Pset4

1a.

i)

Log likelihood for topology 1 is -2746.80

Log likelihood for topology 2 is -2923.72

Log likelihood for topology 3 is -2923.72

Topology 1 is the maximum likelihood tree

ii)

Amplify branch length for branch 1-4 by 1000* thus, branch length will be:

 $0.20843*1000,\, 0.03397*1000,\, 0.03497*1000,\, 0.24952*1000,\, 0.00000,$

0.20843*1000, 0.03397*1000, 0.03497*1000, 0.24952*1000, 0.00000),ncol = 5, nrow = 3,byrow = TRUE)

Outcomes of the modified branch length:

Log likelihood for topology 1 is -5140.38

Log likelihood for topology 2 is -5140.38

Log likelihood for topology 3 is -5140.38

The log likelihood of the three trees are exactly the same, making it less likely to distinguish the best tree.

As the branch 5 getting shorter, the difference between the trees are smaller and it is harder to find the maximum likelihood tree.

iii)

Or we can test in some other genes/longer coding region except for APOE. The optimal tree topology should be consistent with the outcomes of APOE.

For the fixed species, we can also test with Sankoff's algrothm by minimizing the base transitions in each node and get the best tree topology.

1b. output:

root:

ATGAAGGTTCTGTGGGCTGCTGCTGGTCACATTCCTGGCAGGATGCCAGGCCGAGGT GGAGCTGGAGA human:

ATGAAGGTTCTGTGGGCTGCGTTGCTGGTCACATTCCTGGCAGGATGCCAGGCCAAGGTGGAGCTGGAGA

mouse:

ATGAAGGCTCTGTGGGCCGTGCTGTTGGTCACATTGCTGACAGGATGCCTAGCCGAGGGA GAGCAGGTGA

rat:

ATGAAGGCTCTGTGGGCCCTGTTGGTCCCATTGCTGACAGGATGCCTGGCCGAGGGA
GAGCAGGTGA

dog:

ATGAAGGTTCTGTGGGCTGCGCTGGTGGTCACGCTCCTGGCAGGATGCTGGGCCGATGTGCAGCCGGAGC

root:

CAGAGCAGACCGAGTGGCAGAGCGGCCAGCCCTGGGAGCTGGCACTGGGCCGCTTCTG
GGATTACCTGCG

human:

CAGAGCAGACCGAGTGGCAGAGCGGCCAGCGCTGGGAACTGGCACTGGGTCGCTTTTGG
GATTACCTGCG

mouse:

CAGATCAGCTCGAGTGGCAAAGCAACCCTGGGAGCAGGCCCTGAACCGCTTCTGG
GATTACCTGCG

rat:

CAGATCAGCTCCCAGGGCAAAGCGACCAACCCTGGGAGCAGGCCCTGAACCGCTTCTGG
GATTACCTGCG

dog:

CGGAGCAGGCCGGGTGGCAGACTGGCCAGCCCTGGGAGGCGGCGCTGGCCCGCTTCTG
GGATTACCTGCG

root:

CTGGGTGCAGACGCTGTCTGACCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCC
AGGAACTGACG

human:

CTGGGTGCAGACACTGTCTGAGCAGGTGCAGGAGGAGCTGCTCAGCTCCCAGGTCACCC
AGGAACTGAGG

mouse:

CTGGGTGCAGACGCTGTCTGACCAGGTCCAGGAAGAGCTGCAGAGCTCCCAAGTCACACAAGAACTGACG

rat:

CTGGGTGCAGACGCTTTCTGACCAGGTCCAGGAAGAGCTGCAGAGCTCCCAAGTCACACAGGAACTGACG

dog:

CTGGGTGCAGACGCTGTCTGACCAGGTGCAAGAGGGGCGTGCTCAACACCCAGGTCACCC
AGGAACTGACG

root:

GCGCTGATGGACGAGACCATGAAGGAGGTGAAGGCCTACAAATCGGAGCTGGAGGAACA GCTGGGCCCGG

human:

GCGCTGATGGACGAGCCATGAAGGAGTTGAAGGCCTACAAATCGGAACTGGAGGAACAA CTGACCCCGG

mouse:

GCACTGATGGAGGACACTATGACGGAAGTAAAGGCTTACAAAAAGGAGCTGGAGGAACAG CTGGGTCCAG

rat:

GTACTGATGGAGGACACTATGACGGAAGTAAAGGCATACAAAAAGGAGCTGGAGGAACAG CTGGGCCCAG dog:

GCGCTGATGGATGAGGCCATGAAGGAGGTGAAGGCCTACAAGGCGGAGCTGGACGAGCA GCTGGGCCCCA

root:

