

11\_k8\_D

%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

ribribed\_unitary\_pseudogene

TR\_C\_gene

IG\_D\_gene

TR\_V\_pseudogene

IG\_J\_gene

TR\_J\_pseudogene

prime\_overlapping\_ncRNA

macro\_lncRNA

scRNA

sRNA