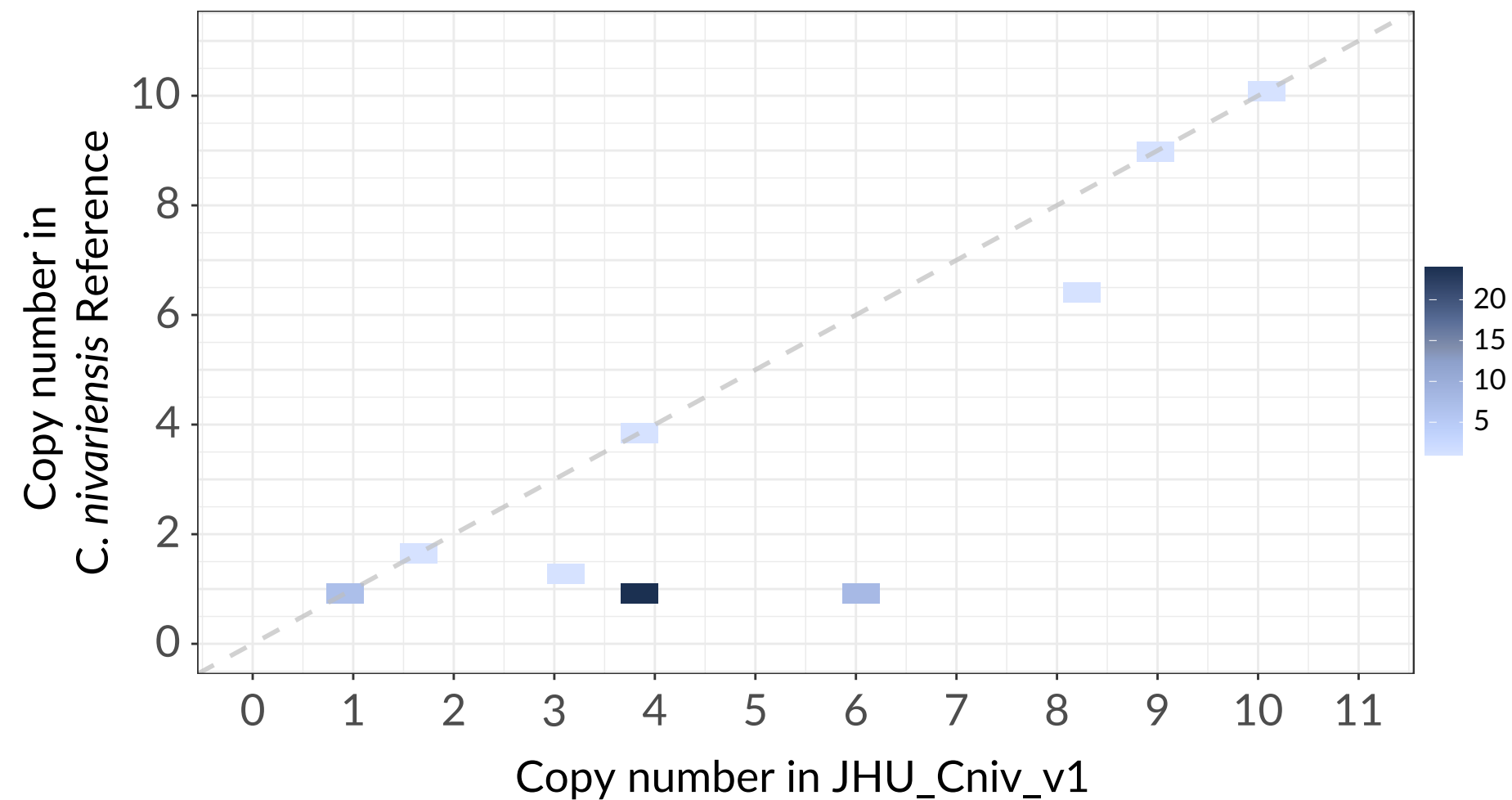
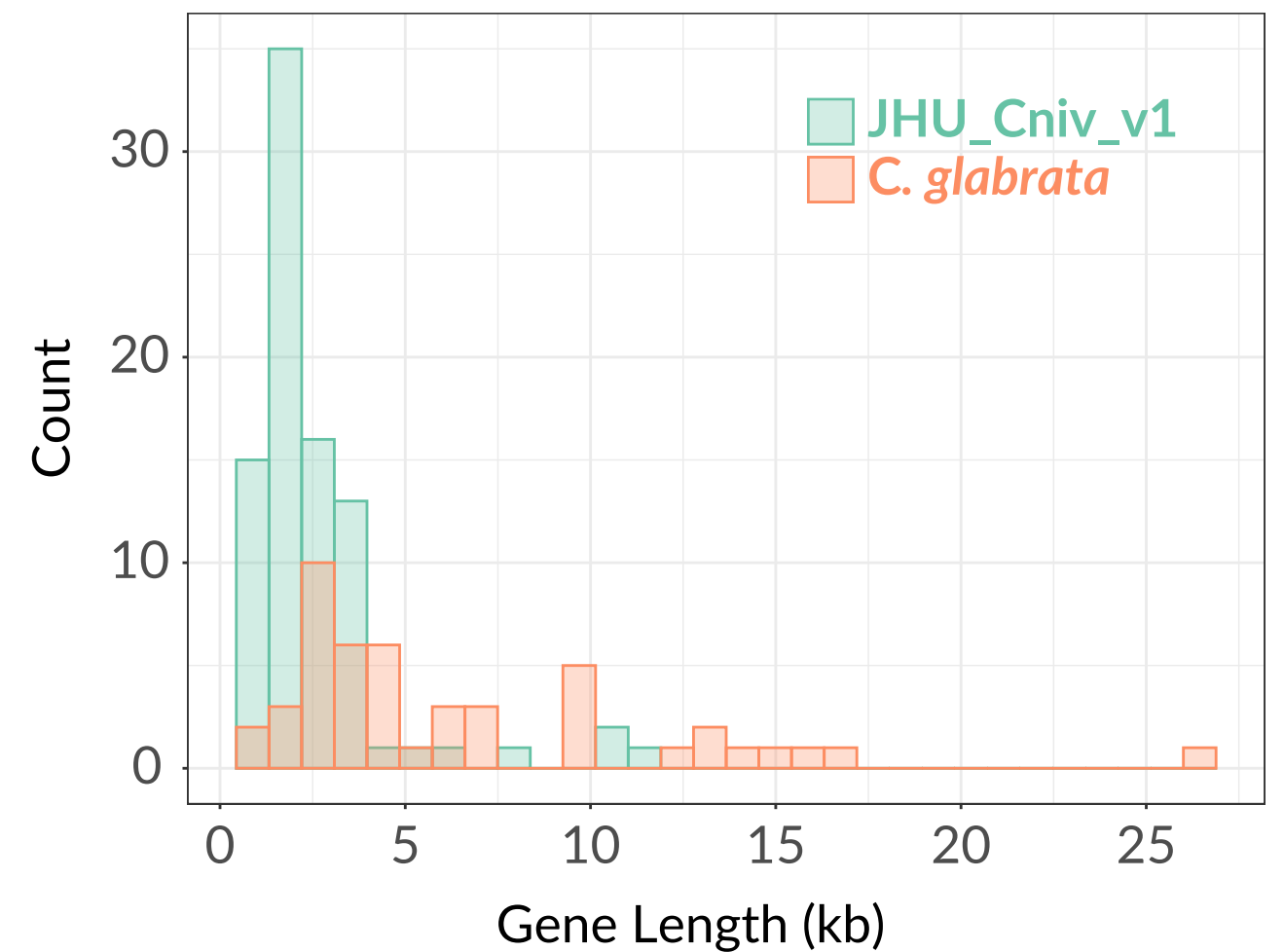


ACopy number of *glabrata* subtelomere gene homologs**B**

Length of predicted GPI-CWPs

**C**

GPI-CWP Max Hit Lengths

