```
Executing file "input.nexus.compute"
  UNIX line termination
  Longest line length = 2472
  Parsing file
  Expecting NEXUS formatted file
   Reading data block
     Allocated taxon set
     Allocated matrix
     Defining new matrix with 10 taxa and 2460 characters
     Data is Dna
     Data matrix is not interleaved
     Gaps coded as -
      Taxon 1 -> SUP35 Kla
      Taxon 2 -> SUP35 Agos
      Taxon 3 -> SUP35_Sc_
      Taxon 4 -> SUP35 Sbou
      Taxon 5 -> SUP35 Sc
      Taxon 6 -> SUP35 Spar
      Taxon 7 -> SUP35 Smik
      Taxon 8 -> SUP35 Sarb
      Taxon 9 -> SUP35 Skud
     Taxon 10 -> SUP35 Seub
     Successfully read matrix
     Setting default partition (does not divide up characters)
     Setting model defaults
     Seed (for generating default start values) = 1571757829
      Setting output file names to "input.nexus.compute.run<i>.<p|t>"
  Exiting data block
  Reading mrbayes block
      Setting autoclose to yes
     Setting nowarnings to yes
     Setting Nst to 6
     Setting Rates to Invgamma
     Setting Nucmodel to 4by4
     Set state frequency prior to default
     Successfully set likelihood model parameters
      Setting number of generations to 100000
      Setting sample frequency to 10
     Running Markov chain
     MCMC stamp = 1376438509
      Seed = 969200195
      Swapseed = 1571757829
     Model settings:
         Data not partitioned --
            Datatype = DNA
            Nucmodel = 4bv4
            Nst
                        Substitution rates, expressed as proportions
                        of the rate sum, have a Dirichlet prior
                        (1.00, 1.00, 1.00, 1.00, 1.00, 1.00)
            Covarion = No
            # States = 4
                        State frequencies have a Dirichlet prior
                        (1.00, 1.00, 1.00, 1.00)
                      = Invgamma
            Rates
                        The distribution is approximated using 4 categories.
                        Likelihood summarized over all rate categories in each
generation.
                        Shape parameter is exponentially
                        distributed with parameter (1.00).
                        Proportion of invariable sites is uniformly dist-
                        ributed on the interval (0.00,1.00).
     Active parameters:
         Parameters
         Revmat
```

```
Statefreq
                      3
  Shape
                      4
  Pinvar
                      5
  Ratemultiplier
                      6
  Topology
                      7
  Brlens
  1 -- Parameter = Revmat
        Type = Rates of reversible rate matrix
        Prior
                  = Dirichlet(1.00,1.00,1.00,1.00,1.00)
        Parameter = Pi
        Type = Stationary state frequencies
        Prior
                  = Dirichlet
        Parameter = Alpha
        Type = Shape of scaled gamma distribution of site rates
        Prior
                 = Exponential(1.00)
        Parameter = Pinvar
        Type = Proportion of invariable sites
        Prior
                 = Uniform(0.00, 1.00)
        Parameter = Ratemultiplier
        Type = Partition-specific rate multiplier
        Prior
                 = Fixed(1.0)
  6 --
        Parameter = Tau
                 = Topology
        Type
                 = All topologies equally probable a priori
        Prior
        Subparam. = V
  7 --
        Parameter = V
        Type = Branch lengths
Prior = Unconstrained:GammaDir(1.0,0.1000,1.0,1.0)
The MCMC sampler will use the following moves:
  With prob. Chain will use move
     0.93 %
              Dirichlet(Revmat)
     0.93 %
              Slider(Revmat)
     0.93 %
              Dirichlet(Pi)
     0.93 %
              Slider(Pi)
     1.85 %
              Multiplier(Alpha)
     1.85 %
              Slider(Pinvar)
     9.26 %
              ExtSPR(Tau,V)
     9.26 %
              ExtTBR(Tau,V)
     9.26 %
              NNI(Tau,V)
     9.26 %
              ParsSPR(Tau,V)
    37.04 %
              Multiplier(V)
    12.96 %
              Nodeslider(V)
     5.56 %
              TLMultiplier(V)
Division 1 has 614 unique site patterns
Initializing conditional likelihoods
Using standard SSE likelihood calculator for division 1 (single-precision)
Initializing invariable-site conditional likelihoods
Initial log likelihoods and log prior probs for run 1:
  Chain 1 -- -13505.482159 -- 36.653401
  Chain 2 -- -12959.497688 -- 36.653401
  Chain 3 -- -13220.200392 -- 36.653401
  Chain 4 -- -13033.906167 -- 36.653401
Initial log likelihoods and log prior probs for run 2:
  Chain 1 -- -13454.486125 -- 36.653401
  Chain 2 -- -13801.708580 -- 36.653401
```

```
Chain 3 -- -13324.366119 -- 36.653401
Chain 4 -- -12408.577175 -- 36.653401
```

