

A user's guide
to
PHYLIP drawtree and drawgram

Introduction

PHYLIP (the PHYLogeny Inference Package) is a package of programs for inferring phylogenies (evolutionary trees). It is available free over the Internet, and written to work on as many different kinds of computer systems as possible.

The installation process is simple:

```
$ sudo apt-get install phylip
```

Once it installed, the package can be accessed in the following way:

```
$ phylip
```

```
Usage: /usr/bin/phylip <program>
```

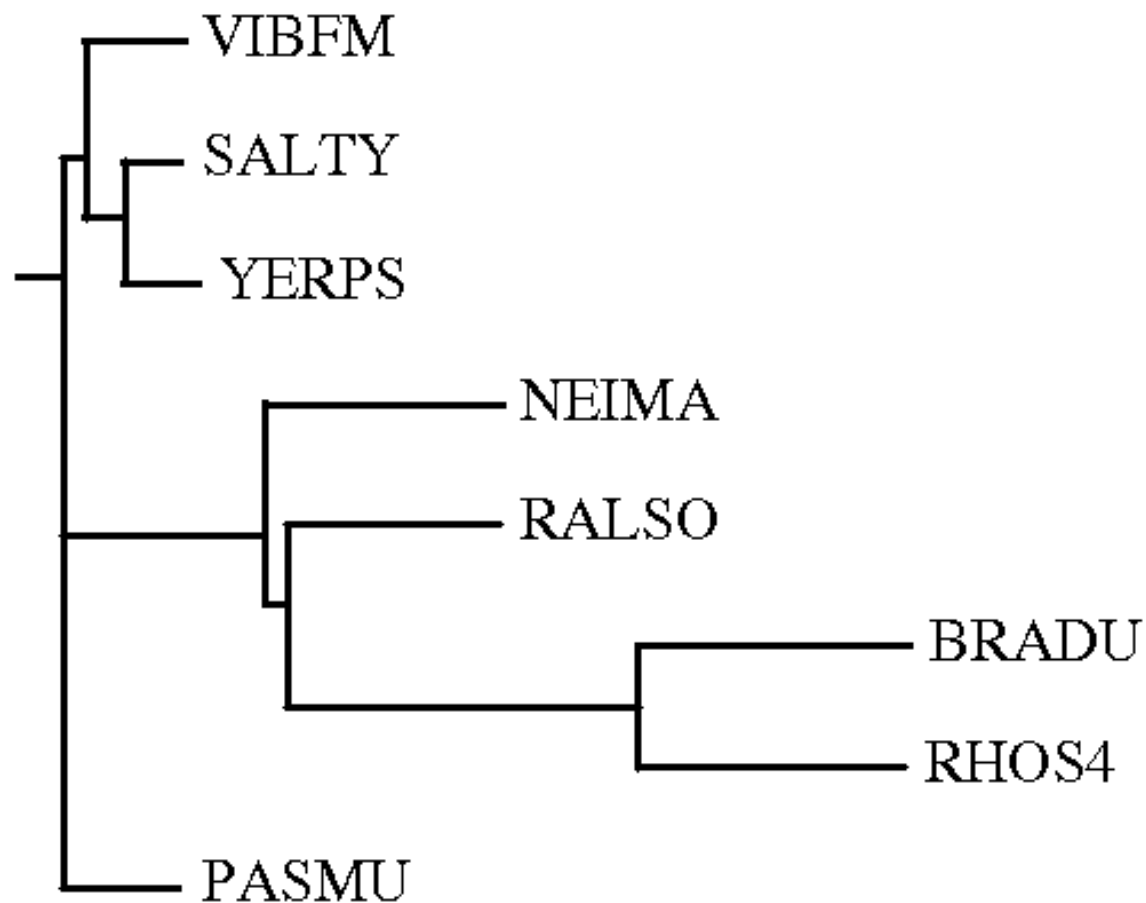
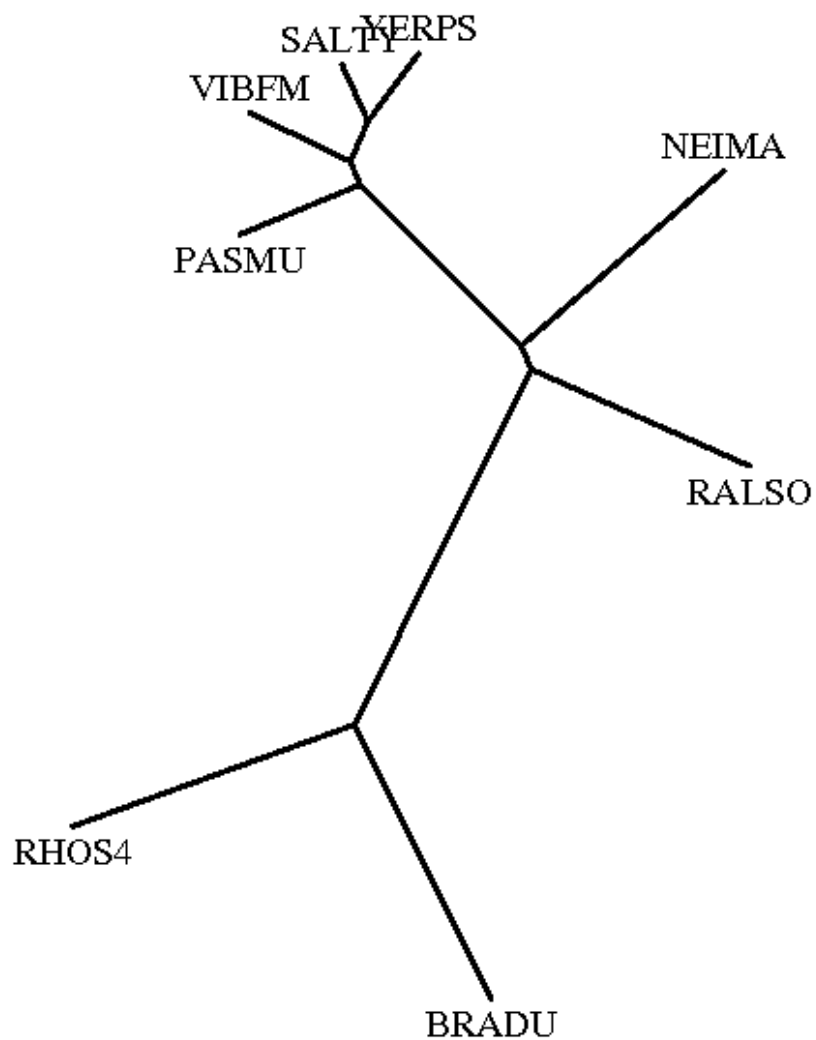
```
Existing programs are:
```

clique	contrast	dnainvar	dnamove	dollop	drawgram	fitch
mix	pars	promlk	restdist	seqboot	consense	dnacomp
dnaml	dnapars	dolmove	drawtree	gendist	move	penny
protdist	restml	treedist	contml	dnadist	dnamlk	dnapenny
dolpenny	factor	kitsch	neighbor	proml	protpars	retree

We are interested in programs drawtree and drawgram.

Trees and trees

drawtree plots **unrooted** trees...



... whereas drawgram plots **rooted** trees.

