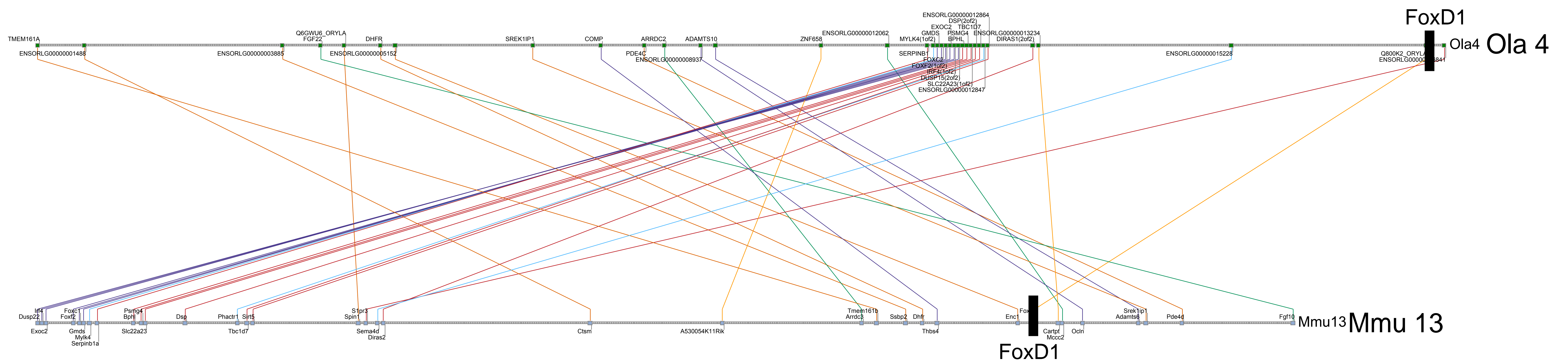
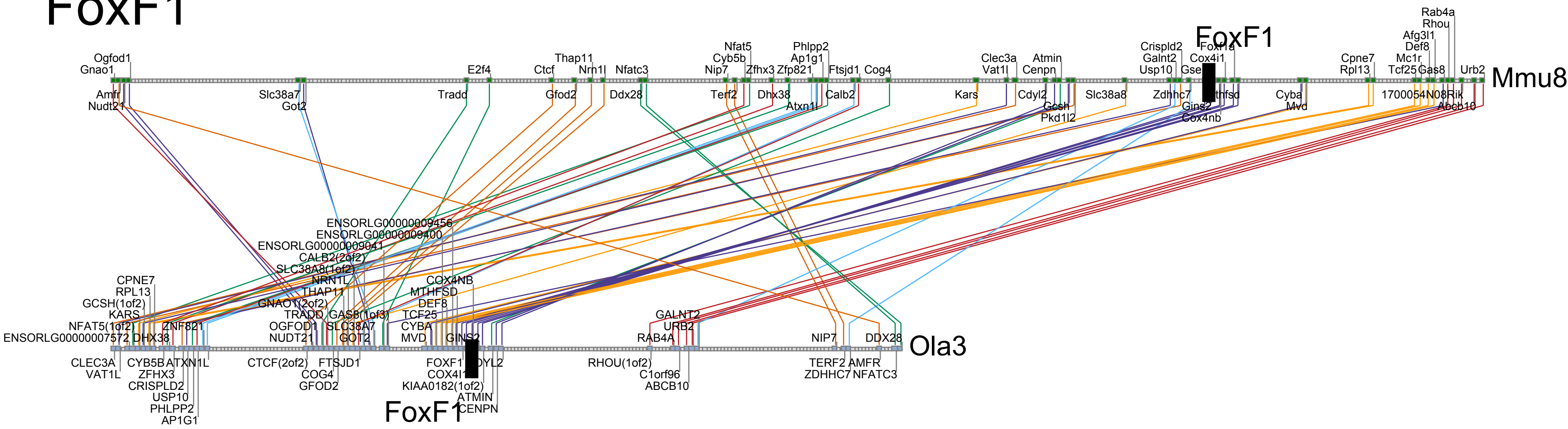


Supplemental Fig.4. Synteny analyses of partial of fox genes were performed using The Synteny Database (Catchen et al. 2009) from mouse and medaka. Generally, 30 Mb around the mouse FOX genes were taken into account (50-gene sliding window). The orthologs location was marked by black bar.

