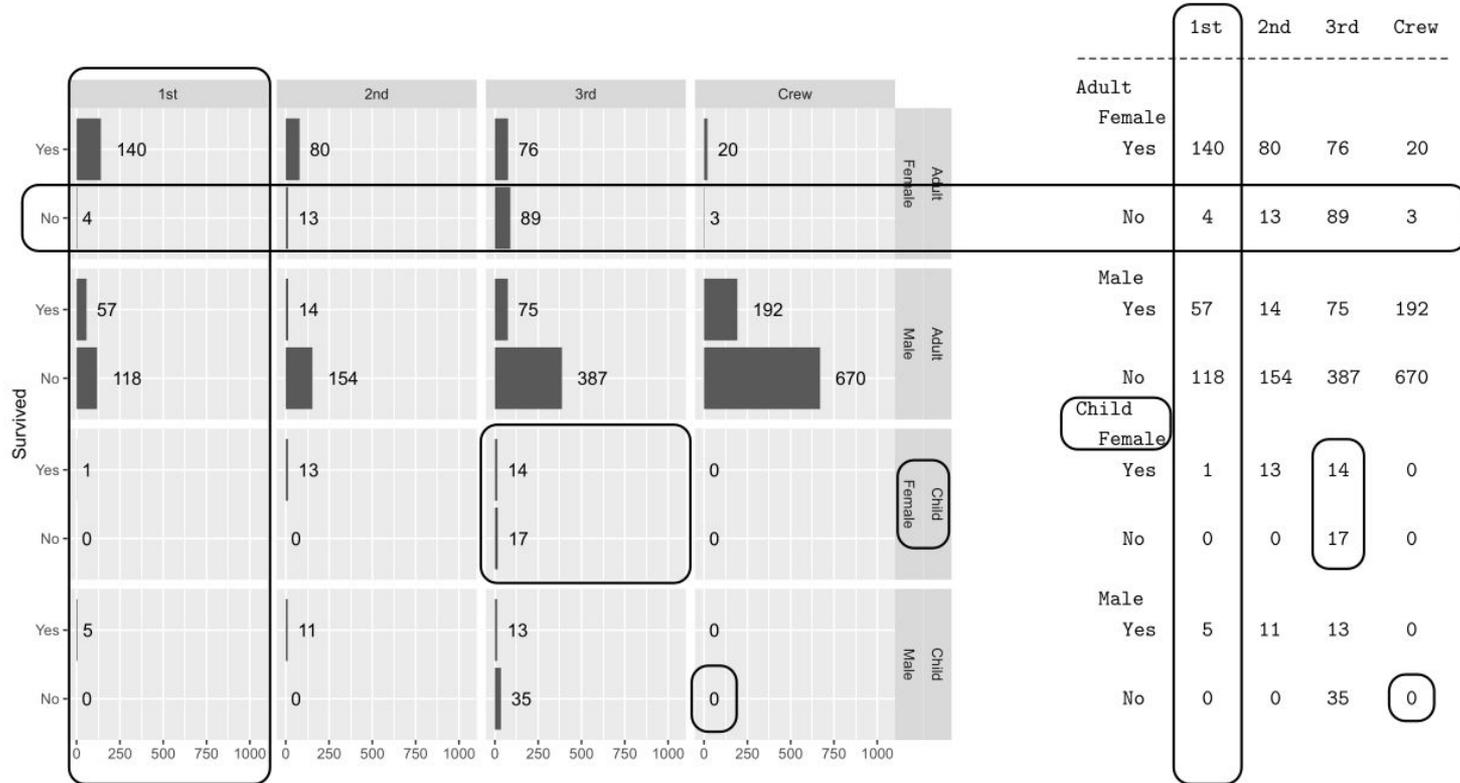
Calculating cell values, right?

The First Step In Creating a Table

Calculating cell values, right?



Reporting Tables Are Faceted Data Visualizations



Imagine Manually Subsetting Facet Data When Using
`ggplot2` (or `lattice`)



Subsetting data and calculating facet statistics

Humans



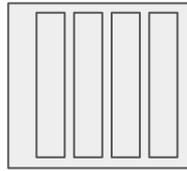
Computers



Basics of rtables



layout
faceting, value
derivation, formatting
instructions



data.frame
Unaggregated data
(variables + obs)

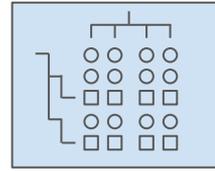


table object
cell values,
table structure,
format instructions,
paths



	ASM A		ASM B	
	M	F	M	F
US				
n	80	60	20	23
Avg (age)	20	21	21	22
CEM				
n	45	34	60	55
Avg (age)	18	19	20	21

formatted table
such as ASCII, pdf,
rtf

Basics of rtables

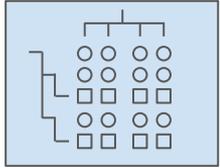
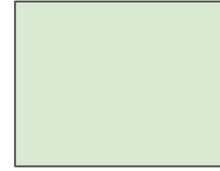
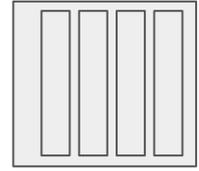


table object
cell values,
table structure,
format instructions,
paths

`<-build_table(`



layout
faceting, value
derivation, formatting
instructions



data.frame
Unaggregated data
(variables + obs)

)

Basics of rtables

print(

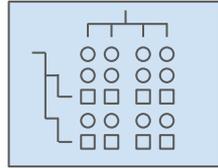


table object
cell values,
table structure,
format instructions,
paths

	ARM A		ARM B	
	M	F	M	F
UB				
n	80	60	20	23
Avg (age)	20	21	21	22
CIN				
n	40	34	60	55
Avg (age)	18	19	20	21

formatted table
here ASCII

Basics of rtables

```
library(rtables)

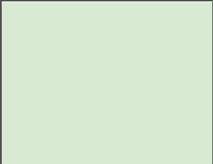
lyt <- basic_table() |>
      split_cols_by("ARM") |>
      analyze("AGE", mean, format = "xx.xx")

tbl <- build_table(lyt, ex_adsl)

print(tbl)
```

Basics of rtables

```
library(rtables)  
  
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  analyze("AGE", mean, format = "xx.xx")
```



layout
faceting, value
derivation, formatting
instructions

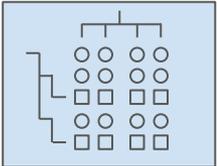
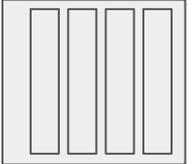


table object
cell values,
table structure,
format instructions,
paths

```
tbl <- build_table(lyt, ex_adsl)  
  
print(tbl)
```

	ARM A		ARM B	
	M	F	M	F
US	80	60	20	23
n	20	21	21	22
Avg (age)				
GB	40	34	60	55
n	18	19	20	21
Avg (age)				

formatted table
here ASCII



data.frame
Unaggregated data
(variables + obs)



Basics of rtables

```
> library(rtables)
Loading required package: magrittr
Loading required package: formatters
>
> lyt <- basic_table() |>
+   split_cols_by("ARM") |>
+   analyze("AGE", mean, format = "xx.xx")
>
> tbl <- build_table(lyt, ex_adsl)
>
> print(tbl)
```

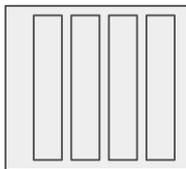
	A: Drug X	B: Placebo	C: Combination
mean	33.77	35.43	35.43

```
> |
```

now let's get cracking



So whats next...



data.frame
Unaggregated data
(variables + obs)



layout
faceting, value
derivation, formatting
instructions

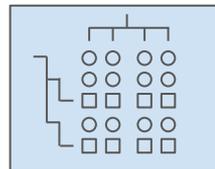
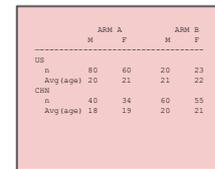


table object
cell values,
table structure,
format instructions,
paths

A diagram representing a formatted table with a pink background and a table structure. The table has columns for 'ARM A' and 'ARM B', each with sub-columns for 'M' and 'F'. The rows represent different variables: 'n', 'Avg (age)', and 'n' again, with values for each combination of arm and gender.

	ARM A		ARM B	
	M	F	M	F
US				
n	80	60	20	23
Avg (age)	20	21	21	22
CEM				
n	45	34	60	55
Avg (age)	18	19	20	21

formatted table
such as ASCII, pdf,
rtf

Layouts declare tables (pre-data)

Table layouts are declared pre-data (symbolically describe the table structure)

- Faceting (row and column)
 - `split_rows_by()` (and sibling funs), `split_cols_by()` (and sibling funs)
- cell value derivation
 - Via `analyze()` and `summarize_row_groups()`
- every layout starts with `basic_table()`
 - Metadata (titles, footer)
 - Display of column counts

Deriving cell values with analyze()

```
lyt <- basic_table() |>  
  analyze("AGE")
```

	all obs
Mean	34.88

```
build_table(lyt, ex_adsl)
```

```
fivenum_afun <- function(x) {  
  in_rows(n = sum(!is.na(x)),  
    "mean (sd)" = c(mean(x), sd(x)),  
    median = median(x),  
    "min - max" = range(x),  
    .formats = c(n = "xx",  
      "mean (sd)" = "xx.x (xx.x)",  
      median = "xx.x",  
      "min - max" = "xx.x - xx.x"))  
}
```

```
lyt2 <- basic_table() %>% analyze("AGE", fivenum_afun)
```

```
build_table(lyt2, ex_adsl)1
```

	all obs
n	400
mean (sd)	34.9 (7.4)
median	34.0
min - max	20.0 - 69.0

Deriving cell values with analyze()

```
lyt <- basic_table() |>  
  analyze("AGE")
```

	all obs
Mean	34.88

```
build_table(lyt, ex_adsl)
```

```
fivenum_afun <- function(x) {  
  in_rows(n = sum(!is.na(x)),  
    "mean (sd)" = c(mean(x), sd(x)),  
    median = median(x),  
    "min - max" = range(x),  
    .formats = c(n = "xx",  
      "mean (sd)" = "xx.x (xx.x)",  
      median = "xx.x",  
      "min - max" = "xx.x - xx.x"))  
}
```

	all obs
n	400
mean (sd)	34.9 (7.4)
median	34.0
min - max	20.0 - 69.0

```
lyt2 <- basic_table() %>% analyze("AGE", fivenum_afun)
```

```
build_table(lyt2, ex_adsl)
```

Analysis - cell value derivation

So far we have seen how layouts are used to define facets.

- Analyses define how the data facet should be summarized and displayed
- The two main analyses functions are
 - `analyze`
 - `summarize_row_groups`
- An analysis can return cells for multiple rows with `in_rows()`
- Cell value formatting can be done with `rcell`, and the various format arguments

Analyzing More Than One Variable Within a Facet

- analyze calls can be called sequentially
 - nested = TRUE (the default) combines them within facets
 - nested = FALSE turns this off and generates a new top-level subtable
- analyze can be applied to multiple variables
 - analyze accepts a list of analysis functions in this case

They are equivalent (by default).

Analysis Functions - Additional Arguments

- When deriving count & percentages one needs access to the column population N
- Analysis functions can optionally accept a number of arguments:
 - `.N_col` for column count
 - `.N_total` for total count
 - `.spl_context` for row-faceting context (see `?spl_context`)
 - `.var` for the name of the variable being analyzed
 - And others (see `?analyze`)



advanced topic, covered in pt 2

Percentages

```
pct_afun <- function(x, .N_col) {  
  rcell(  
    sum(!is.na(x)) * c(1, 1/.N_col),  
    format = "xx (xx.x%)"  
  )  
}
```

```
lyt <- basic_table() |>  
  analyze("AGE", pct_afun)
```

```
build_table(lyt, ex_adsl)
```

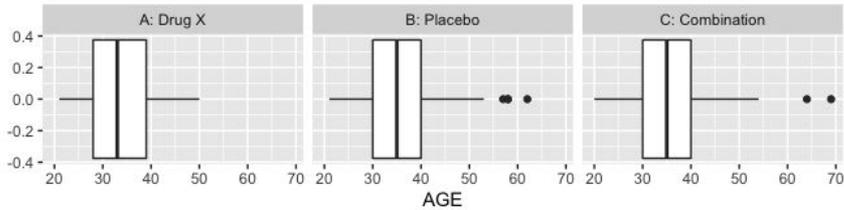
	all obs
pct_afun	400 (100.0%)

Declaring Facets

- `split_rows_by()` (and sibling funcs) add row faceting structure
- `split_cols_by()` (and sibling funcs) add column faceting structure
- Column and row facet structure declared independently
 - As in `facet_grid(rows = , cols =)`

Column Faceting - ggplot2 and rtables

```
ggplot(ex_adsl, mapping = aes(x = AGE)) +  
  geom_boxplot() +  
  facet_grid(cols = vars(ARM))
```

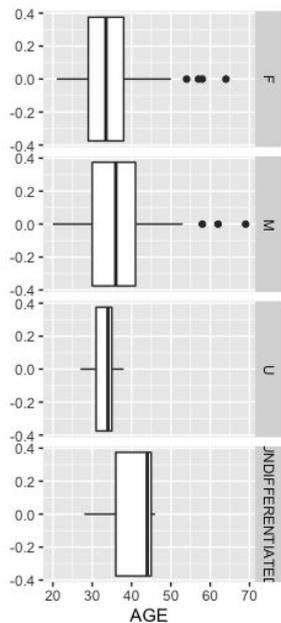


```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  analyze("AGE", range, format = "xx.xx - xx.xx")  
  
build_table(lyt, ex_adsl)
```

	A: Drug X	B: Placebo	C: Combination
range	21.00 - 50.00	21.00 - 62.00	20.00 - 69.00

Row Faceting - ggplot2 and rtables

```
ggplot(ex_adsl, mapping = aes(x = AGE)) +  
  geom_boxplot() +  
  facet_grid(rows = vars(SEX))
```



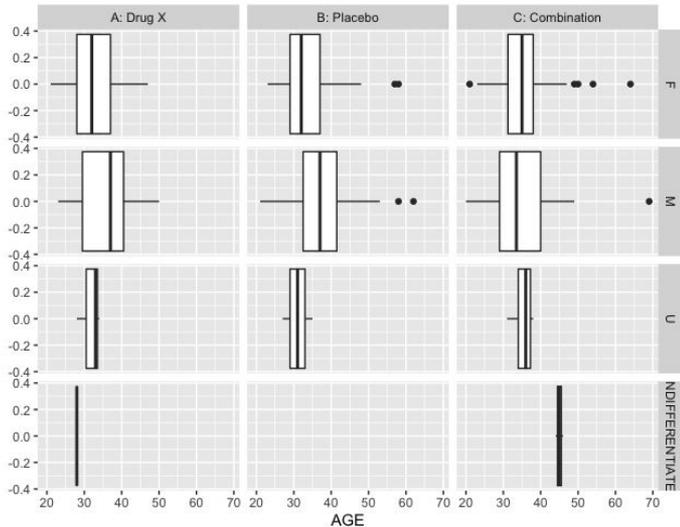
```
lyt2 <- basic_table() |>  
  split_rows_by("SEX") |>  
  analyze("AGE", range,  
    format = "xx.xx - xx.xx")
```

```
build_table(lyt2, ex_adsl)
```

	all obs
F	
range	21.00 - 64.00
M	
range	20.00 - 69.00
U	
range	27.00 - 38.00
JNDIFFERENTIATED	
range	28.00 - 46.00

