

IBD comparison of HTG/GoSfSnJ

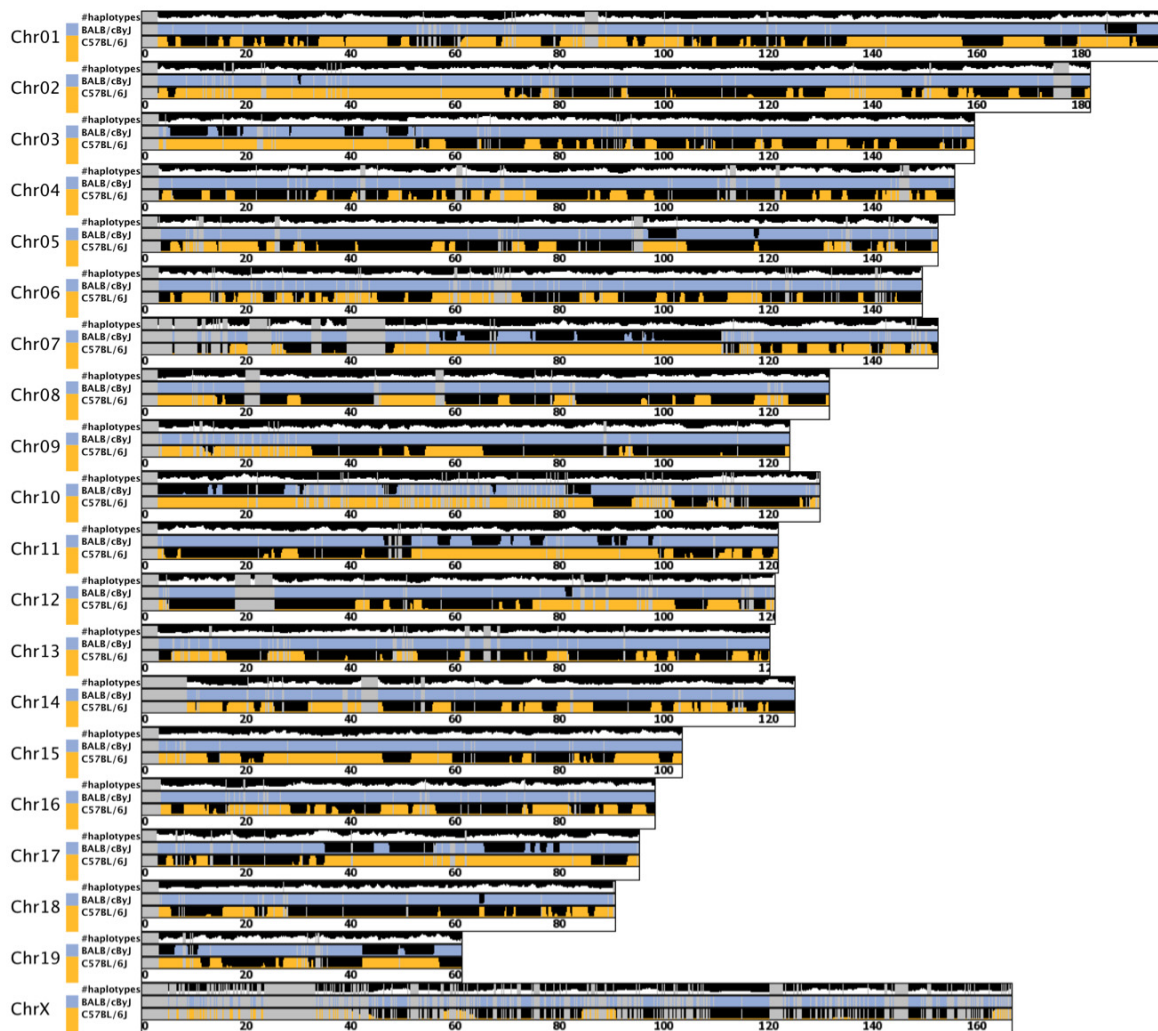


FIGURE S3.—A genome-wide visualization of shared segments between HTG/GoSfSnJ and their predicted ancestors, BALB/cByJ and C57BL/6J, which together covers more than 99.9% of the genome by shared segments. Each colored region represents an IBD region between HTG/GoSfSnJ and either BALB/cByJ or C57BL/6J. A black region represents a region that is not IBD and gray regions represent ambiguous regions.