

Phylogenetic tree showing relationships between 16 species based on 16S rDNA. The tree is rooted at the top left. Bootstrap values are indicated at the nodes.

- MARMOSSET (1)
- MACAQUE (12)
- PIG (2)
- COW (7)
- SHEEP (8)
- DOG (4)
- BAT (5)
- HORSE (6)
- MOUSE (3)
- RAT (15)
- TREE_SHREW (16)
- BONOB0 (9)
- CHIMP (10)
- HUMAN (11)
- GORILLA (14)
- ORANGUTAN (13)

```

/----- MARMOSSET (1)
|
|-- MACAQUE (12)
|   |
|   /----- PIG (2)
|   |
|   /---+----- COW (7)
|   | \-----+
|   |         |-- SHEEP (8)
|   |
|   /-----+----- DOG (4)
|   |
|   |----- BAT (5)
|   |
|   \----- HORSE (6)
|
|/-----+----- MOUSE (3)
|       |
|       /-----+----- RAT (15)
|       |
|       \----- TREE_SHREW (16)
|
\+----- BONOBO (9)
    |
    /+- CHIMP (10)
    ||
    /+\- HUMAN (11)
    ||
    \--+-- GORILLA (14)
        |
        \-- ORANGUTAN (13)

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Calculating tree probabilities...

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50 % credible set contains 28 trees
90 % credible set contains 226 trees
95 % credible set contains 301 trees
99 % credible set contains 361 trees
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