Using a relative burnin of 25.0 % for diagnostics

```
Chain results (100000 generations requested):
```

```
0 -- [-13505.482] (-12959.498) (-13220.200) (-13033.906) * [-13454.486]
(-13801.709) (-13324.366) (-12408.577)
      1000 -- (-10169.643) [-10036.903] (-10105.472) (-10174.701) * (-10056.718)
(-10116.461) (-10109.360) [-9998.675] -- 0:03:18
      2000 -- (-9981.771) [-9938.997] (-9963.056) (-10027.612) * (-9851.835)
(-10034.404) (-9931.038) [-9866.335] -- 0:02:27
      3000 -- (-9887.454) (-9858.486) [-9835.588] (-9919.424) * (-9791.591) (-9933.898)
(-9864.404) [-9782.990] -- 0:02:09
      4000 -- (-9847.149) (-9866.943) [-9755.632] (-9820.259) * (-9771.618) (-9863.980)
(-9842.024) [-9764.692] -- 0:02:24
      5000 -- (-9788.358) (-9816.309) [-9753.982] (-9773.840) * [-9750.462] (-9822.029)
(-9763.682) (-9753.890) -- 0:02:32
     Average standard deviation of split frequencies: 0.129448
      6000 -- (-9746.315) (-9804.057) (-9752.041) [-9760.005] * [-9753.191] (-9785.074)
(-9748.291) (-9754.065) -- 0:02:21
      7000 -- (-9744.799) (-9780.068) [-9747.566] (-9745.425) * (-9745.150) [-9756.949]
(-9747.098) (-9748.586) -- 0:02:26
      8000 -- (-9748.895) (-9764.636) (-9745.409) [-9746.802] * (-9744.312) [-9749.340]
(-9747.217) (-9740.613) -- 0:02:29
      9000 -- (-9738.777) (-9754.385) (-9746.663) [-9744.340] * (-9745.948) (-9747.066)
(-9744.465) [-9750.538] -- 0:02:21
      10000 -- (-9748.432) (-9750.635) (-9744.833) [-9746.272] * (-9745.572) (-9747.051)
(-9748.077) [-9745.525] -- 0:02:15
     Average standard deviation of split frequencies: 0.020950
     11000 -- [-9738.878] (-9749.749) (-9742.779) (-9750.032) * (-9747.315) [-9747.482]
(-9748.964) (-9741.283) -- 0:02:17
     12000 -- (-9743.649) (-9756.744) [-9738.781] (-9743.105) * (-9742.062) (-9744.986)
(-9753.372) [-9740.068] -- 0:02:12
     13000 -- (-9740.209) (-9746.054) (-9754.782) [-9741.778] * (-9748.818) (-9745.438)
[-9749.680] (-9751.027) -- 0:02:13
     14000 -- (-9742.763) (-9742.655) [-9744.972] (-9757.903) * (-9748.045) [-9739.307]
(-9746.226) (-9762.098) -- 0:02:09
     15000 -- (-9750.750) [-9753.964] (-9746.635) (-9756.314) * (-9745.194) (-9747.546)
(-9743.303) [-9741.620] -- 0:02:10
     Average standard deviation of split frequencies: 0.018892
     16000 -- [-9741.092] (-9750.283) (-9743.587) (-9750.737) * (-9737.870) (-9747.565)
(-9746.092) [-9745.937] -- 0:02:06
      17000 - [-9744.230] (-9750.143) (-9755.905) (-9751.141) * (-9743.577) (-9748.706)
(-9755.109) [-9744.863] -- 0:02:11
     18000 -- (-9748.253) (-9749.798) [-9743.788] (-9744.202) * (-9747.962) (-9749.314)
(-9745.579) [-9754.935] -- 0:02:21
     19000 -- (-9741.493) (-9751.559) [-9740.865] (-9753.470) * (-9743.693) [-9742.380]
(-9744.798) (-9744.216) -- 0:02:29
     20000 -- [-9744.995] (-9753.312) (-9747.063) (-9739.786) * (-9744.351) [-9748.140]
(-9744.130) (-9741.149) -- 0:02:36
     Average standard deviation of split frequencies: 0.038983
