# *Drosophila fulvimacula*

**Taxonomic lineage:** Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > Drosophila subgenus Drosophila > repleta group > repleta subgroup

**Assembly:** GCA_035045555.1_ASM3504555v1_genomic

**Dro_ful1 | JAWNOX010000203.1:6591128-6592042 (+) | 915 nt | lncRNA:noe consensus e-value: 5e-170**

5' motif: GCGGT  3' motif: ATCGC  Internal max. Poly-T: 3nt  Trailing-T: 7nt  PSE: 0.86

Upstream promoter alignment showing conserved regions. Sequences include Pol III promoters (noeCR34335, Arthropod_7SK, 7SL, RNase_MRP, RNaseP_nuc, U6, U6atac, tRNAsec, snoRNAMe18SA1806) and Pol II promoters (U1, U2, U3, U4, U4atac, U5, U7, U8, U11, U12, OrCD1). Conserved positions are highlighted in the alignment.