Dear Welkin et al,

Annotation of Squid was fairly straightforward and was performed last year (Spring 2013). I apologize for not getting the finalized version to you sooner. As it was done last school year, the old style notes are present, but I assure you that all NKFs were checked in class on BLASTp and HHpred. Those gps listed as NKF had no functional hits and anything of possible interest is listed for them. My only concerns are the very start and end of the genome as I was not sure how to record the gap properly between the first and last gene. I was also unable to BLAST a gene that we inserted using DNAmaster (the following error kept appearing: HTTP/1.1 404 not found) so I went to NCBI and did the BLASTn on the DNA sequence there and got the hit recorded for 1:1 alignment with gp67 of Fang. All other genes have all BLAST info included on DNAmaster.

Thanks and let me know if you need anything else,

Deb

p.s. This year’s annotation is wrapping up and will be sent out by May 15th.