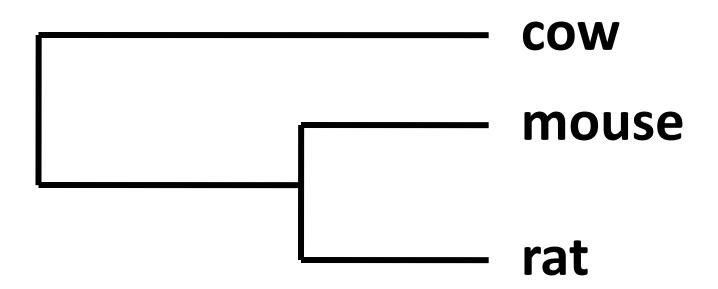
Google as a taxonomic engine

Alexey Shipunov

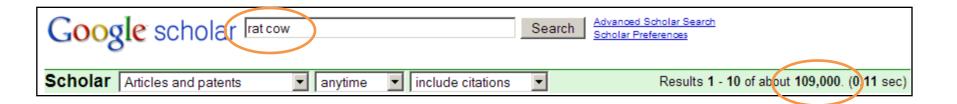
Background: tree



Background: Google Scholar







- Extract phyla names
- Obtain numbers of joint hits
- Do some magic
- Calculate similarity
- Make clusters
- Same, with "-ecology"
- Same, with classes names

Numbers of joint hits

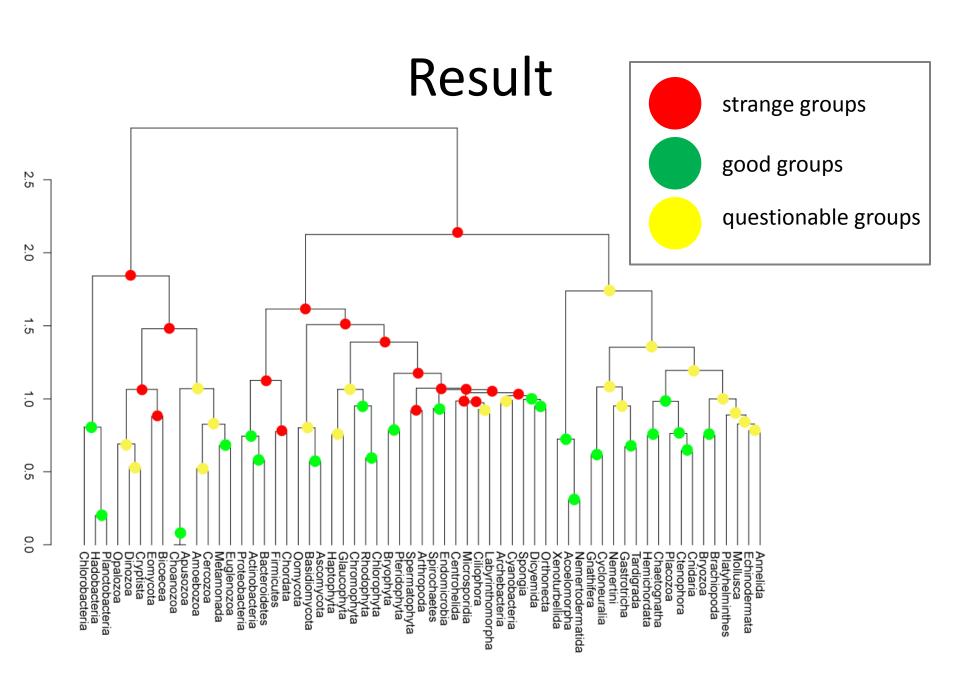
- Phyla names from my "synat" classification
- R script to make command-line queries
- Links text browser to make textual output
- UNIX text tools (sed, grep) to clean results
- Comma-delimited file for import into R

Magic with numbers

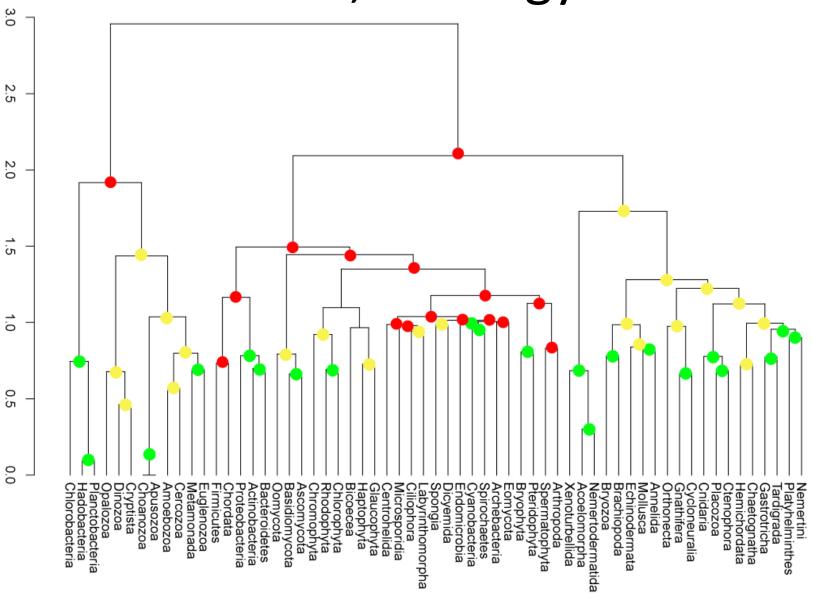
- Some names appear much often than others
 - found numbers of individual hits
 - make weights
 - multiple numbers of joint hits by geometric means of individual hits from each taxon in pair
- Convert table of three column into square matrix
- Convert similarities to dissimilarities

