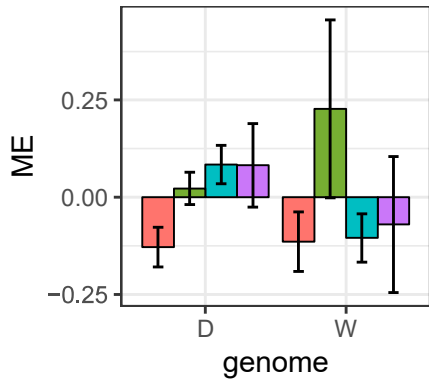
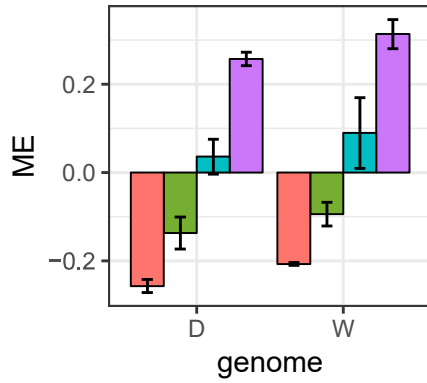


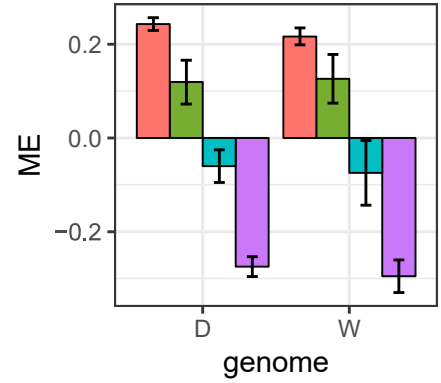
ME0 grey, P=0.3844



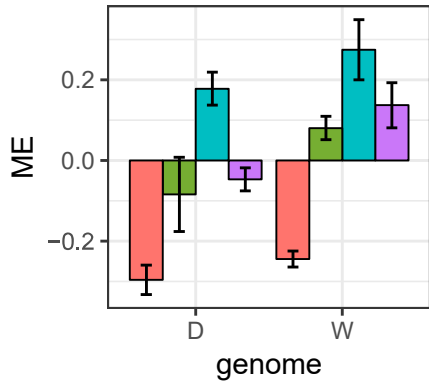
ME1 turquoise, P=0



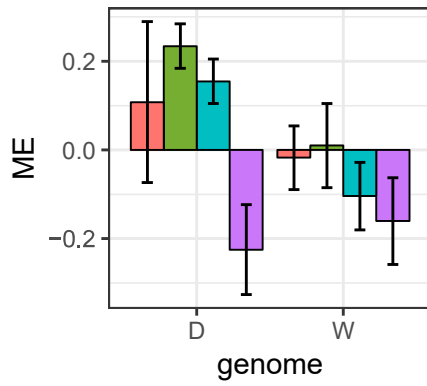
ME2 blue, P=0



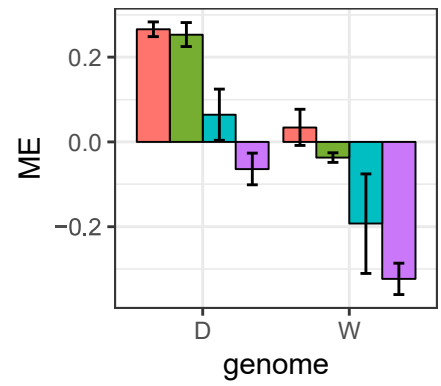