     21000 -- [-9740.074] (-9746.470) (-9743.349) (-9748.883) * [-9746.861] (-9754.067)
(-9751.640) (-9746.388) -- 0:02:45
     22000 -- (-9744.290) (-9744.565) [-9741.835] (-9756.286) * (-9750.986) [-9739.526]
(-9755.131) (-9743.488) -- 0:02:53
     23000 -- (-9744.107) [-9747.558] (-9752.468) (-9761.541) * (-9751.413) [-9750.352]
(-9746.378) (-9745.858) -- 0:02:54
     24000 -- [-9742.942] (-9745.324) (-9750.012) (-9751.290) * (-9753.680) (-9747.034)
```

```
10/22/2019
             www.phylogeny.fr/get result.cgi?task id=d3ac4e07f383653d73d6494b0a0195d0&results in list=9&raw=1&file=...
 (-9756.738) [-9741.258] -- 0:02:51
       25000 -- [-9750.013] (-9745.494) (-9744.562) (-9749.366) * (-9749.780) [-9744.510]
 (-9745.761) (-9756.885) -- 0:02:47
       Average standard deviation of split frequencies: 0.037095
       26000 -- (-9754.533) [-9744.863] (-9742.732) (-9747.753) * (-9747.068) (-9750.543)
 (-9746.222) [-9746.950] -- 0:02:50
       27000 -- (-9744.946) [-9741.405] (-9749.939) (-9751.822) * (-9747.066) (-9739.655)
 (-9749.844) [-9746.684] -- 0:02:53
       28000 -- (-9750.160) (-9742.732) [-9740.951] (-9740.912) * (-9744.585) (-9747.063)
 [-9744.236] (-9749.881) -- 0:02:54
       29000 -- (-9741.272) (-9746.734) [-9741.546] (-9759.673) * (-9742.620) (-9741.399)
 (-9745.178) [-9743.230] -- 0:02:53
       30000 -- (-9744.276) [-9748.688] (-9750.301) (-9750.030) * [-9747.336] (-9747.800)
 (-9749.920) (-9746.957) -- 0:02:50
       Average standard deviation of split frequencies: 0.029005
       31000 -- (-9747.906) [-9744.082] (-9746.397) (-9750.241) * (-9744.622) (-9743.943)
 (-9749.364) [-9745.687] -- 0:02:49
       32000 -- (-9744.502) (-9748.815) [-9744.392] (-9758.713) * (-9745.068) (-9739.127)
 [-9751.696] (-9748.022) -- 0:02:45
       33000 -- (-9743.947) (-9747.094) (-9745.276) [-9753.184] * [-9741.767] (-9744.211)
 (-9742.809) (-9746.801) -- 0:02:42
       34000 -- (-9740.705) [-9746.345] (-9748.576) (-9749.601) * (-9744.711) [-9750.398]
 (-9749.774) (-9745.777) -- 0:02:37
       35000 -- [-9740.497] (-9752.389) (-9746.294) (-9748.158) * (-9751.483) (-9759.155)
 (-9750.306) [-9745.041] -- 0:02:34
       Average standard deviation of split frequencies: 0.022664
       36000 -- (-9742.451) (-9742.726) [-9745.849] (-9753.740) * (-9747.584) [-9744.826]
 (-9756.138) (-9752.411) -- 0:02:31
       37000 -- (-9742.296) [-9744.882] (-9749.795) (-9749.165) * (-9744.524) (-9749.443)
 [-9751.614] (-9739.898) -- 0:02:26
       38000 -- (-9747.463) [-9745.167] (-9744.518) (-9746.000) * (-9744.073) [-9742.325]
 (-9747.270) (-9751.091) -- 0:02:23
       39000 -- [-9741.839] (-9748.788) (-9743.811) (-9741.751) * [-9741.065] (-9740.261)
 (-9745.637) (-9748.800) -- 0:02:20
       40000 -- (-9747.815) [-9754.552] (-9741.467) (-9746.833) * [-9746.834] (-9747.289)
 (-9748.703) (-9747.213) -- 0:02:18
       Average standard deviation of split frequencies: 0.015119
