

# MF3R5\_GFP2

%features

60  
50  
40  
30  
20  
10  
0

■ % in genome  
▨ detected  
■ % in sample

3.2  
2.7  
2.2  
1.6  
1.1  
0.5  
0

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