ME3 brown, P=0



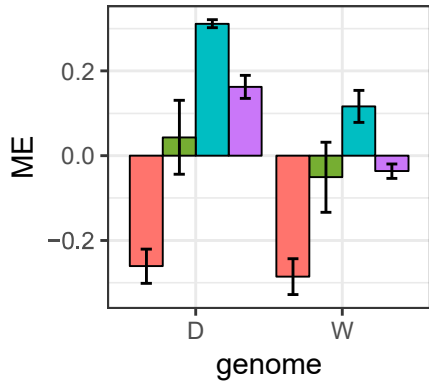
ME4 yellow, P=0.0527



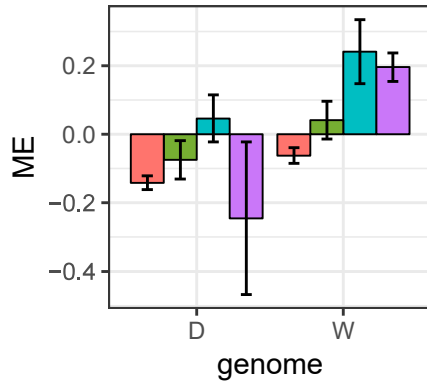
ME5 green, P=0



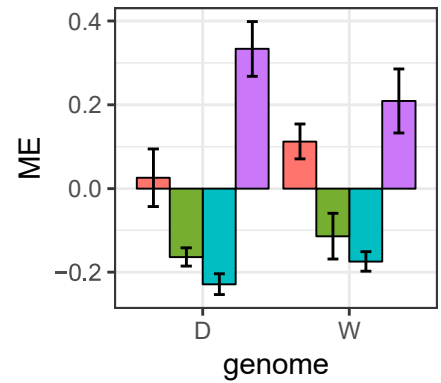
ME6 red, P=0

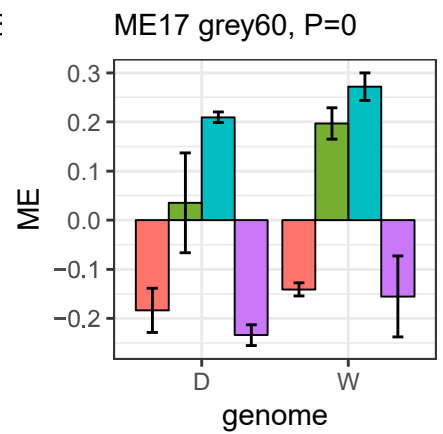
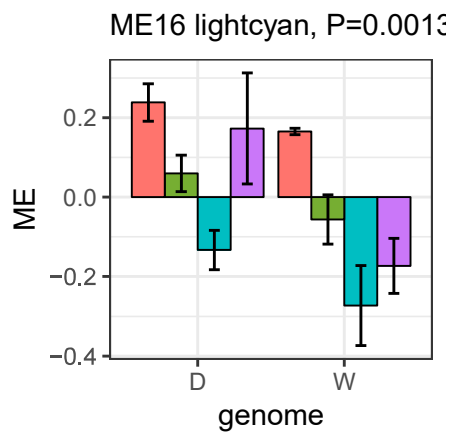
