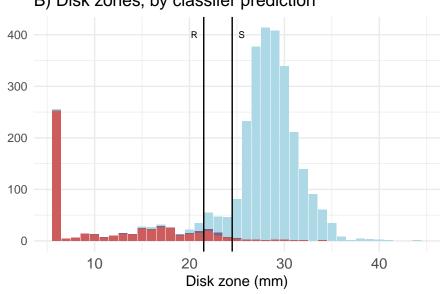


B) Disk zones, by classifer prediction



C) Observed phenotype distribution, per genotype profile