Part 1. drawtree

drawtree dialog

Being called, the program enters interactive mode and asks to enter a tree in **Newick** format:

```
Please enter a new file name> tree.tre
```

Next come the settings with **default** values:

```
0  Screen type (IBM PC, ANSI)?  ANSI
P      Final plotting device:  Postscript printer
B      Use branch lengths:  Yes
L      Angle of labels:  branch points to Middle of label
R      Rotation of tree:  90.0
I      Iterate to improve tree:  Equal-Daylight algorithm
D  Try to avoid label overlap?  No
S      Scale of branch length:  Automatically rescaled
C  Relative character height:  0.3333
F      Font:  Times-Roman
M      Horizontal margins:  1.65 cm
M      Vertical margins:  2.16 cm
#      Page size submenu:  one page per tree
```

Y to accept these or type the letter for one to change

drawtree settings

The **tree** can be obtained in such a way: **multiple alignment** is put through protdist to calculate **distance matrix** and then through neighbor to make a tree in Newick format.

Basic settings (almost all of the list) for an average user are described in this book.

⓪ Screen type (IBM PC, ANSI)? ANSI

Some settings are disregarded due to their unnecessary of changing from defaults.

P Final plotting device: Postscript printer

Postscript printer returns a .ps image. It can be printed to .png with ps2png program in tth-common package:

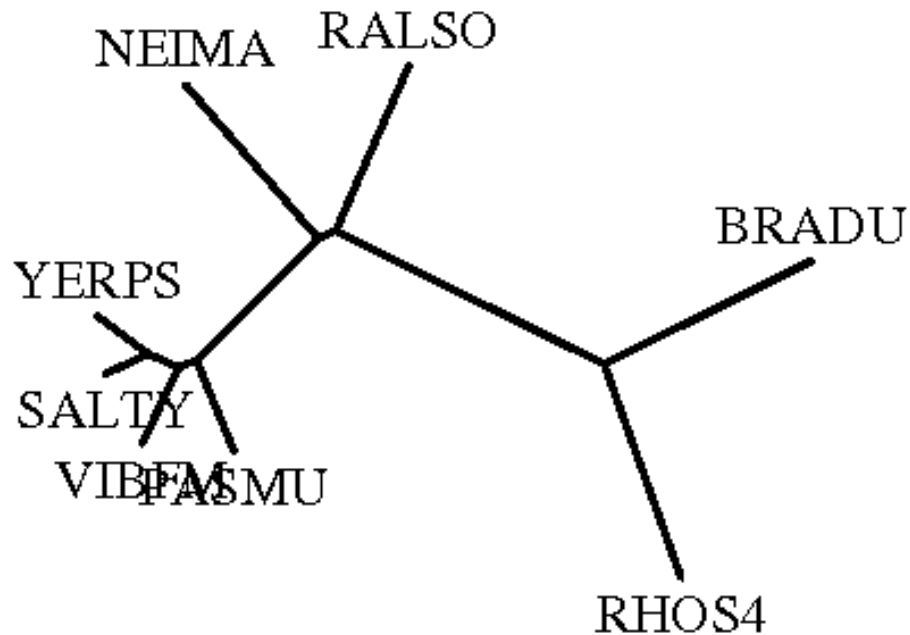
```
$ ps2png image.ps image.png
```

The device can be changed to Windows bitmap so that output images are .bmp files.

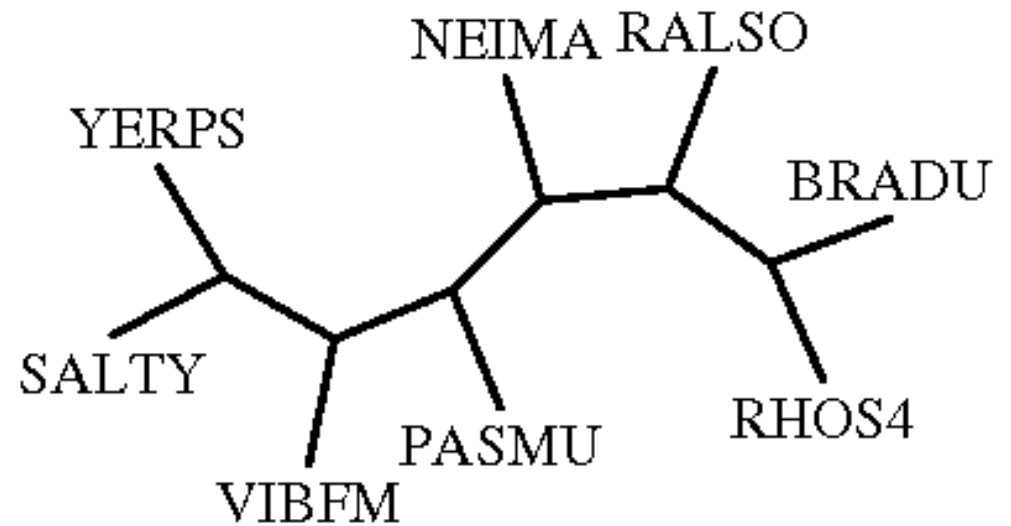
Branch length

B Use branch lengths: Yes

By **default**, branches are calculated based on lengths from Newick formula.



Not using branch length results in equidistant tree.



The equidistant tree looks prettier due to nice layout and may be used in case the length is unnecessary.

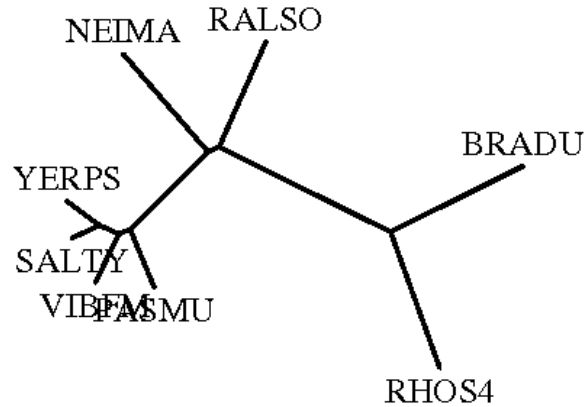
Label positioning

L

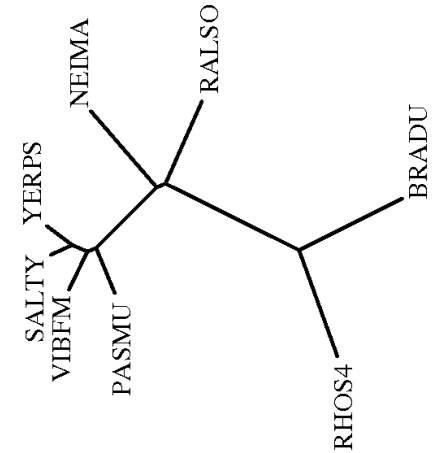
Angle of labels: branch points to Middle of label

Allows to fine-tune label position.

