Pherbellia schoenherri

Taxonomic lineage: Brachycera > Muscomorpha > Sciomyzoidea > Sciomyzidae > Pherbellia

Assembly: GCA 965653165.1 idPheScho1.hap1.1 genomic

Phe_sch1 | OZ289137.1:162234124-162234519 (+) | 396 nt | IncRNA:noe consensus e-value: 1.7e-07

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 8nt PSE: 0.92



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.