

Additional file 6: LD decay within physical distance on each autosome (1-29) in a window of 500 kb. Dotted horizontal lines are the LD background (computed on a subset of non-syntenic SNPs).

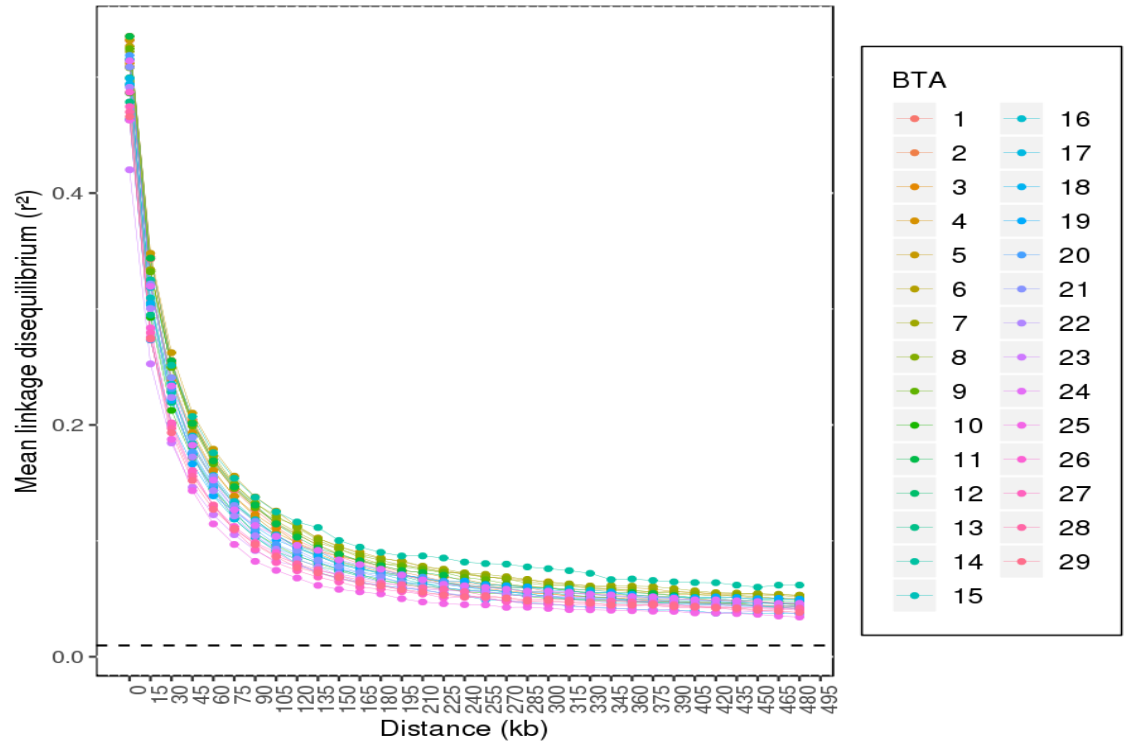


Figure S6: Distribution of average r^2 values for CHA breed with respect to physical distance (kb) in a window of 500 kb

