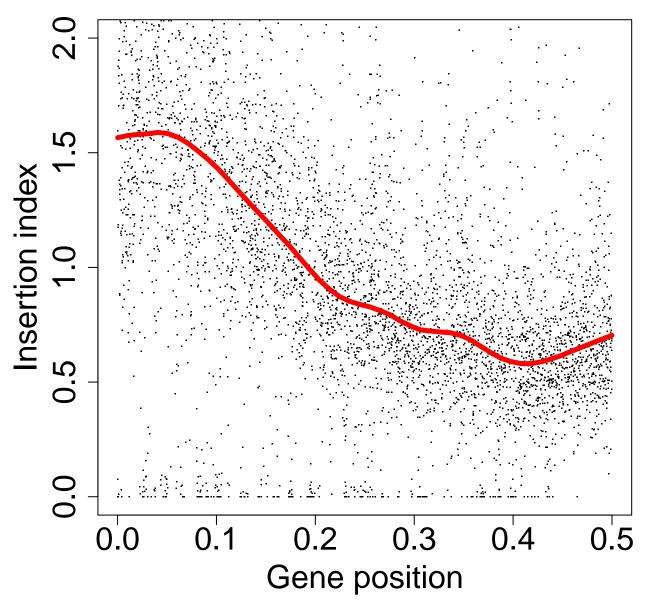
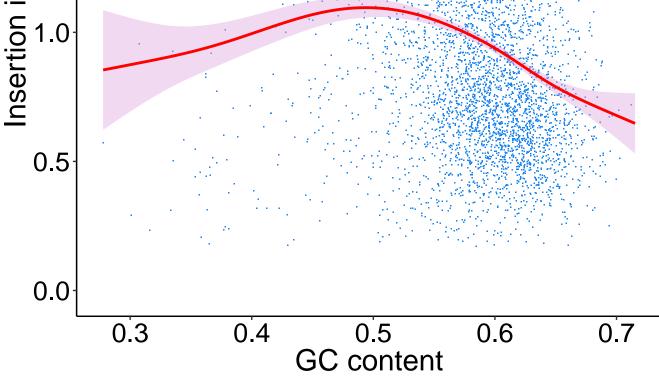


Distance bias - Klebsiella pneumoniae Ec

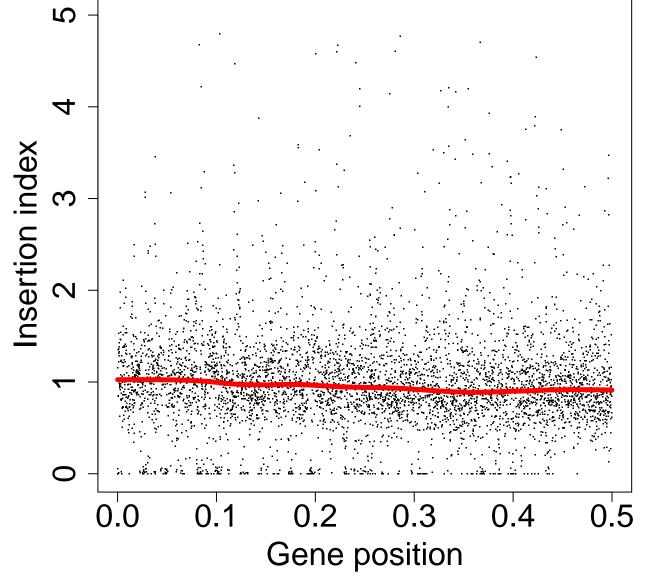


GC bias - Klebsiella pneumoniae Ecl8 2.0 1.5 Insertion index 0.5 0.0 0.4 0.6 0.3 0.5 GC content

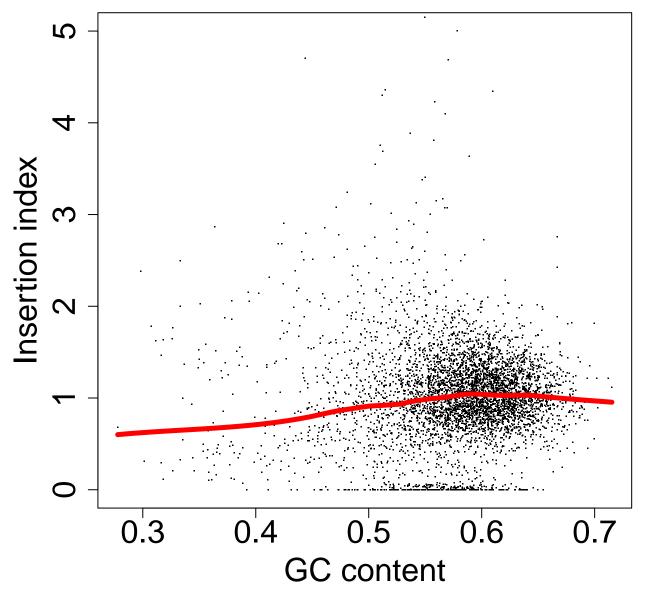
GC bias - Klebsiella pneumoniae Ecl8 2.0 1.5 Insertion index

