

# CyVerse DNA Subway



## Classroom-friendly Bioinformatics



Cold  
Spring  
Harbor  
Laboratory



# Challenge – bringing students into the fold



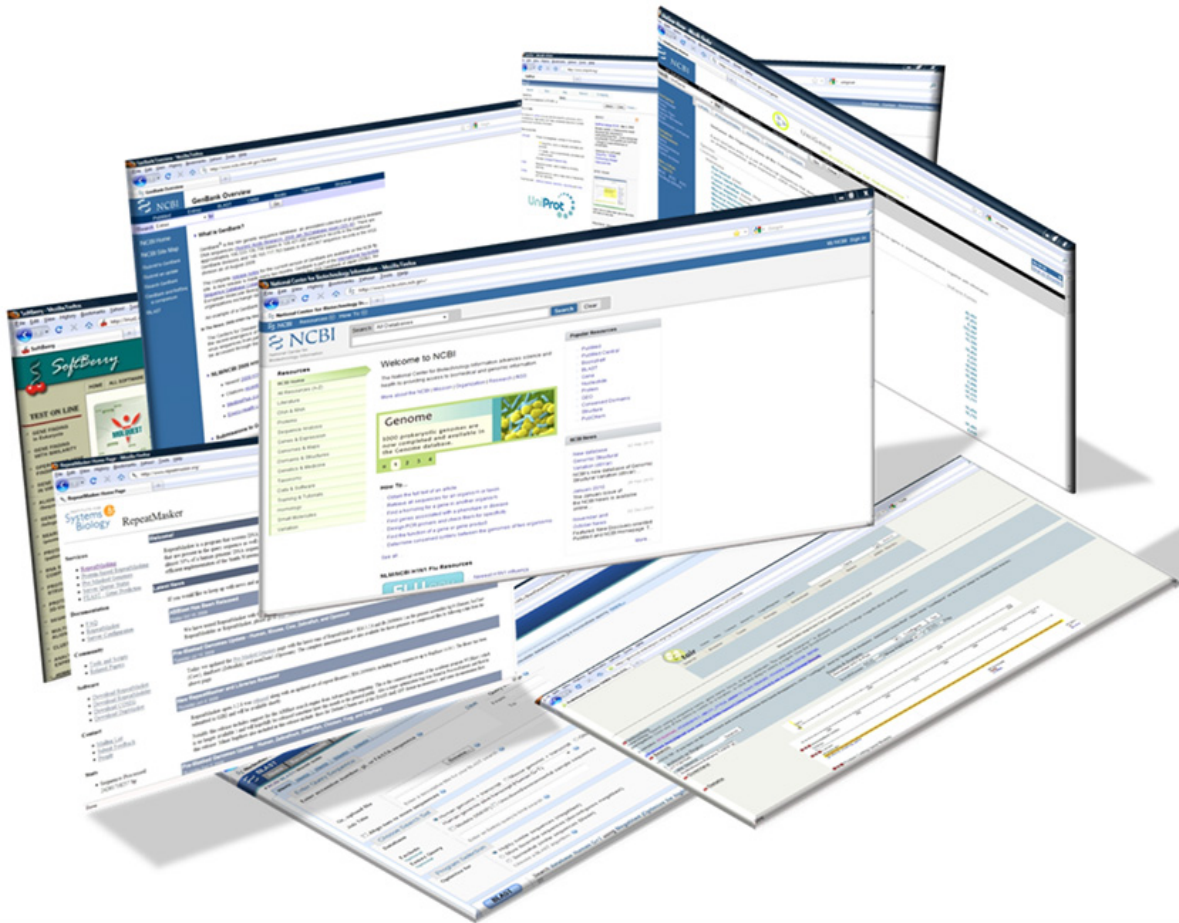
Research

Education

Students can work with the same data at the same time and with the same tools as research scientists.



