

%features

60  
50  
40  
30  
20  
10  
0

protein\_coding  
processed\_pseudogene  
lincRNA  
antisense  
TEC  
miRNA  
snRNA  
snoRNA  
processed\_transcript  
processed\_pseudogene  
misc\_RNA  
rRNA  
sense\_intronic  
antisense\_intronic  
processed\_pseudogene  
unprocessed\_pseudogene  
pseudogene  
antisense\_intronic  
IG\_V\_gene  
scaRNA  
polymorphic\_pseudogene  
TR\_J\_gene  
IG\_V\_pseudogene  
TR\_V\_gene  
ribozyme  
unitary\_pseudogene  
sense\_intronic  
IG\_C\_gene  
TR\_V\_pseudogene  
unprocessed\_pseudogene  
TR\_C\_gene  
TR\_J\_pseudogene  
IG\_D\_gene  
antisense\_intronic  
IG\_J\_gene  
scRNA  
sRNA

% in genome  
detected  
% in sample

3.9  
3.3  
2.6  
2  
1.3  
0.7  
0

UNK