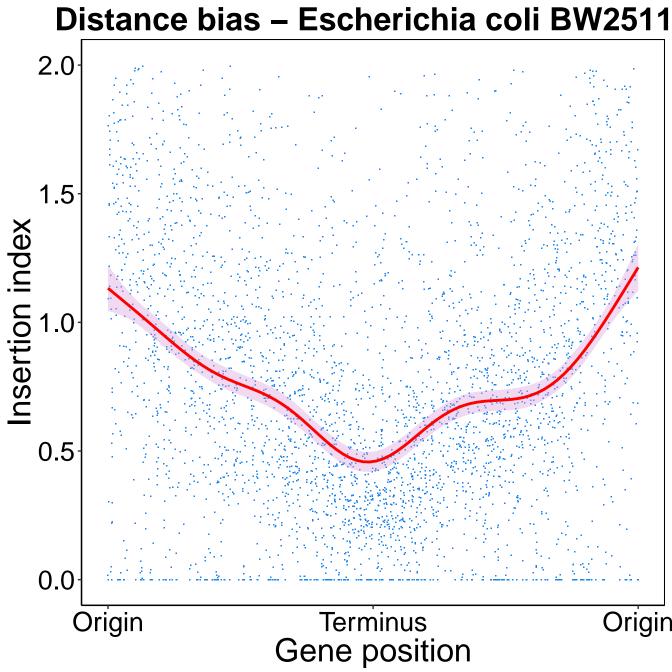


(lebsiella pneumoniae Ecl8 – normalised dist

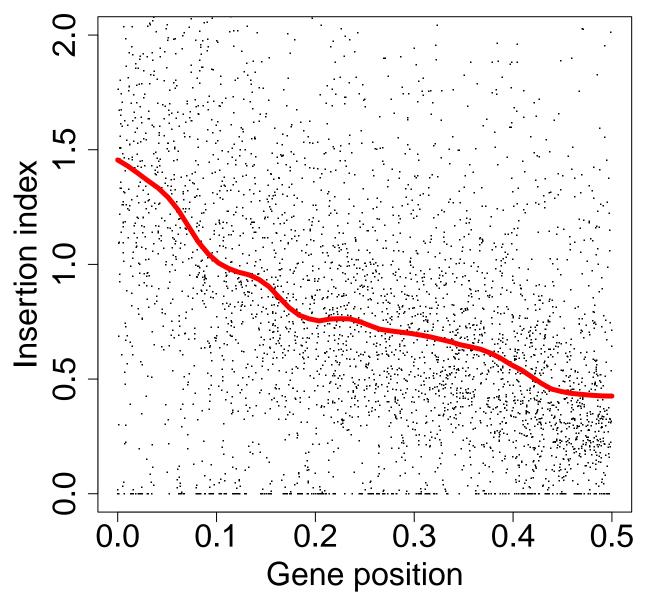


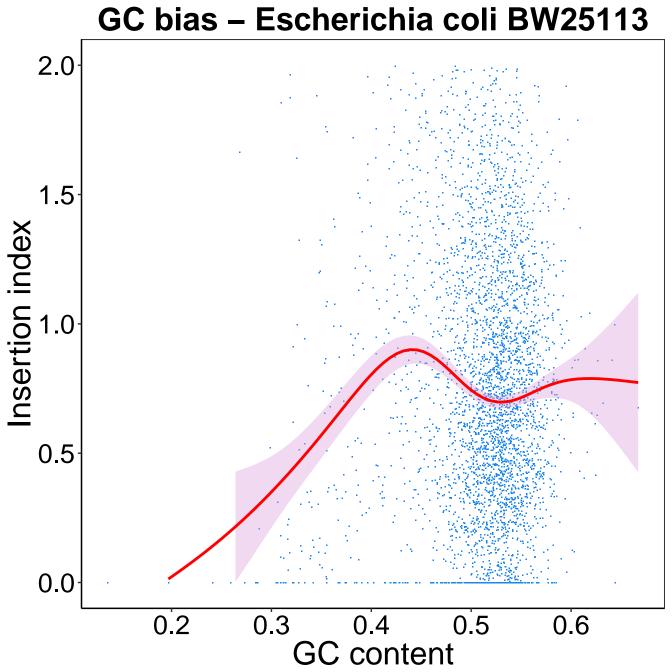
Klebsiella pneumoniae Ecl8 – normalised C

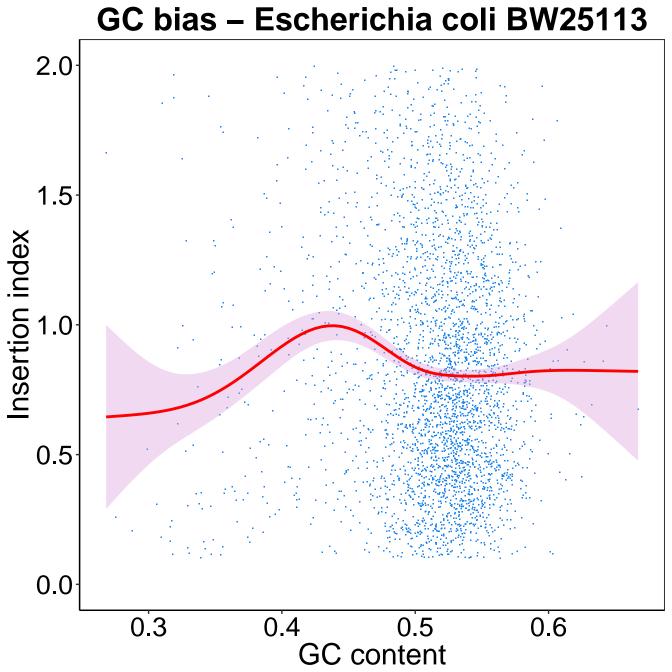




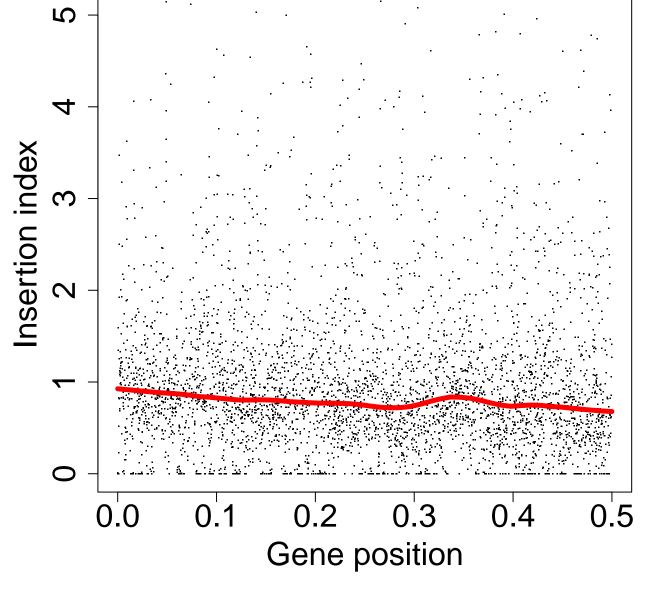
Distance bias – Escherichia coli BW25113



