## 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.