Grid Faceting - ggplot2 and rtables

```
ggplot(ex_adsl, mapping = aes(x = AGE)) +  
  geom_boxplot() +  
  facet_grid(rows = vars(SEX),  
             cols = vars(ARM))
```



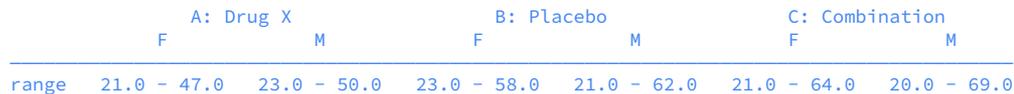
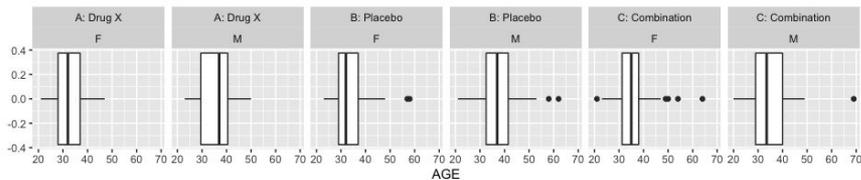
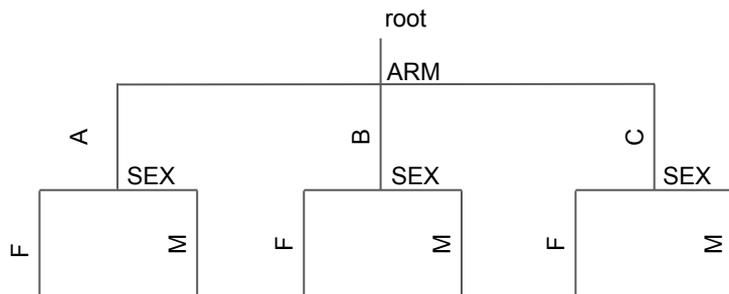
```
lyt3 <- basic_table() |>  
  split_cols_by("ARM") |>  
  split_rows_by("SEX") |>  
  analyze("AGE", range, format = "xx.xx - xx.xx")
```

```
build_table(lyt3, ex_adsl)
```

	A: Drug X	B: Placebo	C: Combination
F			
range	21.00 - 47.00	23.00 - 58.00	21.00 - 64.00
M			
range	23.00 - 50.00	21.00 - 62.00	20.00 - 69.00
U			
range	28.00 - 34.00	27.00 - 35.00	31.00 - 38.00
UNDIFFERENTIATED			
range	28.00 - 28.00	Inf - -Inf	44.00 - 46.00

Nested Faceting Structure

Consecutive splits give nested facet structure, same as giving multiple variables in one dim to `facet_grid()`



Sneak peak into table objects

```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  split_cols_by("B1HL") |>  
  split_rows_by("SEX") |>  
  analyze("AGE", function(x) "")
```

```
tbl <- build_table(lyt, ex_adsl3)
```

```
col_paths_summary(tbl)
```

```
row_paths_summary(tbl)
```

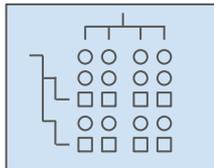
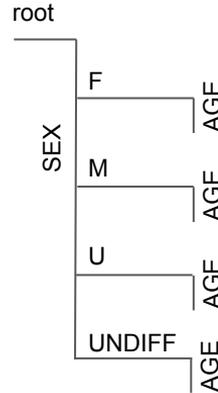
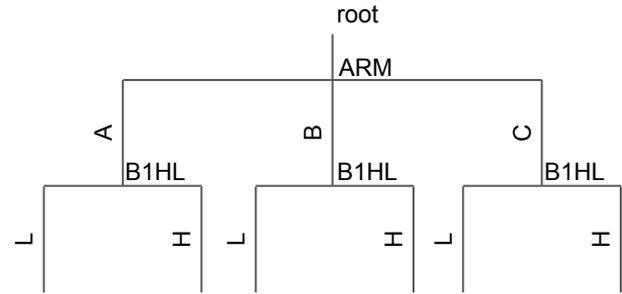
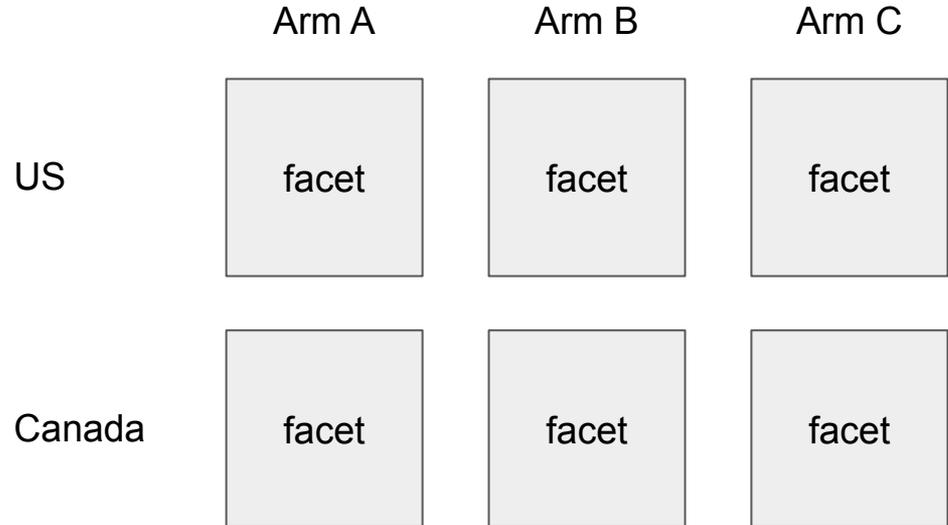


table object
numbers, strings
paths



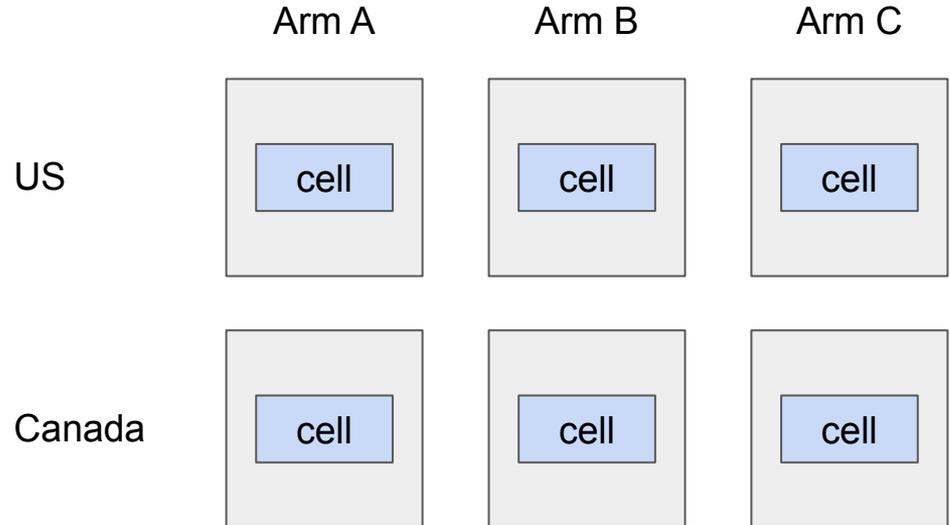
Analyze revisited

- The analysis function is applied *within each facet* declared by the splitting it is nested within, generating **one or more cell values** via `in_rows()`



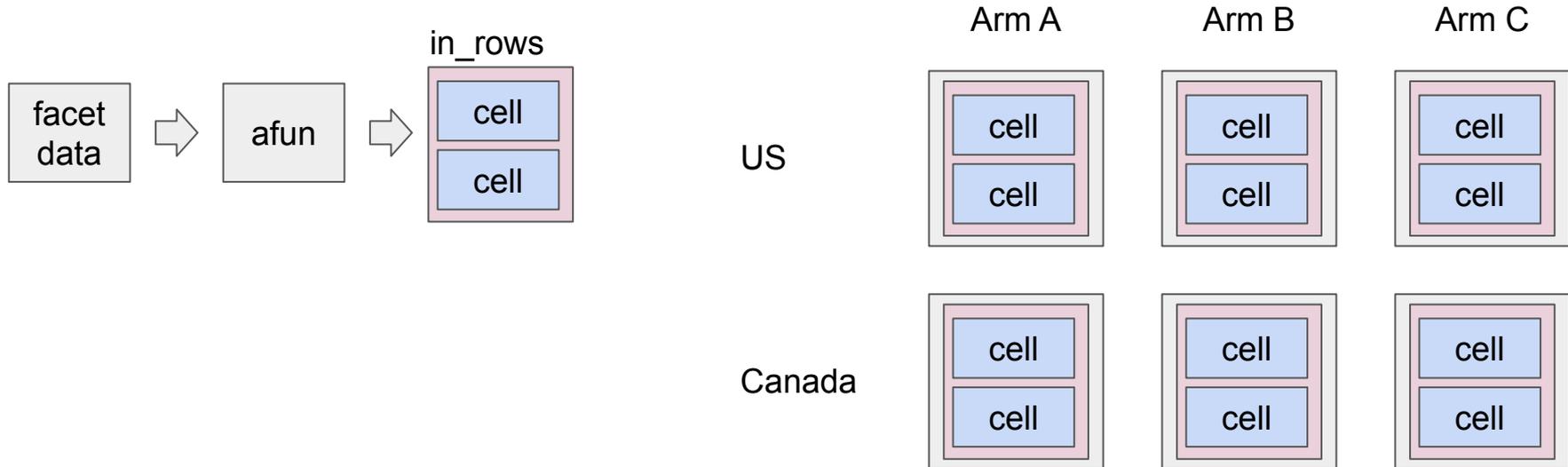
Analyze revisited

- The analysis function is applied *within each facet* declared by the splitting it is nested within, generating **one or more cell values** via `in_rows()`



Analyze revisited

- The analysis function is applied *within each facet* declared by the splitting it is nested within, generating **one or more cell values** via `in_rows()`



Analyze revisited

- The analysis function is applied *within each facet* declared by the splitting it is nested within, generating **one or more cell values** via `in_rows()`

```
lyt2 <- basic_table() %>%
  split_cols_by("ARM") %>%
  split_rows_by("STRATA1") %>%
  analyze("AGE", afun = function(x) {
    in_rows(n = sum(!is.na(x)),
            "mean (sd)" = c(mean(x), sd(x)),
            "median" = median(x),
            "min - max" = range(x),
            .formats = c(n = "xx",
                          "mean (sd)" = "xx.x (xx.x)",
                          median = "xx.x",
                          "min - max" = "xx.x - xx.x"))
  })
build_table(lyt2, ex_adsl)
```

```
> build_table(lyt2, ex_adsl)
```

	A: Drug X	B: Placebo	C: Combination
<hr/>			
A			
n	38	44	40
mean (sd)	33.1 (5.7)	35.1 (7.9)	34.2 (6.2)
median	32.0	32.5	35.0
min - max	24.0 - 46.0	23.0 - 62.0	20.0 - 47.0
B			
n	47	45	43
mean (sd)	33.9 (7.2)	36.0 (9.1)	36.3 (8.4)
median	33.0	36.0	35.0
min - max	23.0 - 48.0	21.0 - 58.0	21.0 - 64.0
C			
n	49	45	49
mean (sd)	34.2 (6.6)	35.2 (6.6)	35.6 (8.2)
median	34.0	35.0	35.0
min - max	21.0 - 50.0	23.0 - 51.0	24.0 - 69.0

Analyze revisited

- The analysis function is applied *within each facet* declared by the splitting it is nested within, generating **one or more cell values** via `in_rows()`

```
lyt2 <- basic_table() %>%
  split_cols_by("ARM") %>%
  split_rows_by("STRATA1") %>%
  analyze("AGE", afun = function(x) {
    in_rows(n = sum(!is.na(x)),
            "mean (sd)" = c(mean(x), sd(x)),
            median = median(x),
            "min - max" = range(x),
            .formats = c(n = "xx",
                         "mean (sd)" = "xx.x (xx.x)",
                         median = "xx.x",
                         "min - max" = "xx.x - xx.x"))
  })
build_table(lyt2, ex_adsl)
```

```
> build_table(lyt2, ex_adsl)
```

	A: Drug X	B: Placebo	C: Combination
<hr/>			
A			
n	38	44	40
mean (sd)	33.1 (5.7)	35.1 (7.9)	34.2 (6.2)
median	32.0	32.5	35.0
min - max	24.0 - 46.0	23.0 - 62.0	20.0 - 47.0
B			
n	47	45	43
mean (sd)	33.9 (7.2)	36.0 (9.1)	36.3 (8.4)
median	33.0	36.0	35.0
min - max	23.0 - 48.0	21.0 - 58.0	21.0 - 64.0
C			
n	49	45	49
mean (sd)	34.2 (6.6)	35.2 (6.6)	35.6 (8.2)
median	34.0	35.0	35.0
min - max	21.0 - 50.0	23.0 - 51.0	24.0 - 69.0

Core Types Of Split

Variable Split

```
split_rows_by("<var>"), split_cols_by("<var>")
```

- Used for *categorical variables*
- One facet per **variable level**
 - Including empty levels for factors
- Facet data is incoming data subset to `<var> == <single level>`
- Most common, basic split

Multivar Split

```
split_rows_by_multivar(<vector of varnames>),  
split_cols_by_multivar(<vector of varnames>)
```

- One facet per ***variable***
- **Facet data is full incoming data**
- Most common/useful in *column space*
- Useful when
 - Incoming data has precalculated statistics in columns
 - E.g., *model summary display*
 - Incoming data is in (very) wide form
 - E.g., different columns for measurements from Visit 1, Visit 2, ...
- **Must use `analyze_colvars()` for cell content derivation**
 - Analysis function can be different for different variables (passed as list)

Multivar Split Example

```
lyt <- basic_table() %>%  
  split_cols_by_multivar(c("RACE", "AGE"),  
                        varlabels = c("Ethn. Present", "Ave. Age")) %>%  
  analyze_colvars(afun = list(RACE = function(x) length(unique(x)),  
                             AGE = function(x) rcell(mean(x), format = "xx.x")))
```

```
> build_table(lyt, DM)  
Ethn. Present Ave. Age  
-----  
3           34.2
```

Static Cut Splits

```
split_rows_by_cuts("<var>", cuts = <>, cumulative = <>),  
split_cols_by_cuts("<var>", cuts = <>, cumulative = <>)
```

- Used to split on values of *numeric/continuous variables*
- One facet per **discretized value** of <var>
- Cut points for discretization are *fixed, data independent*
- Categories can be cumulative (cumulative = TRUE)

Static Cut Split Examples

```
lyt_scut <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "30-40", "40-50", "50+")) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut, DM)
```

	<30 (N=122)	30-40 (N=178)	40-50 (N=45)	50+ (N=11)
Mean	5.98	5.63	6.18	6.68

```
lyt_scut_cum <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "<40", "<50", "All"),  
    cumulative = TRUE) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut_cum, DM)
```

	<30 (N=122)	<40 (N=300)	<50 (N=345)	All (N=356)
Mean	5.98	5.77	5.83	5.85

Static Cut Split Examples

```
lyt_scut <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "30-40", "40-50", "50+")) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut, DM)
```

	<30 (N=122)	30-40 (N=178)	40-50 (N=45)	50+ (N=11)
Mean	5.98	5.63	6.18	6.68

```
lyt_scut_cum <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "<40", "<50", "All"),  
    cumulative = TRUE) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut_cum, DM)
```

