

# Data Transformation with dplyr :: CHEAT SHEET



**dplyr** functions work with pipes and expect **tidy data**. In tidy data:



&



**pipes**

Each **variable** is in its own **column**

Each **observation**, or **case**, is in its own **row**

$x \%>\% f(y)$  becomes  $f(x, y)$

## Summarise Cases

These apply **summary functions** to columns to create a new table of summary statistics. Summary functions take vectors as input and return one value (see back).

**summary function**



**summarise(.data, ...)**  
Compute table of summaries.  
*summarise(mtcars, avg = mean(mpg))*



**count(x, ..., wt = NULL, sort = FALSE)**  
Count number of rows in each group defined by the variables in ... Also **tally()**.  
*count(iris, Species)*

### VARIATIONS

**summarise\_all()** - Apply funs to every column.  
**summarise\_at()** - Apply funs to specific columns.  
**summarise\_if()** - Apply funs to all cols of one type.

## Group Cases

Use **group\_by()** to create a "grouped" copy of a table. dplyr functions will manipulate each "group" separately and then combine the results.



*mtcars %>%  
group\_by(cyl) %>%  
summarise(avg = mean(mpg))*

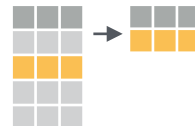
**group\_by(.data, ..., add = FALSE)**  
Returns copy of table grouped by ...  
*g\_iris <- group\_by(iris, Species)*

**ungroup(x, ...)**  
Returns ungrouped copy of table.  
*ungroup(g\_iris)*

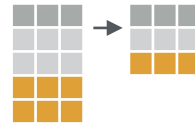
## Manipulate Cases

### EXTRACT CASES

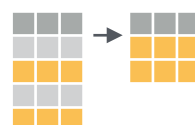
Row functions return a subset of rows as a new table.



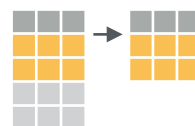
**filter(.data, ...)** Extract rows that meet logical criteria. *filter(iris, Sepal.Length > 7)*



**distinct(.data, ..., .keep\_all = FALSE)** Remove rows with duplicate values.  
*distinct(iris, Species)*



**sample\_frac(tbl, size = 1, replace = FALSE, weight = NULL, .env = parent.frame())** Randomly select fraction of rows.  
*sample\_frac(iris, 0.5, replace = TRUE)*



**sample\_n(tbl, size, replace = FALSE, weight = NULL, .env = parent.frame())** Randomly select size rows. *sample\_n(iris, 10, replace = TRUE)*

**slice(.data, ...)** Select rows by position.  
*slice(iris, 10:15)*

**top\_n(x, n, wt)** Select and order top n entries (by group if grouped data). *top\_n(iris, 5, Sepal.Width)*

### Logical and boolean operators to use with filter()

<	<=	is.na()	%in%		xor()
>	>=	!is.na()	!	&	

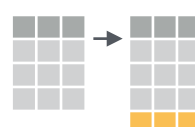
See **?base::Logic** and **?Comparison** for help.

### ARRANGE CASES



**arrange(.data, ...)** Order rows by values of a column or columns (low to high), use with **desc()** to order from high to low.  
*arrange(mtcars, mpg)*  
*arrange(mtcars, desc(mpg))*

### ADD CASES



**add\_row(.data, ..., .before = NULL, .after = NULL)**  
Add one or more rows to a table.  
*add\_row(faithful, eruptions = 1, waiting = 1)*

## Manipulate Variables

### EXTRACT VARIABLES

Column functions return a set of columns as a new vector or table.