       41000 -- (-9742.803) (-9751.058) (-9752.446) [-9742.898] * (-9745.209) (-9747.421)
 (-9747.073) [-9742.830] -- 0:02:13
       42000 -- (-9743.816) (-9748.807) (-9744.616) [-9740.721] * (-9740.646) (-9749.899)
 (-9749.066) [-9740.271] -- 0:02:11
       43000 -- (-9751.875) [-9739.792] (-9744.413) (-9742.397) * [-9744.658] (-9744.241)
 (-9738.097) (-9743.004) -- 0:02:07
       44000 -- (-9747.471) [-9743.598] (-9741.220) (-9747.244) * (-9743.429) (-9739.708)
 [-9748.289] (-9745.216) -- 0:02:04
       45000 -- (-9740.142) (-9745.187) (-9745.619) [-9743.522] * (-9757.743) (-9738.796)
 [-9742.329] (-9751.342) -- 0:02:01
       Average standard deviation of split frequencies: 0.019392
       46000 -- (-9747.185) [-9747.277] (-9754.475) (-9741.696) * [-9743.877] (-9741.333)
 (-9745.263) (-9744.020) -- 0:01:58
       47000 -- (-9748.589) [-9744.437] (-9751.451) (-9740.499) * (-9757.249) (-9748.074)
 [-9743.045] (-9754.704) -- 0:01:55
       48000 -- (-9741.600) (-9744.106) [-9743.421] (-9748.869) * (-9755.698) (-9748.003)
 (-9742.929) [-9741.145] -- 0:01:51
       49000 -- (-9739.902) [-9741.106] (-9754.492) (-9738.703) * (-9748.738) [-9745.165]
 (-9748.040) (-9740.431) -- 0:01:49
       50000 -- (-9751.856) (-9748.382) (-9752.431) [-9747.175] * (-9746.691) (-9741.153)
 (-9741.380) [-9747.402] -- 0:01:46
```

```
Average standard deviation of split frequencies: 0.017060
```

```
51000 -- (-9741.655) (-9743.602) (-9759.079) [-9741.660] * (-9744.426) (-9745.011)
[-9736.639] (-9750.248) -- 0:01:43
     52000 -- (-9750.170) (-9743.302) (-9748.074) [-9744.759] * (-9748.305) (-9742.016)
(-9743.875) [-9744.882] -- 0:01:41
     53000 -- (-9745.508) (-9747.910) (-9748.560) [-9741.666] * (-9743.735) (-9743.442)
(-9747.778) [-9746.585] -- 0:01:38
     54000 -- (-9752.481) (-9749.354) (-9754.525) [-9741.874] * (-9743.213) (-9748.837)
(-9742.719) [-9746.458] -- 0:01:36
     55000 -- (-9742.724) [-9738.782] (-9751.522) (-9756.618) * (-9741.626) (-9753.921)
(-9749.677) [-9744.025] -- 0:01:33
     Average standard deviation of split frequencies: 0.006784
     56000 -- (-9754.106) [-9743.156] (-9756.682) (-9754.886) * (-9749.134) (-9745.650)
[-9740.666] (-9742.858) -- 0:01:31
     57000 -- (-9747.318) [-9747.984] (-9764.372) (-9744.914) * (-9745.825) (-9742.659)
(-9744.871) [-9746.320] -- 0:01:29
     58000 -- (-9743.079) [-9747.741] (-9742.561) (-9751.893) * [-9742.923] (-9750.317)
(-9747.734) (-9745.528) -- 0:01:26
     59000 -- (-9749.513) [-9753.410] (-9749.605) (-9746.561) * (-9744.996) [-9739.287]
(-9746.199) (-9747.201) -- 0:01:24
     60000 -- [-9743.403] (-9753.506) (-9752.752) (-9748.364) * (-9741.553) (-9752.546)
[-9740.675] (-9745.374) -- 0:01:21
     Average standard deviation of split frequencies: 0.008025
     61000 -- (-9760.771) (-9755.219) [-9738.883] (-9750.287) * [-9747.189] (-9747.655)
(-9749.326) (-9745.279) -- 0:01:19
     62000 -- (-9756.707) (-9760.217) (-9748.471) [-9750.281] * (-9751.345) [-9744.625]
