

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Hirtodrosophila > quadrivittata group > confusa subgroup
Assembly: GCA_037075165.1_ASM3707516v1_genomic

[illegible][illegible][illegible]

	Hir_his1	Hir_his2	Hir_his3
Hir_his1	-	43	39
Hir_his2		-	60
Hir_his3			-

Pairwise sequence identity matrix (%) showing similarity between lncRNA sequences.

[illegible]

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.