

Figures show genome averages for matches to *Drosophila melanogaster* proteome measured with tBLASTn, with 95% confidence intervals. Species include three outgroups (*C.elegans*, crustacean *Daphnia pulex*, *Anopheles gambia*), and twelve *Drosophila* species, Dmel .. Dgri, for CAF1 assemblies with two alternates (dpse2r, dyak3r).

