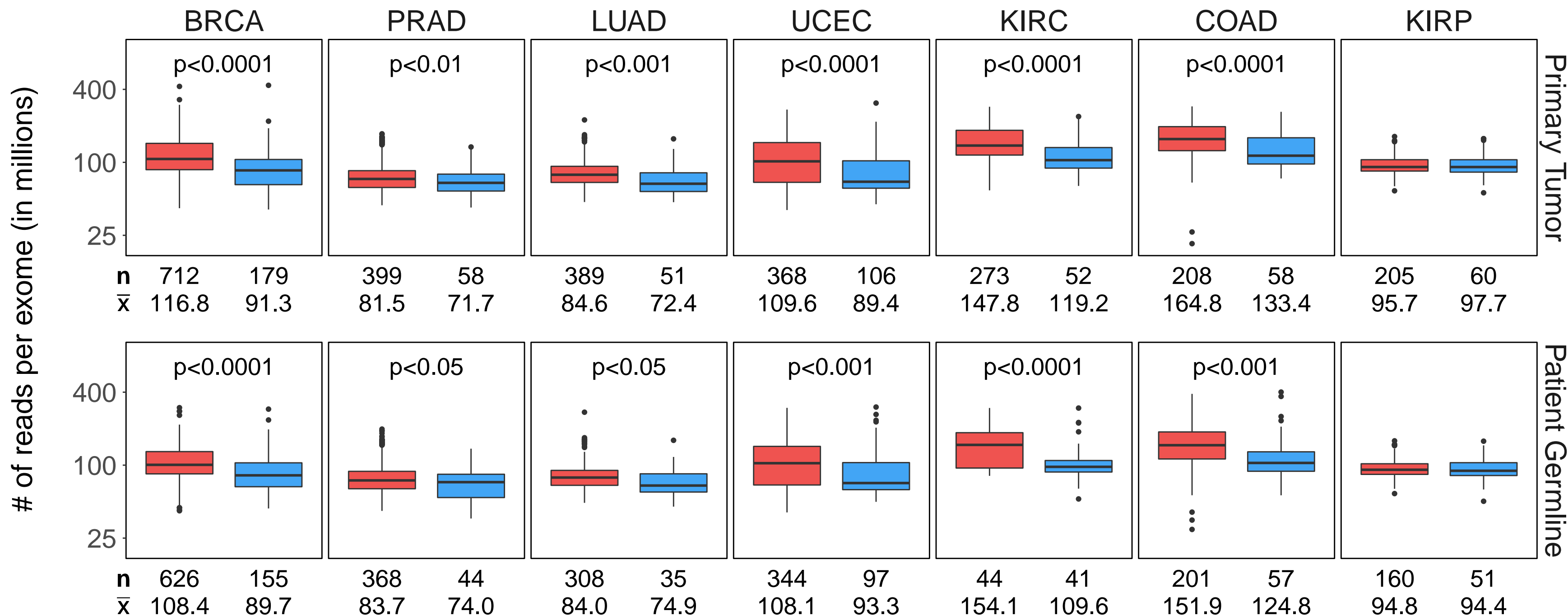
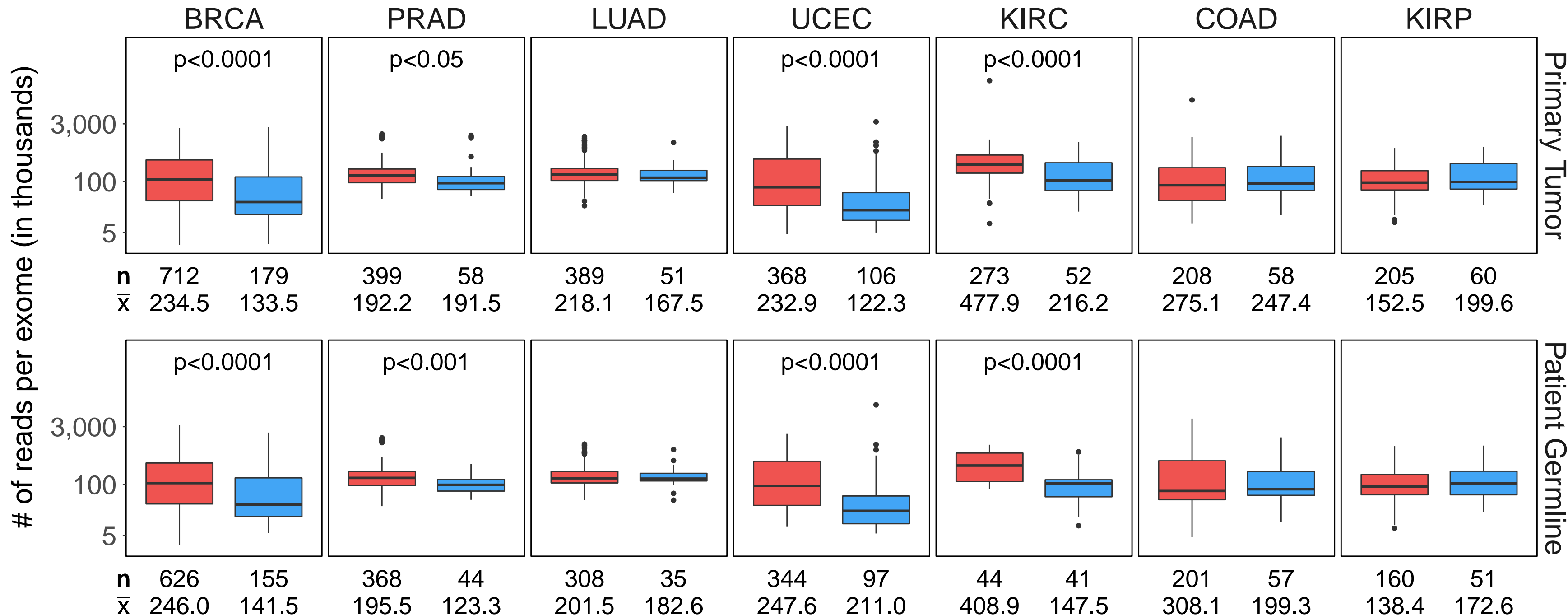


A**Total # of reads per exome****B****# of unmapped reads per exome**

White Black