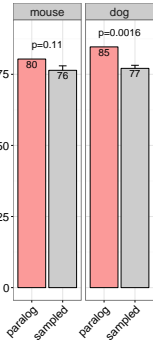
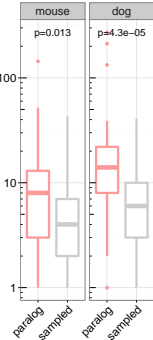


**A**

Ortholog pairs in same TAD [%]

**B**

Hi-C contacts of orthologs

**C**

Pairs conserved in same TAD [%]

