

Sorted Enrichment of Variations in Gene Length vs Total Length

Z-score of Enrichment

10.0
7.5
5.0
2.5
0.0

BLACE
MIR3689A
LINC00459
TRD-GTC2-8
FAM30A
SNAR-G1
TRR-TCG1-1
TRK-CTT3-1
tRNA-Ile-AAT-2-1
TRC-GCA11-1
RCCD1-AS1
tRNA-Arg-CCT-3-1
TRG-TCC1-1
tRNA-Val-CAC-1-3
tRNA-Met-CAT-2-1
TRT-AGT1-2
TRA-TGC4-1
tRNA-Glu-TTC-4-1
tRNA-Ile-AAT-1-1
TRV-TAC1-1
SNAR-G2
TRP-TGG2-1
tRNA-Asn-GTT-3-2
tRNA-Met-CAT-6-1
TRE-TTC2-1
SYT15-AS1
SNAR-A1
Inc-SLCO4A1-8
TRV
TRM-CAT2-1
tRNA-Pro-AGG-2-4
TRG-CCC2-1
RNU5E-1
tRNA-Leu-CAG-2-2
tRNA-Val-CAC-3-1
RNU5D-1
tRNA-Lys-TTT-1-1
tRNA-Lys-TTT-5-1
TRN-GTT2-4
TRE-TTC3-1
tRNA-Asn-GTT-4-1
tRNA-Ala-TGC-3-1
TRA-AGC20-1
TRQ-TTG2-1
TRL-TAG2-1
RNU6-9
SNORD13
TRK-TTT7-1
tRNA-Asp-GTC-1-1
tRNA-Pro-TGG-1-1
TRR-ACG2-3
TRI-AAT4-1
TRC-GCA4-1
TRT-TGT5-1
TRA-AGC15-1
TRP-AGG1-1
TRF-GAA2-1
TRV-AAC5-1
tRNA-Ala-CGC-3-1
TRV-CAC1-5
SNAR-F
tRNA-Ile-AAT-12-1
tRNA-Leu-TAG-3-1
tRNA-Ala-TGC-2-1
TRG1
tRNA-Pro-TGG-3-3
TRR-TCG3-1
RNU5F-1
tRNA-Arg-ACG-1-2
TRL-AAG2-4
TRK-TTT3-2
TRA-AGC11-1
TRQ-CTG6-1
TRA-AGC5-1
TRG-CCC6-1
SNORD118
tRNA-Lys-TTT-6-1
SNAR-C3
TRF-GAA1-4
TRI-AAT5-1
tRNA-Thr-CGT-6-1
tRNA-Lys-CTT-5-1
TRE-TTC5-1
LINC01671
tRNA-Val-AAC-3-1
tRNA-Lys-CTT-1-2
tRNA-Ser-TGA-2-1
tRNA-Ser-CGA-2-1
tRNA-Ala-CGC-1-1
tRNA-Gly-TCC-3-1
tRNA-Gln-TTG-3-1
TRL-TAA4-1
TRK-TTT4-1
TRA-AGC1-1
SNAR-B2
tRNA-Arg-CCT-2-1
tRNA-Ser
TRA-AGC4-1
SNAR-C4
tRNA-Ser-GCT-2-1
TRG-CCC5-1
TRS-AGA2-6
TRA-CGC2-1
TRA-CGC4-1
tRNA-Asn-GTT-1-1
tRNA-Gly-CCC-4-1
tRNA-Thr-TGT-4-1
tRNA-Gln-CTG-2-1
tRNA-Thr-CGT-5-1
TRM-CAT3-2
TRX-CAT1-5
TRA-CGC5-1-001
tRNA-Gly-GCC-2-4
tRNA-Lys-CTT-2-1
tRNA-Glu-CTC-1-2
tRNA-Val-CAC-2-1
tRNA-Thr-TGT-1-1
tRNA-Arg-TCG-4-1
TRT-TGT2-1
Val-tRNA
MIR4538

Gene