



49-Lrh30.faa

41-Lrh20.faa

45-Lrh26.faa

20-Lc 705.faa

44-Lrh23.faa

8-ATCC 8530.faa

35-Lrh10.faa

21-LOCK900.faa

11-CLS17.faa

7-ASCC 3029.faa

28-LR231.faa

46-Lrh27.faa

23-Lr044.faa

36-Lrh11.faa

17-L31.faa

5-ASCC 3016.faa

2-40f.faa

16-HN001.faa

10-BFE5264.faa

60-WQ2.faa

24-Lr071.faa

56-Lrh46.faa

52-Lrh38.faa

38-Lrh14.faa

53-Lrh39.faa

50-Lrh32.faa

40-Lrh18.faa

34-Lrh9.faa

33-Lrh7.faa

15-HCT70.faa

9-ATCC 53103.faa

1-4B15.faa

31-Lrh5.faa

26-Lr108.faa

47-Lrh28.faa

37-Lrh13.faa

14-E800.faa

32-Lrh6.faa

30-Lrh4.faa

19-L35.faa

18-L34.faa

48-Lrh29.faa

43-Lrh22.faa

55-Lrh45.faa

27-Lr140.faa

29-Lrh1.faa

51-Lrh33.faa

25-Lr073.faa

3-2166.faa

54-Lrh44.faa

58-R19-3.faa

12-CRL1505.faa

57-Lrh47.faa

42-Lrh21.faa

59-R709.faa

39-Lrh15.faa

13-DSM 20021.faa

6-ASCC 3018.faa

22-LOCK908.faa

4-ASCC 290.faa

0 0.0129592259996750 0.02591845199935 0.0388776779990250 0.0518369039987

MYA (CORE_PHYLOGENY)