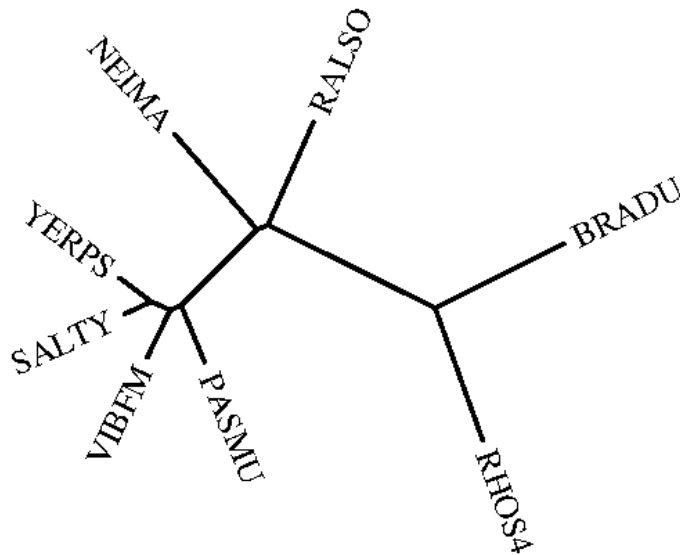
The **default** value points to the **middle** of labels.



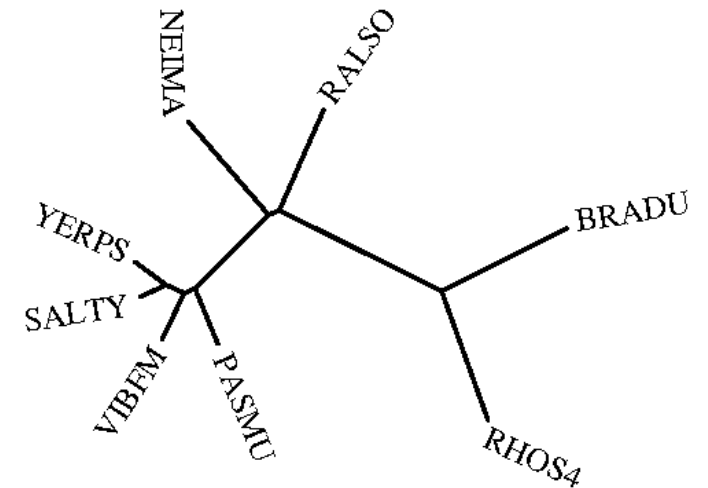
Custom angle can be set, like 90 degrees.



Along placement. Possibly the **best** value of angle.



Radial placement:

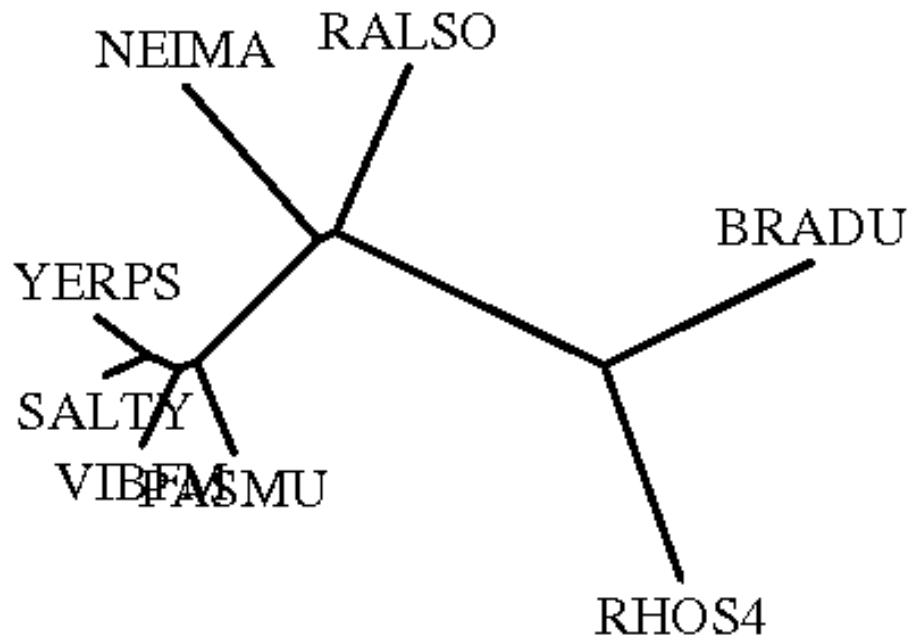


Tree rotation

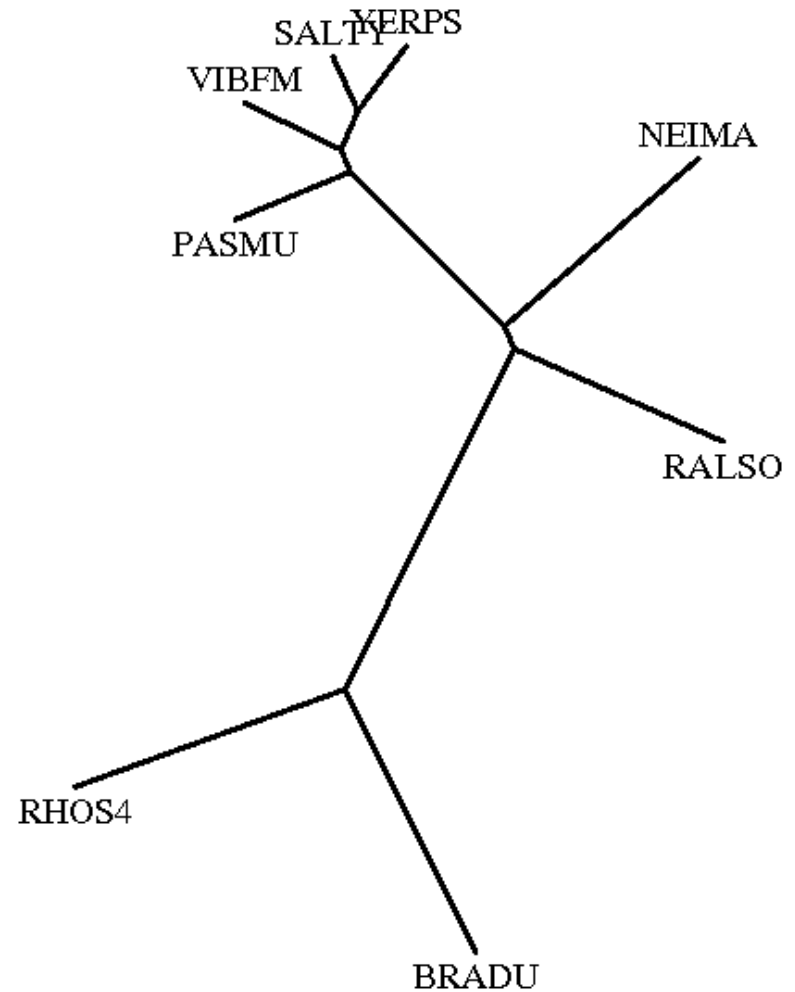
R

Rotation of tree: 90.0

Default, the tree is rotated on **90** degrees.



But you can set **any angle** from -90 to 90. Here is 0.



