[Taxid dump files](ftp://ftp.ncbi.nih.gov/pub/taxonomy/) downloaded off NCBI ftp site, changed formatting

it046050:taxon patricklab$ perl -pi -e "s/\t\|\t/\t/smgi" names.txt

it046050:taxon patricklab$ perl -pi -e "s/\t\|\t/\t/smgi" nodes.txt

it046050:taxon patricklab$ perl -pi -e "s/\t\|\n/\n/smgi" names.txt

it046050:taxon patricklab$ perl -pi -e "s/\t\|\n/\n/smgi" nodes.txt

[prokaryote summary](http://www.ncbi.nlm.nih.gov/genomes/lproks.cgi) (3 sheets were downloaded)