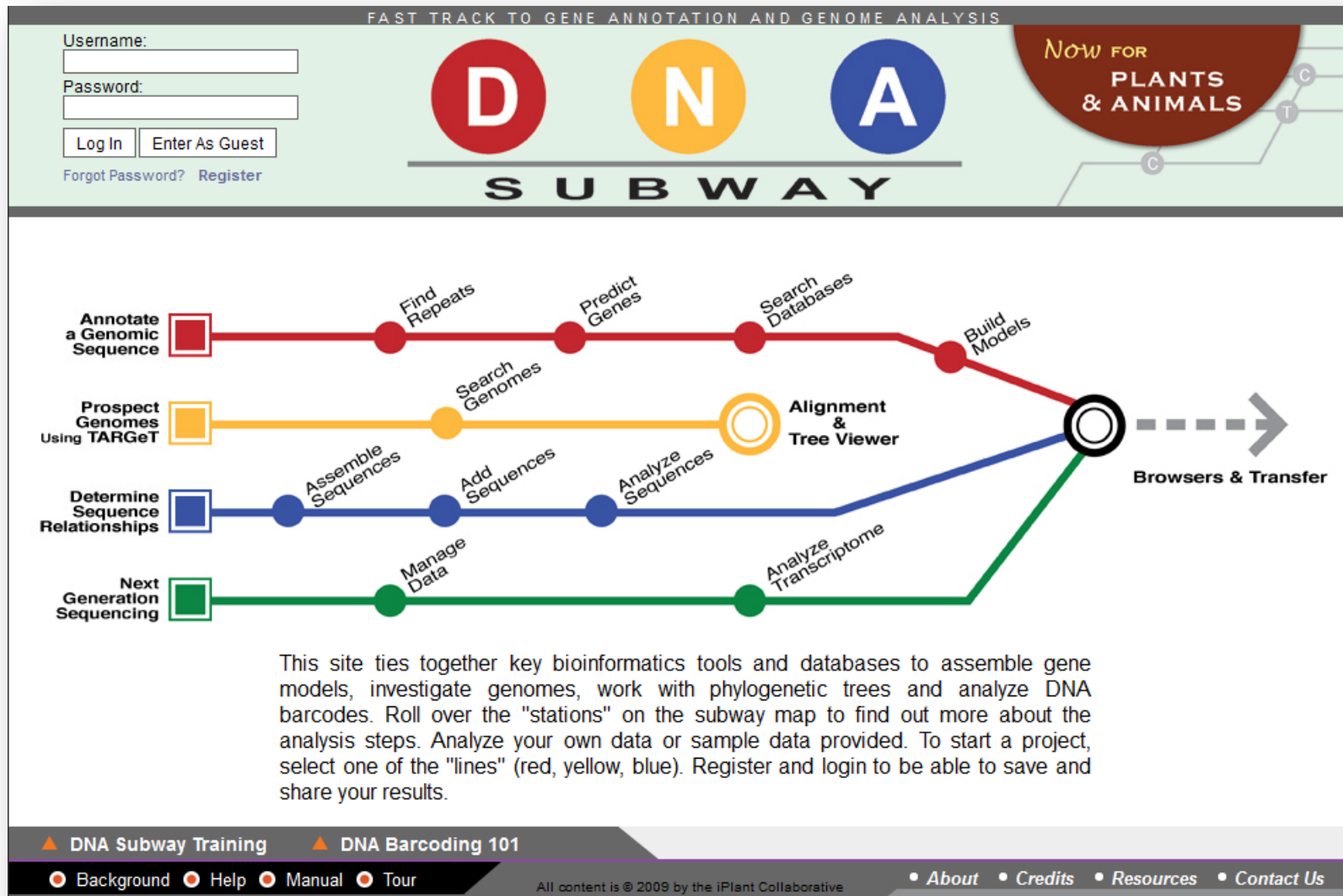
# Can you navigate the tools?



What are your challenges in teaching bioinformatics in the classroom?



