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Presentation available at github.com/bryanhanson/PANIC2017 Additional references & resources on last slide



- R is a free software "environment" for statistical computing and graphics
- The Ecosystem:
 - Base R via "R-Core"
 - Add-on packages from many authors
 - Comprehensive R Archival Network (aka CRAN) (>10,000 packages)
 - Bioconductor (>1,300 packages)
 - Unofficial repositories: Github. Gitlab. SourceForge etc.
 - Support forums
 - User guides galore!



The Ecosystem: Support Resources

- Official Documentation
- Focused, Topical Task Views
- R-Bloggers: over 600 R-oriented bloggers
- Stack Overflow: over 160K questions on use of R
- Hundreds of "Intro to R" documents on the web
- Dozens of free R books on the web
- Many packages have a "vignette" or user guide.
- More resources on last slide.



Features of R

- Written by statisticians
- " ...a rather unlikely linguistic cocktail ... " 1
- Cross-Platform: Windows, Linux, Mac OS
- Infrastructure: ready integration, interactive options
- Interfaces to many other languages, programs
 - SAS, SPSS, python, JavaScript, MATLAB, C++ etc.
- Several ways of running in parallel, using multiple cores
- Command line, or several GUI options



R is Open Source

Free!

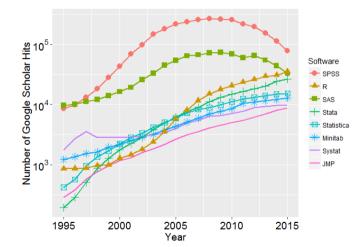
What is R7

- Transparent: All code readily available for inspection
- "Given enough eyeballs, all bugs are shallow" Linus Torvalds
- Many parts of the ecosystem are community driven

"Open source means everyone can see my stupid mistakes. Version control means everyone can see every stupid mistake I've ever made "

Karl Broman





²Bob Muenchen r4stats.com/articles/popularity/



Who Uses R?

- AirBnB
- 7illow³
- Etsy
- NYT
- Twitter
- Facebook



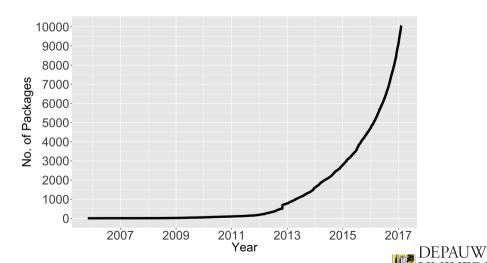


ChemoSpec

Demo

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User Contributed Packages⁴



Automation of Workflow:

data
$$\rightarrow$$
 analysis code + explanatory text \rightarrow figures + tables + text = report

- Many resources for reproducible research
- Several possible input formats
- Typical output formats are pdf files and web pages
- This presentation written with LATEX and R via the knitr package.



- ChemoSpec = Chemometrics + Spectroscopy
- Tools for exploratory data analysis
- No attempt to duplicate functions available on the spectrometer



ChemoSpec: Design Goals

- User friendly design
- Helpful error messages
- Reliable results
- High quality plots
- Consistent plot appearance
- Provide access to a wide range of chemometric operations
- Extensibility
- Developed with metabolomics and IR, NMR & Raman in mind



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What Can ChemoSpec Do?

- Data Cleaning & Prep
 - Import data
 - Remove samples
 - Drop frequency ranges
 - Baseline correction
 - Signal alignment
 - Normalization
 - Savitzky-Golay filters

- Exploratory Data Analysis
 - Plotting & surveying
 - Hierarchical cluster analysis (HCA)
 - Principal component analysis (PCA)
 - PCA diagnostics
 - Score & loading plots
 - ANOVA-PCA
 - Empirical clustering