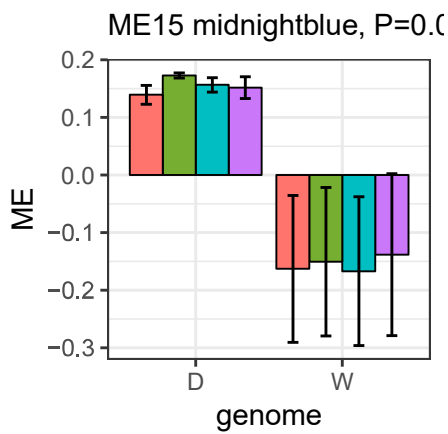
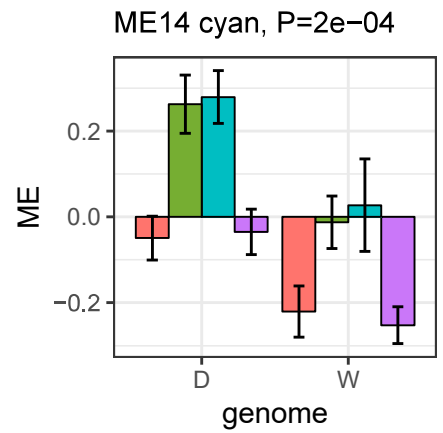
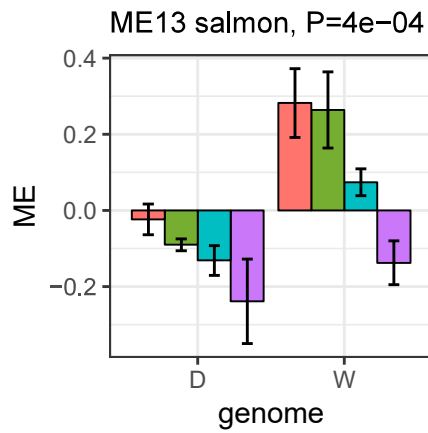
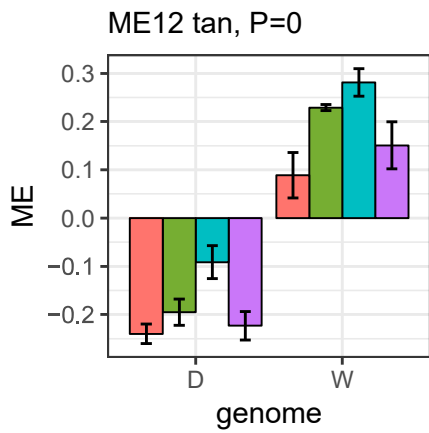
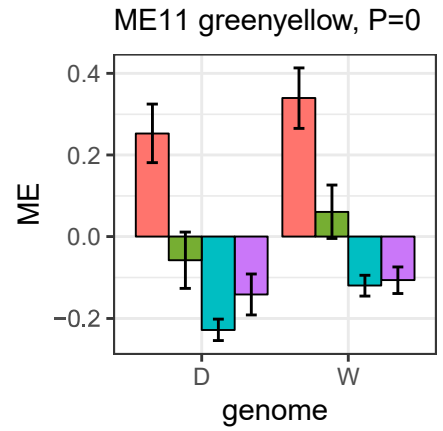
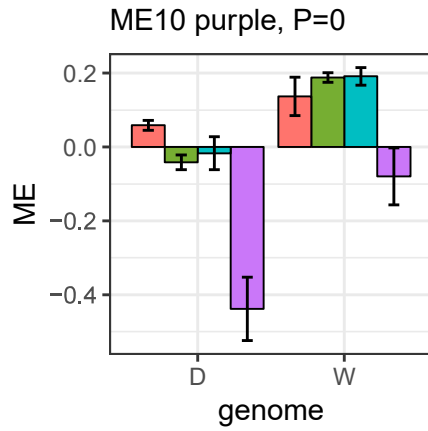
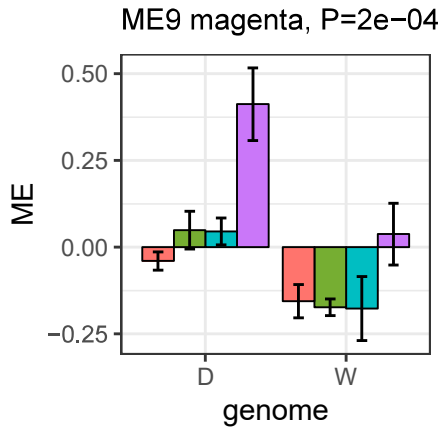