	<30 (N=122)	<40 (N=300)	<50 (N=345)	All (N=356)
Mean	5.98	5.77	5.83	5.85

Static Cut Split Examples

```
lyt_scut <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "30-40", "40-50", "50+")) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut, DM)
```

	<30 (N=122)	30-40 (N=178)	40-50 (N=45)	50+ (N=11)
Mean	5.98	5.63	6.18	6.68

```
lyt_scut_cum <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cuts("AGE", cuts = c(1, 30, 40, 50, 1000),  
    cutlabels = c("<30", "<40", "<50", "All"),  
    cumulative = TRUE) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_scut_cum, DM)
```

	<30 (N=122)	<40 (N=300)	<50 (N=345)	All (N=356)
Mean	5.98	5.77	5.83	5.85

Dynamic Cut Splits

```
split_rows_by_cutfun("<var>", cutfun = <>, cumulative = <>),  
split_cols_by_cutfun("<var>", cutfun = <>, cumulative = <>)
```

- Used to split on values of *numeric/continuous variables*
- One facet per ***discretized value*** of <var>
- Cut points are *dynamic, depend on data*
 - Cut points are calculated once, by ***applying cut fun to data for full table***
 - Cuts identical applications of this split, ***not dependent on any splitting it is nested within***
- Can be cumulative (cumulative = TRUE)

Cut Functions and CutLabel Functions

Cut Function

- Accepts (full-table) data vector for variable
- Returns vector of break points (including lower and upper bound)
 - Like `quantile()`
 - Can be named, if so, labels for upper limit of each cut is used
 - Ignored if `cutlabel` function is specified

CutLabel Function

- Accepts output of `cutfun`
- Returns vector of labels to use

Dynamic Cut Examples

```
helper <- function(vec) paste(round(vec, 1),  
                             collapse = " - ")
```

```
mycutfun <- function(x) {  
  ret <- quantile(x, c(0, .33, .66, 1))  
  forlab <- floor(ret)  
  names(ret) <- c("",  
                 helper(ret[1:2]),  
                 helper(ret[2:3]),  
                 helper(ret[3:4]))  
  ret  
}
```

```
mycumlabfun <- function(x) c(helper(x[1:2]),  
                             helper(x[c(1,3)]),  
                             "All")
```

```
lyt_dyncut<- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cutfun("AGE", cutfun = mycutfun) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_dyncut, DM)
```

	20 - 30 (N=122)	30 - 36 (N=114)	36 - 60 (N=120)
Mean	5.98	5.67	5.90

```
lyt_dyncut_cum <- basic_table(show_colcounts = TRUE) %>%  
  split_cols_by_cutfun("AGE", cutfun = mycutfun,  
                       cutlabelfun = mycumlabfun,  
                       cumulative = TRUE) %>%  
  analyze("BMRKR1")
```

```
> build_table(lyt_dyncut_cum, DM)
```

	20 - 30 (N=122)	20 - 36 (N=236)	All (N=356)
Mean	5.98	5.83	5.85

Mix and Match

All of these split types can be

- Used in both column and row space
 - Though again, multivar doesn't make much sense in row space
- Nested within each other
 - Though nesting splits inside a multivar doesn't make much sense



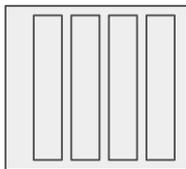
The main points to take away by now are ...

- rables is a sophisticated end 2 end table framework
- tables are faceted visualizations
- rtables tables are created with layouts and data
- layouts declare facets (split_* functions), analyses (analyze function)
- you can read the following code and predict the table structure:

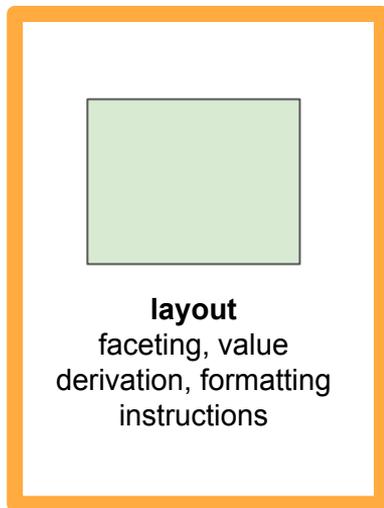
```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  split_cols_by("SEX") |>  
  split_rows_by("STRATA1") |>  
  analyze("AGE", range, format = "xx.xx - xx.xx")  
  
build_table(lyt, ex_adsl)
```



So whats next...



data.frame
Unaggregated data
(variables + obs)



layout
faceting, value
derivation, formatting
instructions

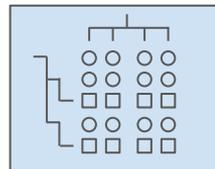
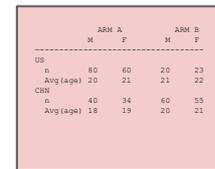


table object
cell values,
table structure,
format instructions,
paths



	ARM A		ARM B	
	M	F	M	F
US				
n	80	60	20	23
avg (age)	20	21	21	22
CEU				
n	45	34	60	55
avg (age)	18	19	20	21

formatted table
such as ASCII, pdf,
rtf

**We have one more topic to cover
for the layouts section**



Group summaries

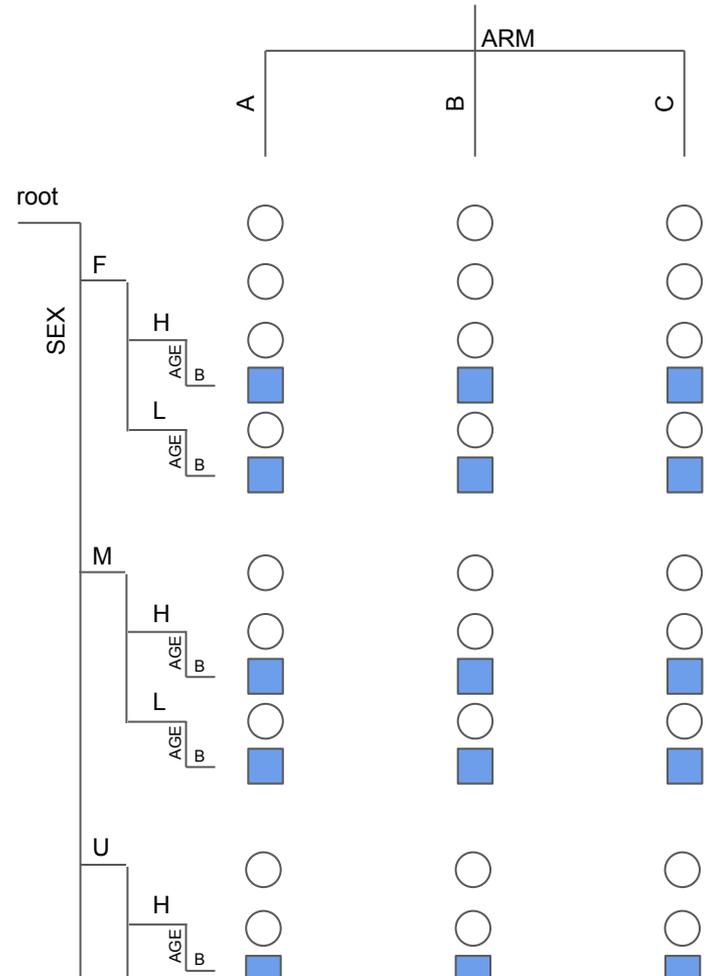
```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \(x) list(B = "a"))

build_table(lyt, ex_adsl3)
```

Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \ (x) list(B = "a"))

build_table(lyt, ex_adsl3)
```

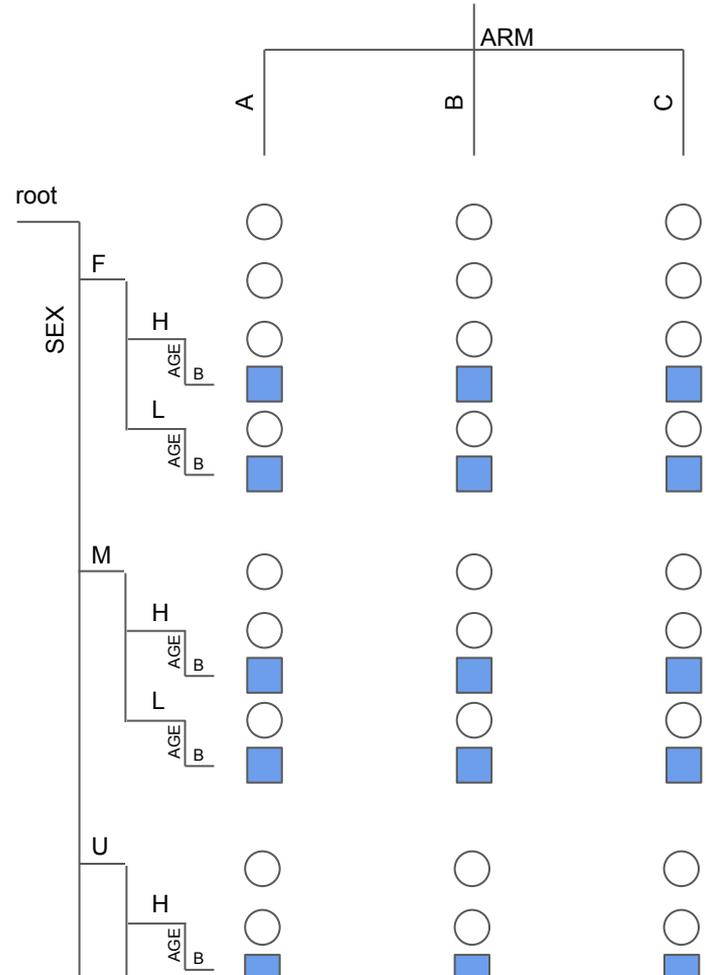


Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \ (x) list(B = "a"))
```

```
build_table(lyt, ex_adsl3)
```

Note, analyze can return multiple rows with `in_rows()`

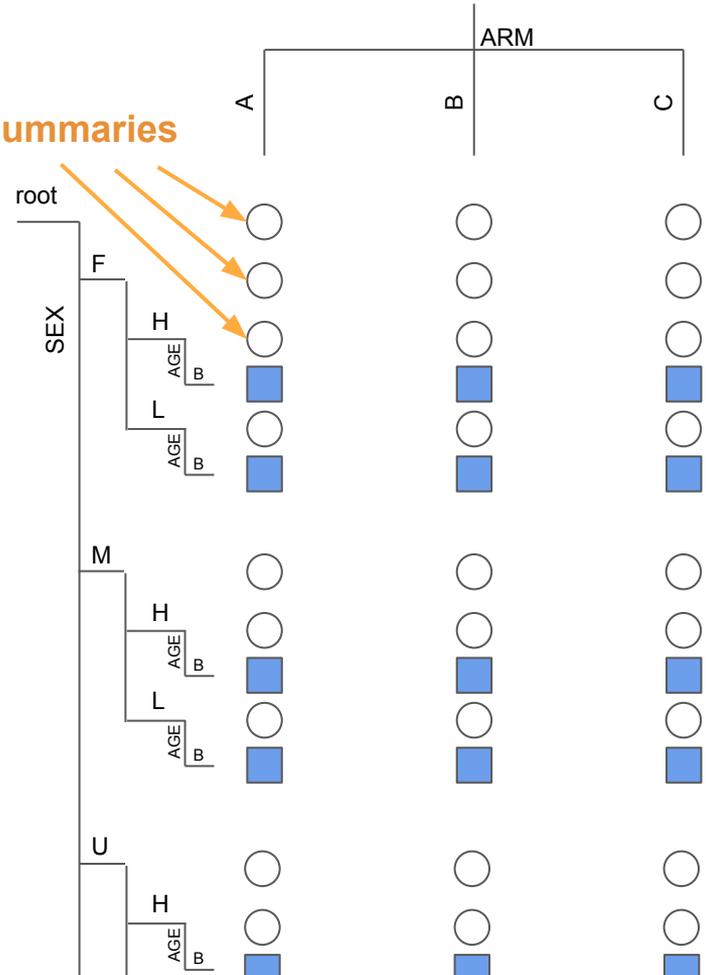


Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \ (x) list(B = "a"))

build_table(lyt, ex_adsl3)
```

3 levels of group summaries

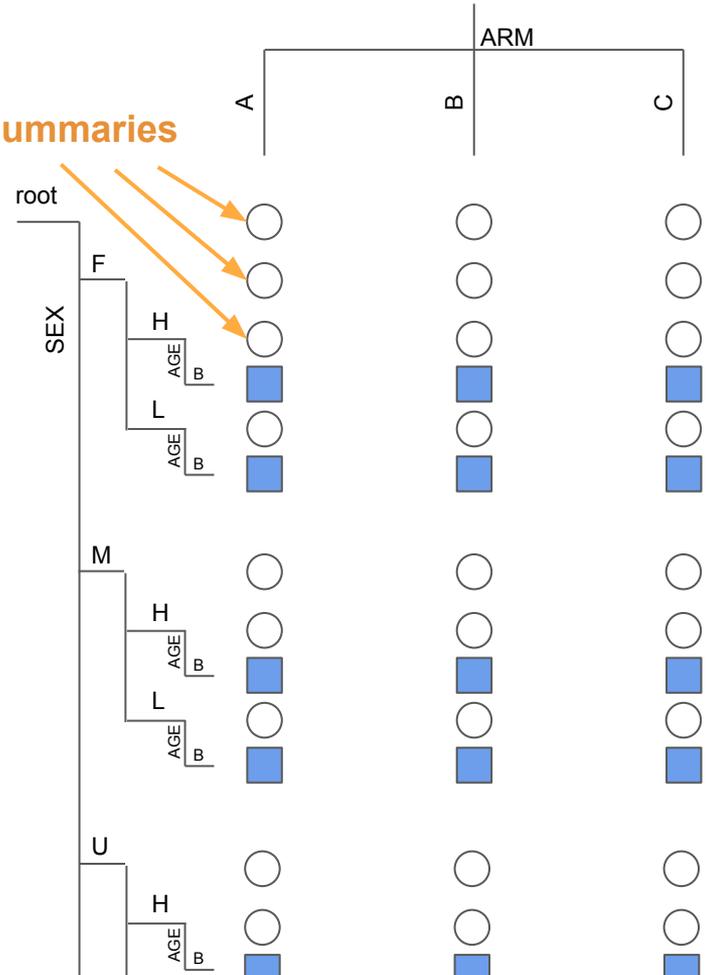


Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \ (x) list(B = "a"))
```

