

%features

60  
40  
20  
0

■ % in genome  
▨ detected  
■ % in sample

3.3  
2.2  
1.1  
0

protein\_coding  
processed\_pseudogene  
lincRNA  
TEC  
antisense  
miRNA  
snRNA  
snoRNA  
processed\_transcript  
unprocessed\_pseudogene  
misc\_RNA  
sense\_intronic  
processed\_pseudogene  
rRNA  
unprocessed\_pseudogene  
pseudogene  
sectional\_promoter\_lncRNA  
scaRNA  
IG\_V\_gene  
polymorphic\_pseudogene  
IG\_V\_pseudogene  
TR\_J\_gene  
sense\_overlapping  
TR\_V\_gene  
ribozyme  
unitary\_pseudogene  
IG\_C\_gene  
tribed\_unitary\_pseudogene  
TR\_C\_gene  
IG\_D\_gene  
TR\_V\_pseudogene  
IG\_J\_gene  
TR\_J\_pseudogene  
prime\_overlapping\_ncRNA  
macro\_lncRNA  
scRNA  
sRNA