

AE3R8_siApoA1-BAF-2

%features

60
50
40
30
20
10
0

% in genome
detected
% in sample

5.7
4.7
3.8
2.8
1.9
0.9
0

protein_coding
pseudogene
lincRNA
antisense
miRNA
misc_RNA
snRNA
snoRNA
sense_intronic
processed_transcript
sense_overlapping
rRNA
polymorphic_pseudogene
TR_J_gene
IG_V_pseudogene
3prime_overlapping_ncrna
IG_V_gene
TR_V_gene
TR_C_gene