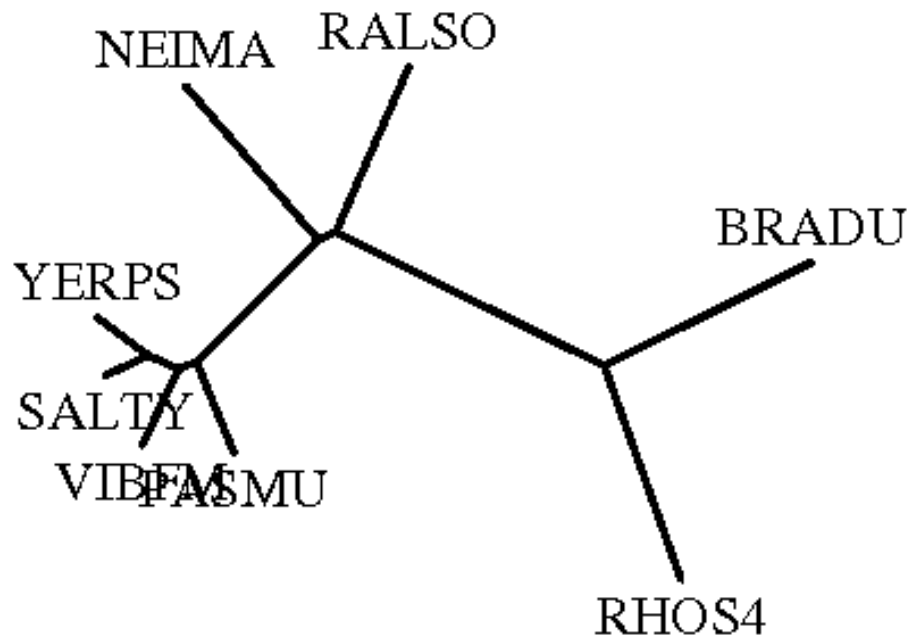
Spin it round.

Tree improvement

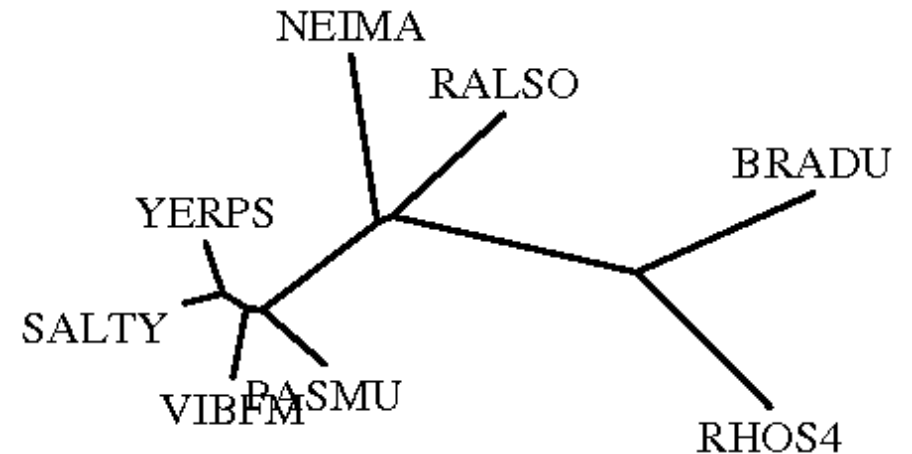
I Iterate to improve tree: Equal-Daylight algorithm

This is used to improve branch angles and prevent lines crossing.

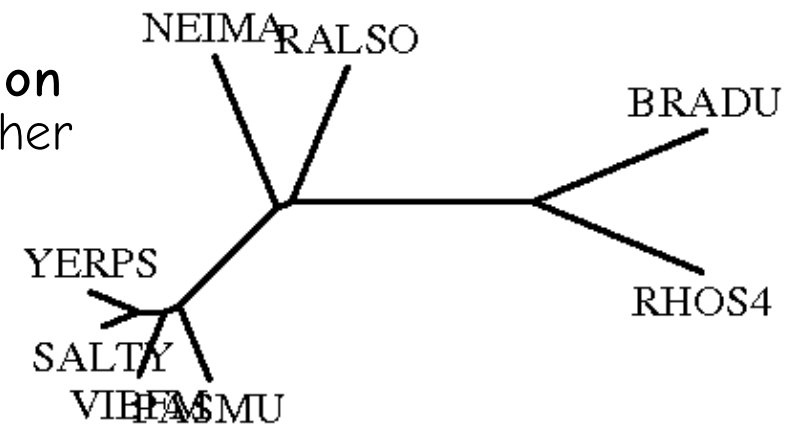
Default is **Equal-Daylight** algorithm.



The **n-Body** looks different.



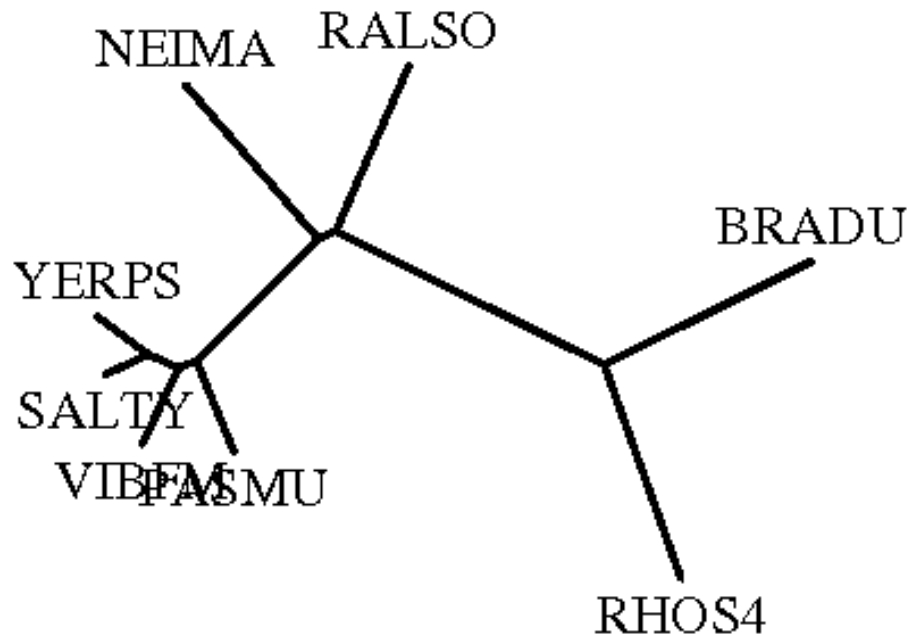
No iteration
uses another
algorithm
from
PLOTREE.



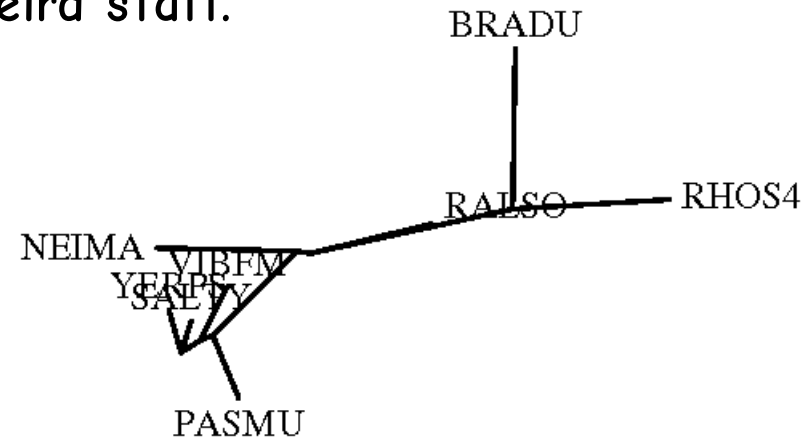
Overlapping labels

D Try to avoid label overlap? No

Default, labels may overlap. But it's almost okay.




When **avoiding**, the program does some weird staff.



Do **not avoid** overlap. Better use positioning of labels.