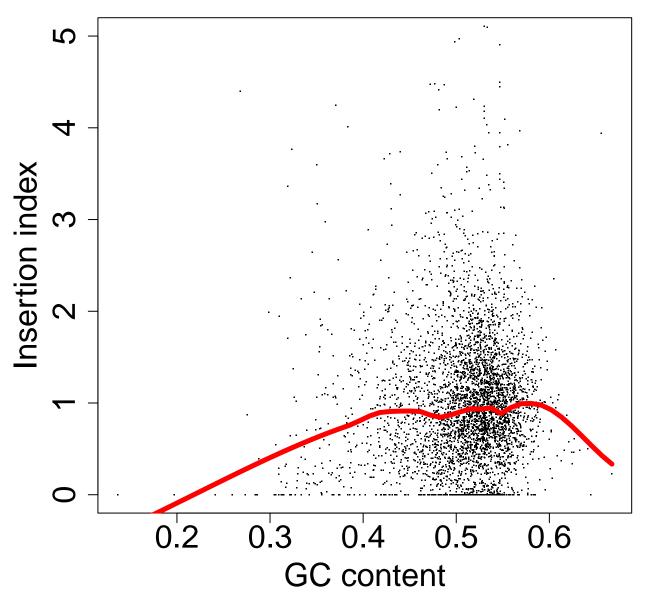


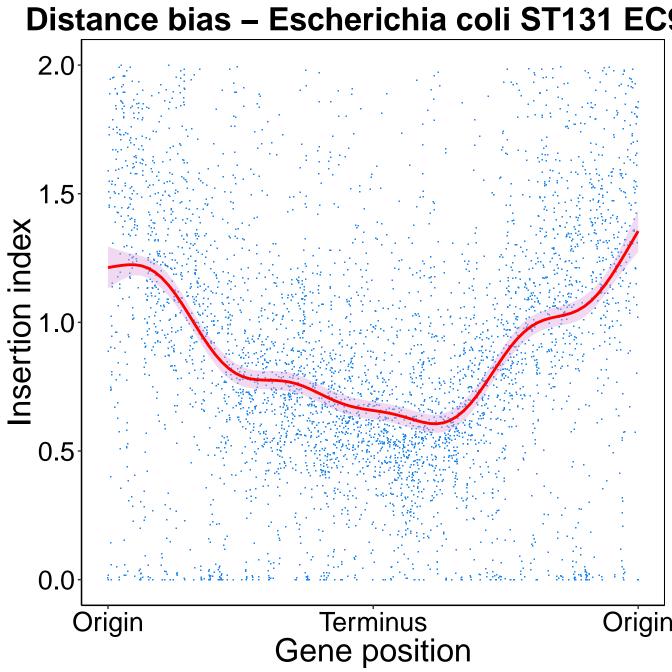


Escherichia coli BW25113 – normalised dista

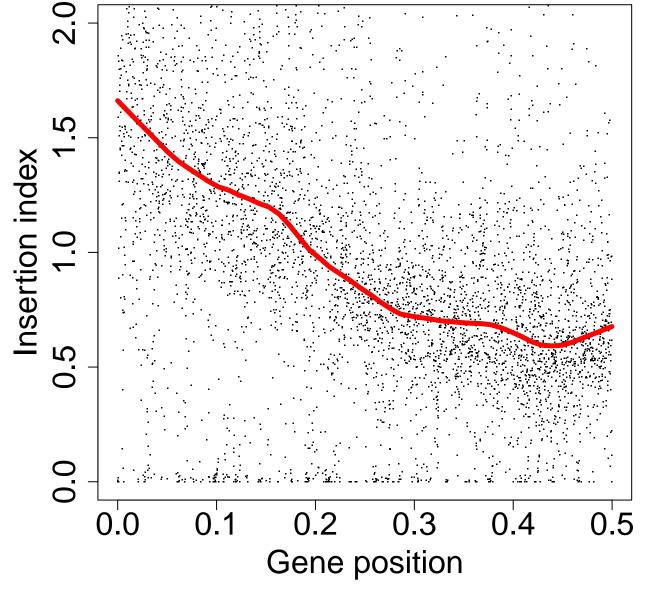


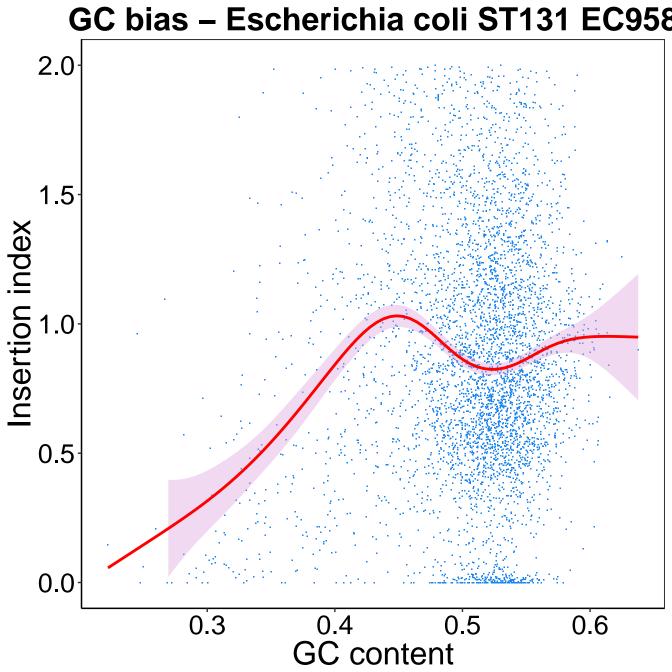
Escherichia coli BW25113 – normalised G

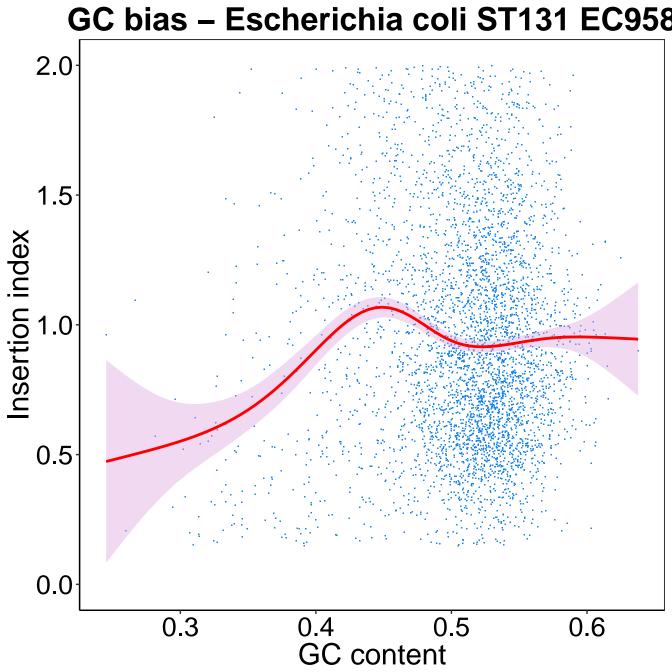




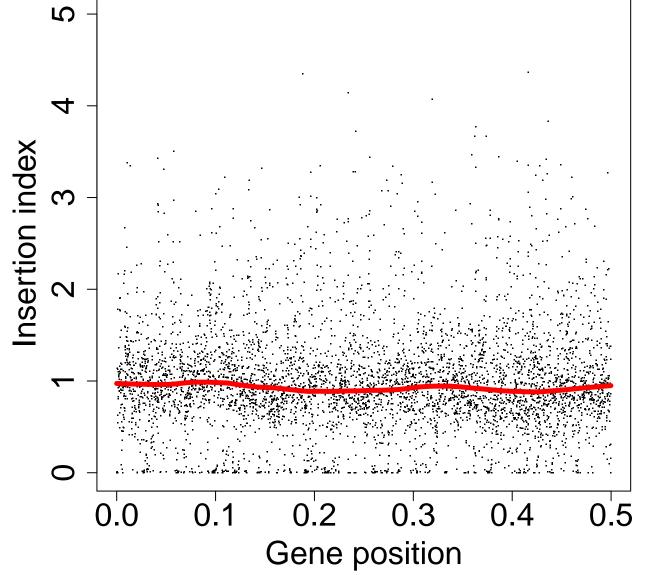