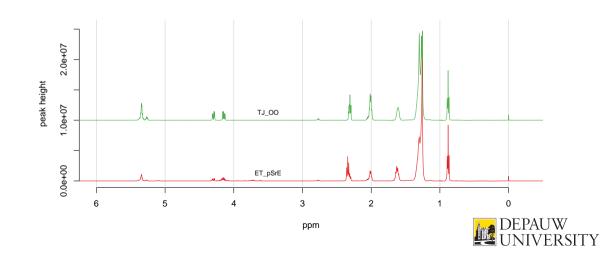


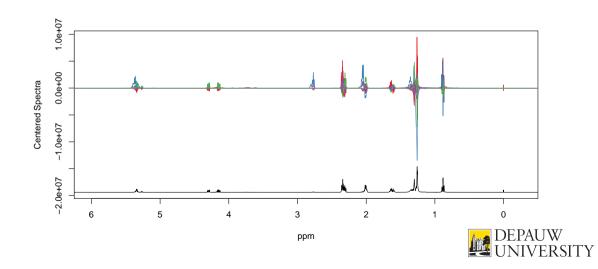
Demonstration Data Set: Saw Palmetto Caps

- Retail samples of Serenoa repens gel caps
- 500 MHz ¹H NMR in CDCl₃
- 4 samples were pure according to the label
- 10 samples have another oil present per label
- 2 outliers: olive oil, and evening primrose oil
- Serenoa repens extracts mainly fatty acids
- Outliers mainly triglycerides

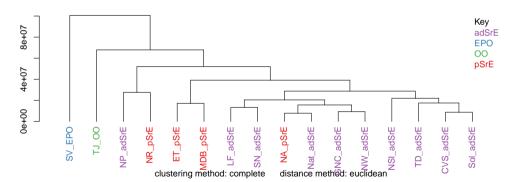


Representative ¹H NMR Spectra



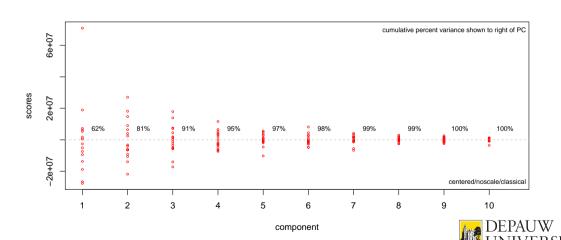


Hierarchical Clustering

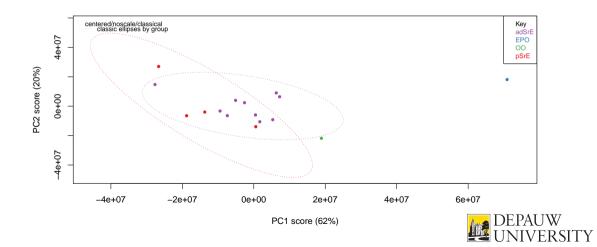




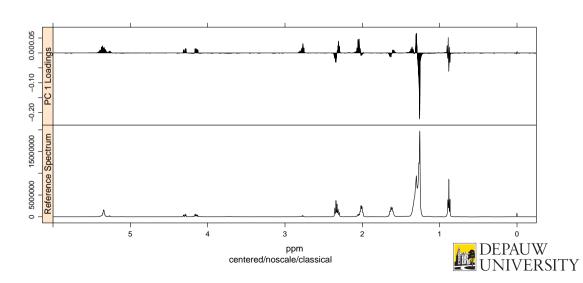
Principal Component Analysis: Scree Plot



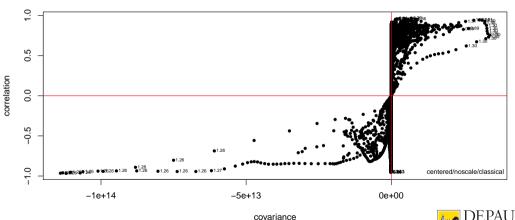
Principal Component Analysis: Score Plot



Principal Component Analysis: Loadings Plot



Principal Component Analysis: "S" Plot





Acknowledgements

- Thanks for your attention!
- Kristie Adams for the invite
- Sabbatical Support, DePauw University



Additional References & Resources

- R Project Home Page
- Selected Topical Task Views
 - Chemometrics & Computational Physics
 - Clinical Trials
 - Experimental Design
 - Pharmacokinetics
 - Machine Learning
 - Reproducible Research
- Bioconductor Home Page

