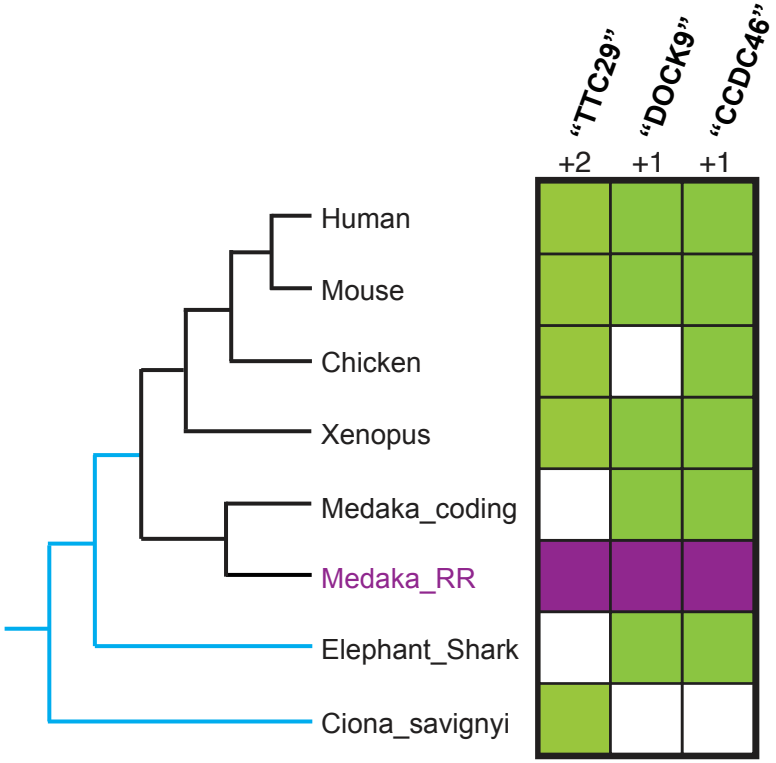


A



Green box = Open reading frame present in the sequence
Purple box = STOP codon present in RF of the sequence
White box = orthologous sequence not found in this species
Blue line = Outgroup species of the tetrapod/ teleost branch

B

