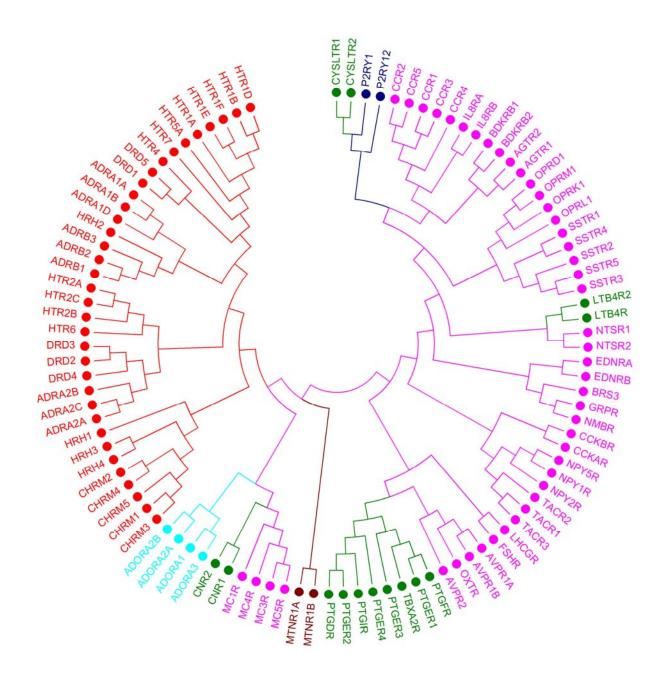
Phylogenetic tree based on the 7TM domain sequence alignment



Phylogenetic tree based on 30 selected residues described in Surgand et al.