And also use **autoresizing** of branch length if you want to discern anything.

S Scale of branch length: Automatically rescaled

1 cm per branch unit. 

Other settings

These are better left unchanged.

C	Relative character height:	0.3333
F	Font:	Times-Roman
M	Horizontal margins:	1.65 cm
M	Vertical margins:	2.16 cm
#	Page size submenu:	one page per tree

Part 2. drawgram

drawgram dialog

The program also requires a tree file. Settings list is somehow different from drawtree.

```
0  Screen type (IBM PC, ANSI):  ANSI
P      Final plotting device:  Postscript printer
H      Tree grows:  Horizontally
S      Tree style:  Phenogram
B      Use branch lengths:  Yes
L      Angle of labels:  90.0
R      Scale of branch length:  Automatically rescaled
D      Depth/Breadth of tree:  0.53
T      Stem-length/tree-depth:  0.05
C      Character ht / tip space:  0.3333
A      Ancestral nodes:  Weighted
F      Font:  Times-Roman
M      Horizontal margins:  1.65 cm
M      Vertical margins:  2.16 cm
#      Pages per tree:  one page per tree
```

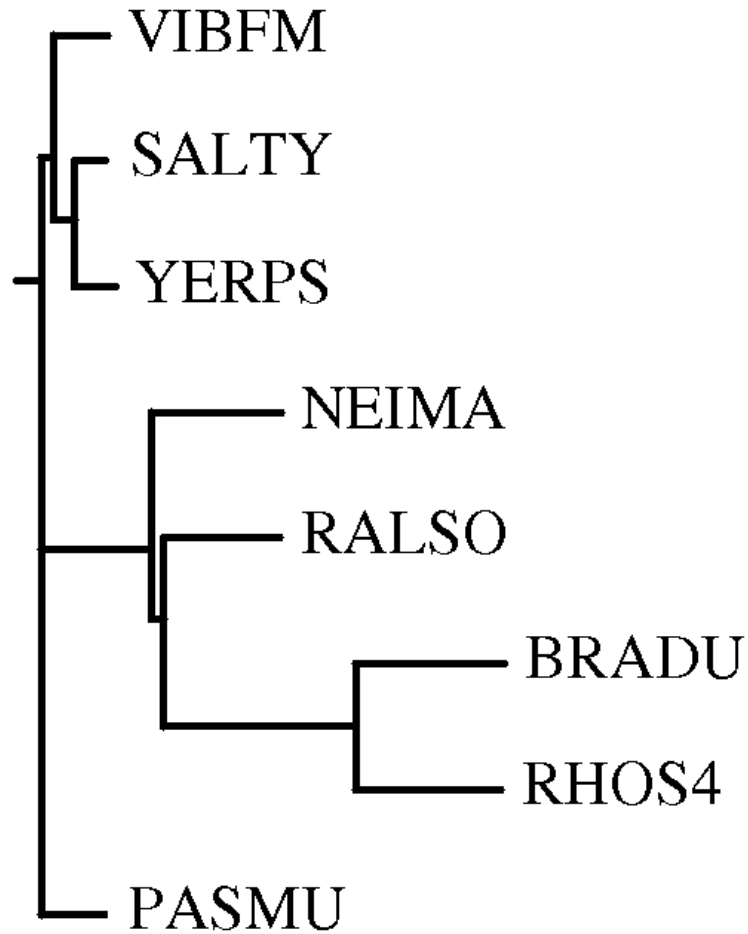
Y to accept these or type the letter for one to change

—

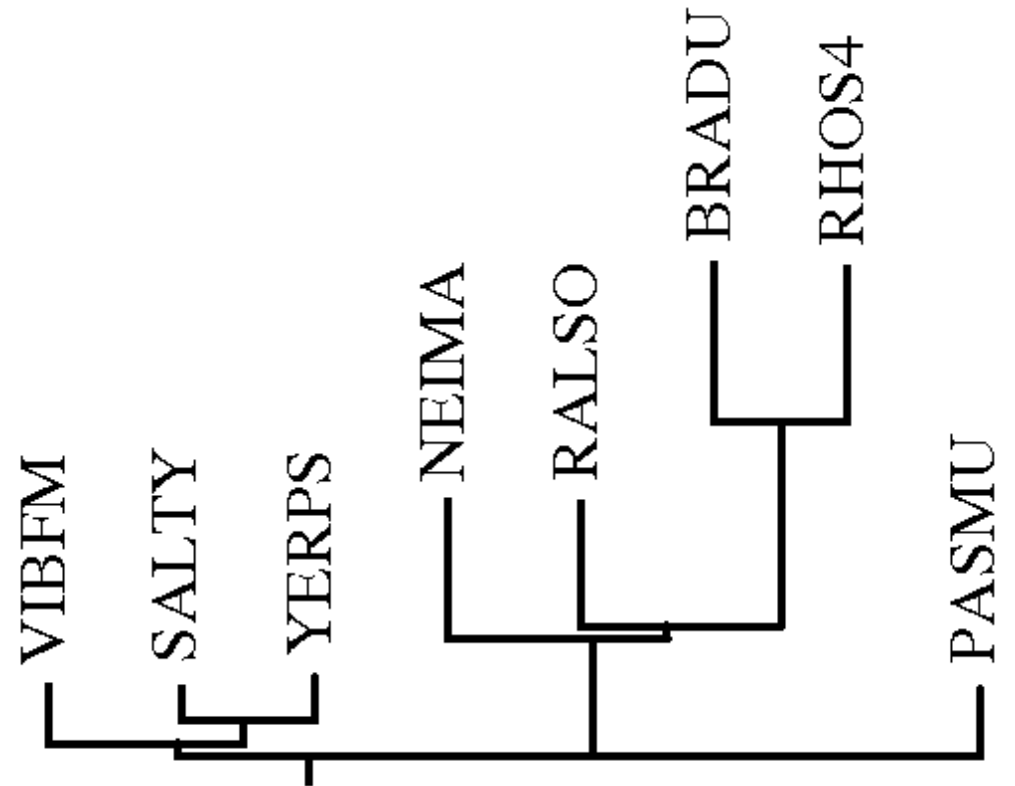
Tree direction

H Tree grows: Horizontally

The tree can grow horizontally.



And vertically.

