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Figure S1. Crossing scheme for hemiclonal analysis. The figure illustrates the crosses needed to (A) sample, (B) amplify and (C) express individual haploid genomes (chromosomes X, II and III) from the four experimental populations. Chromosomes X/Y, II and III are shown left to right. The co-segregating translocation of chromosomes II and III carried by the clone-generator (CG) stock is shown as horizontally linked chromosomes II and III; compound X chromosomes are shown as linked in a horizontal V-shape. Chromosomes forming a target genome are denoted by a T. The males and females shown at the bottom of panel C are those that enter the fitness assays.