

## DK3R3r\_C3

%features

60  
40  
20  
0

■ % in genome  
▨ detected  
■ % in sample

1.7  
1.1  
0.6  
0

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