TGGCGGAGGACGCGGGCCCGGCTGCCAAGGAGCTGCAGGCGGCGCAGGCCCGGC
TGGGCGCGGACAT

human:

TGGCGGAGGACGCGGCACGGCTGTCCAAGGAGCTGCAGGCGGCGCAGGCCCGGCTGGCCGGCGCGGACAT

mouse:

TGGCGGAGGAGACACGGCCAGGCTGGGCAAAGAGGTGCAGGCGGCACAGGCCCGACT CGGAGCCGACAT

rat:

TGGCGGAGGACACGGCCAGGCTGGCTAAAGAGGTGCAGGCGGCACAGGCCCGTCT GGGAGCTGACAT

dog:

TGACCTCGGAGACGCCGGCGCGCGTGGCCAAGGAGCTGCAGGCGGCGCAGGCCCGGCTGCGCTCGGACAT

root:

GGAGGACGTGCGAACCGCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTGGGC CAGAGCACCGAG

human:

GGAGGACGTGTGCGCCCTGGTGCAGTACCGCGGCGAGGTGCAGGCCATGCTCGGC CAGAGCACCGAG

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GGAGGATCTACGCAACCGACTCGGGCAGTACCGCAACGAGGTGCACACCATGCTGGGCC AGAGCACAGAG

rat:

GGAGGATCTACGCAACCGACTCGGGCAGTACCGCAACGAGGTAAACACCATGCTGGGCC AGAGCACAGAG

dog:

GGAGGACGTGCGCAACCGCCTGACGCAGTACCGCGGCGAGCTGCAGGCCATGCTGGGC CAGAGCAGCGAG

root:

human:

GAGCTGCGGGTGCGCCTCCCCACCTGCGCAAGCTGCGTAAGCGGCTCCTCCGCGA
TGCCGATGACC

mouse:

GAGATACGGGCGCGCTCTCCACACCCTGCGCAAGATGCGCAAGCGCTTGATGCGGGA
TGCCGAGGATC

rat:

GAGCTGCGGTCGCCCTCTCCACACCCTGCGCAAGATGCGCAAGCGCCTGATGCGGGA
TGCGGATGATC

dog:

GAGCTGCGGGCGCGCTTCGCCTCCCACATGCGCAAGCTGCGCAAGCGGGTGCTGCGGGACCCGAGGACC

root:

human:

mouse:

TGCAGAAGCGCCTAGCTGTACAAGGCAGGGGCACGCGAGGGCGCCGAGCGCGGTGT GAGTGCCATCCG

rat:

TGCAGAAGCGCCTGGCGGTGTACAAGGCCGGGGCACAGGAGGGCGCCGAGCGCGGTGT GAGTGCTATCCG

dog:

TGCAGAGGCGCCTGGCCGTCTACAAGGCCGGCGTGCGCGAGGGTGCCGAGCGCAGCGT GAGCAGCATCCG

root:

human:

CGAGCGCCTGGGGCCCCTGGTGGAACAGGGCCGCGTGCGGGCCGCCACTGTGGGCTCC
CTGGCCGGCCAG

mouse:

rat:

TGAGCGCCTGGGGCCACTGGTGGAGCAGGGTCGTCAGCGCACAGCCAACCTAGGCGCTGGCGCCCCAG

dog:

CGAGCGCCTCTGGCCGCTGCTGGAGCAGGCCCGCGAGCGCAACGCCAAGGTGGGCGCC CTGGCCACGCAG

root:

human:

mouse:

CCTCTGCGCGATCGCGCCCAGGCTTTTGGTGACCGCATCCGAGGGCGGCTGGAGGAAGT GGGCAACCAGG

rat:

CCCCTGCGCGATCGCGCCCAGGCTTTGAGTGACCGCATCCGAGGGCGGCTGGAGGAAGT GGGCAACCAGG

dog:

CCGCTGCTCGAGCGGGCCGACGCCCTGGGCCAGCAGCTGCGCGGGCAGCTGGAGGAGA
TGAGCAGCCGGG

root:

CCCGCGACCGCCTGGAGGAGGTGCGGGAGCAGATGGAGGAGGTGCGCCCAAGATGGA GGAGCAGGCCCA

human:

CCCGCGACCGCCTGGACGAGGTGAAGGAGCAGGTGGCGGAGGTGCGCCCAAGCTGGA GGAGCAGGCCCA

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CCCGTGACCGCCTAGAGGAGGTGCGTGAGCACATGGAGGAGGTGCGCTCCAAGATGGAGGAACAGACCCA

rat:

CCCGAGACCGCCTAGAGGAGGTGCGTGAGCAGATGGAGGAGGTGCGCTCCAAGATGGAGGAGCAGACCCA

dog:

CCCGCGGCCACCTGGAGGAGATGCGCGAGCAGATACAGGAGGTGCGGGTGAAGATGGA GGAGCAGGCCGA

root:

GCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAGAGCTGGTTCGAGCCCC
TGGTGGAAGAC

human:

GCAGATACGCCTGCAGGCCGAGGCCTTCCAGGCCCGCCTCAAGAGCTGGTTCGAGCCCC TGGTGGAAGAC

mouse:

GCAAATACGCCTGCAGGCGGAGATCTTCCAGGCCCGCCTCAAGGGCTGGTTCGAGCCAAT
AGTGGAAGAC

rat:

GCAGATACGCCTGCAGGCCGAGATCTTCCAGGCCCGCATCAAGGGCTGGTTCGAGCCGC TAGTGGAAGAC

dog:

CCAGATACGCCAAAAGGCCGAGGCCTTCCAGGCGCGCCTCAAGAGCTGGTTCGAGCCCC TGCTGGAAGAC root:

ATGCAGCGCCAGTGGGCCGGGCTGGTGGAGAAGGTGCAGGCTGCCGTGGCCACCAGCCCCCCCTGTGC

human:

ATGCAGCGCCAGTGGGCCGGGCTGGTGGAGAAGGTGCAGGCTGCCGTGGGCACCAGCGCCCCCTGTGC

mouse:

ATGCATCGCCAGTGGGCAAACCTGATGGAGAAGATACAGGCCTCTGTGGCTACCAACCCC
ATCCCAGTGG

rat:

ATGCAGCGCCAGTGGGCAAACCTAATGGAGAAGATACAGGCCTCTGTGGCTACCAACTCC ATCACAGTGC

dog:

ATGCAGCGCCAGTGGGACGGCTGGTGGAGAAGGTGCAGGCGGCCGTGGCCACCATCC
CCACCTCTAAGC

root: CCATGGAGAATCAATGA

human: CCAGCGACAATCACTGA mouse: CCCAGGAGAATCAATGA

rat: CCCTGGAGAATCAATGA dog: CTGTGGAGGAACCATGA

1c.

Note: I highlighted the most of the difference by hand...

root:

MKVLWAALLVTFLAGCQA<mark>E</mark>VELETEQTEWQSGQPWELALGRFWDYLRWVQTLS<mark>D</mark>QVQEELL SSQVTQEL<mark>T</mark> human:

MKVLWAALLVTFLAGCQA<mark>K</mark>VELETEQTEWQSGQRWELALGRFWDYLRWVQTLS<mark>E</mark>QVQEELL SSQVTQEL<mark>R</mark>

root:

ALMDETMKEVKAYKSELEEQL<mark>G</mark>PVAEETRARL<mark>A</mark>KELQAAQARLGADMEDV<mark>RN</mark>RLVQYRGEVQ AMLGQSTE

human:

ALMDETMKELKAYKSELEEQL<mark>T</mark>PVAEETRARL<mark>S</mark>KELQAAQARLGADMEDV<mark>CG</mark>RLVQYRGEVQ AMLGQSTE

root:

ELRARLASHLRKLRKRLLRDADDLQKRLAVYKAGAREGAERG<mark>V</mark>SAIRERLGPLVEQGRERTA<mark>N</mark> VG<mark>A</mark>LA<mark>A</mark>Q

human:

ELRVRLASHLRKLRKRLLRDADDLQKRLAVYQAGAREGAERG<mark>L</mark>SAIRERLGPLVEQGRVRAA<mark>T</mark> VG<mark>S</mark>LA<mark>G</mark>Q

root:

PL<mark>H</mark>ERAQA<mark>L</mark>GERLR<mark>G</mark>R<mark>L</mark>EEMGSR<mark>A</mark>RDRL<mark>E</mark>EV<mark>R</mark>EQ<mark>ME</mark>EVRAK<mark>M</mark>EEQAQQIRLQAEAFQARLKS WFEPLVED

human:

PL<mark>Q</mark>ERAQA<mark>W</mark>GERLR<mark>A</mark>R<mark>M</mark>EEMGSR<mark>T</mark>RDRL<mark>D</mark>EV<mark>K</mark>EQ<mark>VA</mark>EVRAK<mark>L</mark>EEQAQQIRLQAEAFQARLK SWFEPLVED root: MQRQWAGLVEKVQAAVATSPTPVPMENQ\$
human: MQRQWAGLVEKVQAAVGTSAAPVPSDNH\$

2a.

[1]

2b.

Use UCSC phyloPng tool: Parameters: Width:300 Height:512 Decimal places:5

