

## Table Browser

Use this program to get the data associated with a track in text format, to calculate intersections between tracks, and to retrieve DNA sequence covered by a track. See [Using the Table Browser](#) for a description of the controls in this form.

**clade:**  **genome:**  **assembly:**   
**group:**  **track:**   
**table:**    
**region:** ☐ genome ☒ position    
**filter:**   
**intersection:**   
**correlation:**   
**output format:**   
**output file:**  (leave blank to keep output in browser)  
**file type returned:** ☒ plain text ☐ gzip compressed

 

To reset **all** user cart settings (including custom tracks), [click here](#).