3 levels of group summaries

```
build_table(lyt, ex_adsl3)
```



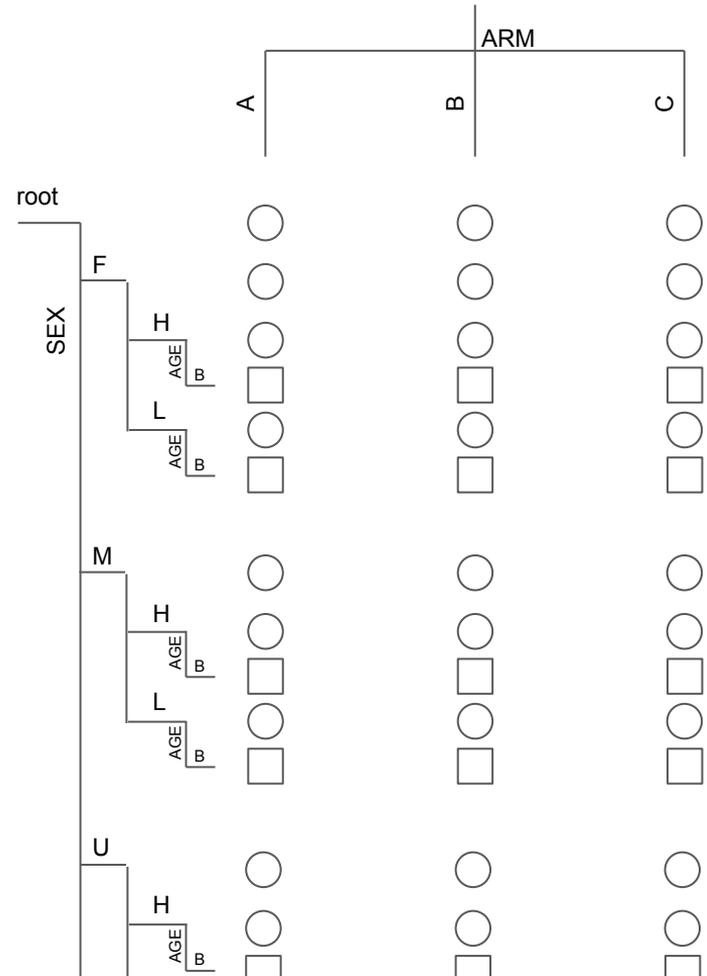


layout

Group summaries

```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  split_rows_by("SEX") |>  
  split_rows_by("B1HL") |>  
  analyze("AGE", \ (x) list(B = "a"))
```

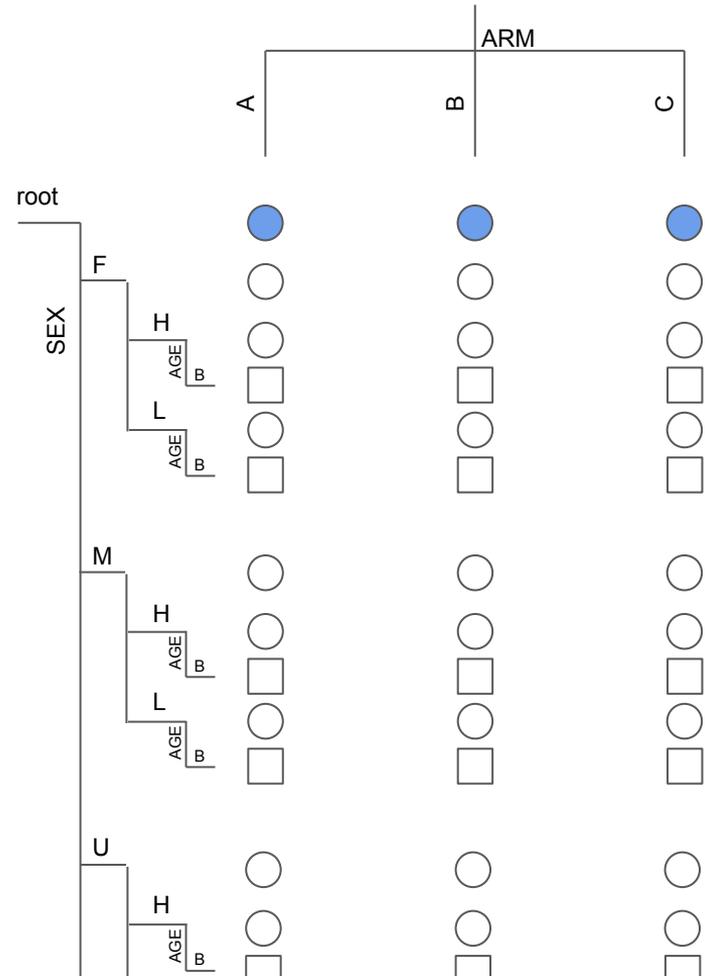
```
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  summarize_row_groups() |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \(x) list(B = "a"))
```

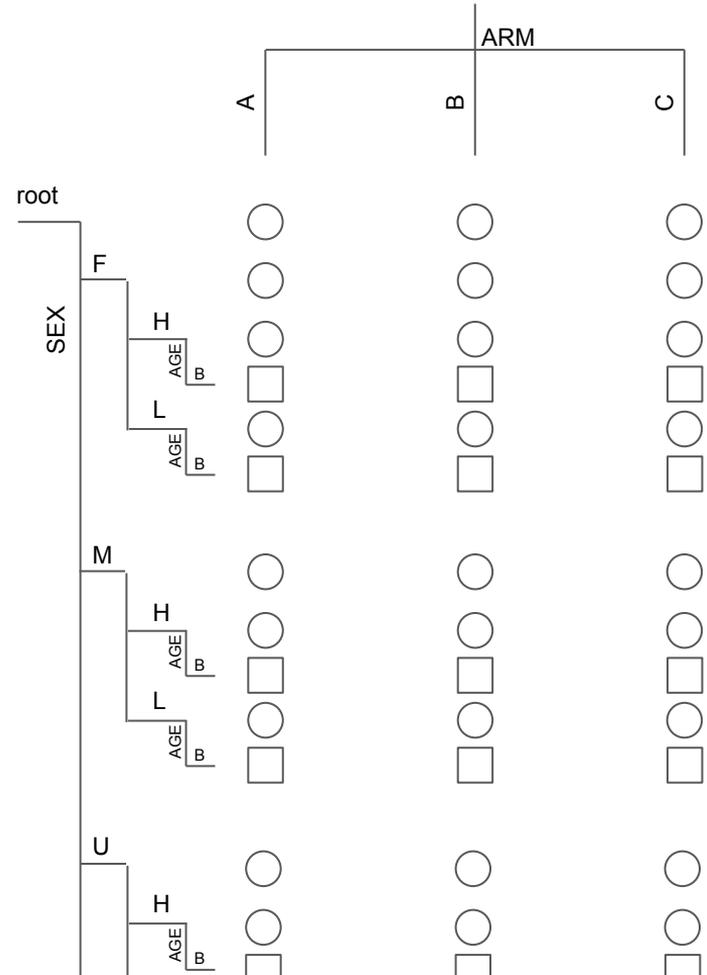
```
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \(x) list(B = "a"))

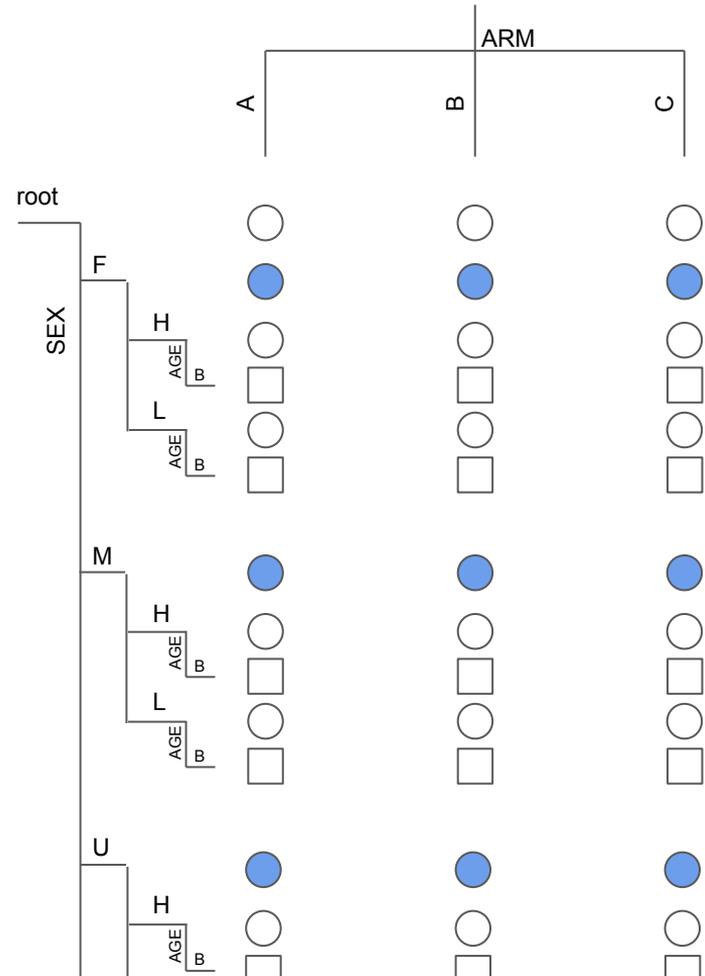
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  summarize_row_groups() |>
  split_rows_by("B1HL") |>
  analyze("AGE", \(x) list(B = "a"))
```

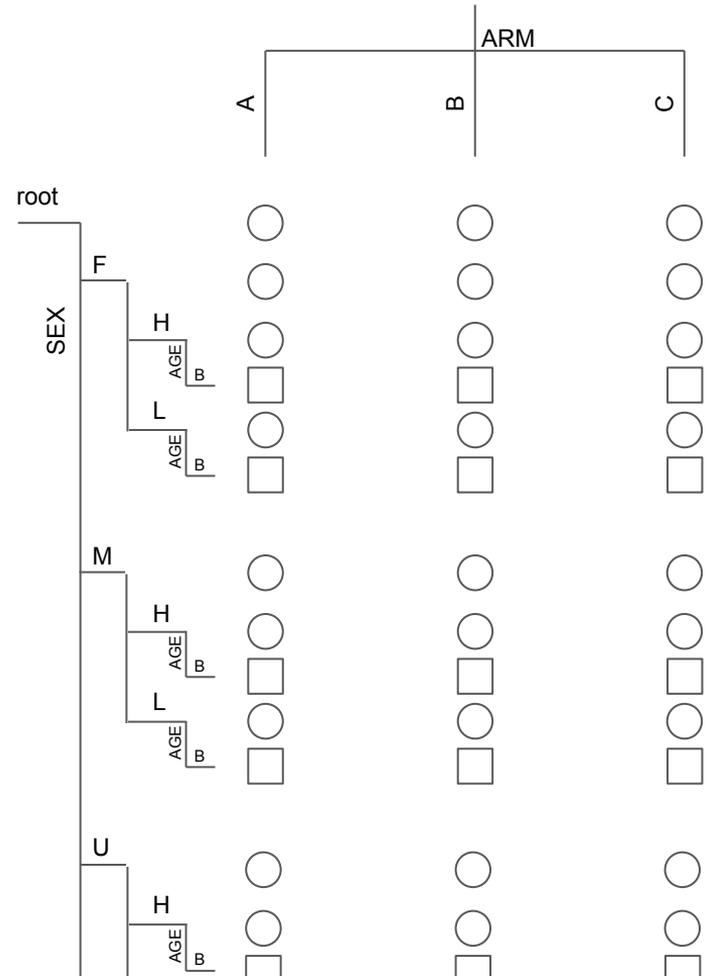
```
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \(x) list(B = "a"))

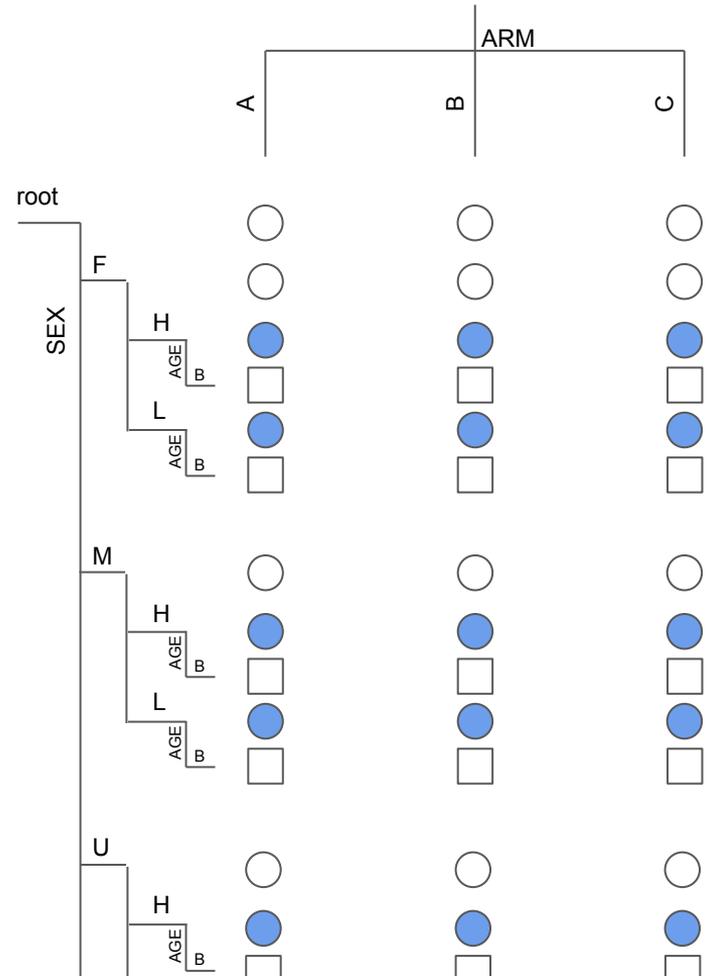
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  summarize_row_groups() |>
  analyze("AGE", \(x) list(B = "a"))
```

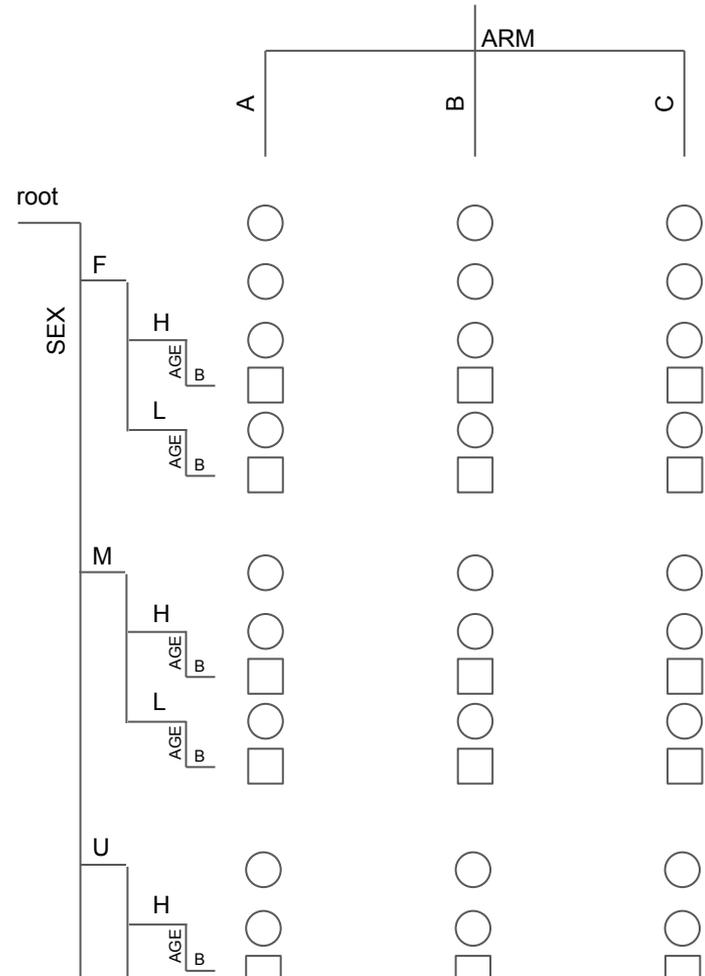
```
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>
  split_cols_by("ARM") |>
  split_rows_by("SEX") |>
  split_rows_by("B1HL") |>
  analyze("AGE", \ (x) list(B = "a"))
```

