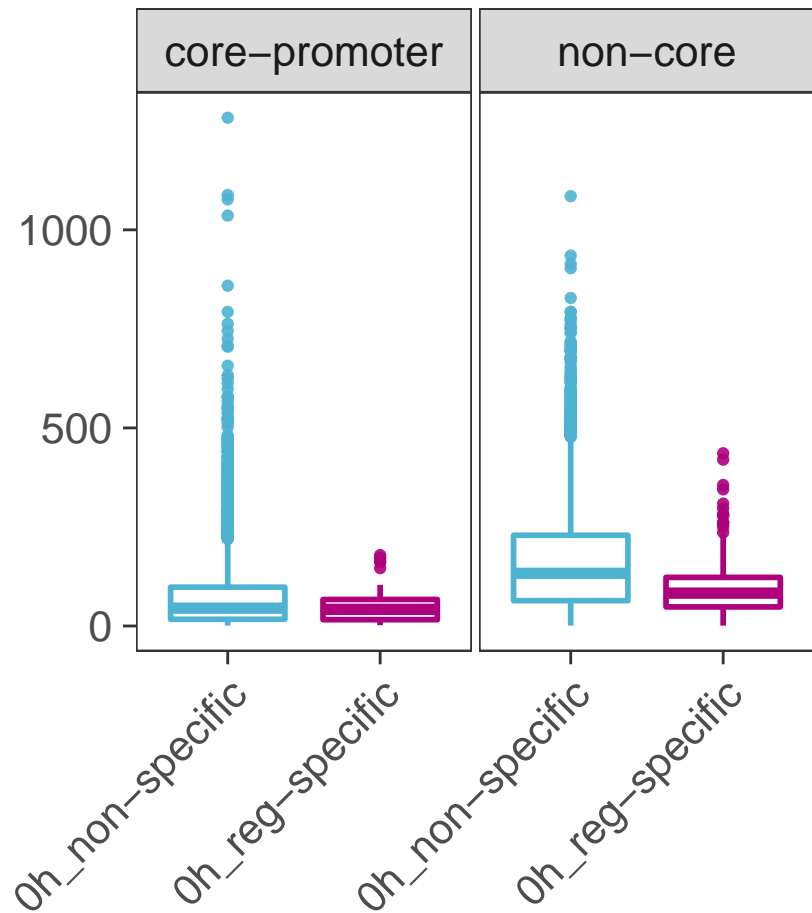


min number of aligned bases



peakGroup

0h_non-specific

0h_reg-specific