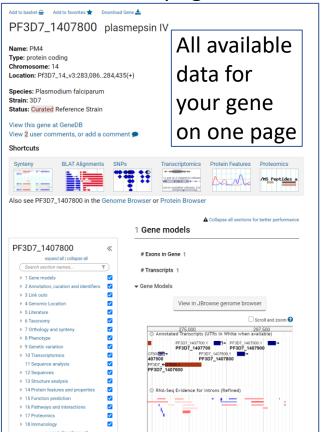
C-1173 VEuPathDB.org: eukaryotic pathogen, vector and, host Omics data mining for everyone

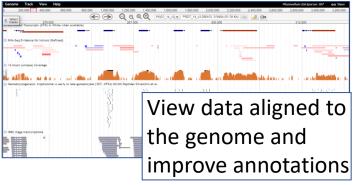
Warrenfeltz S¹, Harb OS², Kissinger JC¹, McDowell MA³, Roos DS², the VEuPathDB Team 1 University of Georgia, 2 University of Pennsylvania, 3 University of Notre Dame

- Free online omics data mining resource
- Over 400 arthropod vectors, eukaryotic pathogens, and fungi supported
- Pre-analyzed omics data is coupled with advanced search capabilities, data visualization and custom analysis tools to facilitate hypothesis testing
- Discover meaningful biological relationships from large volumes of data
- Genome sequence and annotation, transcriptomics, proteomics, epigenomics, metabolomics, population resequencing, clinical data, surveillance data, hostpathogen interactions, and orthology profiles across all integrated organisms

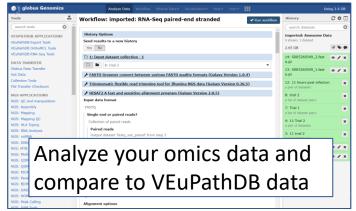


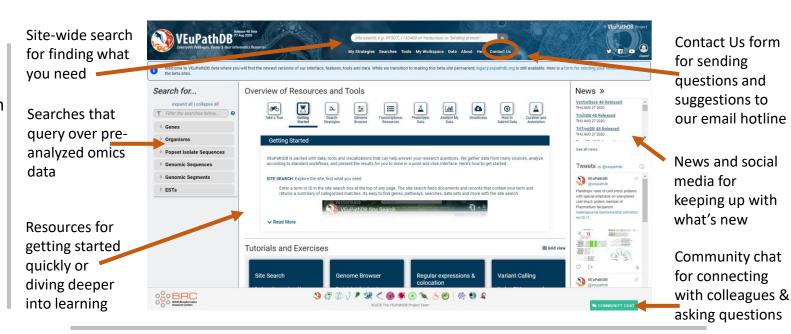


Genome Browser & Apollo

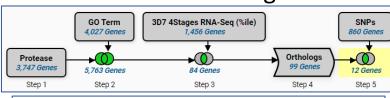


Galaxy Workspace

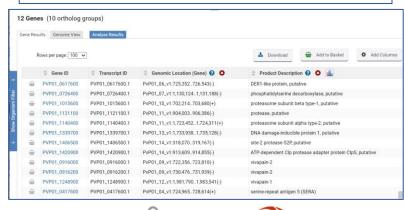




Search Strategies



Probe across many genomes to ask sophisticated questions and find meaningful biological relationships. Relate gene function, expression level and timing, variation and more.



MapVEu:

