

Table S15. Number of matches to the CLAMP PBM motif per Mb on individual chromosomal arms.

<i>D. melanogaster</i>		<i>D. miranda</i>		<i>A. gambiae</i>	
chr2L	91.6	chr2	387.5	chr2L	409.7
chr2R	88.2	chr4	507.3	chr2R	416.5
chr3L	90.6	chr5	136.0	chr3L	334.1
chr3R	91.4	neoX	478.0	chr3R	372.3
chr4	11.9	chrXL	645.4	chrX	870.0
chrX	202.8	chrXR	729.0		