

## SK3R11\_Tdii22\_10m

%features

60  
40  
20  
02  
1.3  
0.7  
0

■ % in genome  
▨ detected  
■ % in sample

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

