

Figure S1 Crossing scheme to generate X<sup>SR</sup> replacement lines. Cross 1: Wild-caught males crossed to lab stock (141.02) and sex-ratio of offspring noted. All lines with skewed sex-ratio were retained. Cross 2: Same wild-caught male as in Cross 1 mated to his daughters. Most offspring from Cross 2 were be female and assuming equal viability, 50% were homozygous for X<sup>SR</sup> and 50% were heterozygous (X<sup>SR</sup>/X<sup>ST</sup>). Cross 3: Virgin females from Cross 2 were mated to standard males (141.02). If these Cross 3 females were heterozygous for the driver, half of sons sired female-biased sex-ratios (3a). If they were heterozygous, all sons sired female-biased sex-ratios (3b). Males from Cross 3b were mated to virgin females from Cross 2 (their aunts) assuring that the male was X<sup>SR</sup>Y and the female was X<sup>SR</sup>YSR. To generate new X<sup>SR</sup>YSR females, virgin X<sup>SR</sup>XSR females were mated to X<sup>SR</sup>Y males. To generate new X<sup>SR</sup>Y males, virgin X<sup>SR</sup>XSR females were mated to X<sup>ST</sup>Y males (141.02). This crossing scheme replaces 50% of the genome each generation with 141.02 genetic material and therefore over several generations, the wild autosomes and Y chromosome are replaced with the lab stock (141.02).