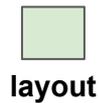
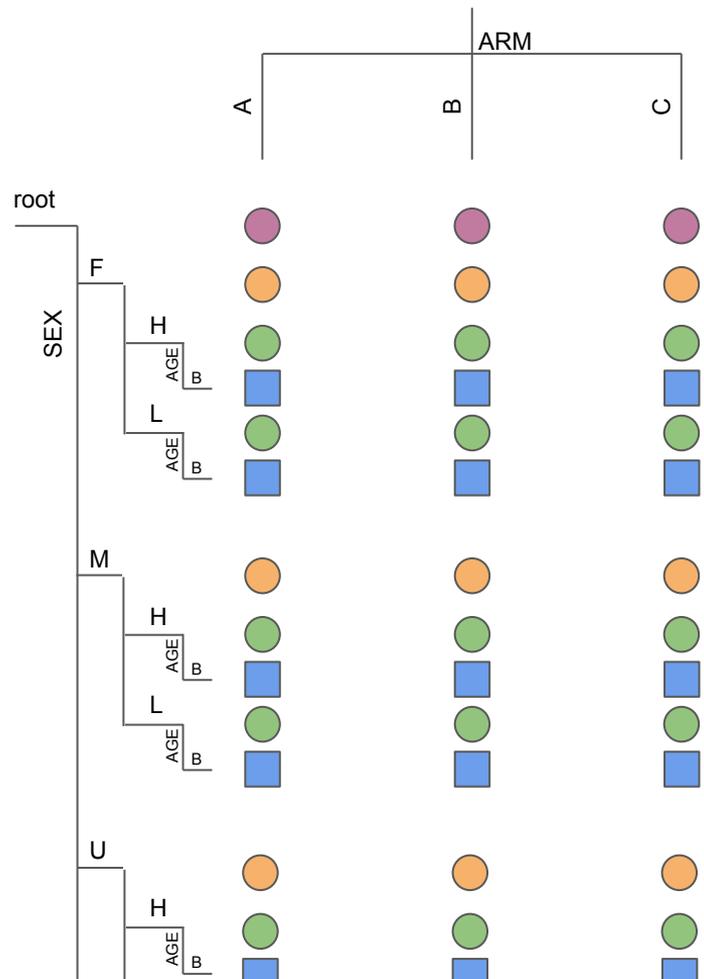
```
build_table(lyt, ex_adsl3)
```



Group summaries

```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  summarize_row_groups() |>  
  split_rows_by("SEX") |>  
  summarize_row_groups() |>  
  split_rows_by("B1HL") |>  
  summarize_row_groups() |>  
  analyze("AGE", afun = \(x) list(B = "a"))
```

```
build_table(lyt, ex_adsl3)
```





layout

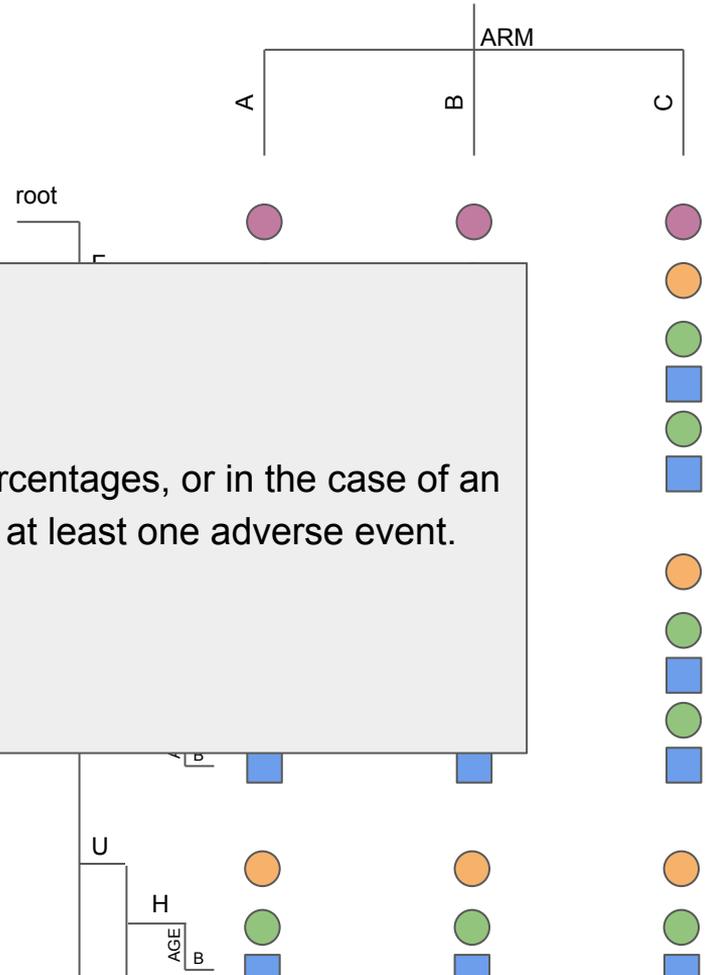
Group summaries

```

lyt <- basic_table() |>
  split_cols_by("ARM") |>
  summarize_row_groups() |>
  split_rows_by("AGE") |>
  summarize_row_groups() |>
  split_rows_by("U") |>
  summarize_row_groups() |>
  analyze("A") |>
  build_table()

```

Usually group summaries hold counts, percentages, or in the case of an adverse events table unique patients with at least one adverse event.



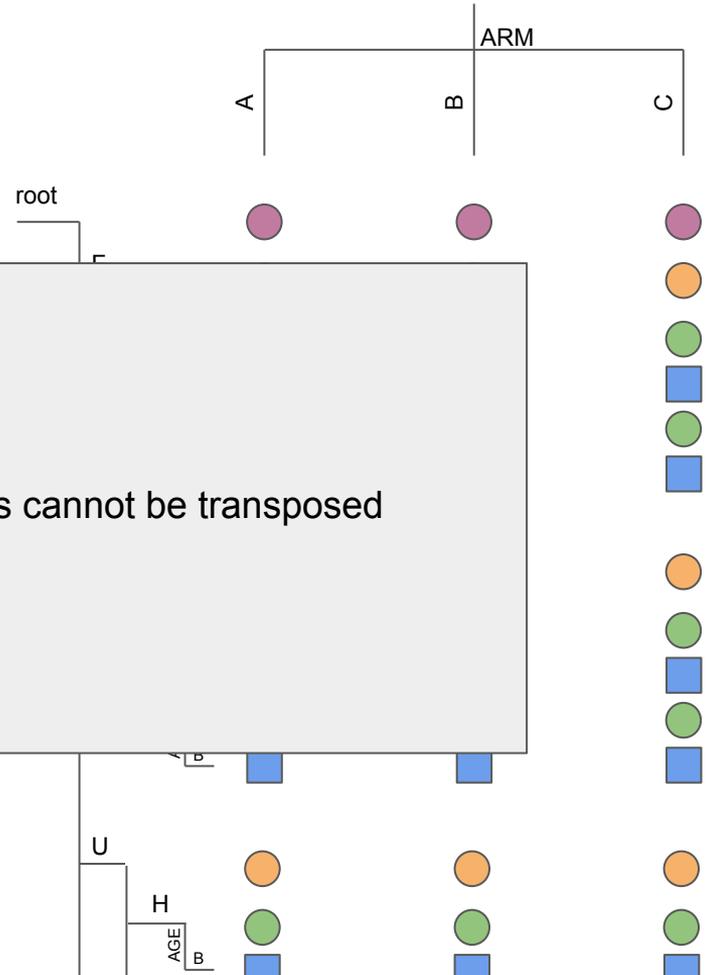


layout

Group summaries

```
lyt <- basic_table() |>  
  split_cols_by("ARM") |>  
  summarize_row_groups() |>  
  split_row_groups() |>  
  summarize_row_groups() |>  
  split_row_groups() |>  
  summarize_row_groups() |>  
  analyze("a") |>  
  build_table()
```

Note: this is one reason why rtables tables cannot be transposed



Group Summary -> Content Table/Rows

The table of rows resulting from a `summary_row_group` layout directive is called that facet's **content table**

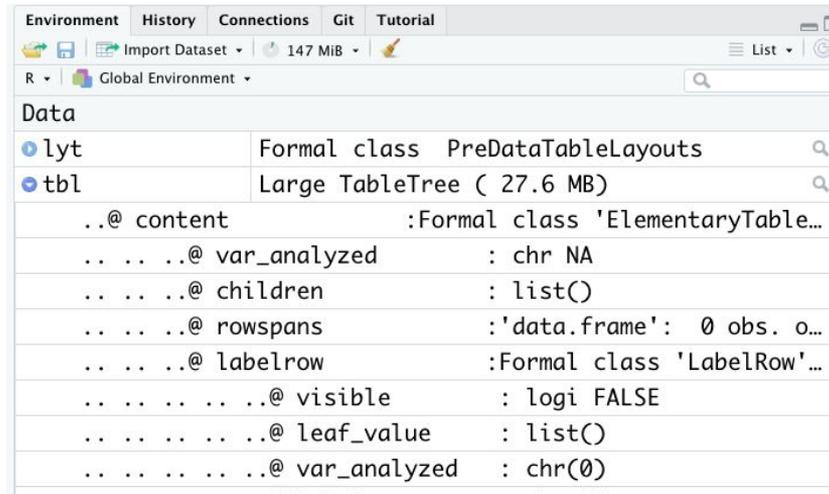
- Artifact from *very* early in the design process
 - Unfortunate
 - Can't be easily changed at this stage



Table Structure

Table objects

- rtables table objects are implemented with a tree data structure
- Directly inspecting table objects' low-level structure is not going to be helpful



The screenshot shows the RStudio Environment pane. The 'Data' section contains two objects: 'lyt' (Formal class PreDataTableLayouts) and 'tbl' (Large TableTree (27.6 MB)). The 'tbl' object is expanded to show its internal structure, which is a tree of objects. The structure is as follows:

tbl	Large TableTree (27.6 MB)
..@ content	:Formal class 'ElementaryTable...'
..@ var_analyzed	: chr NA
..@ children	: list()
..@ rowspans	:'data.frame': 0 obs. o...
..@ labelrow	:Formal class 'LabelRow'...
..@ visible	: logi FALSE
..@ leaf_value	: list()
..@ var_analyzed	: chr(0)

Table objects

- To learn about the structure of a table object:
 - `table_structure()`, `make_row_df()`, `make_col_df`, `row_paths`, `col_paths`
 - `dim()`, `nrow()`, `ncol()`
 - **NOT** `str()` (I mean it, it will not help you)
- rtables table objects are implemented with a tree data structure
 - You won't need to know this beyond understanding pathing
 - Useful for lots of functionality internally
 - Pagination
 - subsetting

Consider A Non-trivial Table

	ARM1		ARM2	
	Male (N=256)	Female (N=248)	Male (N=248)	Female (N=248)
Caucasian (n)	116 (45.3%)	144 (58.1%)	119 (48.0%)	119 (48.0%)
Level A	37 (14.5%)	48 (19.4%)	42 (16.9%)	35 (14.1%)
Age Analysis				
mean	56.42	55.57	56.19	54.53
median	55.91	55.42	58.40	51.73
Age Analysis redux				
range	40.1 - 69.9	40.8 - 69.7	42.2 - 69.7	41.7 - 69.3
Level B	44 (17.2%)	52 (21.0%)	37 (14.9%)	40 (16.1%)
Age Analysis				
mean	54.28	55.24	54.22	54.92
median	54.71	55.47	54.96	55.17
Age Analysis redux				
range	41.2 - 69.5	40.5 - 69.0	40.0 - 68.5	40.0 - 69.2
African American (n)	140 (54.7%)	104 (41.9%)	129 (52.0%)	129 (52.0%)
Level A	45 (17.6%)	40 (16.1%)	48 (19.4%)	44 (17.7%)
Age Analysis				
mean	55.77	55.33	56.26	54.30
median	55.06	54.39	57.20	53.94
Age Analysis redux				
range	42.7 - 69.6	40.4 - 69.3	41.3 - 69.3	40.7 - 68.9
Level B	45 (17.6%)	29 (11.7%)	44 (17.7%)	55 (22.2%)
Age Analysis				
mean	53.85	56.55	56.80	55.88
median	54.36	57.21	57.39	55.71
Age Analysis redux				
range	40.6 - 68.1	40.4 - 69.3	40.1 - 69.0	40.3 - 69.9
Var3 Counts				
level1	121	136	142	120
level2	135	112	106	128

We render this as a rectangular display, but *model* it with structure that is

1. Semantically meaningful
2. Reflects the layout used to create it

Table Structure

```
> table_structure(tbl3)
[TableTree] root
  [TableTree] RACE
    [TableTree] WHITE [cont: 1 x 4]
      [TableTree] FACTOR2
        [TableTree] A [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
        [TableTree] B [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
      [TableTree] BLACK [cont: 1 x 4]
        [TableTree] FACTOR2
          [TableTree] A [cont: 1 x 4]
            [ElementaryTable] AGE (2 x 4)
            [ElementaryTable] AgeRedux (1 x 4)
          [TableTree] B [cont: 1 x 4]
            [ElementaryTable] AGE (2 x 4)
            [ElementaryTable] AgeRedux (1 x 4)
        [ElementaryTable] VAR3 (2 x 4)
```

Table Structure and Layout

```
> table_structure(tbl3)
[TableTree] root
  [TableTree] RACE
    [TableTree] WHITE [cont: 1 x 4]
      [TableTree] FACTOR2
        [TableTree] A [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
        [TableTree] B [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
      [TableTree] BLACK [cont: 1 x 4]
        [TableTree] FACTOR2
          [TableTree] A [cont: 1 x 4]
            [ElementaryTable] AGE (2 x 4)
            [ElementaryTable] AgeRedux (1 x 4)
          [TableTree] B [cont: 1 x 4]
            [ElementaryTable] AGE (2 x 4)
            [ElementaryTable] AgeRedux (1 x 4)
    [ElementaryTable] VAR3 (2 x 4)
```

```
basic_table(show_colcounts = TRUE) %>%
  split_cols_by("ARM") %>%
  split_cols_by("SEX", "Gender", labels_var = "gend_label") %>%
  split_rows_by("RACE", "Ethnicity",
               labels_var = "ethn_label") %>%
  summarize_row_groups("RACE", label_fstr = "%s (n)") %>%
  split_rows_by("FACTOR2", "Factor2",
               split_fun = remove_split_levels("C"),
               labels_var = "fac2_label",
               label_pos = "hidden") %>%
  summarize_row_groups("FACTOR2") %>%
  analyze("AGE", "Age Analysis",
         afun = function(x) list(mean = mean(x),
                                median = median(x)),
         format = "xx.xx") %>%
  analyze("AGE",
         "Age Analysis redux",
         afun = range,
         format = "xx.x - xx.x",
         table_names = "AgeRedux"
         ) %>%
  analyze("VAR3", "Var3 Counts", afun = list_wrap_x(table),
         nested = FALSE,
         show_labels = "visible")
```

Table Structure and Layout

```
> table_structure(tbl3)
[TableTree] root
 [TableTree] RACE
  [TableTree] WHITE [cont: 1 x 4]
   [TableTree] FACTOR2
    [TableTree] A [cont: 1 x 4]
     [ElementaryTable] AGE (2 x 4)
     [ElementaryTable] AgeRedux (1 x 4)
    [TableTree] B [cont: 1 x 4]
     [ElementaryTable] AGE (2 x 4)
     [ElementaryTable] AgeRedux (1 x 4)
  [TableTree] BLACK [cont: 1 x 4]
   [TableTree] FACTOR2
    [TableTree] A [cont: 1 x 4]
     [ElementaryTable] AGE (2 x 4)
     [ElementaryTable] AgeRedux (1 x 4)
    [TableTree] B [cont: 1 x 4]
     [ElementaryTable] AGE (2 x 4)
     [ElementaryTable] AgeRedux (1 x 4)
 [ElementaryTable] VAR3 (2 x 4)
```

