

A detailed phylogenetic tree showing relationships between various bat species based on mitochondrial cytb sequences. The tree is rooted at the bottom left with 'don_site' and branches upwards. Species names are listed along the branches, including man, imp, boon, caque, rmoset, lago, use, bbit, bat, rew, madillo, nrec, modelphis, lvyus, icken, nopus, tradon, gu, brafish, and don_site. A scale bar at the bottom indicates genetic distance from 0 to 1.25. Two vertical arrows point to specific nodes in the tree.