(-9755.921) (-9740.362) -- 0:01:16
     63000 -- [-9747.315] (-9749.720) (-9742.298) (-9743.123) * [-9745.555] (-9744.783)
(-9752.577) (-9744.665) -- 0:01:14
     64000 -- (-9753.829) (-9743.160) [-9743.454] (-9749.682) * (-9749.241) (-9756.019)
(-9737.592) [-9745.990] -- 0:01:12
     65000 -- (-9746.852) (-9745.989) (-9739.283) [-9754.914] * (-9744.583) [-9741.342]
(-9746.836) (-9745.349) -- 0:01:10
     Average standard deviation of split frequencies: 0.007988
     66000 -- (-9746.131) (-9748.890) (-9746.194) [-9741.758] * (-9746.457) (-9741.877)
[-9741.555] (-9762.908) -- 0:01:07
     67000 - [-9745.266] (-9749.473) (-9749.074) (-9748.775) * (-9746.495) (-9742.070)
[-9739.797] (-9758.110) -- 0:01:05
     68000 -- [-9742.051] (-9748.962) (-9750.133) (-9750.611) * (-9747.269) (-9753.447)
[-9739.890] (-9754.100) -- 0:01:03
     69000 -- (-9745.825) (-9744.055) [-9749.832] (-9742.186) * (-9747.517) (-9746.538)
(-9746.226) [-9747.590] -- 0:01:01
     70000 -- [-9751.054] (-9749.476) (-9749.472) (-9741.703) * (-9741.494) (-9737.344)
(-9742.832) [-9741.285] -- 0:00:59
     Average standard deviation of split frequencies: 0.007204
     71000 - (-9753.473) [-9746.342] (-9758.673) (-9744.756) * (-9749.099) (-9741.280)
[-9746.872] (-9744.062) -- 0:00:56
     72000 -- (-9748.775) (-9749.009) (-9750.059) [-9749.817] * (-9752.485) (-9741.199)
(-9746.906) [-9739.639] -- 0:00:54
     73000 -- [-9749.063] (-9751.903) (-9755.155) (-9751.675) * [-9741.180] (-9752.046)
(-9749.585) (-9746.909) -- 0:00:52
     74000 -- (-9736.990) (-9749.607) (-9745.050) [-9743.570] * (-9746.615) (-9745.090)
(-9742.203) [-9747.345] -- 0:00:50
     75000 -- (-9745.218) (-9757.592) [-9744.405] (-9741.304) * [-9747.241] (-9744.261)
(-9753.933) (-9756.062) -- 0:00:48
     Average standard deviation of split frequencies: 0.007793
     76000 -- (-9742.771) (-9745.154) [-9743.892] (-9746.102) * (-9743.312) (-9746.289)
(-9746.126) [-9745.133] -- 0:00:46
```

```
(last 100)
With prob.
                        chain accepted proposals by move
            ( 23 %)
  20.8 %
                        Dirichlet(Revmat)
  49.8 %
             (43%)
                        Slider(Revmat)
   6.3 %
            ( 5 %)
                        Dirichlet(Pi)
            (13%)
  19.4 %
                        Slider(Pi)
  46.3 %
            ( 44 %)
                        Multiplier(Alpha)
  49.4 %
            (50%)
                        Slider(Pinvar)
            (15%)
  16.7 %
                        ExtSPR(Tau,V)
            (2%)
   8.9 %
                        ExtTBR(Tau,V)
            (10%)
  19.0 %
                        NNI(Tau,V)
  22.3 %
            (27%)
                        ParsSPR(Tau,V)
  28.2 %
            (31%)
                        Multiplier(V)
  19.5 %
            (22%)
                        Nodeslider(V)
  12.3 %
            (17%)
                        TLMultiplier(V)
```

