

Figure S6. Segregating inversions in the 43 DGRP lines. *y* axis shows the relative median DOE measures, INV = inversion karyotype (homozygous), INV / ST = heterozygote for the inversion, ST = standard configuration in a homozygous form. Plot titles show name of the inversion, brackets indicate the number of genotypes in each group. Where the number of lines is low each RAL ID is shown individually.