**pull(.data, var = -1)** Extract column values as a vector. Choose by name or index.  
*pull(iris, Sepal.Length)*



**select(.data, ...)**  
Extract columns as a table. Also **select\_if()**.  
*select(iris, Sepal.Length, Species)*

Use these helpers with **select()**,  
e.g. *select(iris, starts\_with("Sepal"))*

<b>contains(match)</b>	<b>num_range(prefix, range)</b>	⋮, e.g. <i>mpg:cyl</i>
<b>ends_with(match)</b>	<b>one_of(...)</b>	⋮, e.g. <i>-Species</i>
<b>matches(match)</b>	<b>starts_with(match)</b>	

### MAKE NEW VARIABLES

These apply **vectorized functions** to columns. Vectorized funs take vectors as input and return vectors of the same length as output (see back).

**vectorized function**



**mutate(.data, ...)**  
Compute new column(s).  
*mutate(mtcars, gpm = 1/mpg)*



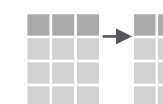
**transmute(.data, ...)**  
Compute new column(s), drop others.  
*transmute(mtcars, gpm = 1/mpg)*



**mutate\_all(.tbl, .funs, ...)** Apply funs to every column. Use with **funs()**. Also **mutate\_if()**.  
*mutate\_all(faithful, funs(log(.), log2(.)))*  
*mutate\_if(iris, is.numeric, funs(log(.)))*



**mutate\_at(.tbl, .cols, .funs, ...)** Apply funs to specific columns. Use with **funs()**, **vars()** and the helper functions for **select()**.  
*mutate\_at(iris, vars(-Species), funs(log(.)))*



**add\_column(.data, ..., .before = NULL, .after = NULL)** Add new column(s). Also **add\_count()**, **add\_tally()**. *add\_column(mtcars, new = 1:32)*



**rename(.data, ...)** Rename columns.  
*rename(iris, Length = Sepal.Length)*

# Vector Functions

## TO USE WITH MUTATE ()

**mutate()** and **transmute()** apply vectorized functions to columns to create new columns. Vectorized functions take vectors as input and return vectors of the same length as output.

**vectorized function**

## OFFSETS

**dplyr::lag()** - Offset elements by 1  
**dplyr::lead()** - Offset elements by -1

## CUMULATIVE AGGREGATES

**dplyr::cumall()** - Cumulative all()  
**dplyr::cumany()** - Cumulative any()  
**cummax()** - Cumulative max()  
**dplyr::cummean()** - Cumulative mean()  
**cummin()** - Cumulative min()  
**cumprod()** - Cumulative prod()  
**cumsum()** - Cumulative sum()

## RANKINGS

**dplyr::cume\_dist()** - Proportion of all values <=   
**dplyr::dense\_rank()** - rank w ties = min, no gaps  
**dplyr::min\_rank()** - rank with ties = min  
**dplyr::ntile()** - bins into n bins  
**dplyr::percent\_rank()** - min\_rank scaled to [0,1]  
**dplyr::row\_number()** - rank with ties = "first"

## MATH

**+**, **-**, **\***, **/**, **^**, **%/%**, **%%** - arithmetic ops  
**log()**, **log2()**, **log10()** - logs  
**<**, **<=**, **>**, **>=**, **!=**, **==** - logical comparisons  
**dplyr::between()** - x >= left & x <= right  
**dplyr::near()** - safe == for floating point numbers

## MISC

**dplyr::case\_when()** - multi-case if\_else()  
*iris %>% mutate(Species = case\_when(  
Species == "versicolor" ~ "versi",  
Species == "virginica" ~ "virgi",  
TRUE ~ Species))*  
**dplyr::coalesce()** - first non-NA values by element across a set of vectors  
**dplyr::if\_else()** - element-wise if() + else()  
**dplyr::na\_if()** - replace specific values with NA  
**pmax()** - element-wise max()  
**pmin()** - element-wise min()  
**dplyr::recode()** - Vectorized switch()  
**dplyr::recode\_factor()** - Vectorized switch() for factors

# Summary Functions

## TO USE WITH SUMMARISE ()

**summarise()** applies summary functions to columns to create a new table. Summary functions take vectors as input and return single values as output.