Acceptance rates for the moves in the "cold" chain of run 2:

```
With prob. (last 100)
                       chain accepted proposals by move
  24.5 %
            (23%)
                        Dirichlet(Revmat)
  49.3 %
            (42%)
                        Slider(Revmat)
   6.6 %
            (5%)
                        Dirichlet(Pi)
  16.6 %
            (20%)
                        Slider(Pi)
  46.6 %
            (40%)
                        Multiplier(Alpha)
  50.4 %
            (44%)
                        Slider(Pinvar)
  16.2 %
            ( 23 %)
                        ExtSPR(Tau, V)
   9.1 %
            ( 8 %)
                        ExtTBR(Tau, V)
  19.1 %
            (20%)
                        NNI(Tau,V)
  22.4 %
            ( 26 %)
                        ParsSPR(Tau,V)
  28.4 %
             ( 27 %)
                        Multiplier(V)
  19.5 %
             (20%)
                        Nodeslider(V)
  13.1 %
             (18%)
                        TLMultiplier(V)
```

Chain swap information for run 1:

	1	2	3	4
1	 	0.76	0.58	0.43
2	16882		0.79	0.61
3	16634	16788		0.79
4	16667	16373	16656	

Chain swap information for run 2:

1	2 3	3 4
2 16714 3 16580 16	0.79 0.66 0.78 642 782 16565	0.62 0.80

Upper diagonal: Proportion of successful state exchanges between chains Lower diagonal: Number of attempted state exchanges between chains

Chain information:

Using relative burnin ('relburnin=yes'), discarding the first 25 % of sampled trees

Writing statistics to files input.nexus.compute.<parts|tstat|vstat|trprobs|con> Examining first file ...

Found one tree block in file "input.nexus.compute.run1.t" with 10001 trees in last block

Expecting the same number of trees in the last tree block of all files

Tree reading status:

0	10	20	30	40	50	60	70	80	90	100
V	V	V	V	V	V	V	V	V	V	V

Read a total of 20002 trees in 2 files (sampling 15002 of them) (Each file contained 10001 trees of which 7501 were sampled)

General explanation:

In an unrooted tree, a taxon bipartition (split) is specified by removing a branch, thereby dividing the species into those to the left and those to the right of the branch. Here, taxa to one side of the removed branch are denoted '.' and those to the other side are denoted '*'. Specifically, the '.' symbol is used for the taxa on the same side as the outgroup.

