

## IT3R39-3343-4

%features

60  
40  
20  
0

■ % in genome  
▨ detected  
■ % in sample

3.1  
2.1  
1  
0

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