**summary function**

## COUNTS

**dplyr::n()** - number of values/rows  
**dplyr::n\_distinct()** - # of uniques  
**sum(!is.na())** - # of non-NA's

## LOCATION

**mean()** - mean, also **mean(!is.na())**  
**median()** - median

## LOGICALS

**mean()** - Proportion of TRUE's  
**sum()** - # of TRUE's

## POSITION/ORDER

**dplyr::first()** - first value  
**dplyr::last()** - last value  
**dplyr::nth()** - value in nth location of vector

## RANK

**quantile()** - nth quantile  
**min()** - minimum value  
**max()** - maximum value

## SPREAD

**IQR()** - Inter-Quartile Range  
**mad()** - median absolute deviation  
**sd()** - standard deviation  
**var()** - variance

# Row Names

Tidy data does not use rownames, which store a variable outside of the columns. To work with the rownames, first move them into a column.

**rownames\_to\_column()**  
Move row names into col.  
*a <- rownames\_to\_column(iris, var = "C")*

**column\_to\_rownames()**  
Move col in row names.  
*column\_to\_rownames(a, var = "C")*

Also **has\_rownames()**, **remove\_rownames()**

# Combine Tables

## COMBINE VARIABLES

**x** + **y** = **A B C A B D**  

a	t	1			
b	u	2			
c	v	3			

 + 

a	t	3			
b	u	2			
d	w	1			

 = 

a	t	1	a	t	3
b	u	2	b	u	2
c	v	3	d	w	1

Use **bind\_cols()** to paste tables beside each other as they are.

**bind\_cols(...)** Returns tables placed side by side as a single table.  
BE SURE THAT ROWS ALIGN.

Use a "Mutating Join" to join one table to columns from another, matching values with the rows that they correspond to. Each join retains a different combination of values from the tables.

**left\_join(x, y, by = NULL, copy=FALSE, suffix=c(".x", ".y"),...)**  
Join matching values from y to x.

**right\_join(x, y, by = NULL, copy = FALSE, suffix=c(".x", ".y"),...)**  
Join matching values from x to y.

**inner\_join(x, y, by = NULL, copy = FALSE, suffix=c(".x", ".y"),...)**  
Join data. Retain only rows with matches.

**full\_join(x, y, by = NULL, copy=FALSE, suffix=c(".x", ".y"),...)**  
Join data. Retain all values, all rows.

Use **by = c("col1", "col2", ...)** to specify one or more common columns to match on.  
*left\_join(x, y, by = "A")*

Use a named vector, **by = c("col1" = "col2")**, to match on columns that have different names in each table.  
*left\_join(x, y, by = c("C" = "D"))*

Use **suffix** to specify the suffix to give to unmatched columns that have the same name in both tables.  
*left\_join(x, y, by = c("C" = "D"), suffix = c("1", "2"))*

## COMBINE CASES

**x** + **y** = **A B C**  

a	t	1
b	u	2
c	v	3

 + 

a	t	3
c	v	3
d	w	4

Use **bind\_rows()** to paste tables below each other as they are.

**bind\_rows(..., .id = NULL)**  
Returns tables one on top of the other as a single table. Set .id to a column name to add a column of the original table names (as pictured)

**intersect(x, y, ...)**  
Rows that appear in both x and y.

**setdiff(x, y, ...)**  
Rows that appear in x but not y.

**union(x, y, ...)**  
Rows that appear in x or y. (Duplicates removed). **union\_all()** retains duplicates.

Use **setequal()** to test whether two data sets contain the exact same rows (in any order).

## EXTRACT ROWS

**x** + **y** = **A B C**  

a	t	1
b	u	2
c	v	3

 + 

a	t	3
b	u	2
d	w	1

Use a "Filtering Join" to filter one table against the rows of another.

**semi\_join(x, y, by = NULL, ...)**  
Return rows of x that have a match in y. USEFUL TO SEE WHAT WILL BE JOINED.

**anti\_join(x, y, by = NULL, ...)**  
Return rows of x that do not have a match in y. USEFUL TO SEE WHAT WILL NOT BE JOINED.