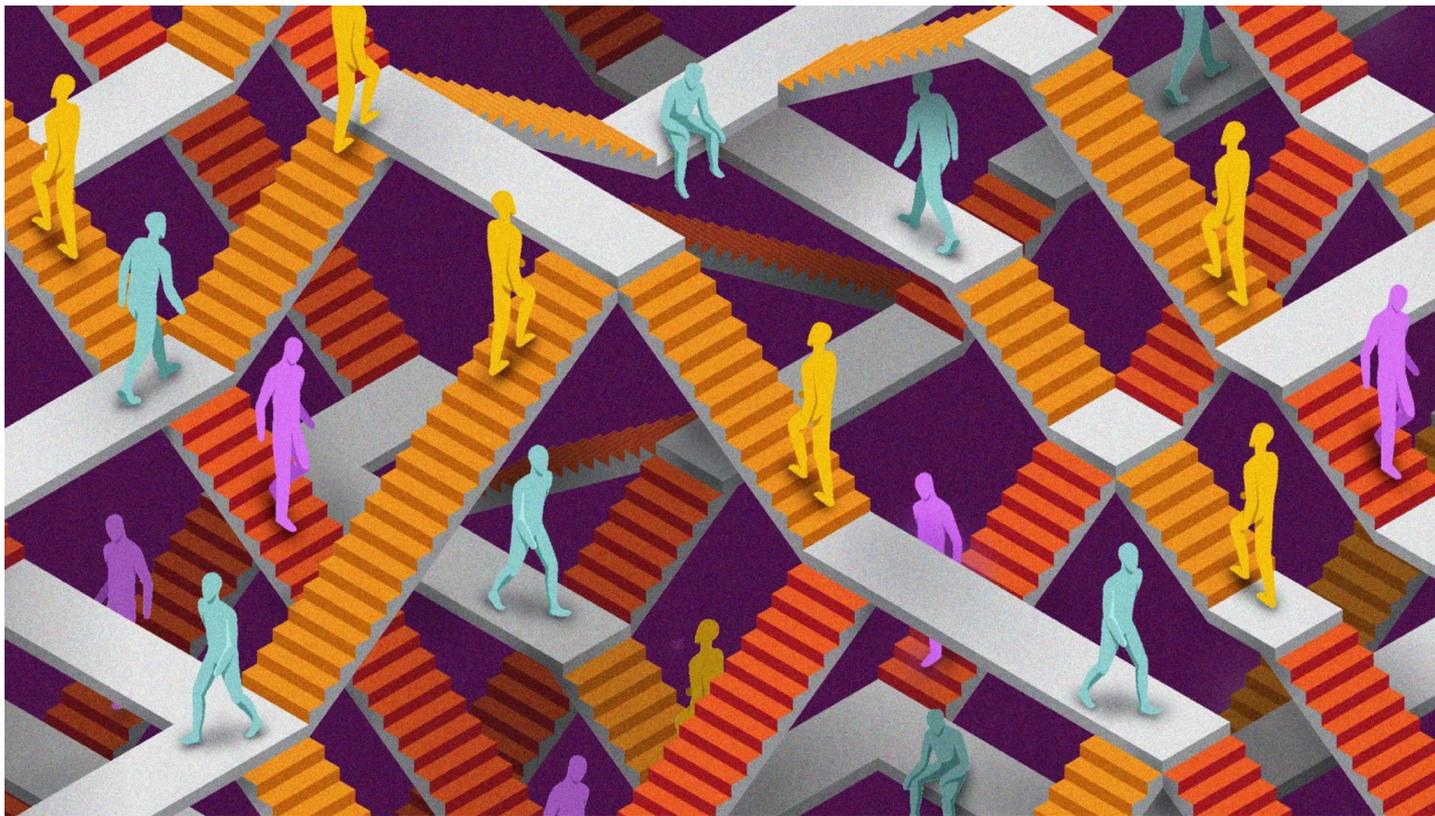
```
basic_table(show_colcounts = TRUE) %>%
  split_cols_by("ARM") %>%
  split_cols_by("SEX", "Gender", labels_var = "gend_label") %>%
  split_rows_by("RACE", "Ethnicity",
               labels_var = "ethn_label") %>%
  summarize_row_groups("RACE", label_fstr = "%s (n)") %>%
  split_rows_by("FACTOR2", "Factor2",
               split_fun = remove_split_levels("C"),
               labels_var = "fac2_label",
               label_pos = "hidden") %>%
  summarize_row_groups("FACTOR2") %>%
  analyze("AGE", "Age Analysis",
         afun = function(x) list(mean = mean(x),
                                median = median(x)),
         format = "xx.xx") %>%
  analyze("AGE",
         "Age Analysis redux",
         afun = range,
         format = "xx.x - xx.x",
         table_names = "AgeRedux"
         ) %>%
  analyze("VAR3", "Var3 Counts", afun = list_wrap_x(table),
         nested = FALSE,
         show_labels = "visible")
```

Table Structure and Layout

```
> table_structure(tbl3)
[TableTree] root
[TableTree] RACE
[TableTree] WHITE [cont: 1 x 4]
[TableTree] FACTOR2
[TableTree] A [cont: 1 x 4]
[ElementaryTable] AGE (2 x 4)
[ElementaryTable] AgeRedux (1 x 4)
[TableTree] B [cont: 1 x 4]
[ElementaryTable] AGE (2 x 4)
[ElementaryTable] AgeRedux (1 x 4)
[TableTree] BLACK [cont: 1 x 4]
[TableTree] FACTOR2
[TableTree] A [cont: 1 x 4]
[ElementaryTable] AGE (2 x 4)
[ElementaryTable] AgeRedux (1 x 4)
[TableTree] B [cont: 1 x 4]
[ElementaryTable] AGE (2 x 4)
[ElementaryTable] AgeRedux (1 x 4)
[ElementaryTable] VAR3 (2 x 4)
```

```
basic_table(show_colcounts = TRUE) %>%
  split_cols_by("ARM") %>%
  split_cols_by("SEX", "Gender", labels_var = "gend_label") %>%
  split_rows_by("RACE", "Ethnicity",
               labels_var = "ethn_label") %>%
  summarize_row_groups("RACE", label_fstr = "%s (n)") %>%
  split_rows_by("FACTOR2", "Factor2",
               split_fun = remove_split_levels("C"),
               labels_var = "fac2_label",
               label_pos = "hidden") %>%
  summarize_row_groups("FACTOR2") %>%
  analyze("AGE", "Age Analysis",
         afun = function(x) list(mean = mean(x),
                                median = median(x)),
         format = "xx.xx") %>%
  analyze("AGE",
         "Age Analysis redux",
         afun = range,
         format = "xx.x - xx.x",
         table_names = "AgeRedux"
        ) %>%
  analyze("VAR3", "Var3 Counts", afun = list_wrap_x(table),
         nested = FALSE,
         show_labels = "visible")
```

Pathing



Pathing

- Semantically descriptive way of describing position in table based on facet structure
 - Individual cells
 - Rows
 - Columns and column groups
 - Subtables
- Enforces valid structure of result when used for subsetting
 - Unlike position-based subsetting

Paths Of Our Table

```
> row_paths_summary(tb13)
```

rowname	node_class	path
—		
Caucasian (n)	ContentRow	root, RACE, WHITE, @content, Caucasian (n)
Level A	ContentRow	root, RACE, WHITE, FACTOR2, A, @content, Level A
Age Analysis	LabelRow	root, RACE, WHITE, FACTOR2, A, AGE
mean	DataRow	root, RACE, WHITE, FACTOR2, A, AGE, mean
median	DataRow	root, RACE, WHITE, FACTOR2, A, AGE, median
Age Analysis redux	LabelRow	root, RACE, WHITE, FACTOR2, A, AgeRedux
range	DataRow	root, RACE, WHITE, FACTOR2, A, AgeRedux, range
Level B	ContentRow	root, RACE, WHITE, FACTOR2, B, @content, Level B
Age Analysis	LabelRow	root, RACE, WHITE, FACTOR2, B, AGE
mean	DataRow	root, RACE, WHITE, FACTOR2, B, AGE, mean
median	DataRow	root, RACE, WHITE, FACTOR2, B, AGE, median
Age Analysis redux	LabelRow	root, RACE, WHITE, FACTOR2, B, AgeRedux
range	DataRow	root, RACE, WHITE, FACTOR2, B, AgeRedux, range
African American (n)	ContentRow	root, RACE, BLACK, @content, African American
(n)		
Level A	ContentRow	root, RACE, BLACK, FACTOR2, A, @content, Level A
Age Analysis	LabelRow	root, RACE, BLACK, FACTOR2, A, AGE
mean	DataRow	root, RACE, BLACK, FACTOR2, A, AGE, mean
median	DataRow	root, RACE, BLACK, FACTOR2, A, AGE, median
Age Analysis redux	LabelRow	root, RACE, BLACK, FACTOR2, A, AgeRedux
range	DataRow	root, RACE, BLACK, FACTOR2, A, AgeRedux, range
Level B	ContentRow	root, RACE, BLACK, FACTOR2, B, @content, Level B
Age Analysis	LabelRow	root, RACE, BLACK, FACTOR2, B, AGE
mean	DataRow	root, RACE, BLACK, FACTOR2, B, AGE, mean
median	DataRow	root, RACE, BLACK, FACTOR2, B, AGE, median
Age Analysis redux	LabelRow	root, RACE, BLACK, FACTOR2, B, AgeRedux
range	DataRow	root, RACE, BLACK, FACTOR2, B, AgeRedux, range
Var3 Counts	LabelRow	root, VAR3
level1	DataRow	root, VAR3, level1

```
> col_paths_summary(tb13)
```

label	path
ARM1	ARM, ARM1
Male	ARM, ARM1, SEX, M
Female	ARM, ARM1, SEX, F
ARM2	ARM, ARM2
Male	ARM, ARM2, SEX, M
Female	ARM, ARM2, SEX, F

Path Introspection

- `row_paths_summary` - primarily for interactive use, returns `data.frame`
- `row_paths` - returns list of paths, useful programmatically
- `make_row_df` - returns larger amount of information in `data.frame`
 - paths in the `path` column

Example of Pathing

```
> table_structure(tbl3)
[TableTree] root
  [TableTree] RACE
    [TableTree] WHITE [cont: 1 x 4]
      [TableTree] FACTOR2
        [TableTree] A [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
        [TableTree] B [cont: 1 x 4]
          [ElementaryTable] AGE (2 x 4)
          [ElementaryTable] AgeRedux (1 x 4)
      [TableTree] BLACK [cont: 1 x 4]
    [TableTree] FACTOR2
      [TableTree] A [cont: 1 x 4]
        [ElementaryTable] AGE (2 x 4)
        [ElementaryTable] AgeRedux (1 x 4)
      [TableTree] B [cont: 1 x 4]
        [ElementaryTable] AGE (2 x 4)
        [ElementaryTable] AgeRedux (1 x 4)
  [ElementaryTable] VAR3 (2 x 4)
```

```
> cell_values(tbl3, c("root", "RACE", "WHITE", "FACTOR2", "B", "AgeRedux"))
$ARM1.M
[1] 41.21257 69.45142
```

```
$ARM1.F
[1] 40.54095 68.99051
```

```
$ARM2.M
[1] 40.03586 68.45793
```

```
$ARM2.F
[1] 40.04115 69.21873
```

```
> tt_at_path(tbl3, c("root", "RACE", "WHITE", "FACTOR2", "B", "AgeRedux"))
```

	ARM1		ARM2	
	Male (N=256)	Female (N=248)	Male (N=248)	Female (N=248)
Age Analysis redux range	41.2 - 69.5	40.5 - 69.0	40.0 - 68.5	40.0 - 69.2

Pathing To Subtables vs Pathing To Content Tables

Selecting subtable

```
> tt_at_path(tbl, c("root", "RACE", "WHITE"))
```

	ARM1		ARM2	
	Male (N=256)	Female (N=248)	Male (N=248)	Female (N=248)
Caucasian (n)	116 (45.3%)	144 (58.1%)	119 (48.0%)	119 (48.0%)
Level A	37 (14.5%)	48 (19.4%)	42 (16.9%)	35 (14.1%)
Age Analysis				
mean	56.42	55.57	56.19	54.53
median	55.91	55.42	58.40	51.73
Age Analysis redux				
range	40.1 - 69.9	40.8 - 69.7	42.2 - 69.7	41.7 - 69.3
Level B	44 (17.2%)	52 (21.0%)	37 (14.9%)	40 (16.1%)
Age Analysis				
mean	54.28	55.24	54.22	54.92
median	54.71	55.47	54.96	55.17
Age Analysis redux				
range	41.2 - 69.5	40.5 - 69.0	40.0 - 68.5	40.0 - 69.2

Selecting content (group summary)

```
> tt_at_path(tbl, c("root", "RACE", "WHITE", "@content"))
```

	ARM1		ARM2	
	Male (N=256)	Female (N=248)	Male (N=248)	Female (N=248)
Caucasian (n)	116 (45.3%)	144 (58.1%)	119 (48.0%)	119 (48.0%)

Layout <-> Structure <-> Pathing

- Layout maps directly (and predictably) to table structure
- Pathing directly describes table structure, therefore
- **Layout maps directly (and predictably) to paths, and vice versa**

Layout <-> Structure <-> Pathing

- Layout maps directly (and predictably) to table structure
- Pathing directly describes table structure, therefore
- **Layout maps directly (and predictably) to paths, and vice versa**



(Semi-) Advanced Topics

Tables With Subsections

Tables Can Have Multiple Top-level Sections

- With no row splitting, this is just multiple `analyze` calls
 - Or one `analyze` call with multiple vars
- With row splitting, this is done by adding new ***non-nested*** layouting instructions
 - `analyze(..., nested = FALSE)`
 - `split_rows_by(..., nested = FALSE)` **or**
 - `split_rows_by(...)` directly after `analyze()`

```

lyt_subtabs <- basic_table() %>%
  split_cols_by("ARM") %>%
  split_rows_by("SEX", split_fun = drop_split_levels) %>%
  analyze("AGE") %>%
  split_rows_by("RACE",
    split_fun = keep_split_levels(c("ASIAN", "WHITE"))) %>%
  split_rows_by("SEX", split_fun = drop_split_levels) %>%
  analyze("AGE")

```

```
> build_table(lyt_subtabs, DM)
```

	A: Drug X	B: Placebo	C: Combination
F			
Mean	33.71	33.84	34.89
M			
Mean	36.55	32.10	34.28
ASIAN			
F			
Mean	33.55	34.00	34.90
M			
Mean	35.03	31.10	34.39
WHITE			
F			
Mean	35.88	38.57	36.50
M			
Mean	44.00	35.29	34.00

```

lyt_subtabs <- basic_table() %>%
  split_cols_by("ARM") %>%
  split_rows_by("SEX", split_fun = drop_split_levels) %>%
  analyze("AGE") %>%
  split_rows_by("RACE",
    split_fun = keep_split_levels(c("ASIAN", "WHITE"))) %>%
  split_rows_by("SEX", split_fun = drop_split_levels) %>%
  analyze("AGE")

```

```

> build_table(lyt_subtabs, DM)
      A: Drug X   B: Placebo   C: Combination