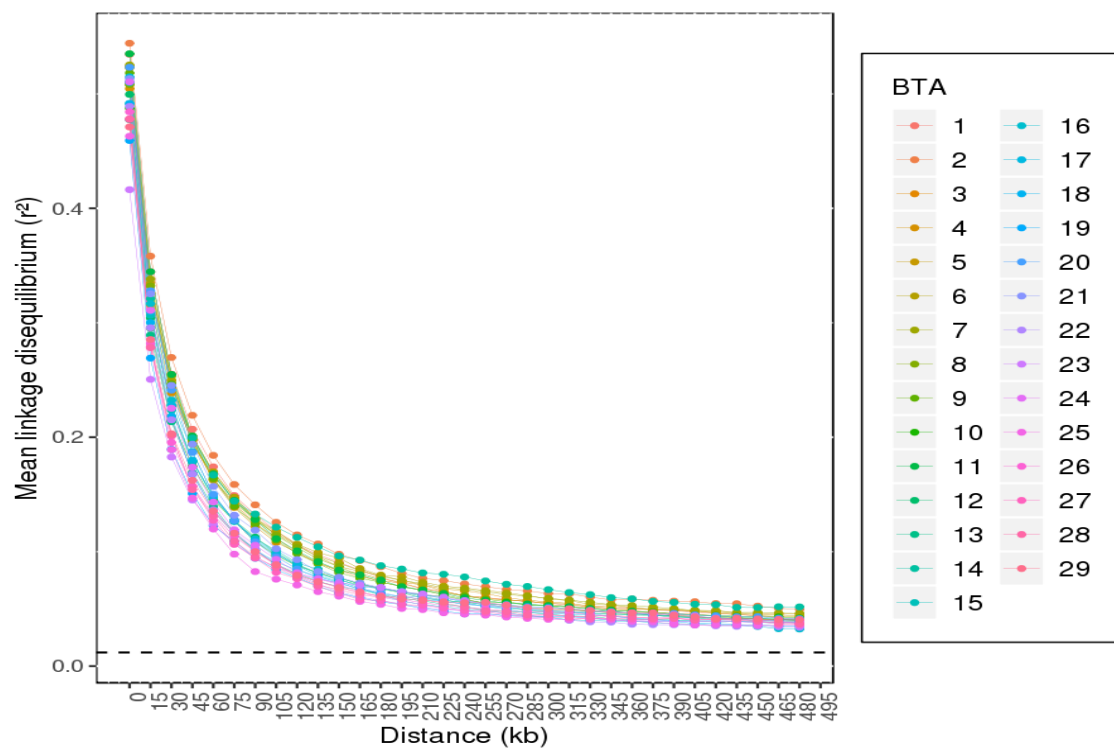


Figure S7: Distribution of average r^2 values for LIM breed with respect to physical distance (kb) in a window of 500 kb

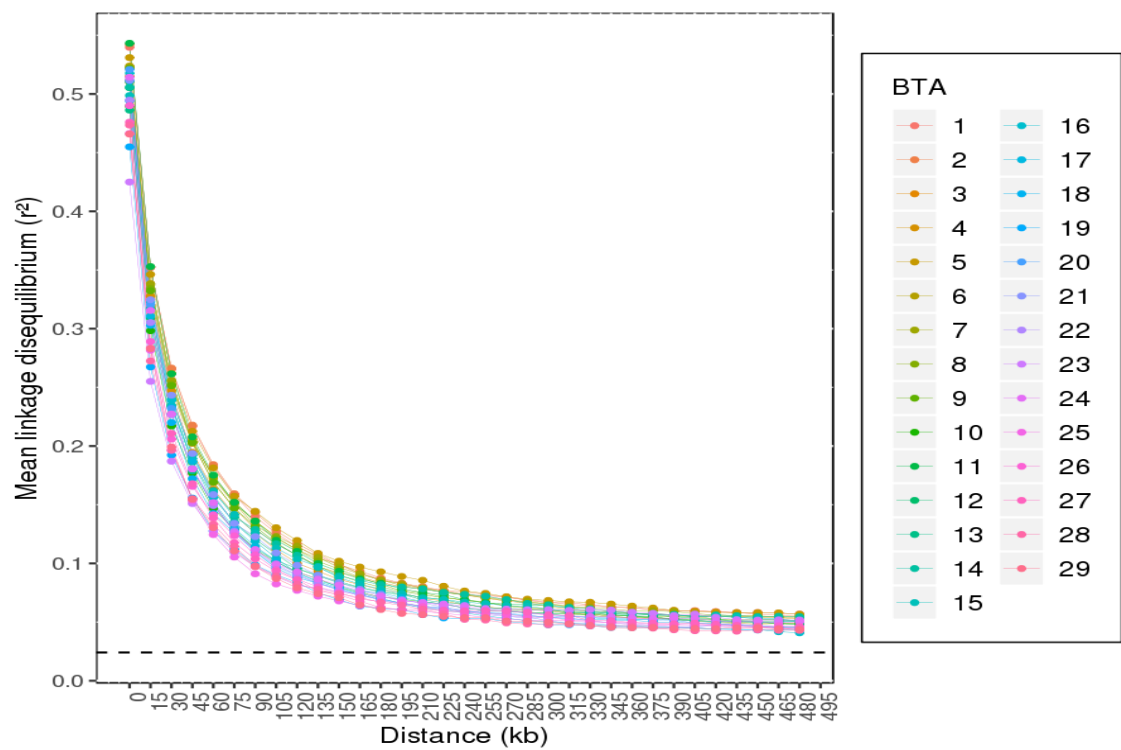


Figure S8: Distribution of average r^2 values for BLA breed with respect to physical distance (kb) in a window of 500 kb