

A phylogenetic tree diagram showing the evolutionary relationships between 16 species. The tree is rooted at the top left and branches downwards. Bootstrap values are indicated at the nodes. The species names are listed on the right, with their corresponding numbers in parentheses.

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

Bootstrap values at the nodes:

- Node between (1) and (2): 100
- Node between (3) and (5): 100
- Node between (13) and (14): 100
- Node between (15) and (16): 100
- Node between (4) and (6): 74
- Node between (7) and (8): 100
- Node between (9) and (11): 98
- Node between (10) and (12): 54
- Node between (8) and (9): 100
- Node between (10) and (11): 99
- Node between (6) and (7): 100
- Node between (8) and (10): 100
- Node between (12) and (11): 99

[illegible]

Calculating tree probabilities...

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50 % credible set contains 14 trees
90 % credible set contains 48 trees
95 % credible set contains 61 trees
99 % credible set contains 85 trees
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