Anatomy of the FungiDB main page

The Eukaryotic Pathogen, Vector and Host Informatics Resources (https://beta.VEuPathDB.org) is comprised of a family of bioinformatics resources including an integrated functional genomics database for fungi and oomycetes - FungiDB.

FungiDB (https://beta.FungiDB.org) is a free online resource for data-mining and functional genomics analysis. It provides an easy-to-use, interactive interface to explore genomes, annotation, functional data (transcriptomics or proteomics), metabolic pathways and results from numerous genome wide analyses (ie. InterPro scan, signal peptide and transmembrane domain predictions, orthology, etc.). FungiDB also contains an expanding number of genomes from species spanning the Oomycetes and Fungi groups including but not limited to plant, animal, and human pathogens.

