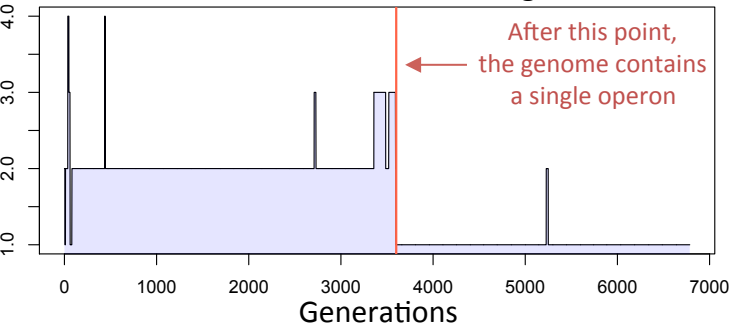
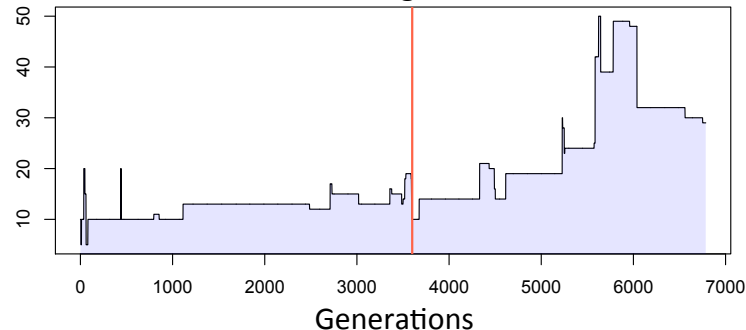
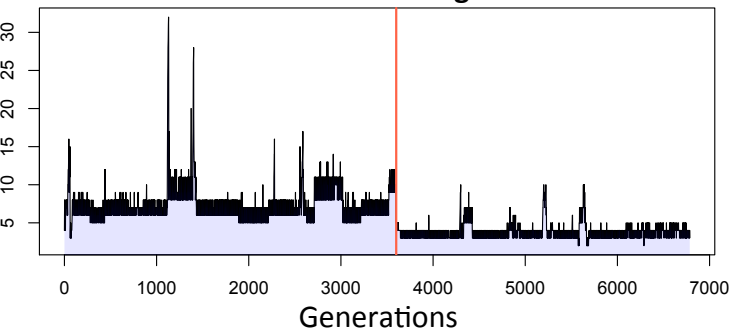
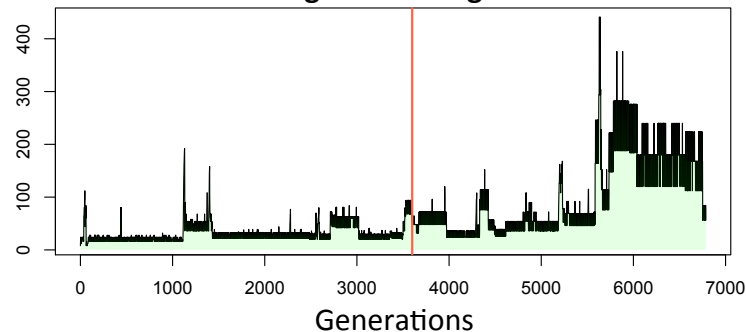
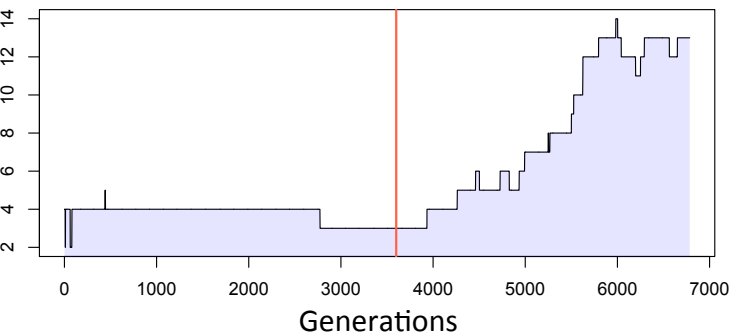


a. Number of functional regions**b. Functional genome size****c. Number of nodes in the regulation network****d. Number of edges in the regulation network****e. Number of nodes in the metabolic network****f. Number of edges in the metabolic network**