

Sheet1_2

Type	#found	% of total	with PAS	with good PAS	mean_covr
all	42822	0.27	0.84	0.75	205.768874128
protein_coding	31381	0.41	0.86	0.77	204.25956967
processed_transcript	5289	0.14	0.79	0.69	174.413102328
retained_intron	3346	0.2	0.78	0.68	268.982709751
nonsense_mediated_decay	2274	0.27	0.8	0.68	224.043801653
lincRNA	111	0.14	0.86	0.74	26
non_coding	95	0.29	0.78	0.69	225.747747748
processed_pseudogene	78	0.01	0.31	0.22	22.125
pseudogene	50	0.05	0.86	0.8	104.509803922
snoRNA	44	0.03	0.2	0.16	6.5777777778
ncrna_host	23	0.79	0.39	0.39	1140.85714286
TEC	20	0.57	0.9	0.85	54.45
snRNA	18	0.01	0.44	0.28	4.7777777778
antisense	12	0.09	0.92	0.83	21.3333333333
miRNA	10	0.01	0.4	0.3	3.1
transcribed_unprocessed_pseudogene	10	0.03	0.7	0.6	27.6
misc_RNA	9	0.01	0.33	0.11	5.3333333333
retrotransposed	8	0.04	0	0	5.4444444444
polymorphic_pseudogene	7	0.24	0.71	0.71	109.222222222
Mt_tRNA	7	0.32	0.29	0	13.25
unprocessed_pseudogene	6	0	0.33	0.17	31.5
scRNA_pseudogene	4	0.01	0.25	0.25	2.5
rRNA_pseudogene	3	0.02	0	0	36.6666666667
ambiguous_orf	3	0.06	1	1	31
IG_C_gene	3	0.17	0.67	0.33	322.666666667
unitary_pseudogene	2	0.01	1	1	5
rRNA	2	0	0	0	2.5
transcribed_processed_pseudogene	2	0.01	0.5	0.5	12
snRNA_pseudogene	1	0.01	1	1	2
IG_J_pseudogene	1	0.33	0	0	5
Mt_rRNA	1	0.5	0	0	6
tRNA_pseudogene	1	0.01	0	0	5
Mt_tRNA_pseudogene	1	0	0	0	3