

## 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:** Vertebrate ▾   **genome:** Human ▾   **assembly:** Mar. 2006 ▾

**group:** Genes and Gene Prediction Tracks ▾   **track:** Conservation ▾

**table:** multiz17way ▾   [describe table schema](#)

**region:**  genome  position chr11:93103000-93108000 [lookup](#)

**filter:** [create](#)

**intersection:** [create](#)

**correlation:** [create](#)

**output format:** MAF - multiple alignment format ▾

**output file:** region.maf (leave blank to keep output in browser)

**file type returned:**  plain text  gzip compressed

[get output](#)   [summary/statistics](#)

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