

*k*-mers

120000

100000


80000

60000

40000

20000

0

 *hist.txt*

1

2

3

4

5

6

7

8

9

10

11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

27

28

29

30

Percentage of Genomes