```

F	Mean	33.71	33.84	34.89
M	Mean	36.55	32.10	34.28
ASIAN				
F	Mean	33.55	34.00	34.90
M	Mean	35.03	31.10	34.39
WHITE				
F	Mean	35.88	38.57	36.50
M	Mean	44.00	35.29	34.00

```

> table_structure(tbl)
[TableTree] root
  [TableTree] SEX
    [TableTree] F
      [ElementaryTable] AGE (1 x 3)
    [TableTree] M
      [ElementaryTable] AGE (1 x 3)
  [TableTree] RACE
    [TableTree] ASIAN
      [TableTree] SEX
        [TableTree] F
          [ElementaryTable] AGE (1 x 3)
        [TableTree] M
          [ElementaryTable] AGE (1 x 3)
    [TableTree] WHITE
      [TableTree] SEX
        [TableTree] F
          [ElementaryTable] AGE (1 x 3)
        [TableTree] M
          [ElementaryTable] AGE (1 x 3)

```

Generally Don't Need rbind

- Modeling the different subtables in the layout is cleaner
 - Nice paths
 - Same details during pagination
- Rare exceptions:
 - Subtables derived from different base data
 - Different column structure
 - *In both cases, this results in a table whose structure doesn't match its presentation*

Overall, using rbind to construct tables should be avoided unless you *actually* need it, which you almost always don't

Controlling Splitting Behavior Via Provided Functions

Split Functions - Generalizing Faceting

- By default `split_*_by` generate faceting which partitions data based on a categorical variable
 - Same as faceting in `ggplot2`, `lattice`
- We can control which facet panes are generated via *split functions*
 - Split functions: `drop_split_levels`
 - Split function Factories: `remove_split_levels(excl=)`,
`trim_levels_in_group(innervar =)`, `add_combo_levels(combosdf =)`,
`add_overall_col(valname =)`

Manipulating Factor Levels

We leave examples of these to the reader

- `drop_split_levels`
- `remove_split_levels(excl = <>)`
- `drop_and_remove_levels(excl = <>)`
- `keep_split_levels(only = <>, reorder = <>)`
- `reorder_split_levels(neworder = <>, newlabels = <>, drlevels = <>)`

Preventing extraneous 0-rows in nested splits

```
trim_levels_in_group(innervar = <>)
```

Drop unobserved levels of variable `innervar` ***independently within each facet generated by this split***

- Used to control levels of `innervar` going into a nested split or analyze
- In practice, this controls (`splitvar`, `innervar`) value pairs

trim_levels_in_group(innervar)

```
> lyt <- basic_table() %>%  
+   split_rows_by("outer_fac") %>%  
+   split_rows_by("inner_fac") %>%  
+   analyze("value", mean, format = "xx.x")  
> build_table(lyt, dat)
```

all obs

A

A1
mean 1.0

A2
mean 1.8

B1
mean NA

B2
mean NA

global
mean 5.4

B

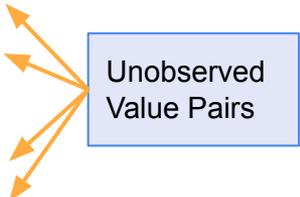
A1
mean NA

A2
mean NA

B1
mean 2.9

B2
mean 4.0

global
mean 4.8



Unobserved
Value Pairs

trim_levels_in_group(innervar)

```
> lyt <- basic_table() %>%  
+   split_rows_by("outer_fac") %>%  
+   split_rows_by("inner_fac") %>%  
+   analyze("value", mean, format = "xx.x")  
> build_table(lyt, dat)  
      all obs
```

```
A  
  A1  
    mean    1.0  
  A2  
    mean    1.8  
  B1  
    mean    NA  
  B2  
    mean    NA  
global  
  mean    5.4  
B  
  A1  
    mean    NA  
  A2  
    mean    NA  
  B1  
    mean    2.9  
  B2  
    mean    4.0  
global  
  mean    4.8
```



Unobserved
Value Pairs

```
> lyt2 <- basic_table() %>%  
+   split_rows_by("outer_fac", split_fun = trim_levels_in_group("inner_fac")) %>%  
+   split_rows_by("inner_fac") %>%  
+   analyze("value", mean, format = "xx.x")  
> build_table(lyt2, dat)  
      all obs
```

```
A  
  A1  
    mean    1.0  
  A2  
    mean    1.8  
global  
  mean    5.4  
B  
  B1  
    mean    2.9  
  B2  
    mean    4.0  
global  
  mean    4.8
```

Full Control Of Nested Facet Value Combinations

```
trim_levels_to_map(map)
```

Fully restrict facet-value-combination space to a pre-specified set of combinations across any number of variables ***regardless of whether a combination is observed in the data***

- Used when some combinations are nonsensical but other combinations are rare but should be reported
 - E.g., adverse events tables with multiple levels of summary

trim_levels_to_map

```
> lyt <- basic_table() %>%  
+   split_rows_by("LBCAT") %>%  
+   split_rows_by("PARAMCD") %>%  
+   analyze("ANRIND")  
> build_table(lyt, ex_adlb)
```

all obs

CHEMISTRY

ALT

HIGH	279
LOW	260
NORMAL	2261

CRP

HIGH	293
LOW	271
NORMAL	2236

IGA

HIGH	0
LOW	0
NORMAL	0

IMMUNOLOGY

ALT

HIGH	0
LOW	0
NORMAL	0

CRP

HIGH	0
LOW	0
NORMAL	0

IGA

HIGH	278
LOW	286
NORMAL	2236

trim_levels_to_map

```
> lyt <- basic_table() %>%  
+   split_rows_by("LBCAT") %>%  
+   split_rows_by("PARAMCD") %>%  
+   analyze("ANRIND")  
> build_table(lyt, ex_adlb)
```

```
all obs
```

CHEMISTRY	
ALT	
HIGH	279
LOW	260
NORMAL	2261
CRP	
HIGH	293
LOW	271
NORMAL	2236
IGA	
HIGH	0
LOW	0
NORMAL	0
IMMUNOLOGY	
ALT	
HIGH	0
LOW	0
NORMAL	0
CRP	
HIGH	0
LOW	0
NORMAL	0
IGA	
HIGH	278
LOW	286
NORMAL	2236

```
> map
```

```
LBCAT PARAMCD ANRIND  
1 CHEMISTRY ALT LOW  
2 CHEMISTRY CRP LOW  
3 CHEMISTRY CRP HIGH  
4 IMMUNOLOGY IGA HIGH
```

```
> lyt2 <- basic_table() %>%  
+   split_rows_by("LBCAT") %>%  
+   split_rows_by("PARAMCD", split_fun = trim_levels_to_map(map = map)) %>%  
+   analyze("ANRIND")  
> build_table(lyt2, ex_adlb)
```

```
all obs
```

CHEMISTRY	
ALT	
LOW	260
CRP	
LOW	271
HIGH	293
IMMUNOLOGY	
IGA	
HIGH	278

trim_levels_to_map

```
> lyt <- basic_table() %>%  
+   split_rows_by("LBCAT") %>%  
+   split_rows_by("PARAMCD") %>%  
+   analyze("ANRIND")  
> build_table(lyt, ex_adlb)  
all obs
```

```
CHEMISTRY  
ALT  
HIGH      279  
LOW       260  
NORMAL    2261  
CRP  
HIGH      293  
LOW       271  
NORMAL    2236  
IGA  
HIGH      0  
LOW       0  
NORMAL    0  
IMMUNOLOGY  
ALT  
HIGH      0  
LOW       0  
NORMAL    0  
CRP  
HIGH      0  
LOW       0  
NORMAL    0  
IGA  
HIGH      278  
LOW       286  
NORMAL    2236
```

```
> map  
LBCAT PARAMCD ANRIND  
1 CHEMISTRY ALT LOW  
2 CHEMISTRY CRP LOW  
3 CHEMISTRY CRP HIGH  
4 IMMUNOLOGY IGA HIGH
```

Allowed Value
Combinations

```
> lyt2 <- basic_table() %>%  
+   split_rows_by("LBCAT") %>%  
+   split_rows_by("PARAMCD", split_fun = trim_levels_to_map(map = map)) %>%  
+   analyze("ANRIND")  
> build_table(lyt2, ex_adlb)  
all obs
```

```
CHEMISTRY  
ALT  
LOW      260  
CRP  
LOW      271  
HIGH     293  
IMMUNOLOGY  
IGA  
HIGH     278
```

Adding Combination Levels

```
add_combo_levels (combo_df) ,  
add_overall_level (valname) (special case)
```

Add combination levels (overall level is a special case w/ convenience function) to existing levels of a split, optionally also excluding some of the original levels

add_combo_levels (combodf)

Combinations are declared in a data.frame/tribble with the following columns:

- valname - name of the combined value (and thus the generated facet)
- label - label for the generated facet
- levelcombo - character vector of values to combine (typically factor levels of underlying var)
- exargs - a list of extra arguments corresponding to the split level (usually empty list)

```
> combodf <- tribble(
+   ~valname, ~label,      ~levelcombo,      ~exargs,
+   "A_B",    "Arms A+B", c("A: Drug X", "B: Placebo"), list(),
+   "A_C",    "Arms A+C", c("A: Drug X", "C: Combination"), list())
>
> lyt <- basic_table(show_colcounts = TRUE) %>%
+   split_cols_by("ARM", split_fun = add_combo_levels(combodf)) %>%
+   analyze("AGE")
>
> build_table(lyt, DM)
```

	A: Drug X (N=121)	B: Placebo (N=106)	C: Combination (N=129)	Arms A+B (N=227)	Arms A+C (N=250)
Mean	34.91	33.02	34.57	34.03	34.73

add_combo_levels (combo_df)

Combinations are declared in a data.frame/tribble with the following columns:

- valname - name of the combined value (and thus the generated facet)
- label - label for the generated facet
- levelcombo - character vector of values to combine (typically factor levels of underlying var)
- exargs - a list of extra arguments corresponding to the split level (usually empty list)

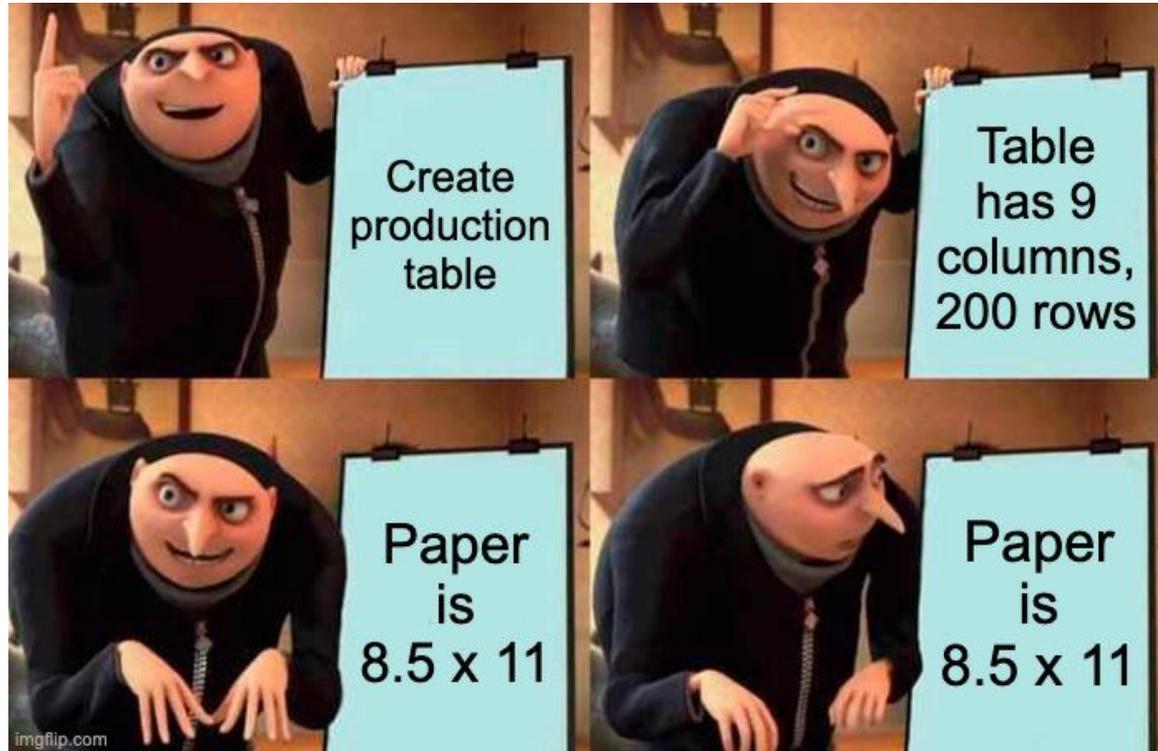
```
> combo_df <- tribble(
+   ~valname, ~label,      ~levelcombo,      ~exargs,
+   "A_B",    "Arms A+B",  c("A: Drug X", "B: Placebo"),  list(),
+   "A_C",    "Arms A+C",  c("A: Drug X", "C: Combination"), list()
+ )
>
> lyt <- basic_table(show_colcounts = TRUE) %>%
+   split_cols_by("ARM", split_fun = add_combo_levels(combo_df)) %>%
+   analyze("AGE")
>
> build_table(lyt, DM)
```

	A: Drug X (N=121)	B: Placebo (N=106)	C: Combination (N=129)	Arms A+B (N=227)	Arms A+C (N=250)
Mean	34.91	33.02	34.57	34.03	34.73

One new level
per row.

N's handled
automatically

Pagination



Pagination Problem

Page 1	Page 2	
Page 3	Page 4	
Page 5	Page 6	

Note some rows, columns and row names need to be repeated for context.

