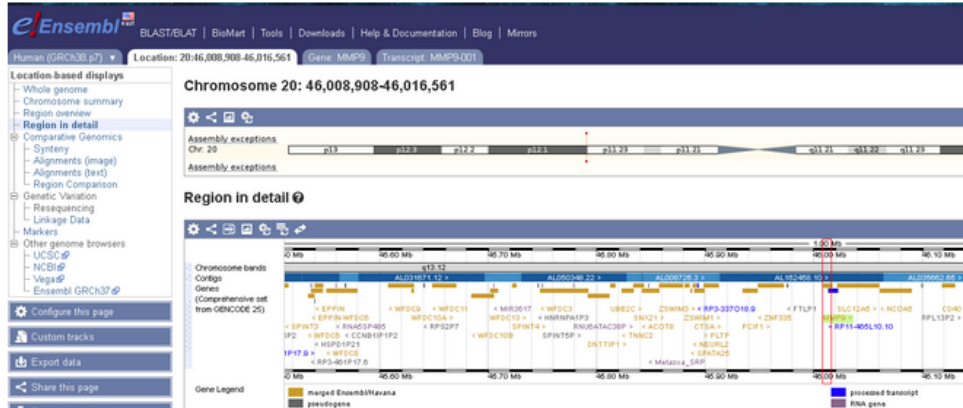


Introduction to Genome Browsers. Part 1 Ensembl



Ensembl provides access to genomic information with a number of visualization tools. By using Ensembl researchers can download data directly (e.g., genomic sequences), visualize many types of data (e.g., structural, variation, regulatory) directly on a genome assembly. In this session will review the basic functionalities and navigation of Ensembl by using specific examples. We will also use BioMart interface to answer questions and retrieve data and information from databases without the need of having any programming expertise.

Please bring your laptop to follow the instructor.



Date: Tuesday, September 19, 2017
Time: 10:30am - 12:00pm
Location: Cushing/Whitney Medical Library, Room 103 TCC, 333 Cedar St, New Haven CT 06520
Campus: Medical School
Categories: Bioinformatics