Distance bias – Escherichia coli ST131 EC9



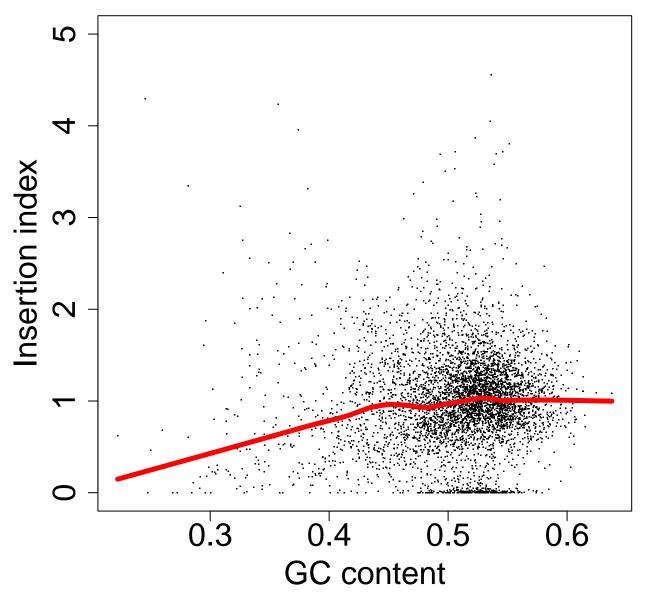


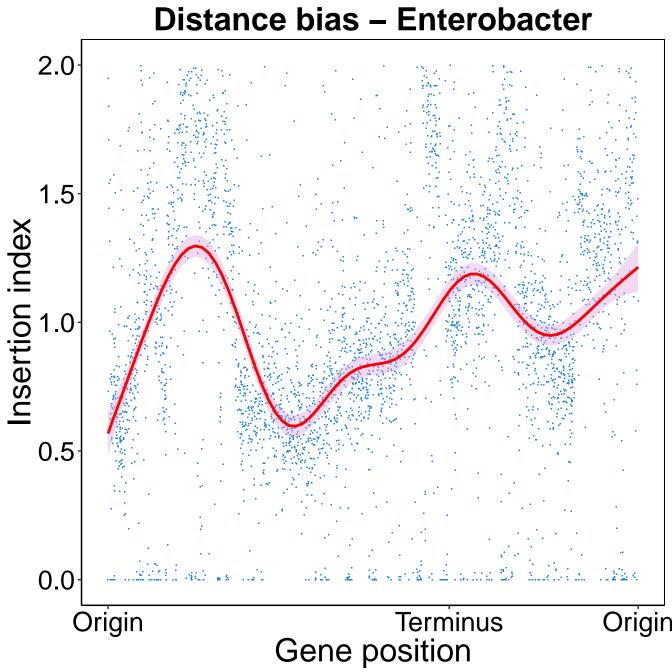


scherichia coli ST131 EC958 – normalised dis

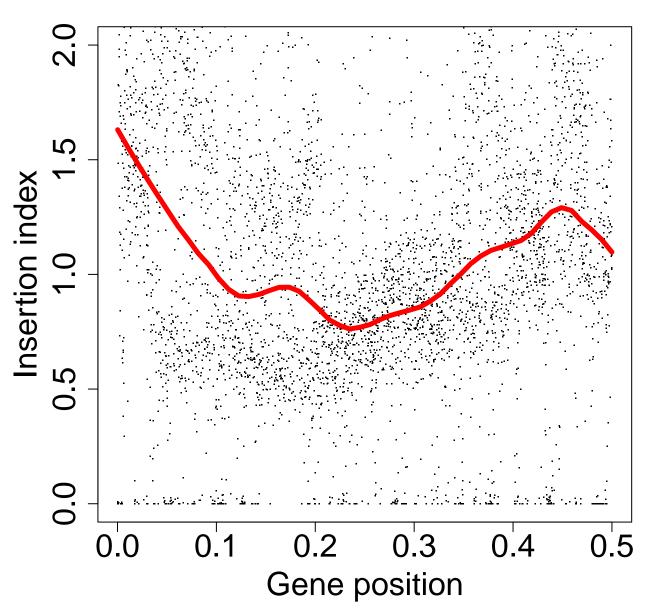


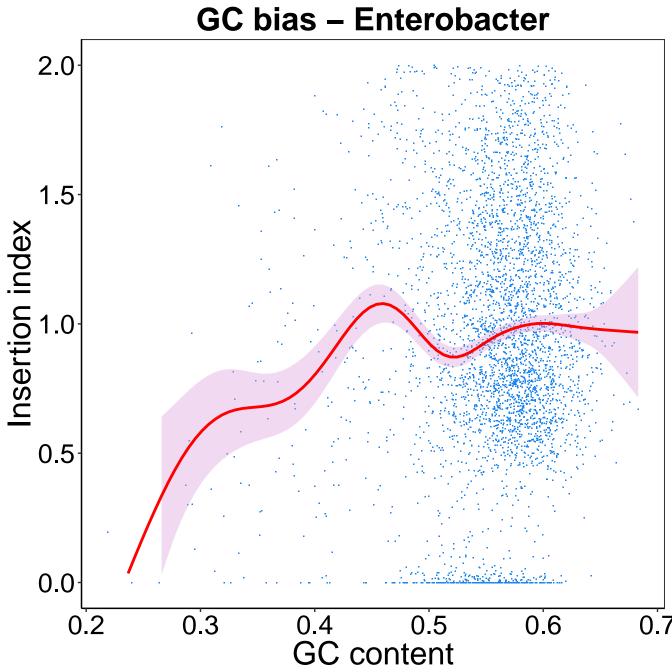
Escherichia coli ST131 EC958 – normalised