ME7 black, P=0.0317

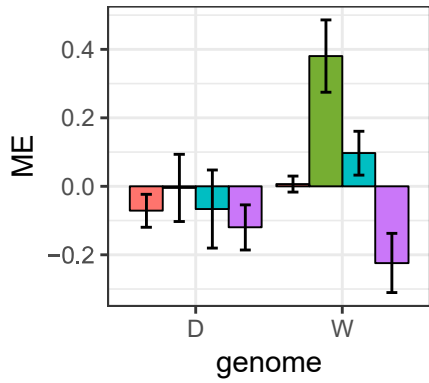


ME8 pink, P=0

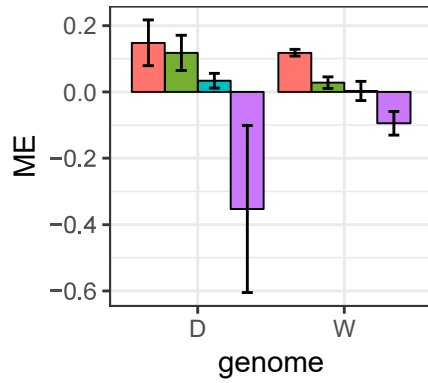




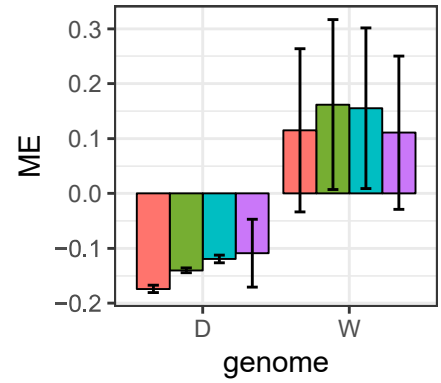
ME18 lightgreen, P=0.003



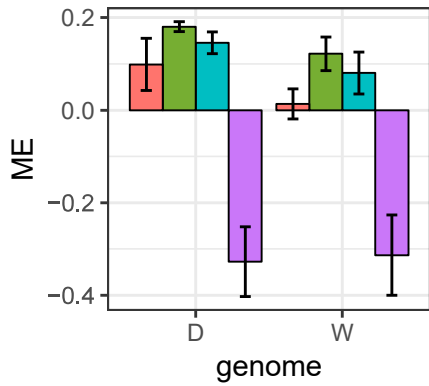
ME19 lightyellow, P=0.036



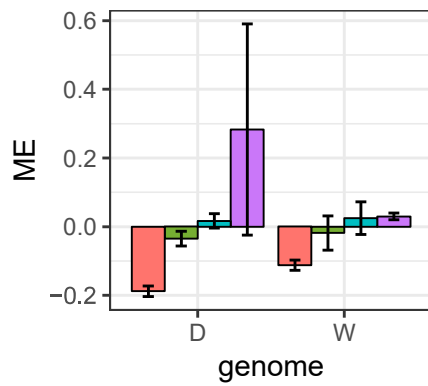
ME20 royalblue, P=0.135



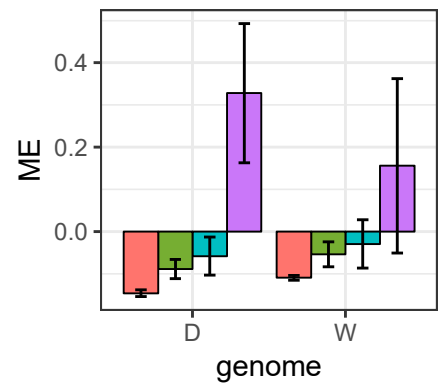
ME21 darkred, P=0



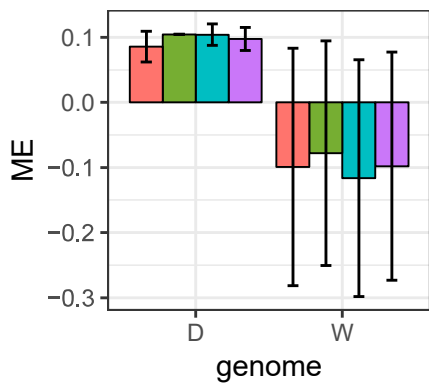