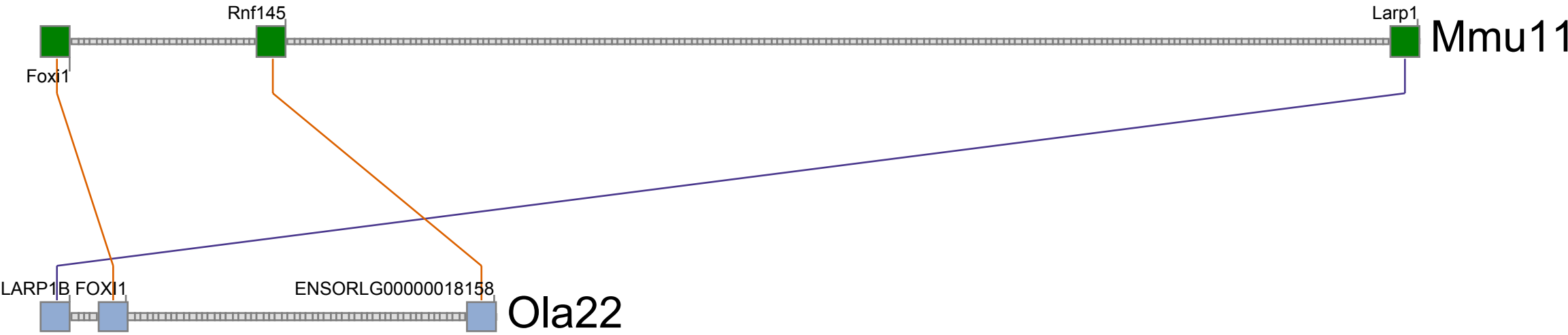
FoxD1



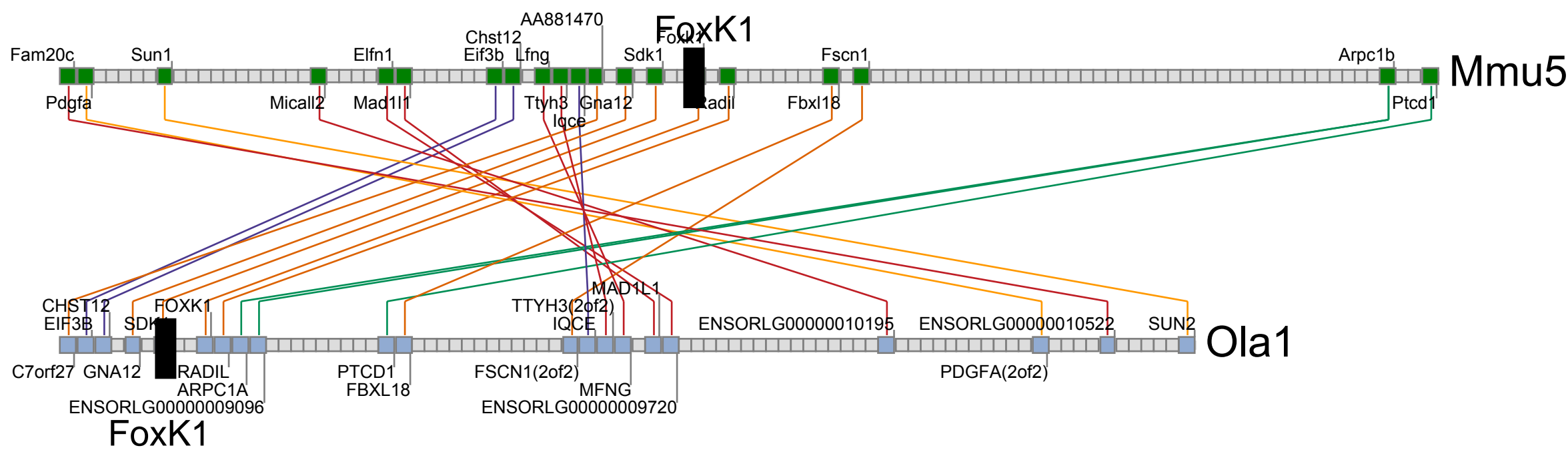
FoxF1



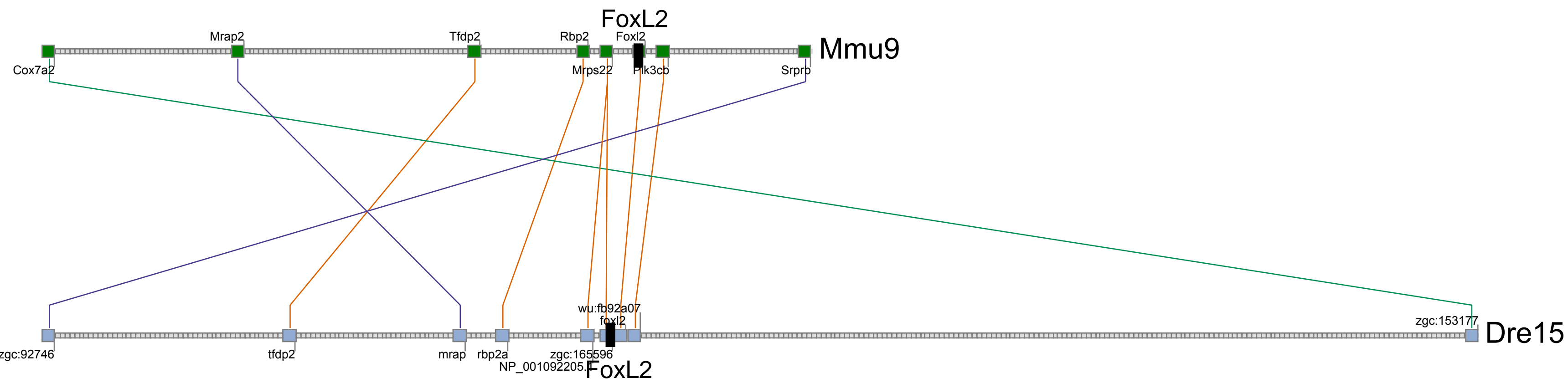
