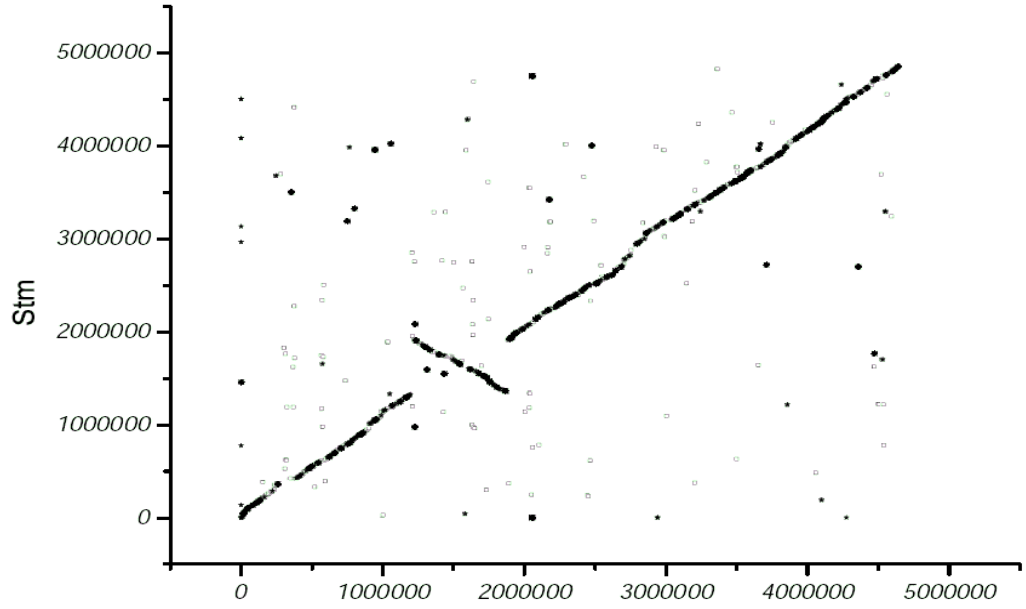
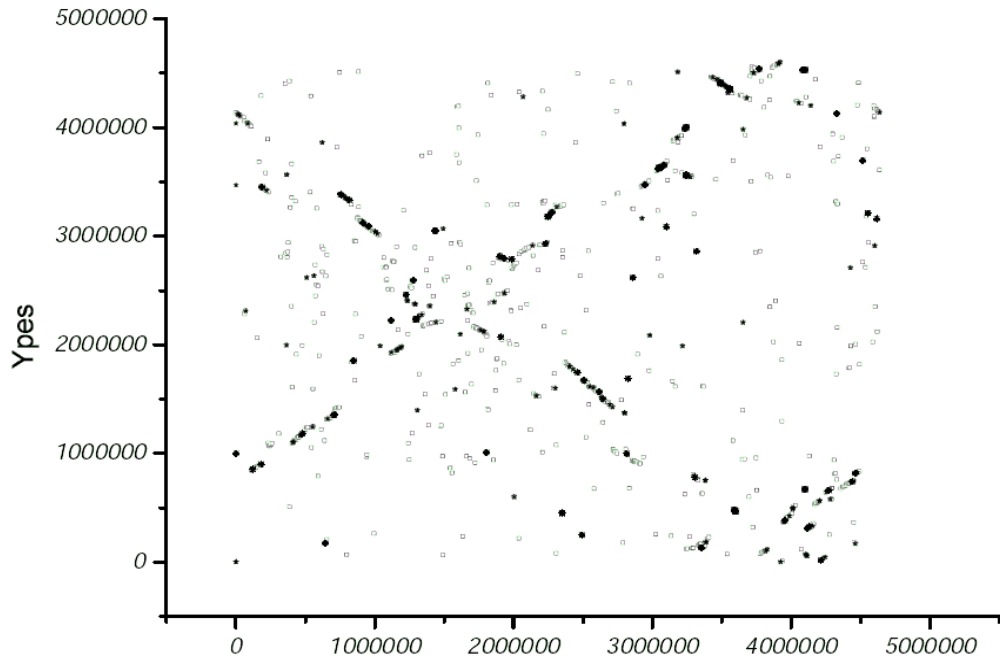


R4: Conservation of regulatory sites and chromosome vicinity of their target genes



A



B

Conserved genes and regulatory sites between the Eco and the Stm (A), and the Eco and the Ypes (B) genomes. Pairs of orthologous genes between each pair of genomes have x and y coordinates that correspond to their positions in each genome. Genes that share the same regulation in both genomes are represented as filled squares; genes that belong to the **other regulon** or **lost link** categories are represented as open squares.