In a rooted or clock tree, the tree is rooted using the model and not by reference to an outgroup. Each bipartition therefore corresponds to a clade, that is, a group that includes all the descendants of a particular branch in the tree. Taxa that are included in each clade are denoted using '*', and taxa that are not included are denoted using the '.' symbol.

The output first includes a key to all the bipartitions with frequency larger or equual to (Minpartfreq) in at least one run. Minpartfreq is a parameter to sumt command and currently it is set to 0.10. This is followed by a table with statistics for the informative bipartitions (those including at least two taxa), sorted from highest to lowest probability. For each bipartition, the table gives the number of times the partition or split was observed in all runs (#obs) and the posterior probability of the bipartition (Probab.), which is the same as the split frequency. If several runs are summarized, this is followed by the minimum split frequency (Min(s)), the maximum frequency (Max(s)), and the standard deviation of frequencies (Stddev(s)) across runs. The latter value should approach 0 for all bipartitions as MCMC runs converge.

This is followed by a table summarizing branch lengths, node heights (if a clock model was used) and relaxed clock parameters (if a relaxed clock model was used). The mean, variance, and 95 % credible interval are given for each of these parameters. If several runs are summarized, the potential scale reduction factor (PSRF) is also given; it should approach 1 as runs converge. Node heights will take calibration points into account, if such points were used in the analysis.

Note that Stddev may be unreliable if the partition is not present in all runs (the last column indicates the number of runs that sampled the partition if more than one run is summarized). The PSRF is not calculated at all if the partition is not present in all runs. The PSRF is also sensitive to small sample sizes and it should only be considered a rough guide to convergence since some of the assumptions allowing one to interpret it as a true potential scale reduction factor are violated in MrBayes.

List of taxa in bipartitions:

- 1 -- SUP35_Kla
- 2 -- SUP35 Agos
- 3 -- SUP35 Sc
- 4 -- SUP35_Sbou
- 5 -- SUP35 Sc
- 6 -- SUP35_Spar
- 7 -- SUP35_Smik
- 8 -- SUP35_Sarb 9 -- SUP35 Skud
- 10 -- SUP35 Seub

Key to taxon bipartitions (saved to file "input.nexus.compute.parts"):

```
ID -- Partition
1 -- .******
2 -- .*.....
3 -- ..*.....
4 -- ...*.....
5 -- ....*....
6 -- .....*....
7 -- .....*...
8 -- .....*..
9 -- .....*.
10 -- .....*
11 -- ...*****...
12 -- ..******
13 -- ..****....
14 -- ..***.....
15 -- .....*.*
16 -- ..****.*.
17 -- ..**.....
18 -- ..*.*....
19 -- .....***
20 -- ...**.....
21 -- ..******.
22 -- ..******
```

Summary statistics for informative taxon bipartitions (saved to file "input.nexus.compute.tstat"):

ID	#obs	Probab.	Sd(s)+	Min(s)	Max(s)	Nruns
11	15002	1.000000	0.000000	1.000000	1.000000	2
12	15002	1.000000	0.000000	1.000000	1.000000	2
13	15002	1.000000	0.000000	1.000000	1.000000	2
14	15002	1.000000	0.000000	1.000000	1.000000	2
15	11280	0.751900	0.024321	0.734702	0.769097	2
16	7851	0.523330	0.004242	0.520331	0.526330	2
17	5875	0.391614	0.015743	0.380483	0.402746	2
18	4651	0.310025	0.014046	0.300093	0.319957	2
19	4594	0.306226	0.002451	0.304493	0.307959	2
20	4476	0.298360	0.001697	0.297160	0.299560	2
21	3334	0.222237	0.023567	0.205573	0.238901	2
22	2541	0.169377	0.001791	0.168111	0.170644	2

⁺ Convergence diagnostic (standard deviation of split frequencies) should approach 0.0 as runs converge.