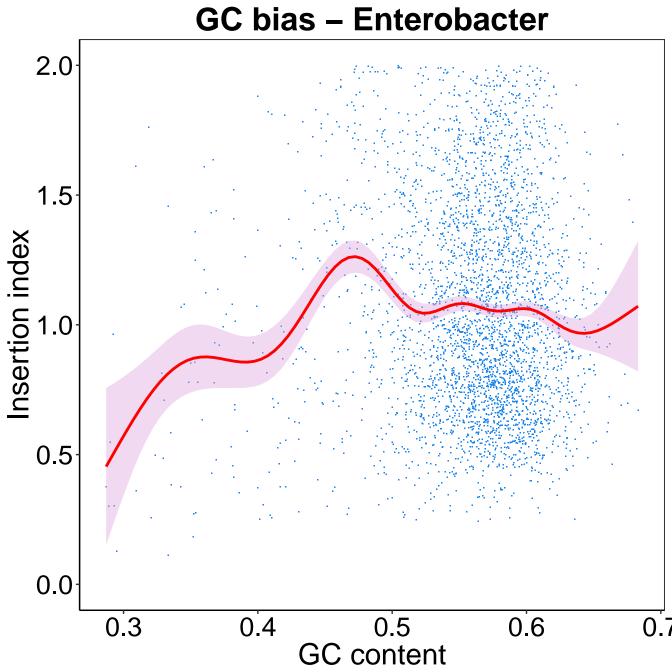




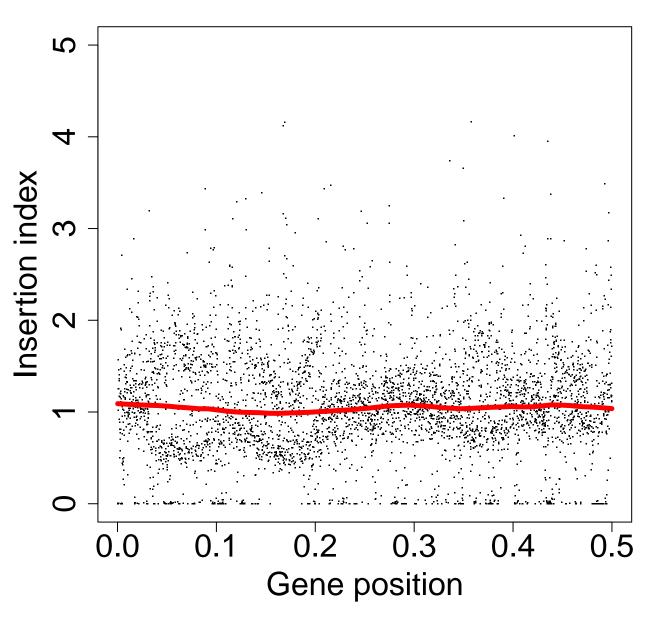
Distance bias – Enterobacter



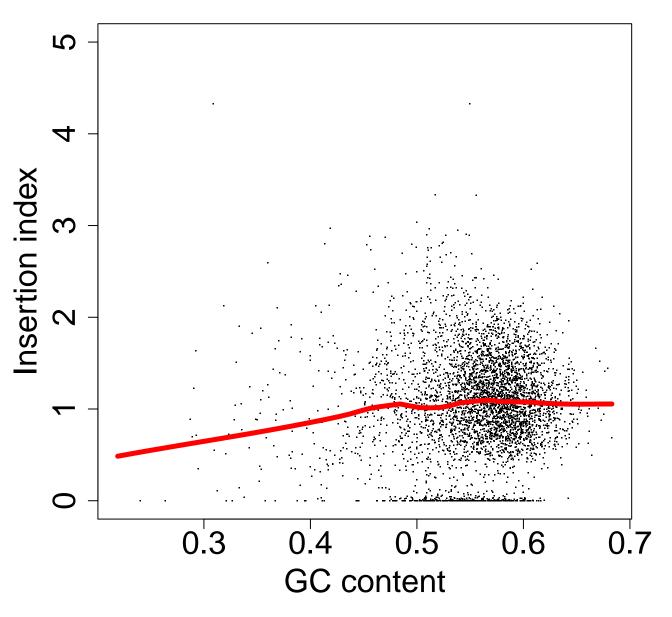


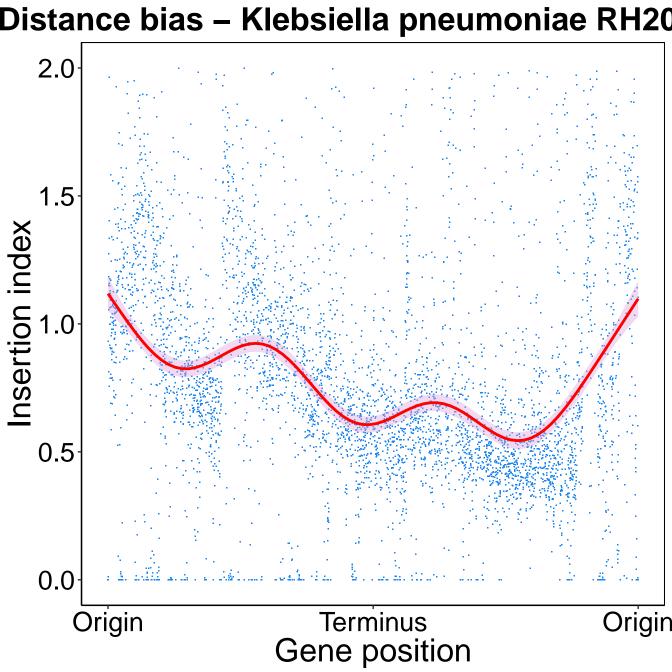


Enterobacter – normalised distance

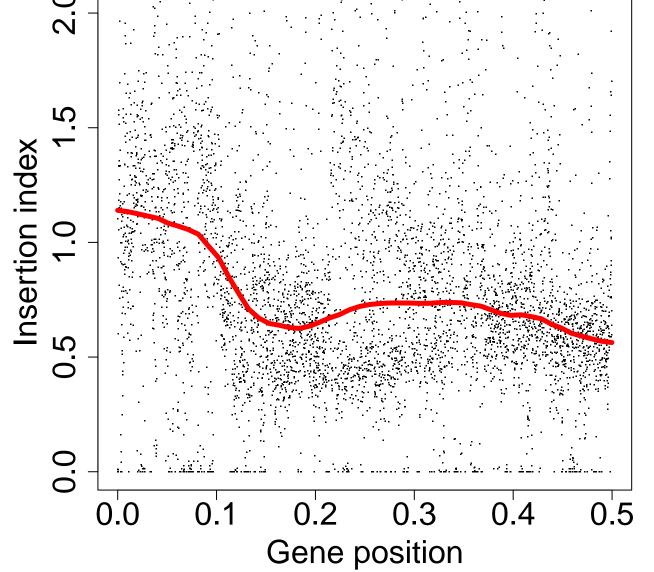


Enterobacter – normalised GC

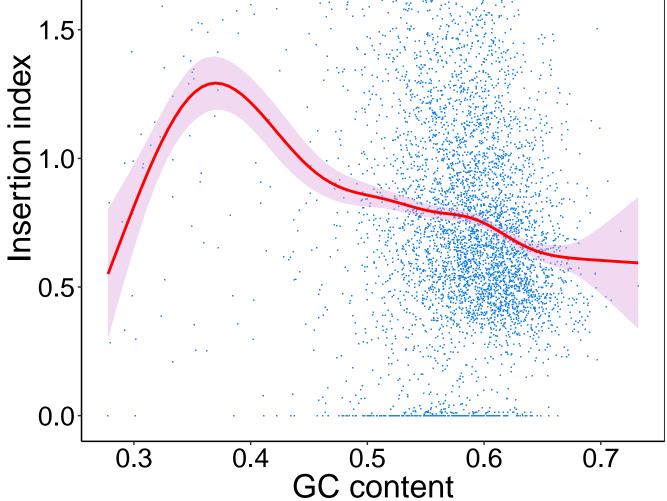




