

## Testing Report

Version of GenMAPP Builder: 2.0b71

Computer on which export was run: Personal computer

Postgres Database name: CT\_KS\_20131119\_32bit\_gmb2b71

UniProt XML filename: Uniprot\_XML\_C.trachomatis\_serovar\_A\_KS\_20131114.xml

- UniProt XML version (The version information can be found at [the UniProt News Page](#) ): UniProt release 2013\_11
  - Original file name from [UniProt site](#) : uniprot-organism%3A315277+keyword%3A181.xml
- Time taken to import: 1.20 min

GO OBO-XML filename: Go\_daily-termdb\_v2\_HD\_20131107.obo-xml

- GO OBO-XML version (The version information can be found in the file properties after the file downloaded from the [GO Download page](#) has been unzipped): 11/06/2013
  - Original file name as listed in [beta.geneontology.org](#): go\_daily-termdb.obo-xml.gz
- Time taken to import: 13.05 min
- Time taken to process: 10.80 min

GOA filename: 22183.C\_trachomatis\_A\_KS\_20131114.goa

- GOA version (News on [this page](#) records past releases; current information can be found in the Last modified field on the [FTP site](#) ): 11/12/13
  - Original file name as listed in the FTP site: 22183.C\_trachomatis\_A.goa
- Time taken to import: 0.04 min
- Name of .gdb file: Ct-Std\_v2\_KS\_20131121.gdb
- Time taken to export .gdb:
  - Start Time: 10:56 AM (approximately)
  - End Time: 11:14:26 AM
- Upload your file and link to it here: [Ct-Std\\_External\\_20131121.gdb](#)

Note: There is one ID "CTA\_0406/CTA\_0407/CTA\_0408" that is a combination of three predicted genes that are actually one gene. The Tally Engine and PostgreSQL count this as one gene resulting in a total gene count of 917. However, through Access and *xmllpipedbmatch*, the ID is separated into three separate genes bringing the total count to 919.