ME22 darkgreen, P=0.240



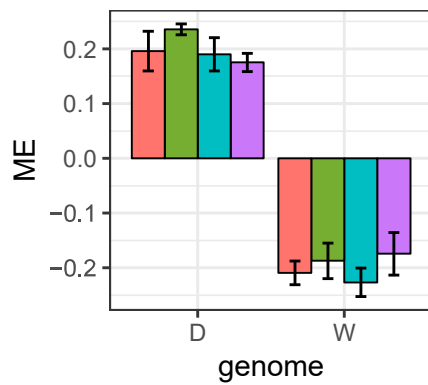
ME23 darkturquoise, P=0.0



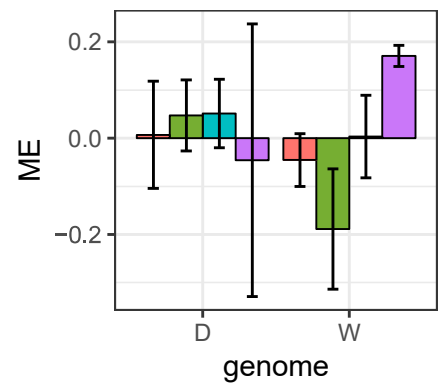
ME24 darkgrey, P=0.6761



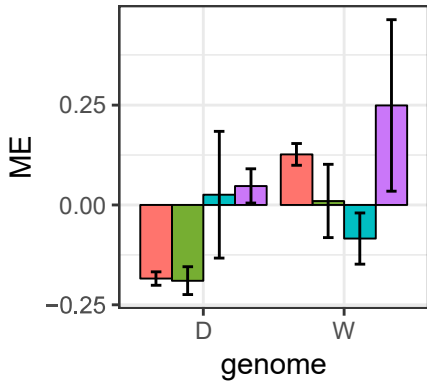
ME25 orange, P=0



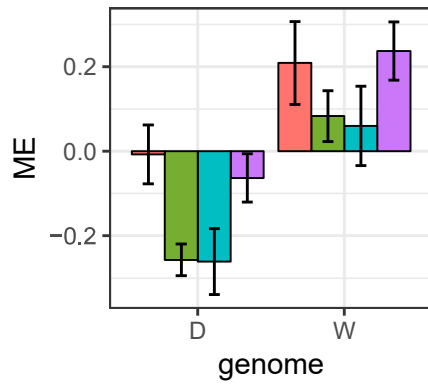
ME26 darkorange, P=0.70



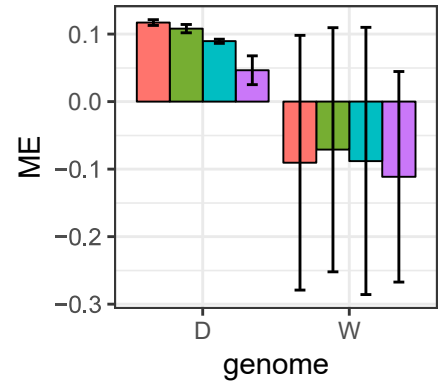
ME27 white, P=0.1099



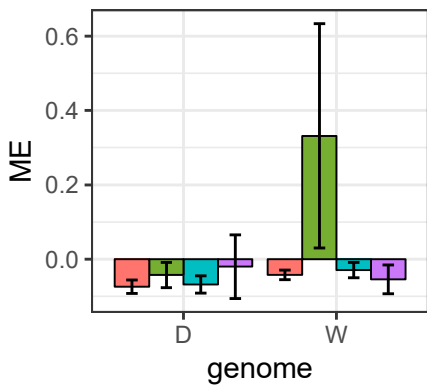
