

Assembly: GCA_005959815.1_ROC_SBull_1.0_genomic

5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 3nt Trailing-T: 10nt PSE: 0.93

5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.98

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	Sar_bul1	Sar_bul2	Sar_bul3
Sar_bul1	-	31	32
Sar_bul2		-	97
Sar_bul3			-

[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.