Distance bias – Klebsiella pneumoniae RH20

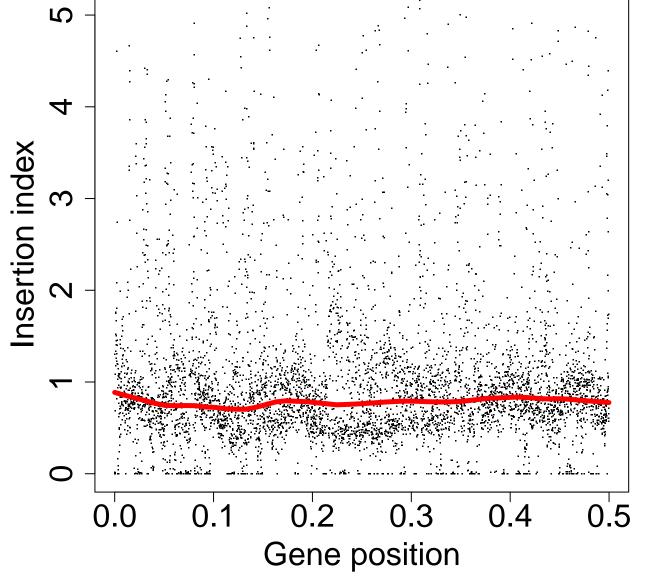


GC bias – Klebsiella pneumoniae RH2012

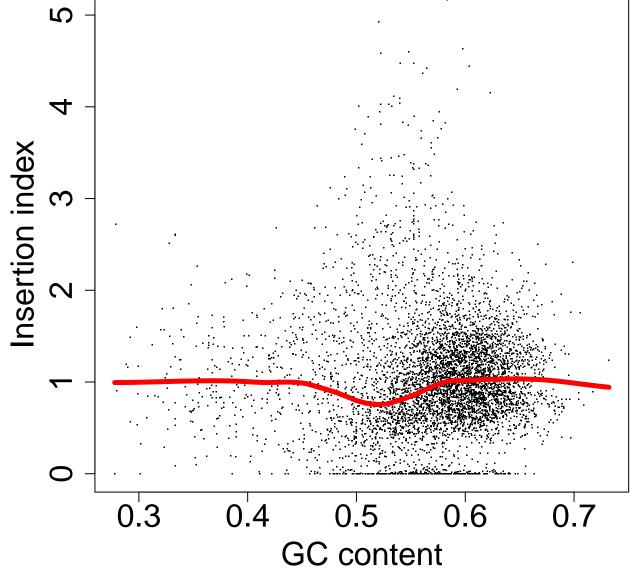


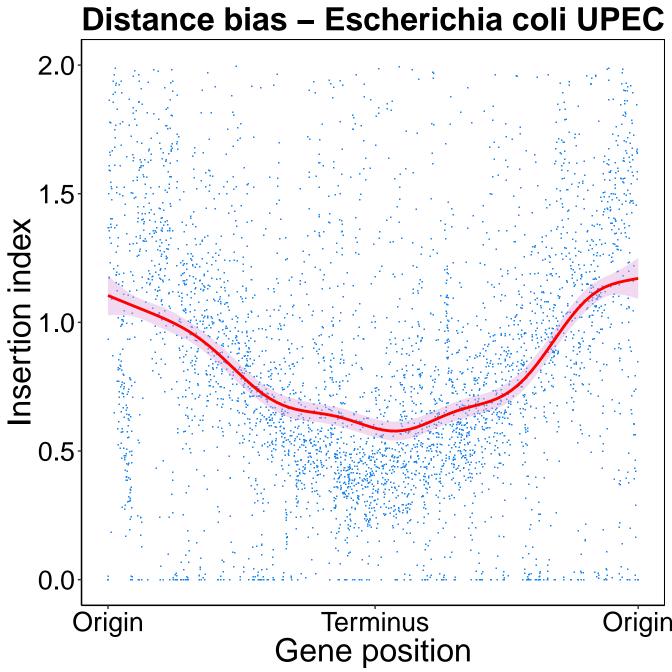
GC bias - Klebsiella pneumoniae RH2012 2.0 1.5 Insertion index 0.5 0.0 0.4 0.5 GC content 0.6 0.3 0.7

