Type	#found	% of total	with PAS	with good PAS	mean_covr
all	17324	0.11	0.89	0.81	18.9066417106
protein_coding	12982	0.17	0.91	0.83	18.4179172351
processed_transcript	1675	0.04	0.83	0.75	19.8174927114
retained_intron	1525	0.09	0.88	0.77	20.0623794212
nonsense_mediated_decay	916	0.11	0.87	0.78	21.3333333333
snoRNA	52	0.03	0.04	0.02	38.5769230769
non_coding	40	0.12	0.85	0.8	9.9090909091
processed_pseudogene	26	0	0.15	0.12	8.4230769231
pseudogene	23	0.03	0.96	0.87	8.347826087
lincRNA	15	0.02	1	0.87	7
misc_RNA	12	0.01	0	0	7.9166666667
snRNA	12	0.01	0.17	0.08	17.7142857143
ncrna_host	9	0.31	1	1	43.2222222222
miRNA	8	0	0	0	5.8888888889
TEC	6	0.17	1	0.83	3
rRNA_pseudogene	6	0.03	0	0	233.833333333
scRNA_pseudogene	4	0.01	0	0	2.5
Mt_tRNA	3	0.14	0.33	0	4.75
IG_C_gene	2	0.11	0.5	0.5	6.5
unprocessed_pseudogene	2	0	0.5	0.5	2.5
transcribed_unprocessed_pseudogene	2	0.01	1	1	4
rRNA	1	0	0	0	2
polymorphic_pseudogene	1	0.03	1	1	22
IG_J_gene	1	0.01	0	0	3
retrotransposed	1	0	0	0	3