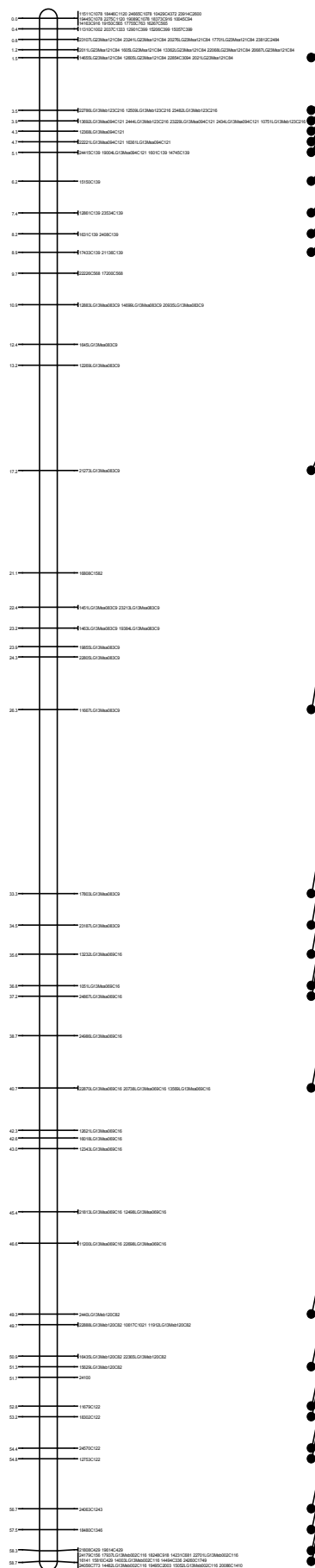
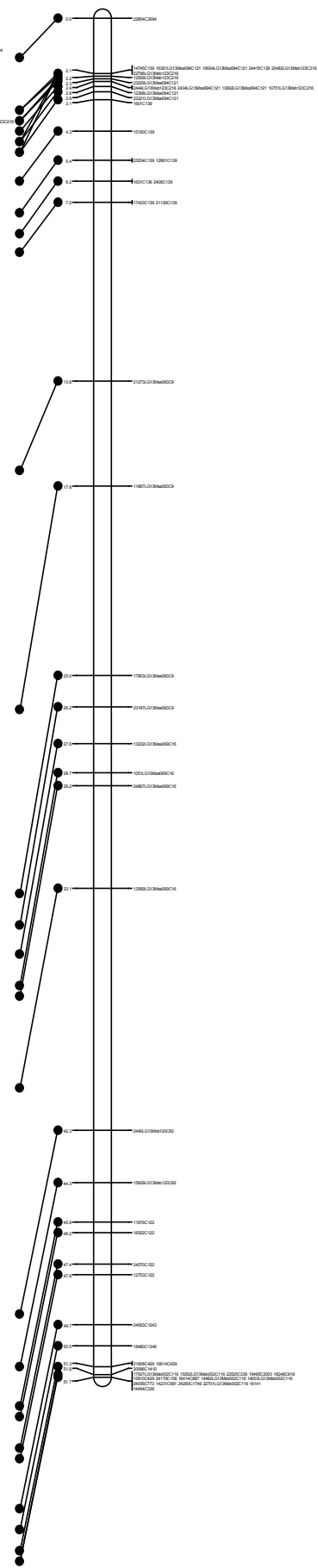


Fig. S1

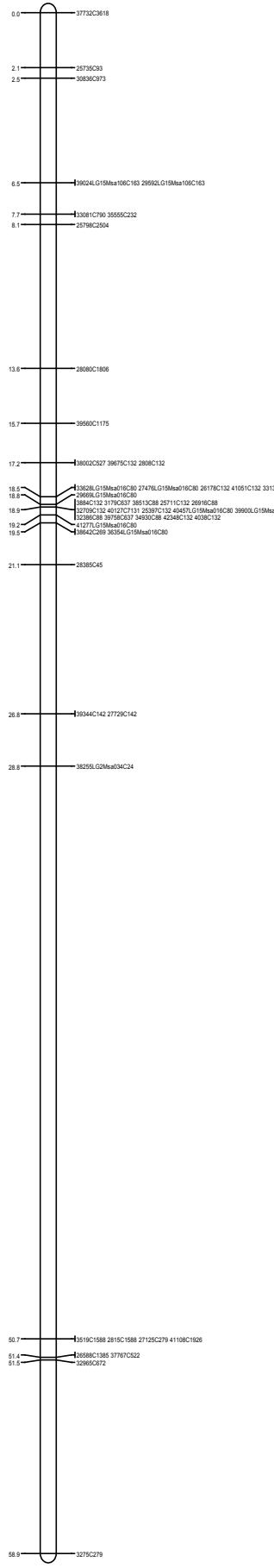
Xma LG13 120 markers ML



Xma LG13 120 markers RM



Xma LG15 110 markers RM



Xma LG15 110 markers ML

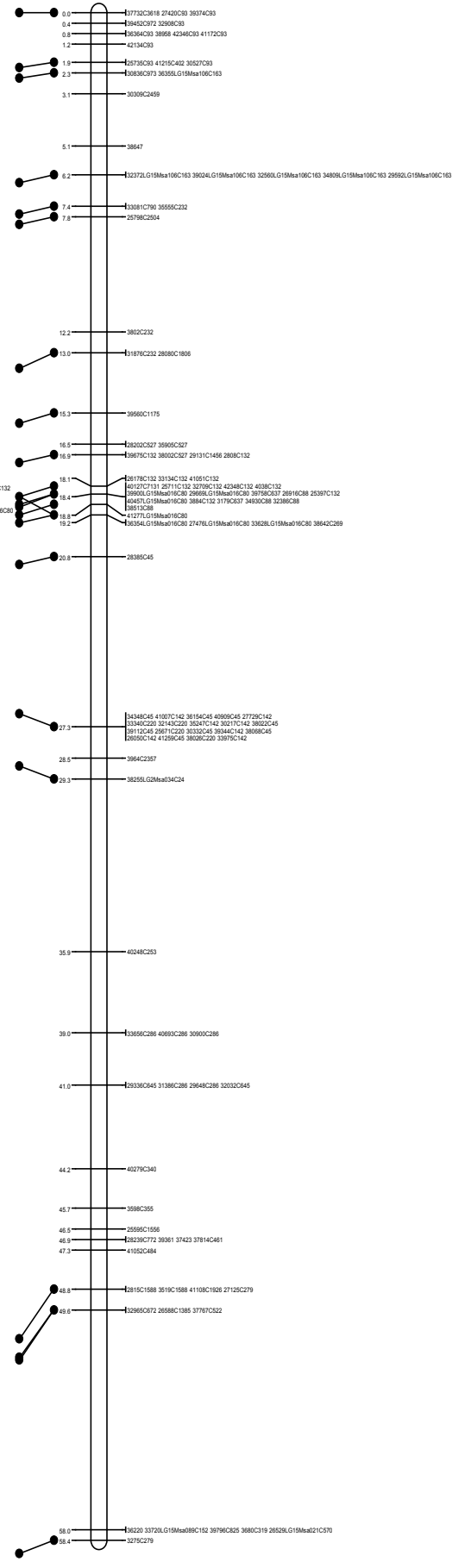


Figure S1 A comparison of marker location along platyfish LG11 using the Maximum Likelihood and the Regression Mapping algorithms in Joinmap 4.0 with a common set of 135 markers. Results show no significant differences.