Similarity

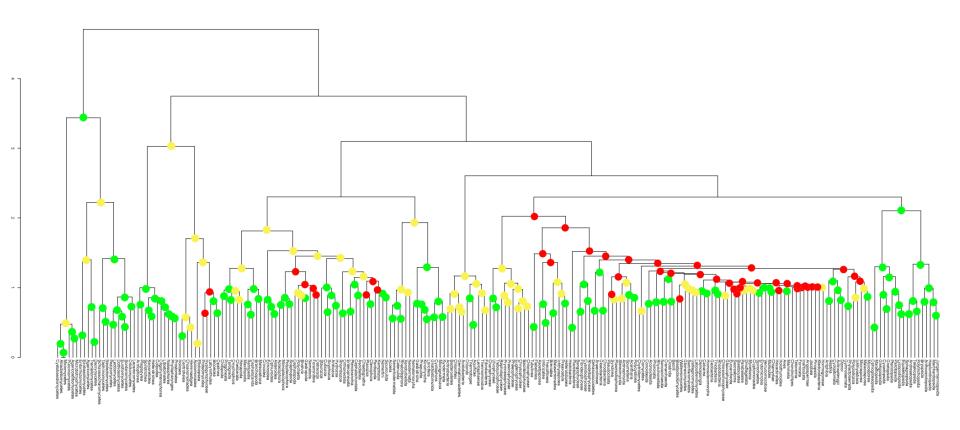
- Calculate Euclidean distances
- Ward's method hierarchical clustering
- Tree of clusters is NOT a phylogenetic tree



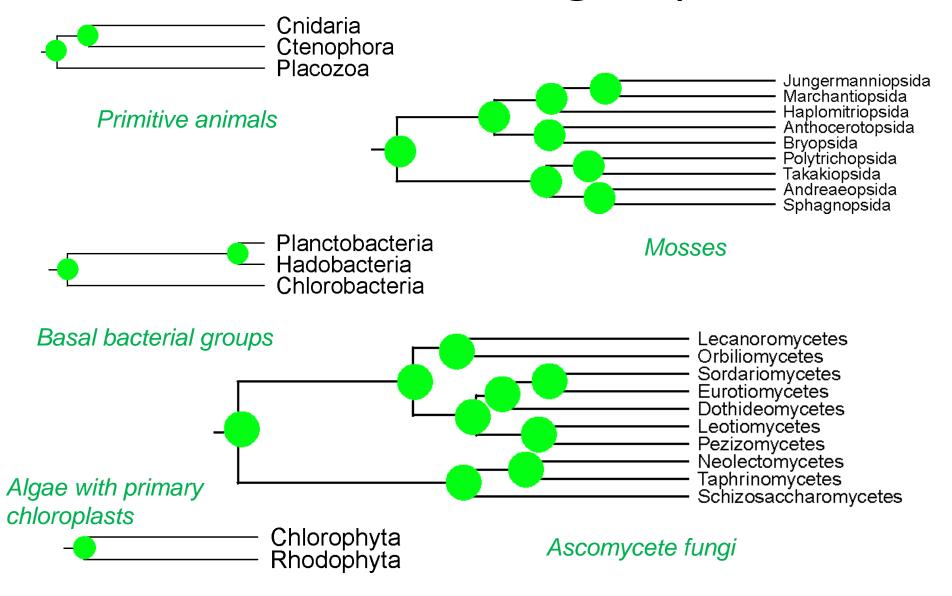
Same, -ecology



Classes



Some reliable groups

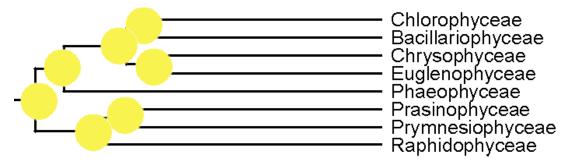


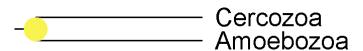
Questionable groups



Spiralians + starfishes

Algal mix





Two distantly related groups, both contain amoebae

Strange groups

Arthropoda
Spermatophyta

Insects + flowering plants

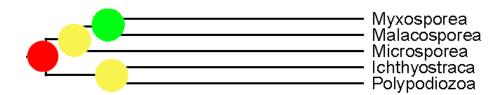
Ostracoda
Foraminifera
Nematoda
Gastropoda
Bivalvia
Polychaeta
Cephalopoda

protists

————— Chordata ————— Firmicutes

Creatures with shells

Vertebrates + Grampositive bacteria



Parasitic creatures, animals and protists

Conclusions

- It is working!
- In most cases, only closest taxa were revealed;
 animals, bacteria and fungi were intermixed
- "-ecology" did not help
- Classes are generally better than phyla

"Best classification is a classification which does not exist, it is a constantly changing product of processing all reliable data available on-line"