



Figure S1 Relative positions of LKAM sequences in relation to the *L. sativa* draft genome assembly 'Lsat_1_v4'. a.- i., Pseudo-chromosomes 1 – 9 respectively. Circularized for illustration, the scale on the inner circle depicts base pair (bp) intervals, the outer circle is labeled with the LKAM markers prefixed LS1_ then the numeric identifier.