Type	#found	% of total	with PAS	with good PAS	mean_covr
all	6407	0.04	0.91	0.83	9.3180902186
protein_coding	4698	0.06	0.92	0.84	9.1972904318
processed_transcript	673	0.02	0.89	0.82	9.302189781
retained_intron	641	0.04	0.91	0.77	9.201863354
nonsense_mediated_decay	305	0.04	0.91	0.85	11.2597402597
non_coding	22	0.07	0.86	0.77	3.0454545455
snoRNA	18	0.01	0.11	0.06	9.1666666667
ncrna_host	9	0.31	1	1	7.4444444444
pseudogene	8	0.01	0.75	0.75	4.125
processed_pseudogene	8	0	0	0	7
snRNA	5	0	0	0	8.6
lincRNA	5	0.01	1	0.8	3.2
rRNA_pseudogene	4	0.02	0	0	101.75
miRNA	4	0	0	0	2.5
misc_RNA	3	0	0	0	4.6666666667
IG_C_gene	2	0.11	0.5	0.5	6.5
IG_J_gene	1	0.01	0	0	3
polymorphic_pseudogene	1	0.03	1	1	3