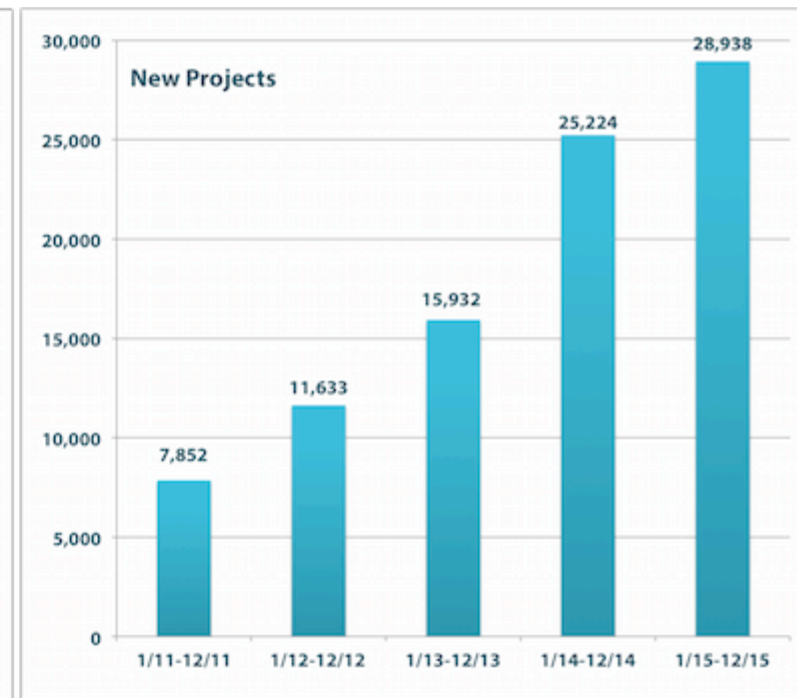
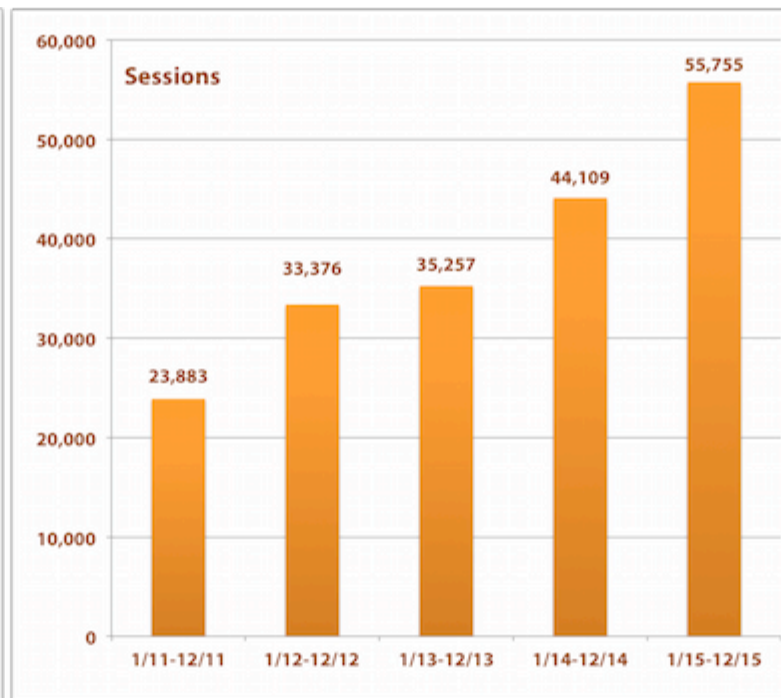
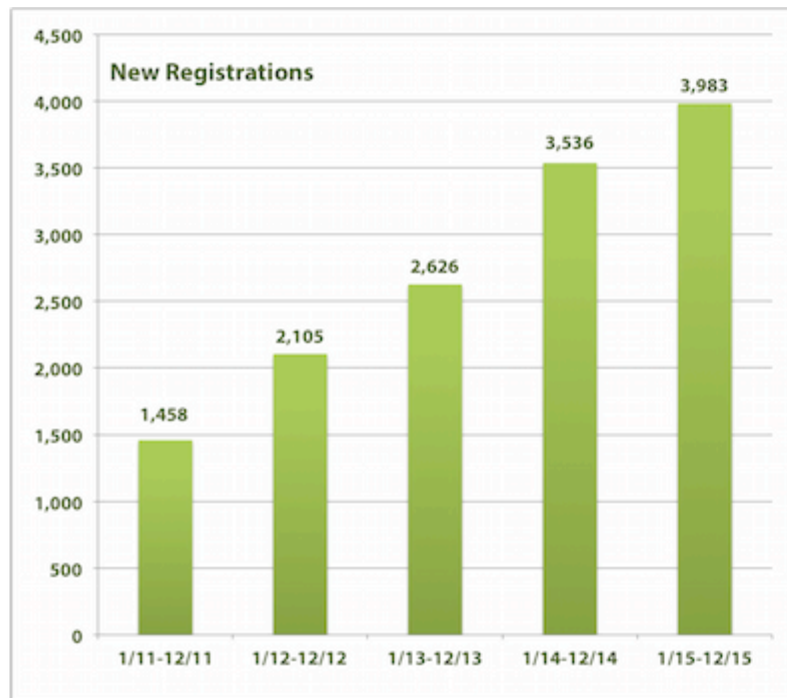
# Take the Subway



