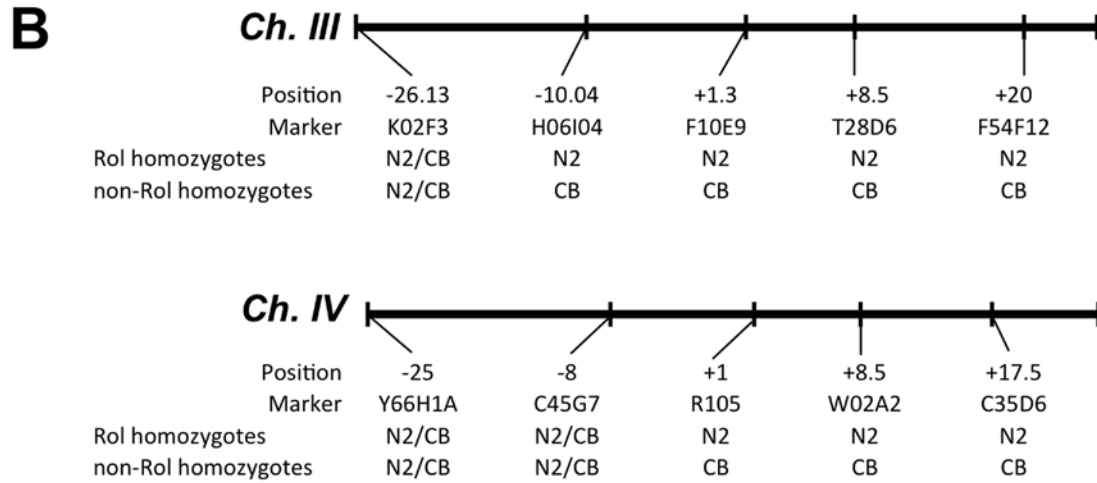


**A**

	Position	Ch. I central	Ch. II central	Ch. III central	Ch. IV central	Ch. V central	X Ch. central
	Marker	K04F10	T13C2	F10E9	R105	F20D6	F45E1
Rol homozygotes		N2/CB	N2/CB	N2	N2	N2/CB	N2/CB
non-Rol homozygotes		N2/CB	N2/CB	CB	CB	N2/CB	N2/CB



**Figure S1 Snip-SNP mapping of the locus of *mels4* (containing the *lacO* array) in AV221.** Summary of the SNP mapping analysis used to identify linkage of *mels4* (containing the *lacO* array) to markers on both chromosomes III and IV, showing the genetic map positions of SNP markers used in this analysis; see Materials and Methods for explanation. In brief, the co-integrated marker *rol-6(su1006)* was used as a surrogate marker to track the presence of *mels4*, which was generated in the N2 genetic background. Heterozygous F1 worms were generated by crossing the AV221 and CB4856 strains, and SNP marker genotypes were assessed both for F2 worms that were homozygous for *mels4* (Rol homozygotes) or that lacked *mels4* (non-Rol homozygotes). “N2/CB” indicates that both alleles were detected, reflecting independent assortment of *mels4* and the SNP marker or crossing over between the array and the SNP marker. “N2” (in the Rol homozygotes) or “CB” (in the non-Rol homozygotes) indicates tight linkage of the SNP marker to *mels4*. The unusual marker segregation pattern observed indicates that *mels4* is associated with a reciprocal translocation between chromosomes III and IV (see Materials and Methods and Figure S2).