ME28 skyblue, P=8e-04



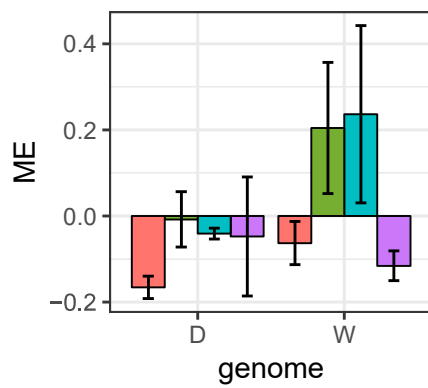
ME29 saddlebrown, P=0.7



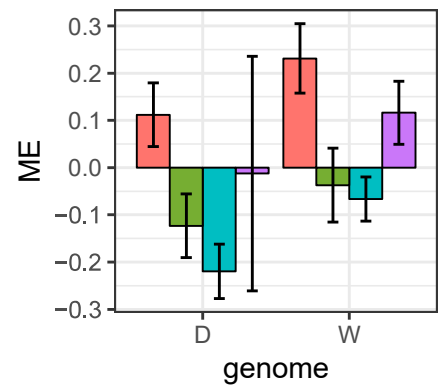
ME30 steelblue, P=0.2626



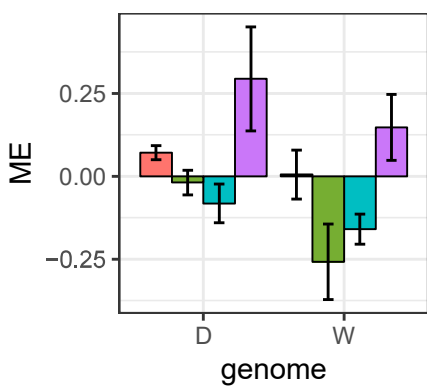
ME31 paleturquoise, P=0.



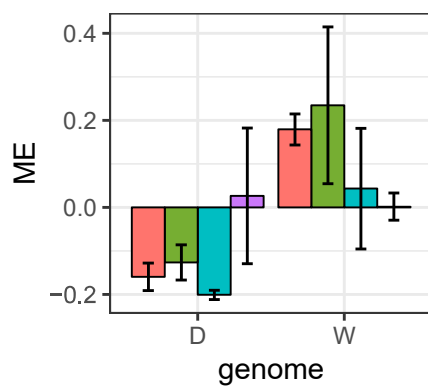
ME32 violet, P=0.148



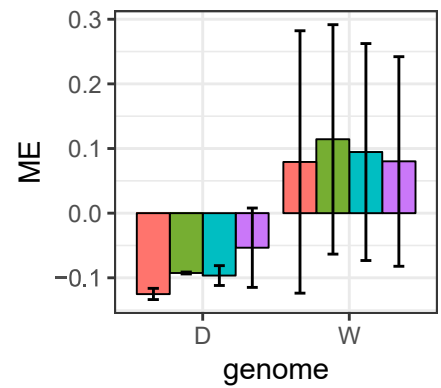
ME33 darkolivegreen, P=



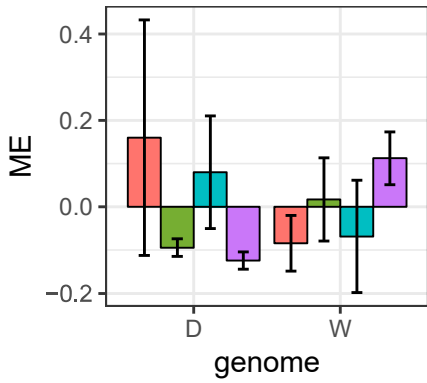
ME34 darkmagenta, P=0.