Faculty identified guiding requirements  
that shaped the development of CyVerse educational platforms:

- *Mix lecture and lab* – have a wet bench “hook”
- *Student-scientist partnerships* – someone has to care about the data
- *Co-investigation* – projects should potentially lead to publications
- *Scale* – platforms should support projects multiple classrooms can join.





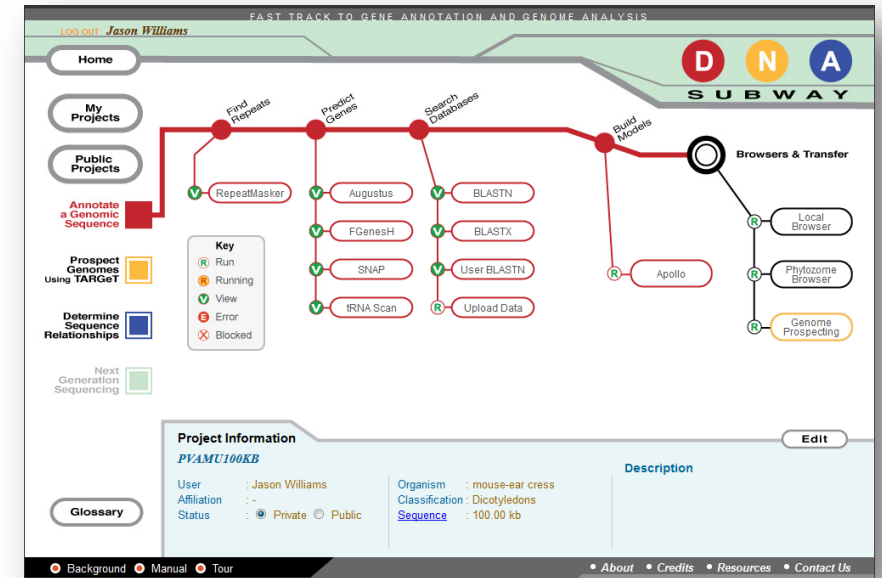
More than 13,000 users

More than 28,000 student projects in 2015



## Red Line

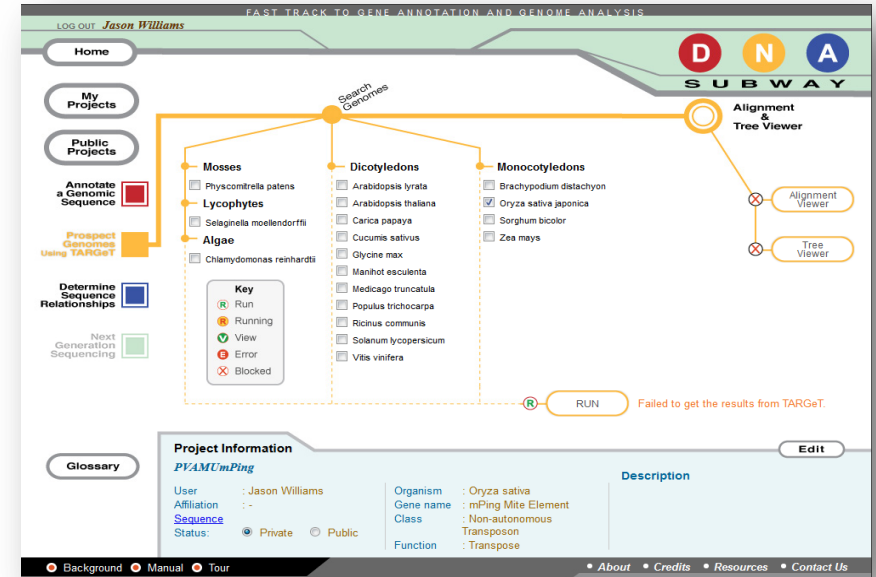
- Analyze up to 150 KB of DNA sequence
- De novo gene prediction
- Construct evidence-based gene models
- Visualize genome sequence in browser