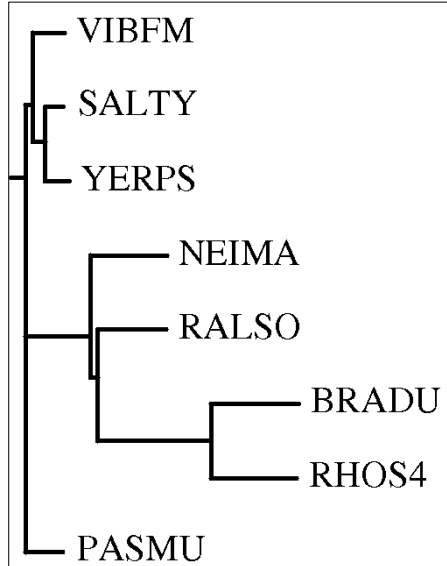


Various styles

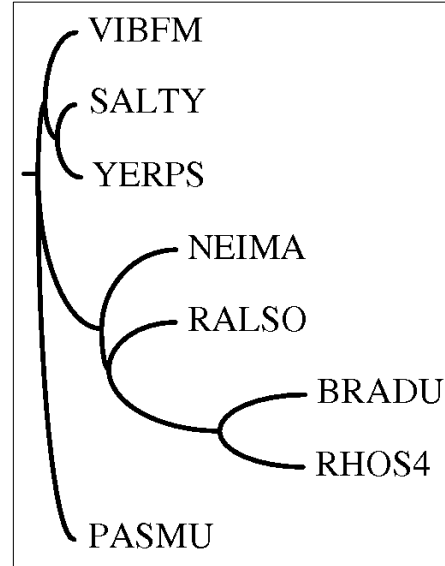
S

Tree style: Phenogram

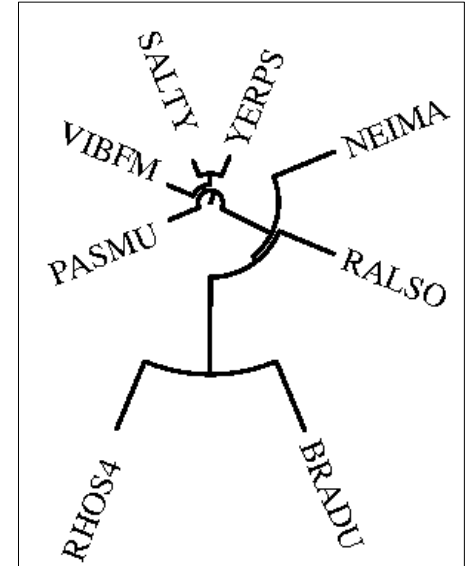
Phenogram



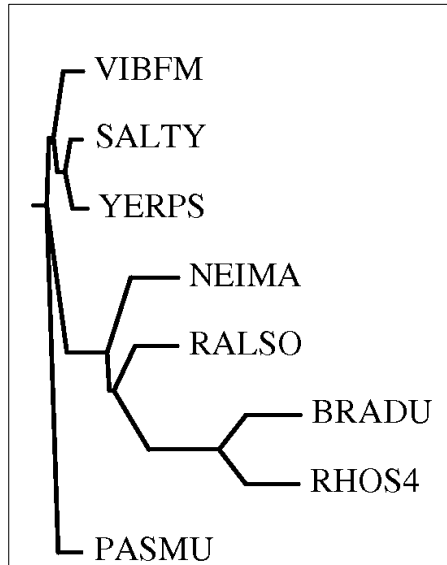
Curvogram



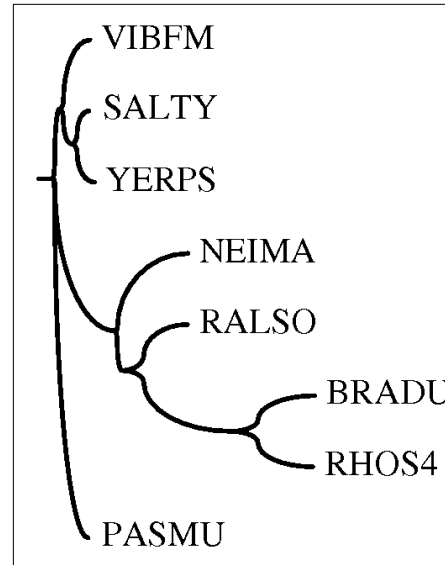
Circular



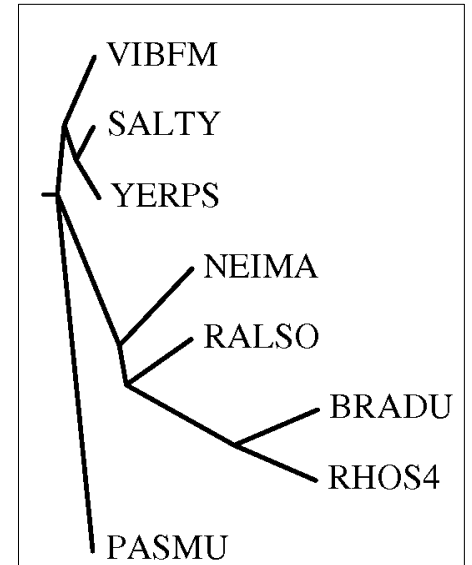
Eurogram



Swoopogram



Cladogram



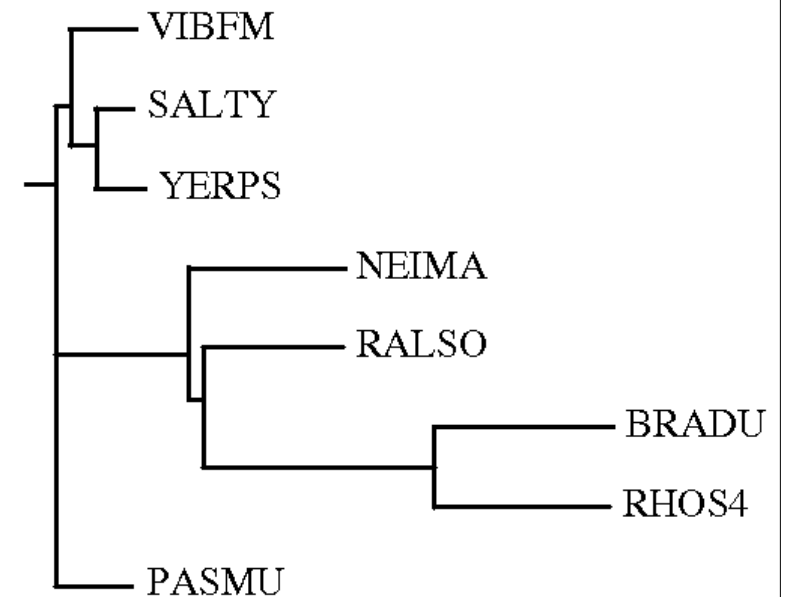
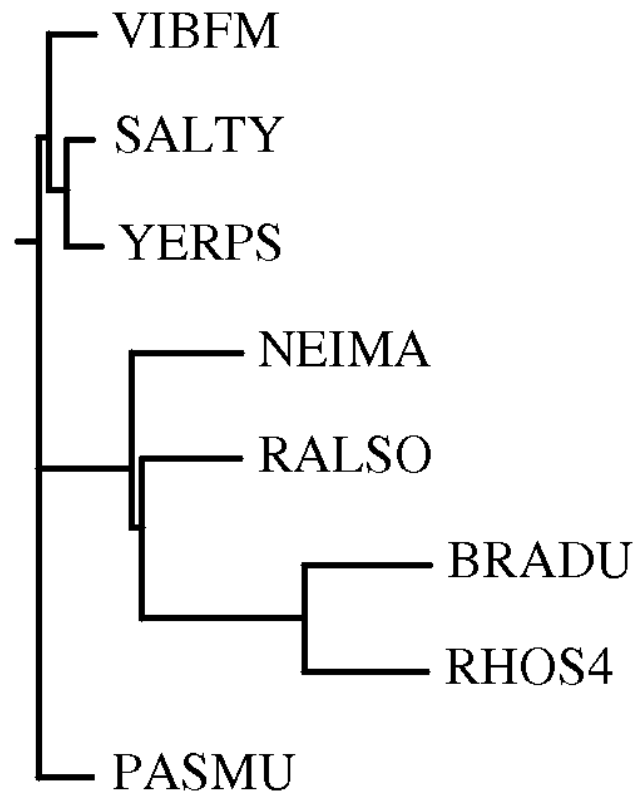
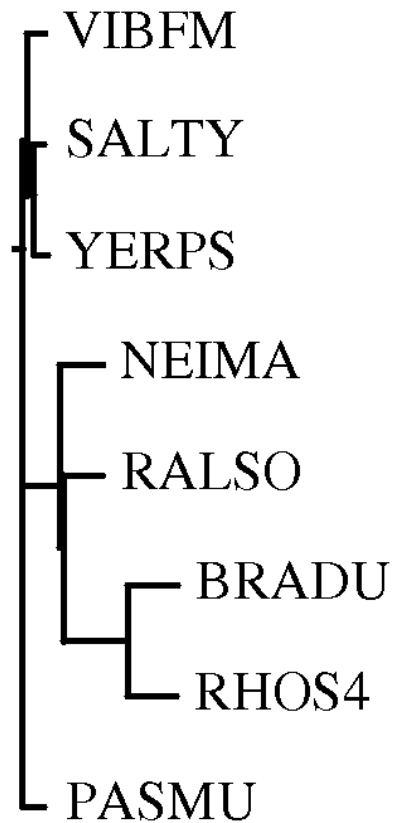
Depth of tree

