

Removed sequence

30%  
20%  
10%  
0%

mSymSyn1#Hap1

mSymSyn1#Hap2

mGorGor1#MATERNAL

Col-CC-v2

GCA\_000597845.1

grch38

mm39

grch38

SGDref

ZH13

SL5

athaliana82

ecoli50

ecoli500

hsapiens90.chr6

mmusculus36.chr19

primates16.chr6

scerevisiae142

soy37

tomato23

Pangenome