



Figure S1 M regions visualization through SGD browser (<http://www.yeastgenome.org/>). M regions are represented as blue arrows on the last track. M regions may correspond to already known repeated elements such as Ty and LTR elements (M14_40) or to multi-copy genes. For example, the *YNL034W* (570,477-572,315) and *YNL033W* (572,999-573,853) loci, covered by M14_43, are duplications of the locus *YNL018C* (601,774-599,936) covered by M14_49, and the locus *YNL019C* (599,230-598,376) covered by M14_48, respectively. The SGD interface includes functional annotations; for example, the meiotic “Double strand break hotspots: Pan et al. (2011)”.