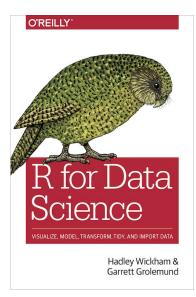
Introduction to data science with R and the tidyverse



Daniel Lundin

http://r4ds.had.co.nz/



Microbial model Systems



The "Excel view" of data, a.k.a. "wide" format

species	order	site 1	site 2	site 3	site 4	site 5
Berberis hispanica	Ranunculales	26	12	37	0	5
Allium ebusitanum	Asparagales	521	0	2	51	485
Anacamptis morio	Asparagales	54	10	0	48	16
Carduus nutans	Asterales	55	8	41	23	8
Colchicum cupanii	Liliales	14	0	0	34	0
Colchicum ritchii	Liliales	0	12	0	3	17





The "data view" of data, a.k.a. "long" format

species	order	site	count
Berberis hispanica	Ranunculales	site 1	26
Allium ebusitanum	Asparagales	site 1	521
Anacamptis morio	Asparagales	site 1	54
Carduus nutans	Asterales	site 1	55
Colchicum cupanii	Liliales	site 1	14
Colchicum ritchii	Liliales	site 1	0
Berberis hispanica	Ranunculales	site 2	12
Allium ebusitanum	Asparagales	site 2	0
Anacamptis morio	Asparagales	site 2	10
Carduus nutans	Asterales	site 2	8
Colchicum cupanii	Liliales	site 2	0
Colchicum			

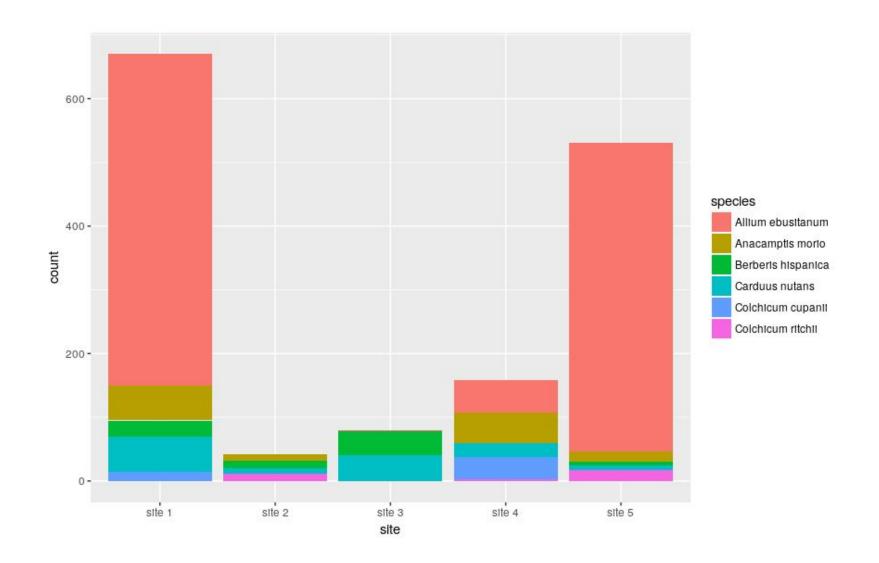
Linnaeus University

Ecology and Evertifies Microbial model Speanisa

Ranunculales site 3

12

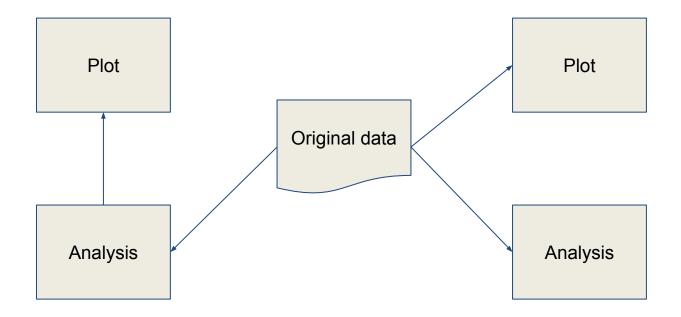








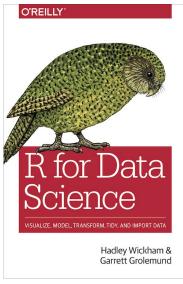
You need to be good at format conversion...











http://r4ds.had.co.nz/

- > install.packages('tidyverse')
- > library(tidyverse)

https://www.tidyverse.org/

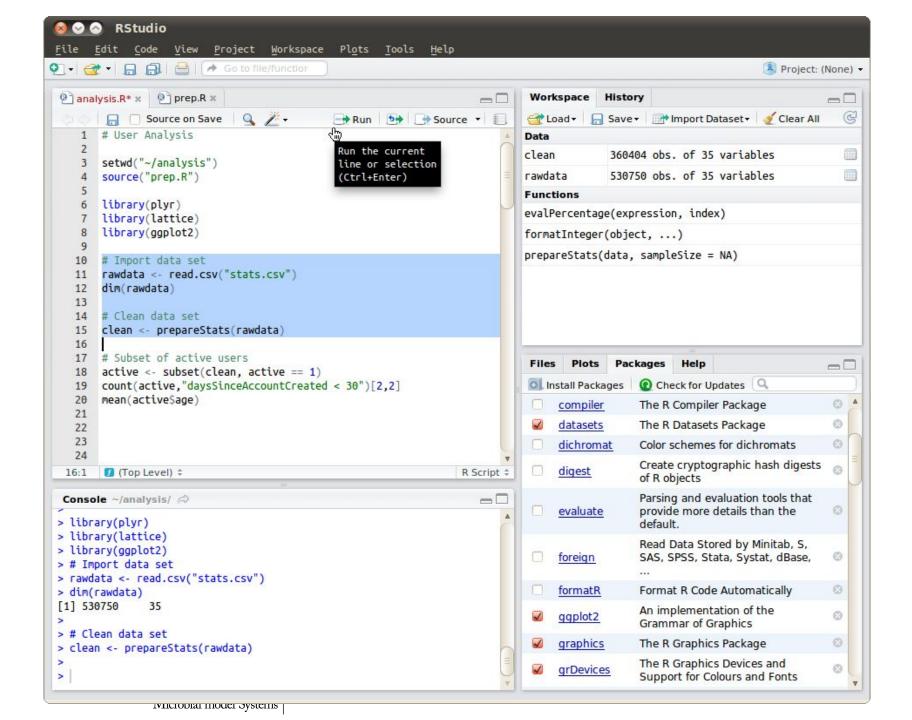
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RStudio



https://www.rstudio.com/

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What I expect from you

- 1. No knowledge of R
- 2. Knowledge of "base" R
- Knowledge of tidyverse R
- 4. You prefer Python, Perl, Ruby, ...

"Daniel, I'm not following!"



What to show (notes to myself)

Centre

```
readr: read_tsv()/write_tsv()
tidyr: gather()/spread()
tidyr: separate()/unite()
dplyr: filter()/select()
dplyr: group_by() and summarise()
dplyr: union()/inner_join() et al.
ggplot2: ggplot(): geom_col(), geom_point(), facet_wrap()
```