D Depth/Breadth of tree: 0.53

Less

Default

Greater



The greater variant looks prettier out of them all. But needs other adjustments.

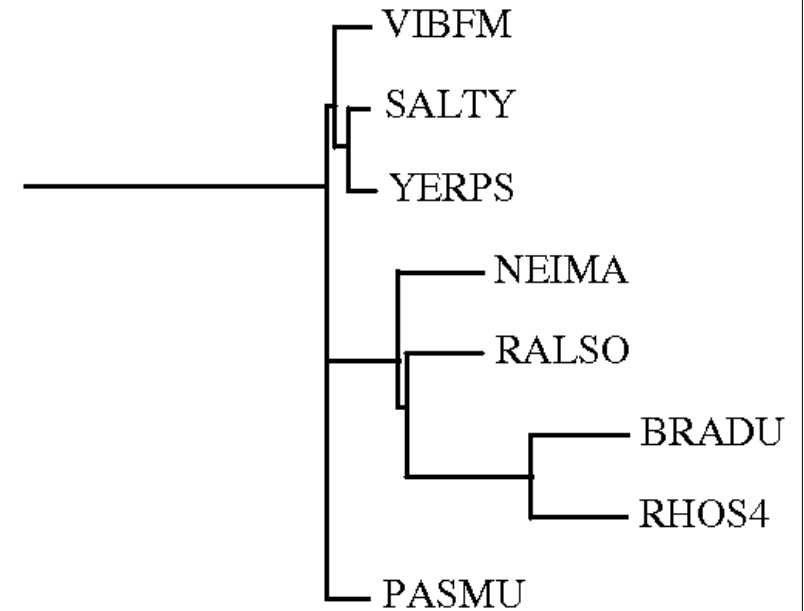
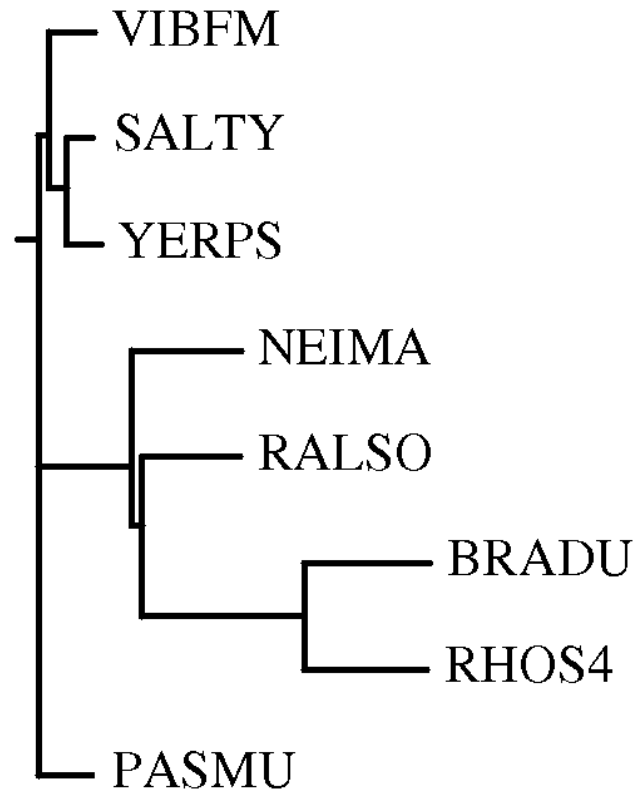
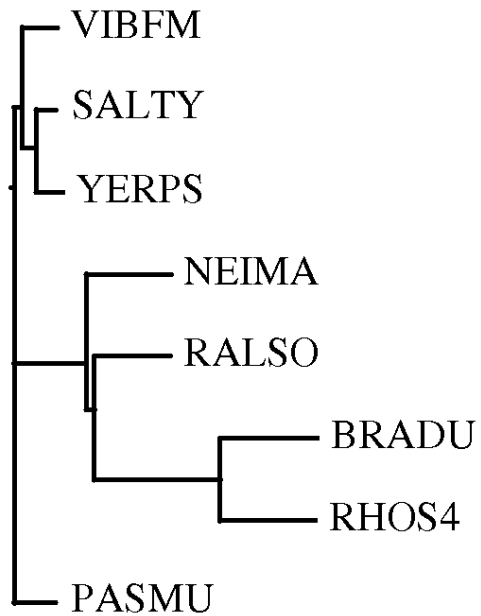
Stem length

T Stem-length/tree-depth: 0.05

Less

Default

Greater



Indeed, root length needs some enlargement.

