

Table S1. Genome resequencing coverage of Drosophila strains and species

Strain	Total Reads	Mapped Reads	Coverage	SNPs and indels modified	New sequence added
<i>D. mel zhr*</i>	69,794,712	66,705,080	30.04	659,661	NA
<i>D. mel z30*</i>	51,969,198	49,523,960	22.31	1,142,005	NA
<i>D. sim tsimbazaza</i>	63,719,674	61,579,018	30.18	1,369,487	12,633,195
<i>D. sechellia (reference)</i>	55,413,474	50,308,326	22.73	56,067	1,614,249

* Genomes from these strains were reported in Coolon *eg al.*, 2012.