Summary statistics for branch and node parameters (saved to file "input.nexus.compute.vstat"):

Nruns	Parameter	Mean	Variance	Lower	Upper	Median	PSRF+
2	length[1]	0.258096	0.001156	0.198801	0.331625	0.256191	1.000
2	length[2]	0.347219	0.001665	0.267526	0.425785	0.345348	1.001
2	length[3]	0.000923	0.000000	0.000008	0.002189	0.000776	1.000
2	length[4]	0.000951	0.000000	0.000018	0.002280	0.000790	1.000
2	length[5]	0.002373	0.000001	0.000604	0.004635	0.002227	1.000

.0/22/20.	19 www.phyli	ogeny.n/get_resu	it.cgiftask_id=d5	ac4e0/13630330/	/300494DUaU1930	iooresuits_iii_list=	=9&IaW=1&IIIe=
2	length[6]	0.032538	0.000030	0.022481	0.043566	0.032235	1.000
2	length[7]	0.074641	0.000075	0.058317	0.091728	0.074363	1.000
2	length[8]	0.077298	0.000080	0.059876	0.095372	0.076621	1.002
2	length[9]	0.096884	0.000121	0.075469	0.118979	0.096610	1.000
2	length[10]	0.096712	0.000147	0.071170	0.119736	0.096729	1.000
2	length[11]	0.030741	0.000060	0.015423	0.045381	0.030718	1.000
2	length[12]	0.323802	0.001652	0.245816	0.400342	0.322102	1.001
2	length[13]	0.035404	0.000040	0.023280	0.047457	0.035118	1.001
2	length[14]	0.046873	0.000037	0.034764	0.058888	0.046724	1.001
2	length[15]	0.015521	0.000042	0.002187	0.027095	0.015583	1.001
2	length[16]	0.012755	0.000044	0.000213	0.024242	0.012441	1.000
2	length[17]	0.000677	0.000000	0.000000	0.001998	0.000490	1.000
2	length[18]	0.000523	0.000000	0.000000	0.001589	0.000365	1.000
2	length[19]	0.010327	0.000037	0.000152	0.021819	0.009545	1.005
2	length[20]	0.000528	0.000000	0.000000	0.001523	0.000369	1.000
2	length[21]	0.014213	0.000074	0.000005	0.029748	0.013257	1.001
2	length[22]	0.010399	0.000051	0.000002	0.023248	0.009298	1.000
2							

⁺ Convergence diagnostic (PSRF = Potential Scale Reduction Factor; Gelman and Rubin, 1992) should approach 1.0 as runs converge. NA is reported when deviation of parameter values within all runs is 0 or when a parameter value (a branch length, for instance) is not sampled in all runs.

Summary statistics for partitions with frequency >= 0.10 in at least one run:
Average standard deviation of split frequencies = 0.007322
Maximum standard deviation of split frequencies = 0.024321
Average PSRF for parameter values (excluding NA and >10.0) = 1.001
Maximum PSRF for parameter values = 1.005

Clade credibility values:

```
\-----+
                  \----- SUP35 Seub (10)
```

Phylogram (based on average branch lengths):

```
/----- SUP35 Kla (1)
------ SUP35 Agos (2)
                                            / SUP35_Sc__ (3)
                                       /----+ SUP35 Sbou (4)
                                    /----+ \ \ SUP35_Sc (5)
                                     \\----- SUP35_Spar (6)
                                   \\----- SUP35 Smik (7)
                                \\----- SUP35_Skud (9)
                                | /----- SUP35 Sarb (8)
                                 \----- SUP35 Seub (10)
```

|-----| 0.100 expected changes per site

Calculating tree probabilities...

Credible sets of trees (26 trees sampled):

- 50 % credible set contains 5 trees
- 90 % credible set contains 11 trees
- 95 % credible set contains 12 trees
- 99 % credible set contains 14 trees

Exiting mrbayes block Reached end of file

Tasks completed, exiting program because mode is noninteractive To return control to the command line after completion of file processing, set mode to interactive with 'mb -i <filename>' (i is for interactive) or use 'set mode=interactive'