osiella pneumoniae RH201207 – normalised o

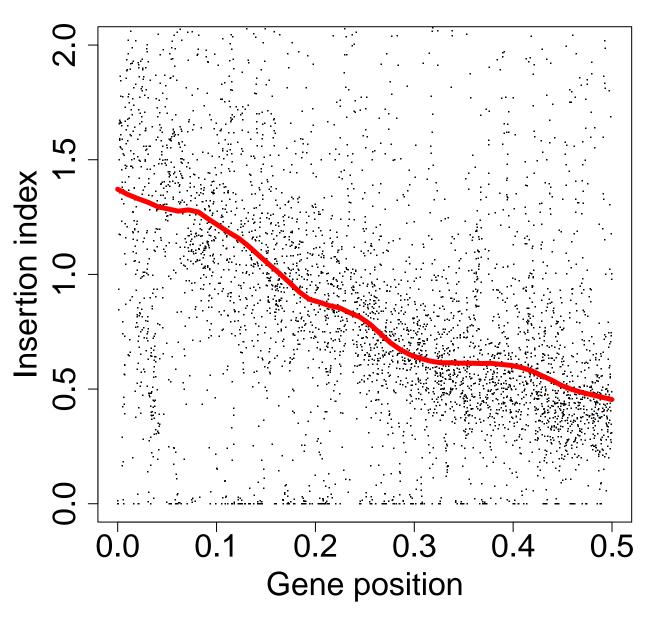


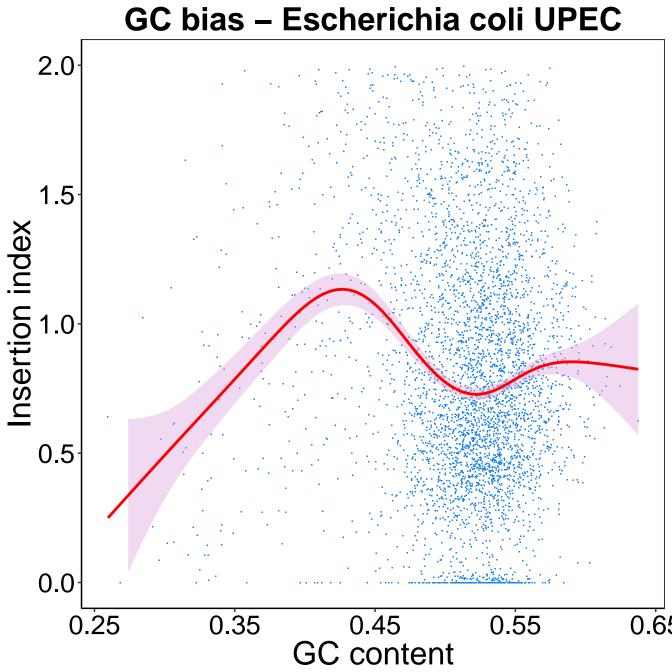
lebsiella pneumoniae RH201207 – normalise

