

Genome averages for tBLASTn matches to three proteomes, with 95% confidence intervals. Species include three outgroups (*C.elegans*, *Daphnia pulex*, *Anopheles gambia*), and twelve *Drosophila* species, Dmel .. Dgri, for CAF1 assemblies. Averages are shown for Fruitfly (blue), Mouse (brown) Worm (green) proteomes.