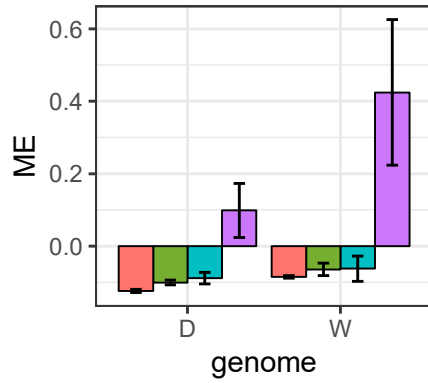
ME35 sienna3, P=0.7332



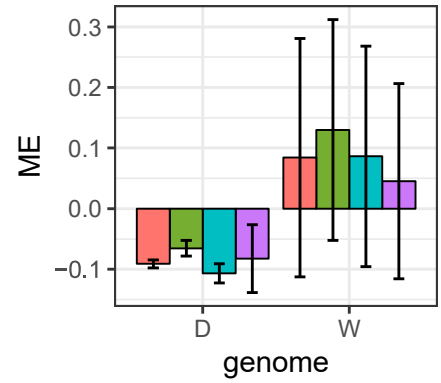
ME36 yellowgreen, P=0.6



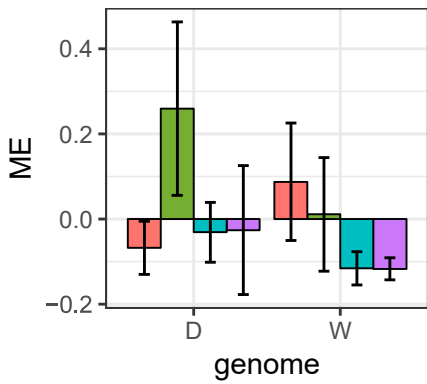
ME37 skyblue3, P=0.0019



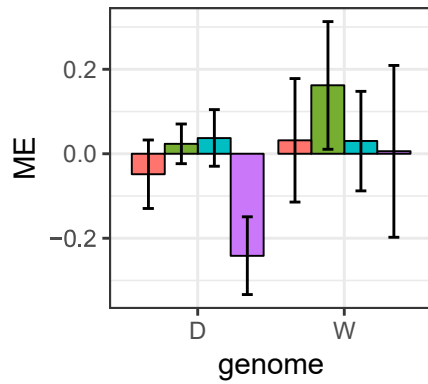
ME38 plum1, P=0.7879



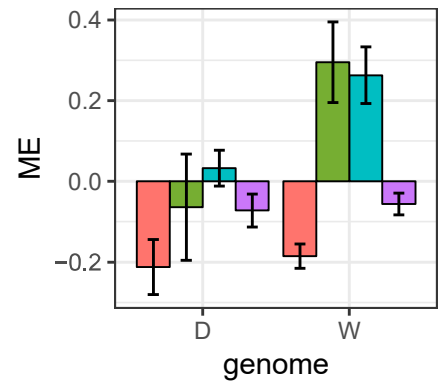
ME39 orangered4, P=0.40



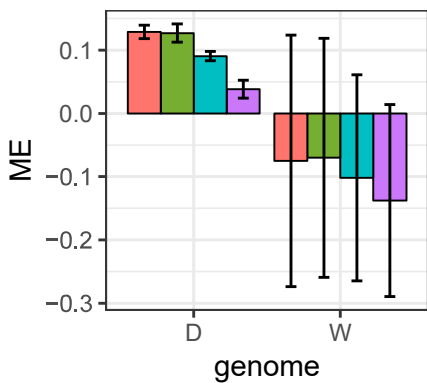
ME40 mediumpurple3, P=



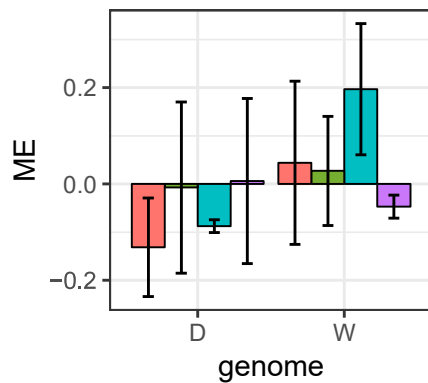
ME41 lightsteelblue1, P=8