## Yellow Line

- Analyze DNA or protein sequence
- Search plant genomes using TARGeT
- Explore gene duplications, transposons, and non-coding sequences not detectable in conventional BLAST searches





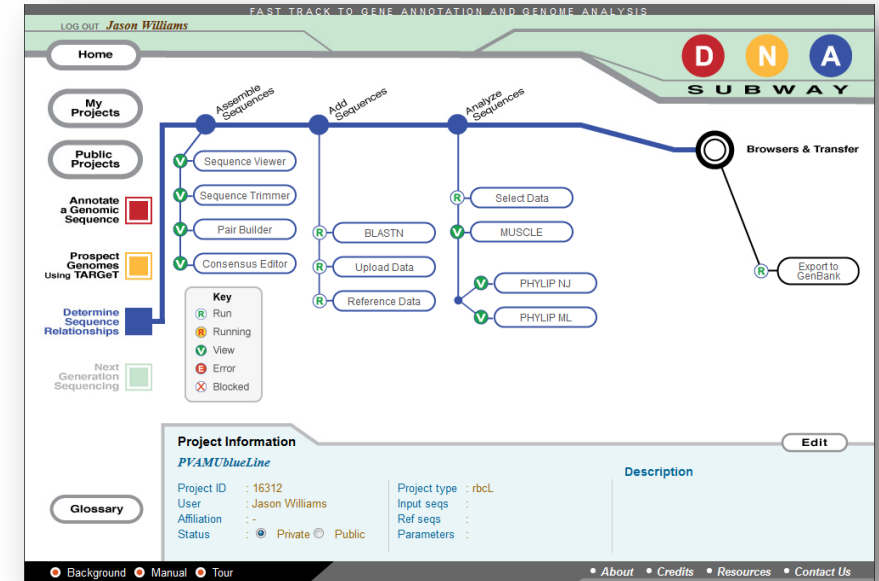
# DNA Subway

SUBWAY Blue Line: DNA barcoding, and phylogenetics



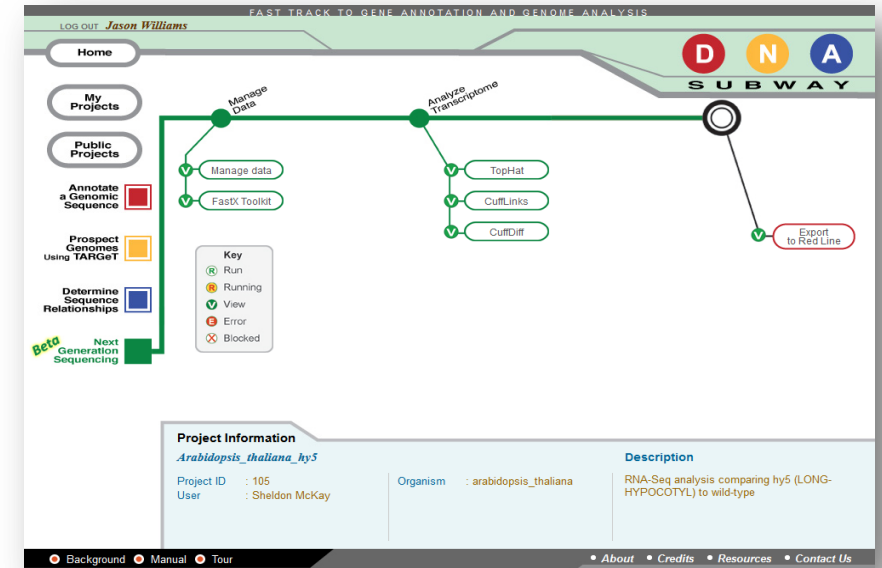
## Blue Line

- Analyze DNA or protein sequence
- Search plant genomes using TARGeT
- Explore gene duplications, transposons, and non-coding sequences not detectable in conventional BLAST searches



## Green Line

- Examine RNA-Seq data for differential expression
- Use High-performance computing to analyze complete datasets
- Generate lists of genes and fold-changes; add results to Red Line projects





Transforming Science Through Data-driven Discovery

## CyVerse Executive Team



**Parker Antin**  
**Nirav Merchant**  
**Eric Lyons**



**Matt Vaughn**



**Doreen Ware**  
**Dave Micklos**



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