

%features

60  
40  
20  
02.4  
1.6  
0.8  
0

■ % in genome  
▨ detected  
■ % in sample

protein\_coding  
processed\_pseudogene  
lincRNA  
antisense  
TEC  
miRNA  
processed\_transcript  
snoRNA  
snRNA  
unprocessed\_pseudogene  
d\_processed\_pseudogene  
misc\_RNA  
sense\_intronic  
rRNA  
unprocessed\_pseudogene  
pseudogene  
sectional\_promoter\_lncRNA  
polymorphic\_pseudogene  
IG\_V\_gene  
scaRNA  
TR\_J\_gene  
sense\_overlapping  
IG\_V\_pseudogene  
unitary\_pseudogene  
IG\_C\_gene  
ribozyme  
ribosomal\_pseudogene  
TR\_C\_gene  
TR\_V\_gene  
TR\_V\_pseudogene  
IG\_J\_gene  
TR\_J\_pseudogene  
IG\_D\_gene  
sRNA  
TR\_D\_gene