Context-Preserving Pagination

title		page nr. *
subtitles		
page titles		
top left	column structure	
row structure	cells	
referential footnotes		
page footer		
global footer		
provenance footer		page nr. *

Repeated on:

	every page
	some pages
	no pages
	depends

* forthcoming feature

Pagination Specifications

- Vertical Pagination repeats context information after page breaks
 - Content and label rows
- Pagination happens *after* wordwrapping
- Horiz pagination is identical across all sections of vert pagination
- Horizontal and Vertical pagination can be done separately, but
 - Both performed when page dimensions are defined
- Pagination assumes monospaced fonts
- ***Use verbose = TRUE for debugging when pagination fails***

Specifying page-size and font

- Page Dimensions

- `page_type`
 - `letter, a4, legal`
 - **Can be used with** `landscape = TRUE`
- `pg_width, pg_height` (in inches)
- `lpp, cpp` (raw height in lines, width in chars)

- Font

- `font_size`
- `font_family` (must be monospaced)

Converted to lines and characters:

```
page_lcpp(page_type = "legal",  
          font_size = 10,  
          font_family = "Courier")
```

```
$cpp  
[1] 84
```

```
$lpp  
[1] 93
```

Invoking Pagination

- **Directly:** `paginate_table`
 - Returns list of tables representing the pages (width first ordering)
- **Indirectly** (`export_as_txt(page_type = "a4")`)
 - Exports file containing paginated version of the table
 - txt, pdf, rtf exporters support pagination

Page-by Row Splits

- Force pagination between levels of a row-split
 - `split_rows_by("varname", page_by = TRUE)`
 - `page_prefix` - appended with split value label to create page titles
- Page-by row splits cannot be nested within non-page_by row splits
- Page-by pagination happens *before* size-based vertical pagination
 - Horizontal pagination is unaffected
 - Vertical pagination is performed separately within each 'page' defined by full set of page-by splits

Page-by Row Splitting

```
lyt <- basic_table(title = "Main Title",
                   subtitles = "Subtitle") |>
  split_cols_by("ARM") |>
  split_rows_by("SEX",
               page_by = TRUE,
               page_prefix = "Sex",
               split_fun = drop_split_levels) |>
  analyze("AGE", mean)

tbl <- build_table(lyt, DM)
```

```
> paginate_table(tbl)
```

```
$F
```

```
Main Title
```

```
Subtitle
```

```
Sex: F
```

	A: Drug X	B: Placebo	C: Combination
mean	33.7	33.8	34.9

```
$M
```

```
Main Title
```

```
Subtitle
```

```
Sex: M
```

	A: Drug X	B: Placebo	C: Combination
mean	36.5	32.1	34.3

Note about Page-by row splits

Currently forced pagination happens *at pagination time*, meaning it won't show up when you print the table without pagination:

```
> tbl
```

```
Main Title
```

```
Subtitle
```

	A: Drug X	B: Placebo	C: Combination
F			
mean	33.7	33.8	34.9
M			
mean	36.5	32.1	34.3

This may change in future releases

Customizing Appearance And Rendering Behavior



Section Dividers

Horizontal dividers (incl. blank line) placed after row groups

```
lyt <- basic_table(title = "Main Title",  
                   subtitles = "Subtitle") %>%  
  split_cols_by("ARM") %>%  
  split_rows_by("STRATA1", section_div = "-") %>%  
  split_rows_by("SEX", section_div = " ",  
               split_fun = drop_split_levels) %>%  
  analyze("AGE", mean, format = "xx.x")  
  
tbl <- build_table(lyt, DM)
```

```
> tbl  
Main Title  
Subtitle
```

		A: Drug X	B: Placebo	C: Combination
A				
F	mean	30.9	32.9	36.0
M	mean	35.1	31.1	35.6

B				
F	mean	34.9	32.9	34.4
M	mean	36.6	32.1	34.4

C				
F	mean	35.2	36.0	34.3
M	mean	37.4	32.8	32.8

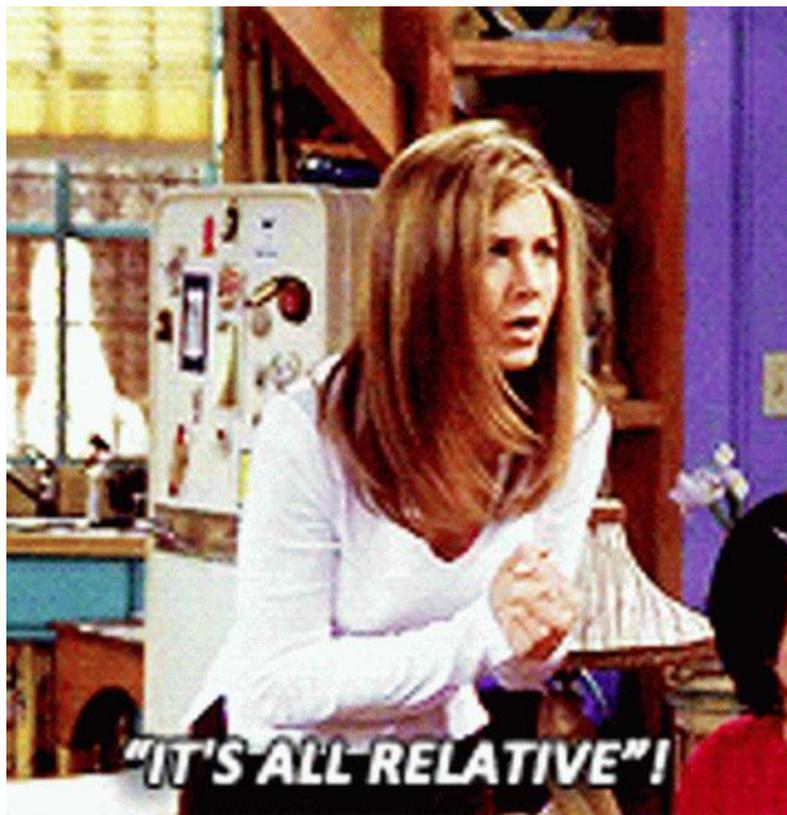
Section dividers

- Can be different for different levels of faceting
- Can be blank lines (use " ")
- When multiple section dividers would apply, only the one for the largest group is printed
- No section divider is ever printed at the end of the table body

Indent Modification

- Indenting happens automatically
 - Including visible label and hidden label cases
- Indent levels can be modified at the layout stage
 - `indent_mod` argument to, well, most things

Understanding indent_mod



Normal Table

```
lyt <- basic_table() %>%  
  split_rows_by("STRATA1") %>%  
  split_rows_by("SEX",  
    split_fun = drop_split_levels) %>%  
  analyze("AGE", mean, format = "xx.x")  
  
tbl1 <- build_table(lyt, DM)
```

```
> tbl1  
      all obs  
-----  
A  
  F  
  mean  33.2  
  M  
  mean  34.5  
B  
  F  
  mean  34.2  
  M  
  mean  34.0  
C  
  F  
  mean  35.1  
  M  
  mean  34.5
```

Indent Mods

```
lyt1 <- basic_table() %>%  
  split_rows_by("STRATA1") %>%  
  split_rows_by("SEX",  
               split_fun = drop_split_levels) %>%  
  analyze("AGE", mean, format = "xx.x")  
  
tbl1 <- build_table(lyt1, DM)
```

Individual
Mod



```
> tbl1  
-----  
all obs  
A  
  F  
  mean 33.2  
  M  
  mean 34.5  
B  
  F  
  mean 34.2  
  M  
  mean 34.0  
C  
  F  
  mean 35.1  
  M  
  mean 34.5
```

Indent Mods

```
lyt2 <- basic_table() %>%
  split_rows_by("STRATA1", indent_mod = 1) %>%
  split_rows_by("SEX",
    split_fun = drop_split_levels) %>%
  analyze("AGE", mean, format = "xx.x")

tbl2 <- build_table(lyt2, DM)
```

Total effective
change

```
+1 = +1
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = +1
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = +1
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
+1 = 0 + (+1)
```

Individual
Mod

```
■ →
■
■
■
■
■ →
■
■
■
■
■ →
■
■
■
■
```

```
> tbl2
      all obs
-----
A
  F
  mean 33.2
  M
  mean 34.5
B
  F
  mean 34.2
  M
  mean 34.0
C
  F
  mean 35.1
  M
  mean 34.5
```

Indent Mods

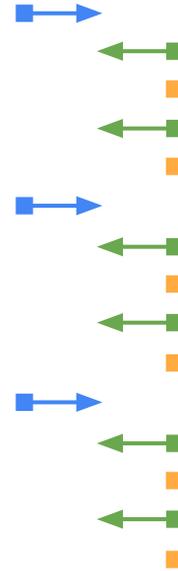
```
lyt3 <- basic_table() %>%
  split_rows_by("STRATA1", indent_mod = 1) %>%
  split_rows_by("SEX", indent_mod = -1,
    split_fun = drop_split_levels) %>%
  analyze("AGE", mean, format = "xx.x")

tbl3 <- build_table(lyt3, DM)
```

Total effective
change

```
+1 = +1
0 = -1 + (+1)
0 = 0 + (0)
0 = -1 + (+1)
0 = 0 + (0)
+1 = +1
0 = -1 + (+1)
0 = 0 + (0)
0 = -1 + (+1)
0 = 0 + (0)
+1 = +1
0 = -1 + (+1)
0 = 0 + (0)
0 = -1 + (+1)
0 = 0 + (0)
```

Individual
Mod



> tbl3

		all obs
A		
F		
mean		33.2
M		
mean		34.5
B		
F		
mean		34.2
M		
mean		34.0
C		
F		
mean		35.1
M		
mean		34.5

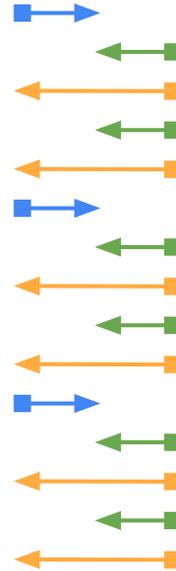
Indent Mods

```
lyt4 <- basic_table() %>%
  split_rows_by("STRATA1", indent_mod = 1) %>%
  split_rows_by("SEX", indent_mod = -1,
    split_fun = drop_split_levels) %>%
  analyze("AGE", mean, format = "xx.x",
    indent_mod = -2)
tbl4 <- build_table(lyt4, DM)
```

Total effective
change

```
+1 = +1
0 = -1 + (+1)
-2 = -2 + (0)
0 = -1 + (+1)
-2 = -2 + (0)
+1 = +1
0 = -1 + (+1)
-2 = -2 + (0)
0 = -1 + (+1)
-2 = -2 + (0)
+1 = +1
0 = -1 + (+1)
-2 = -2 + (0)
0 = -1 + (+1)
-2 = -2 + (0)
```

Individual
Mod



> tbl4
all obs

all obs	
A	
F	
mean	33.2
M	
mean	34.5
B	
F	
mean	34.2
M	
mean	34.0
C	
F	
mean	35.1
M	
mean	34.5

Indent Mods

- You can also apply indent mods to individual rows
 - `.indent_mods` argument in `in_rows()`
 - `indent_mod` argument in `rcell()`
 - First non-zero `indent_mod` promoted to apply to row, indent mods on other cells in the row are ignored
- I leave that as an exercise

Table Inset



Table Inset

```
lyt6 <- basic_table(title = "Main Title",
                    subtitles = "Subtitle",
                    main_footer = "Fooooooter",
                    prov_footer = "le provenance",
                    inset = 10) %>%
  split_cols_by("ARM") %>%
  split_rows_by("SEX",
               split_fun = drop_split_levels) %>%
  analyze("AGE", mean, format = "xx.x")

tbl6 <- build_table(lyt6, DM)
```

Table Inset

```
lyt6 <- basic_table(title = "Main Title",
  subtitles = "Subtitle",
  main_footer = "Fooooooter",
  prov_footer = "le provenance",
  inset = 10) %>%
  split_cols_by("ARM") %>%
  split_rows_by("SEX",
    split_fun = drop_split_levels) %>%
  analyze("AGE", mean, format = "xx.x")

tbl6 <- build_table(lyt6, DM)
```

```
> tbl6
Main Title
Subtitle
```



		A: Drug X	B: Placebo	C: Combination
F	mean	33.7	33.8	34.9
M	mean	36.5	32.1	34.3

Fooooooter

le provenance

Table Inset

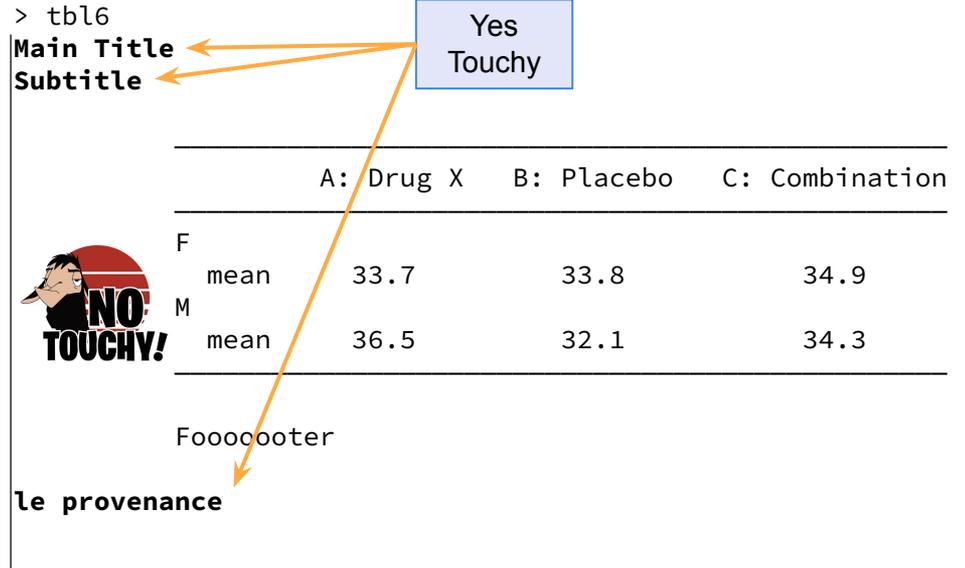
```
lyt6 <- basic_table(title = "Main Title",  
  subtitles = "Subtitle",  
  main_footer = "Foocooooter",  
  prov_footer = "le provenance",  
  inset = 10) %>%  
  split_cols_by("ARM") %>%  
  split_rows_by("SEX",  
    split_fun = drop_split_levels) %>%  
  analyze("AGE", mean, format = "xx.x")  
  
tbl6 <- build_table(lyt6, DM)
```

```
> tbl6
```

		A: Drug X	B: Placebo	C: Combination
F	mean	33.7	33.8	34.9
M	mean	36.5	32.1	34.3

Foocooooter

le provenance



The diagram illustrates the 'inset' parameter in the R code. A blue box labeled 'Yes Touchy' has three orange arrows pointing to the 'Main Title', 'Subtitle', and 'Foocooooter' elements of the table output. A cartoon character with a 'NO TOUCHY!' sign is also present.

Word Wrapping and Width



Title/footer And Column Widths

- `max_width` - controls rendered width of title and footer information
 - Including referential footnotes
 - Currently need `tf_wrap = TRUE`
- `colwidths` (most funcs) - controls rendered width of columns
 - `widths` (`toString`) - same argument, difference in name an unfortunate artifact
 - Includes row labels + `opleft` as first “column”
 - Controls width of column labels
- Pagination machinery (Direct and via import) takes into account word-wrapping
 - **Only** if you specify the above to the paginator, table does not carry this info around

A Wide Table (From Our Tests)

```
> tt_for_wrap
Enough long title to be probably wider than expected
```

	Incredibly long column name to be wrapped	This_should_be_somewhere_split	C: Combination
ASIAN			
AGE			
Mean	32.50	36.68	36.99
EOSDY			
Mean	A very long content to_be_wrapped_and_splitted	A very long content to_be_wrapped_and_splitted	A very long content to_be_wrapped_and_splitted
BLACK OR AFRICAN AMERICAN			
AGE			
Mean	34.27	34.93	33.71
EOSDY			
Mean	A very long content to_be_wrapped_and_splitted	A very long content to_be_wrapped_and_splitted	A very long content to_be_wrapped_and_splitted

Also this seems quite wider than expected initially.

I Make It Fit And Then I Sits

```
> cat(export_as_txt(tt_for_wrap, tf_wrap = TRUE, max_width = 20,  
                  colwidths = c(25, 20, 15, 20)))
```

Enough long title to
be probably wider
than expected

	Incredibly long column name to be wrapped	This_should_be_ somewhere_split	C: Combination
<hr/>			
ASIAN			
AGE			
Mean	32.50	36.68	36.99
EOSDY			
Mean	A very long content to_be_wrapped_and_sp litted	A very long content to_be_w rapped_and_spli tted	A very long content to_be_wrapped_and_sp litted
<hr/>			
BLACK OR AFRICAN AMERICAN			
AGE			
Mean	34.27	34.93	33.71
EOSDY			
Mean	A very long content to_be_wrapped_and_sp litted	A very long content to_be_w rapped_and_spli tted	A very long content to_be_wrapped_and_sp litted

Also this seems
quite wider than
expected initially.



Render the table

Multiple formats are supported

- ASCII Text - `export_as_txt`, `toString` (no pagination)
- PDF - `export_as_pdf`
- RTF - `export_as_rtf` (experimental) via `r2rtf`
- HTML - `as_html`
 - No pagination
- flextable object - `tt_to_flextable`
- PPT, DOCX - indirectly, via `officer` + `flextable`

And That's rtables in ~2.5 Hours

