Analysis of CSV File Reading in R

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Outline

- Background
- ▶ Experiment
- ► ANOVA
- ► Additional Analysis
- Discussion

Background

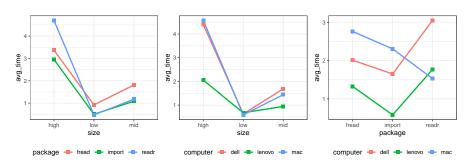
- ▶ Statisticians struggle with reading in large dataset (Matthews, 2021)
- ▶ read.csv, part of base R (R Core Team, 2021), is old and slow.
- ► There are better and more efficient data I/O algorithms in R (Gillespie & Lovelace, 2021)
 - ▶ rio::import() (Chan et al., 2021)
 - ▶ readr::read_csv() (Wickham & Hester, 2021) part of tidyverse (Wickham et al., 2019)
 - ▶ data.table::fread() (Dowle & Srinivasan, 2021)

Experiment

- ▶ 3³ factorial design (Montgomery, 2012)
 - \blacktriangleright File size: low (~0.5GB), medium (~1.2GB), high (~2GB)
 - ► Computer: Lenovo, Dell, MacBook
 - ► Function: rio::import(), readr::read_csv(),
 data.table::fread()
- ► Model: insert formula
- ightharpoonup n = 20 replicates
- ▶ How were the data generated?

Data Understanding

► Interaction plots

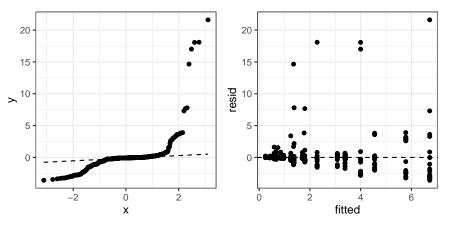


► All model terms are significant...

Term	df	SS	MS	F	p-value
computer	2	118.6155	59.3078	12.8017	0.0000
size	2	908.9852	454.4926	98.1034	0.0000
package	2	38.4974	19.2487	4.1549	0.0162
computer:size	4	136.3553	34.0888	7.3582	0.0000
computer:package	4	114.5762	28.6441	6.1829	0.0001
size:package	4	87.9878	21.9970	4.7481	0.0009
computer:size:package	8	144.3655	18.0457	3.8952	0.0002
Residuals	513	2376.6214	4.6328	NA	NA

► However . . .

► There are issues with both **normality** and **homoscedasticity**



lacktriangle Confirmed by Shapiro-Wilk and Levene tests (both with p-value pprox 0)

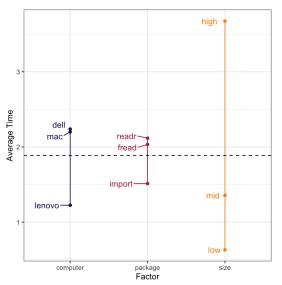
- ► Transforming the data (log, Box-Cox) did not fix issues with model assumptions
- ▶ This leads us to consider a nonparametric approach,
 - ▶ No distribution assumption about the data
 - ▶ Permutation test for a three way factorial designs
 - ▶ asbio::perm.fact.test() (Aho, 2021)

► A permutation test gives significant results

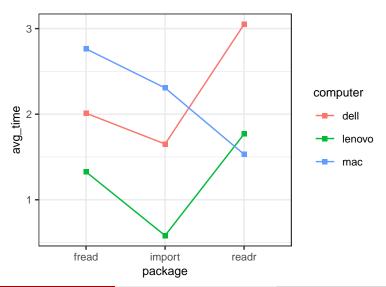
Term	Initial F	df	p-value
size	98.1034	2	0.0001
computer	12.8017	2	0.0001
package	4.1549	2	0.0123
size:computer	7.3582	4	0.0001
size:package	4.7481	4	0.0007
computer:package	6.1829	4	0.0002
size:computer:package	3.8952	8	0.0003
Residual	NA	513	NA

- ▶ Post-hoc analysis: pairwise permutation tests
- ▶ Use False Discovery Rate (Benjamini & Hochberg, 1995) as p-value adjustment method
- ▶ Which pairs differ in mean reading time?
 - ► Speed: all pairs
 - ► Computer: Lenovo-Dell, Lenovo-Mac
 - ▶ Package/Function: data.table::fread() rio::import(), rio::import() - readr::read_csv()

► Univariate effects plot of the factors



▶ Investigate reading method (package/function) for MacBook only



- ► A simulation study
- ▶ Simulate more data, in addition to the existing 3 data files
- Consider 10 file sizes (in GB)
 - ightharpoonup 0.49, 1.22, 1.51, 1.73, 1.96, 2.16, 2.36, 2.61, 2.94, 3.32
- ▶ Use MacBook to read in each file using the 3 functions, and record the time elapsed

► ANCOVA

► Response: time elapsed (in seconds)

► Treatment: function (3 levels)

► Covariate: file size (in GB)

Term	SS	df	F	p-value
(Intercept)	10.5154	1	8.4657	0.0073
meth	187.3580	2	75.4188	0.0000
size	192.4662	1	154.9502	0.0000
Residuals	32.2950	26	NA	NA

▶ Model assumptions are met.

statistic	p.value	method
0.9732	0.6286	Shapiro-Wilk normality test

statistic	p.value	df	df.residual	method
2.3633	0.1133	2	27	Levene test

► Multiple comparisons - Tukey contrasts

contrast	estimate	se	statistic	adj p-value
import - fread	-0.2873	0.4984	-0.5764	0.8338
readr - fread	-5.4391	0.4984	-10.9127	0.0000
readr - import	-5.1518	0.4984	-10.3362	0.0000

▶ readr significantly differs from the other 2 methods when using Mac.

Cheers.

- Acknowledgments
 - ► Lance Davis
 - ▶ Mike Perry
 - ▶ Greg Matthews
- Take advantage of more efficient algorithms!!!
- ► Greg's YouTube video: youtu.be/E5KJkooW4RY
- GitHub: github.com/qntkhvn/read_speed
- Question?

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