

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.