Foxl1



FoxK1



FoxL2

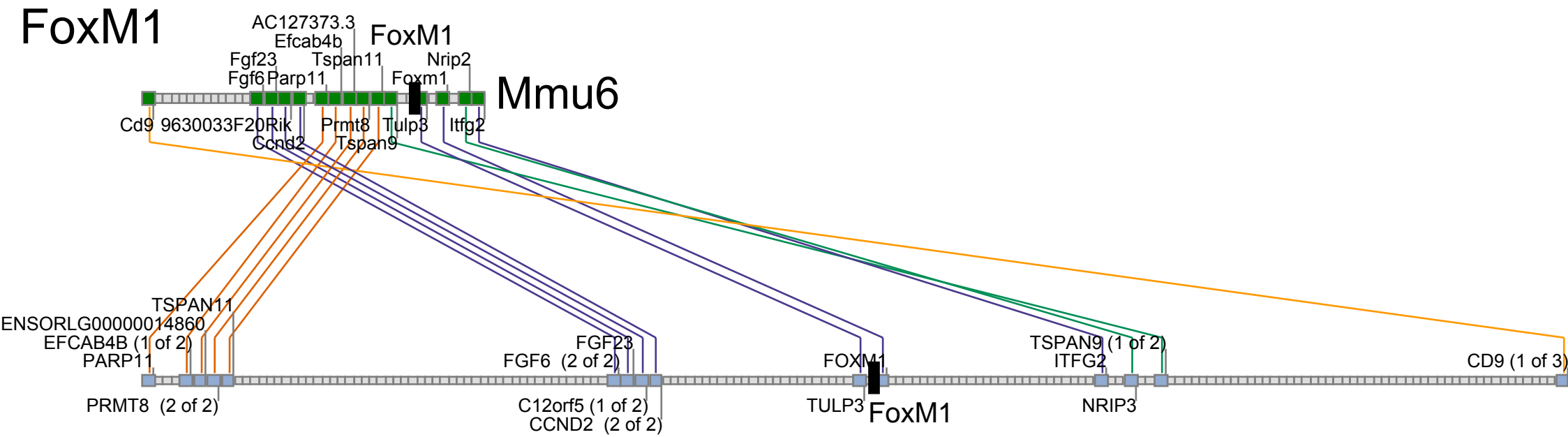


FoxM1

FoxM1

Mmu6

Ola23



FoxP1

