

Sheet1_2

Type	#found	% of total	with PAS	with good PAS	mean_covr
all	28992	0.18	0.89	0.8	69.5013387777
protein_coding	21570	0.28	0.91	0.82	67.8263400108
processed_transcript	3226	0.09	0.85	0.76	65.106738223
retained_intron	2348	0.14	0.86	0.75	83.9606879607
nonsense_mediated_decay	1596	0.19	0.87	0.76	81.355983163
non_coding	61	0.19	0.87	0.77	61.421875
lincRNA	44	0.06	0.8	0.66	14.25
processed_pseudogene	35	0	0.17	0.14	15.4571428571
pseudogene	31	0.03	0.9	0.87	55.7096774194
snoRNA	20	0.01	0.1	0.1	6
TEC	13	0.37	1	0.92	20.4615384615
ncrna_host	9	0.31	1	1	679
snRNA	6	0	0.33	0	3
Mt_tRNA	5	0.23	0.4	0	5.2
transcribed_unprocessed_pseudogene	5	0.02	1	1	15.4
antisense	4	0.03	1	0.75	18.5
rRNA_pseudogene	3	0.02	0	0	7
IG_C_gene	3	0.17	0.67	0.33	322.666666667
retrotransposed	3	0.01	0	0	2.6666666667
misc_RNA	2	0	0	0	5.5
scRNA_pseudogene	2	0	0	0	2.5
rRNA	1	0	0	0	2
polymorphic_pseudogene	1	0.03	1	1	3
IG_J_pseudogene	1	0.33	0	0	5
unprocessed_pseudogene	1	0	0	0	2
ambiguous_orf	1	0.02	1	1	11
transcribed_processed_pseudogene	1	0.01	1	1	9