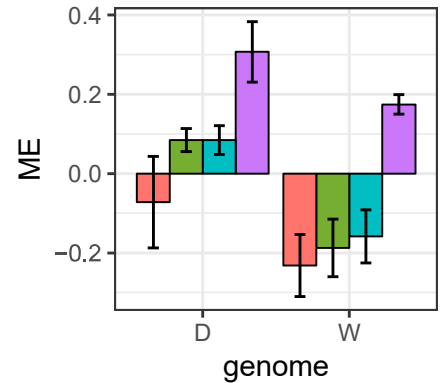
ME42 lightcyan1, P=0.634

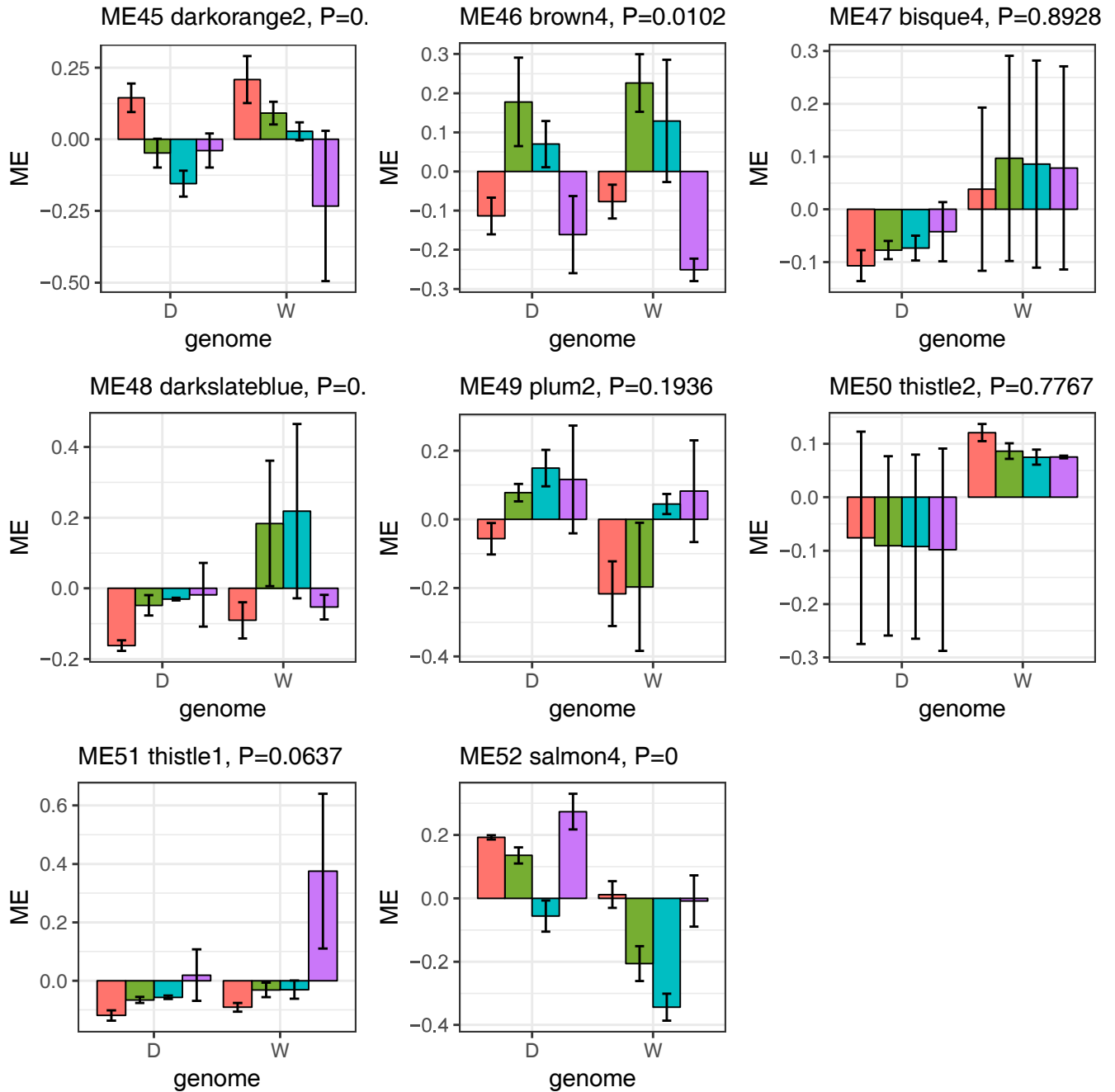


ME43 ivory, P=0.7524



ME44 floralwhite, P=4e-0





**Figure S3.** Average module eigengene values from the homoeologous fiber network across developmental timespoints for wild and domesticated fiber. Generalized expression patterns for the genes within each module are represented by the MEs. Developmental stages: red = 5 dpa, green = 10 dpa, blue = 15 dpa, purple = 20 dpa.