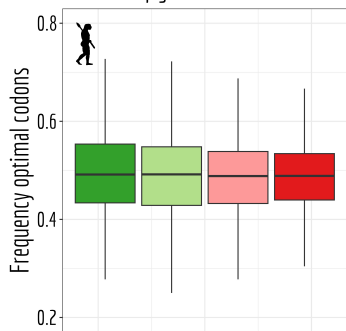
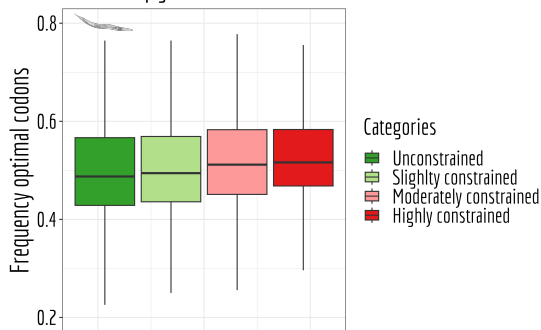


**A**

Number of gene = 766

**B**

Number of gene = 840

**C**

Difference in proportion of optimal codons between the 25% highest and the 25% lowest constrained sites