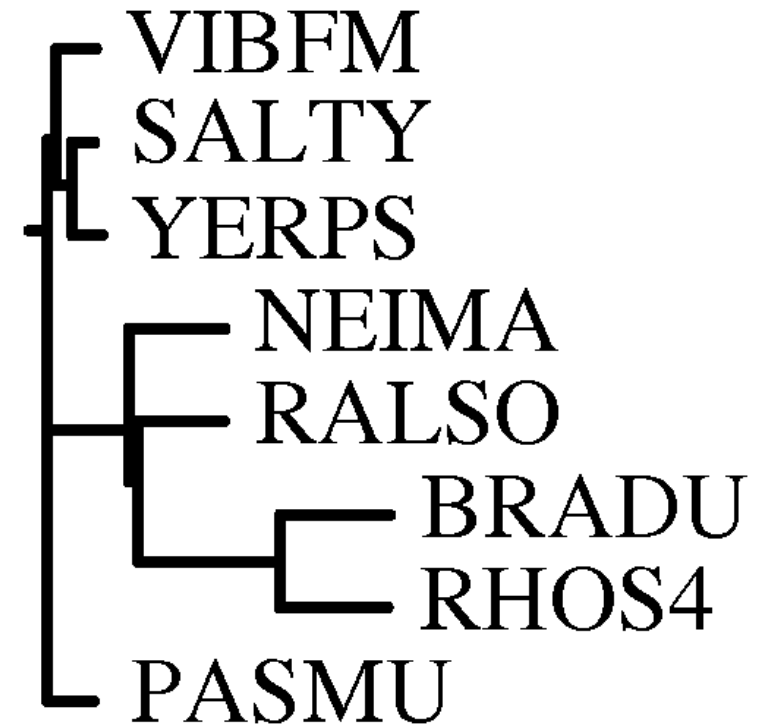
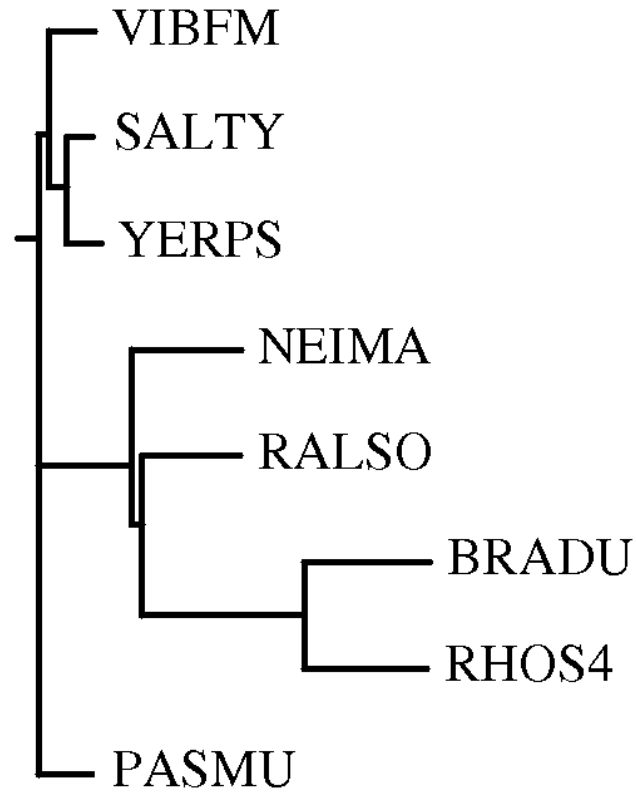
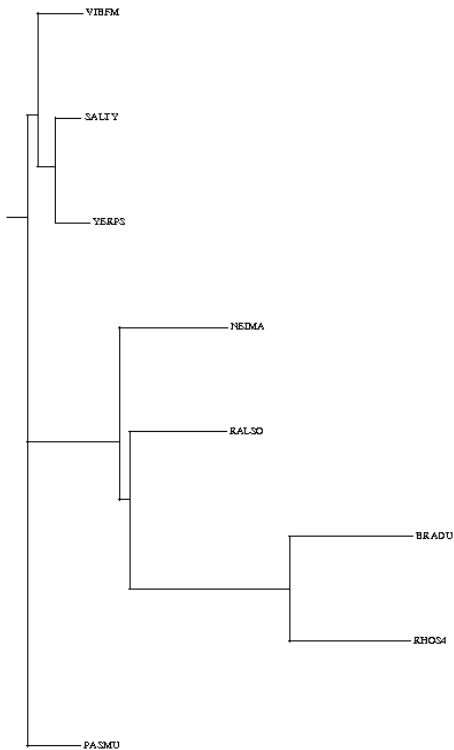
Character size

C Character ht / tip space: 0.3333

Less

Default

Greater



Better left default.

Node positions

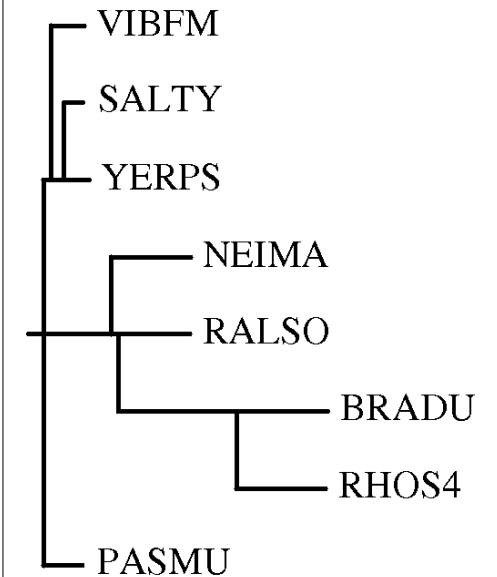
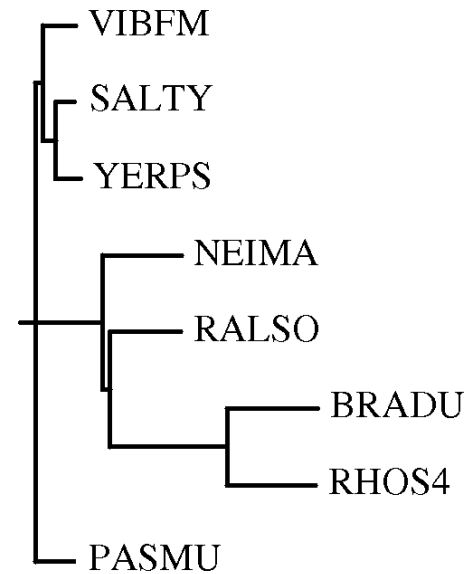
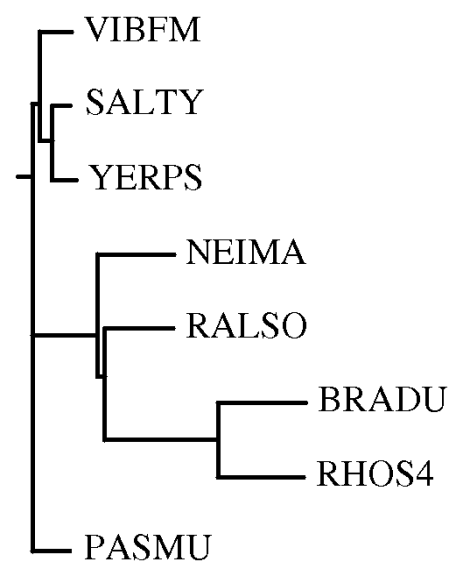
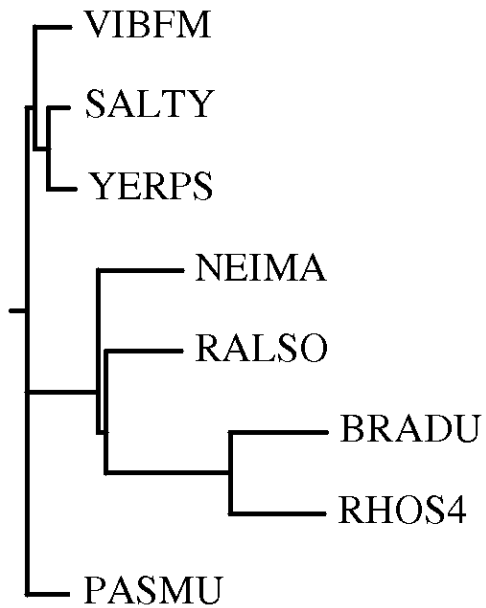
A Ancestral nodes: Weighted

Centered

Weighted

Intermediate

Innermost



Intermediate looks fine but weighted reflects distances between tips.

That's all, folks!

About

Made by Dmitry Mylarshchikov, FBB MSU student.

E-mail: dmitrymyl@gmail.com

Website: kodomo.fbb.msu.ru/~dmitrymyl

Available online on: kodomo.fbb.msu.ru/~dmitrymyl/term04/philip

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