

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](#).