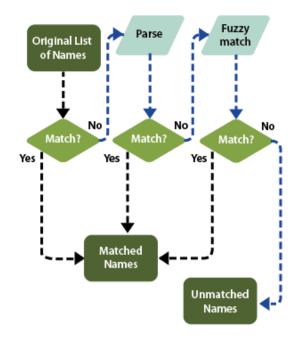
Lessons from Plant Genomics



Taxonomic Name Resolution Service (TNRS)

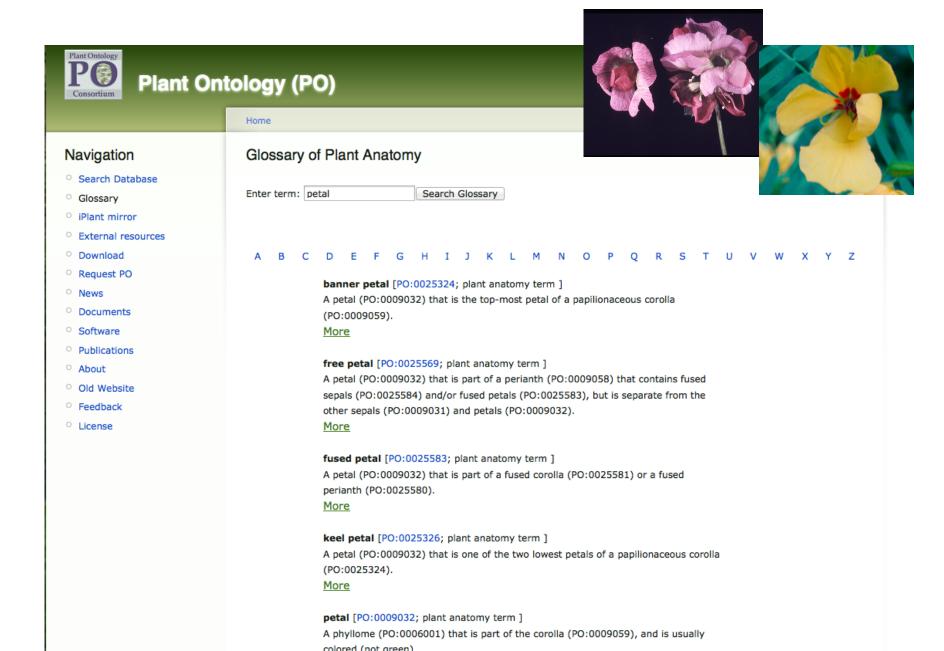
The Taxonomic Name Resolution Service (TNRS) is a tool for the computer-assisted standardization of plant scientific names. TNRS corrects spelling errors and alternative spellings to a standard list of names and converts out-of-date names to the currently accepted name. TNRS can process many names at once, saving hours of tedious and error-prone manual name correction.



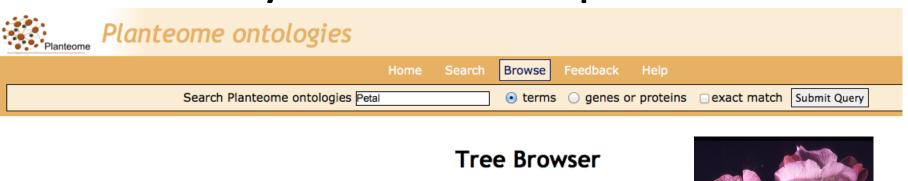


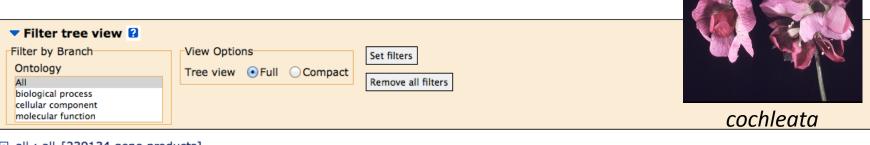
Variety

Variety: Anatomical and morphological



Variety: Genes and proteins





- all : all [239134 gene products]
 - PO:0025131: plant anatomical entity [104456 gene products]
 - - - PO:0025007: collective plant organ structure [86953 gene products]
 - PO:0025023 : collective phyllome structure [56355 gene products]

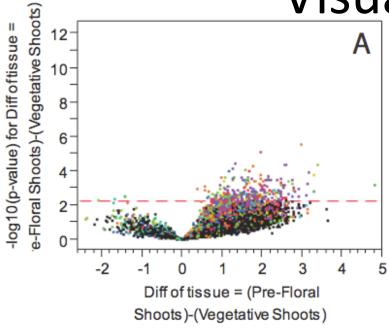
- blade-on-petiole
- - - - → participates_in relationship PO:0025082 : reproductive shoot system [59790 gene products]

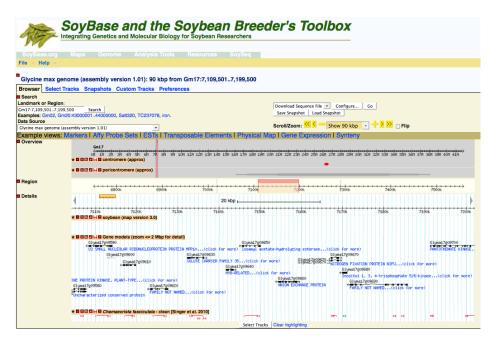


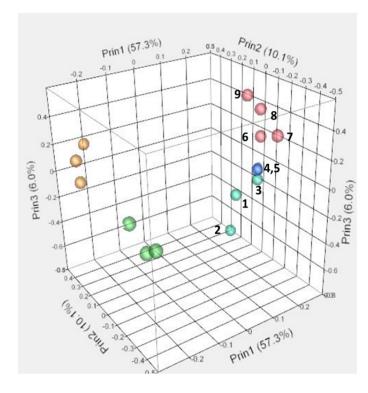
Plantcell Volume 17(5):

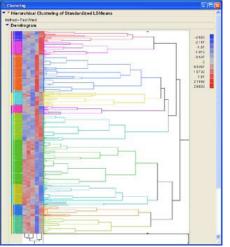
434-1448

Visualization



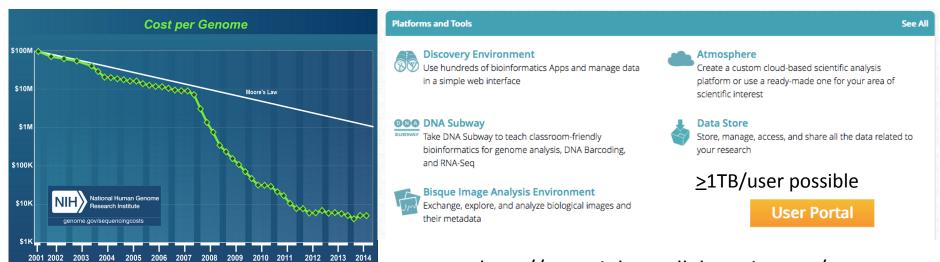






Velocity, Volume, and Veracity

- 2003 Human Genome Project completed after 13 years
- Sanger-based sequencing (average read length=500-600 bases): 6-fold coverage
- 454 sequencing (average read length=300-400 bases): 10-fold coverage
- Illumina and SOLiD sequencing (average read length=50-100 bases): 30-fold coverage



www.genome.gov/sequencingcosts

http://www.iplantcollaborative.org/