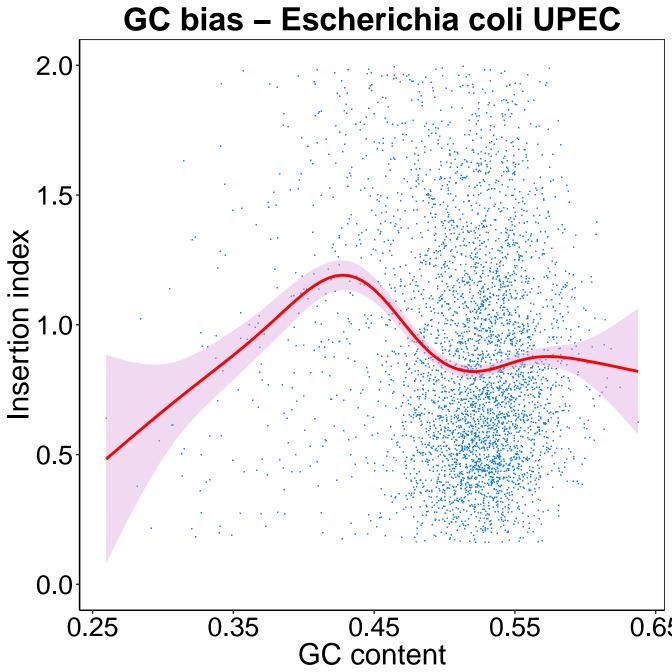




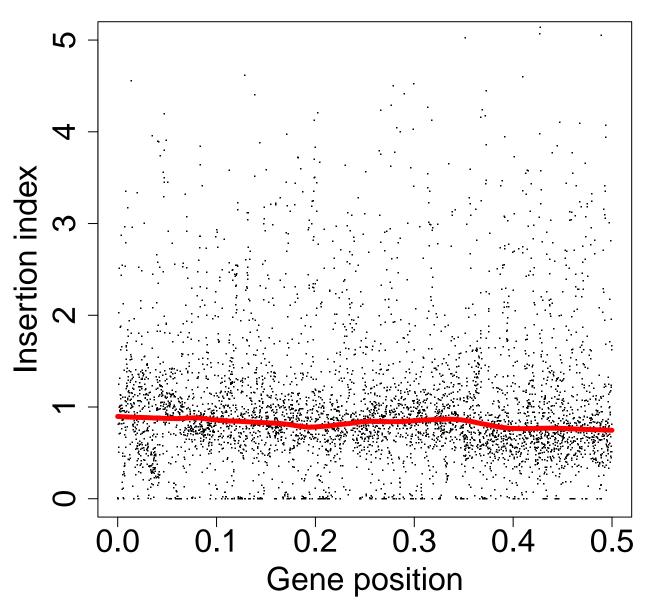
Distance bias - Escherichia coli UPEC



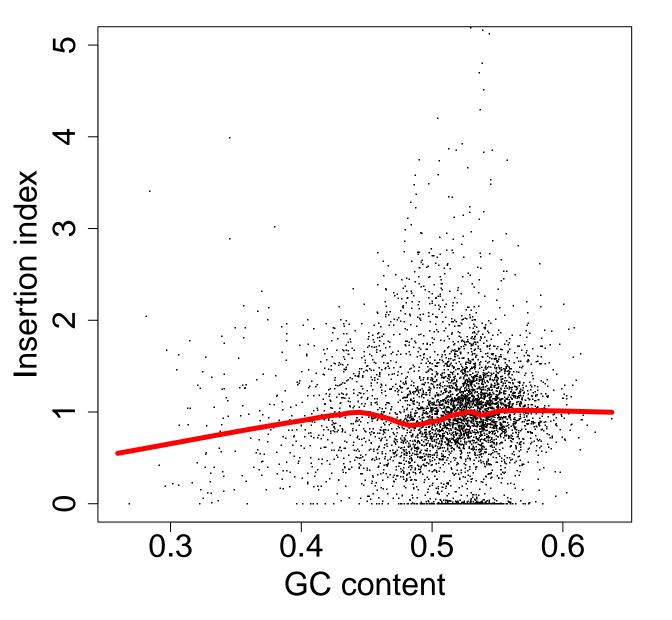


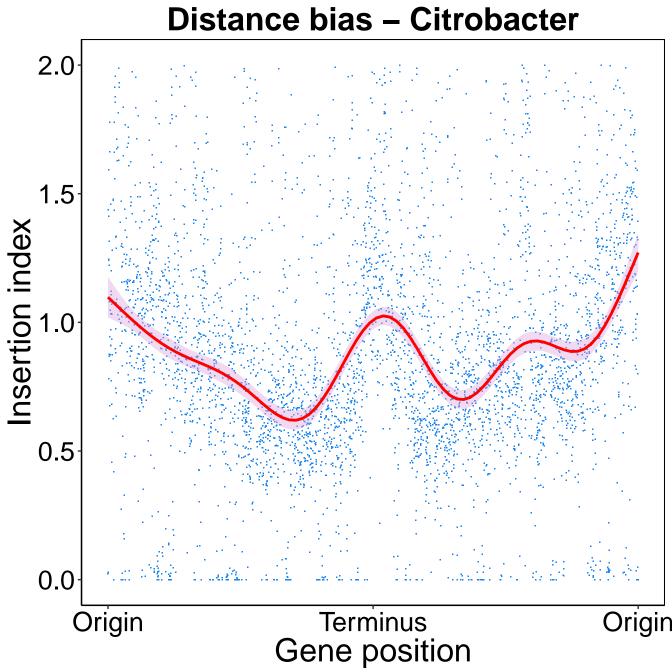


Escherichia coli UPEC - normalised distan

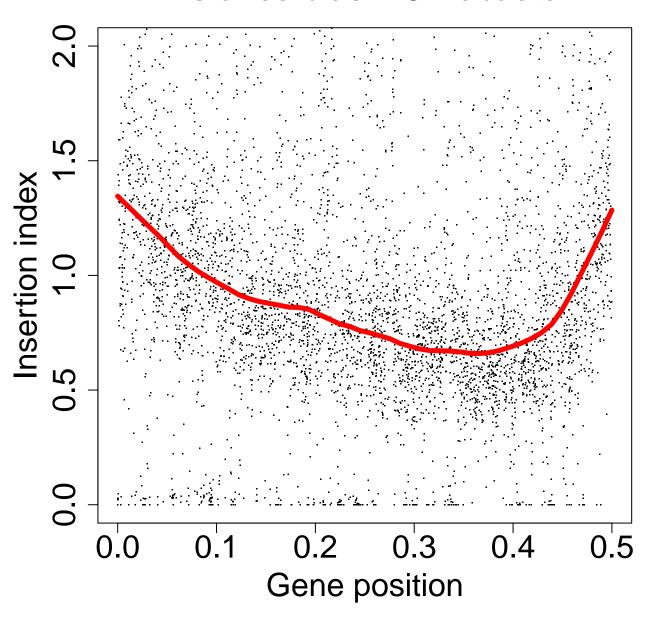


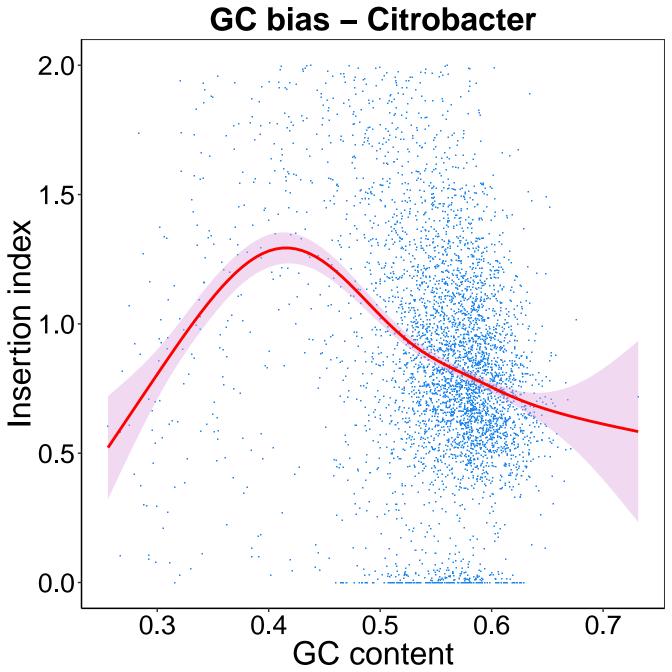
Escherichia coli UPEC – normalised GC

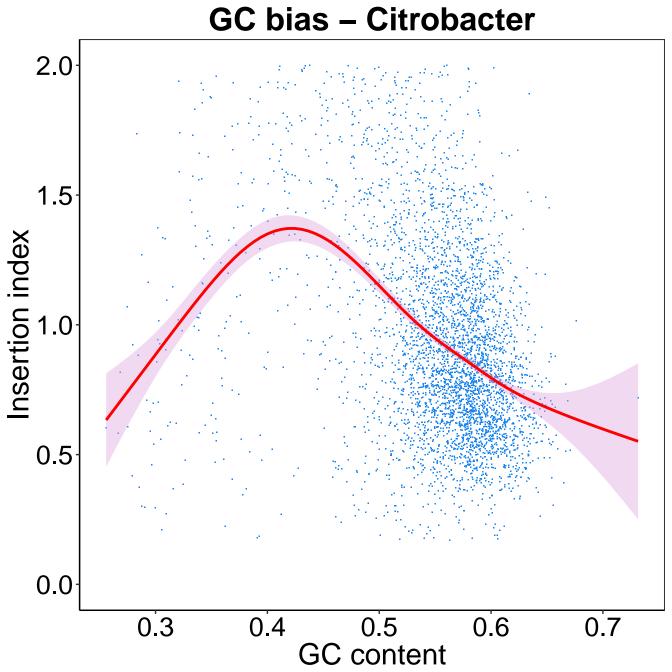




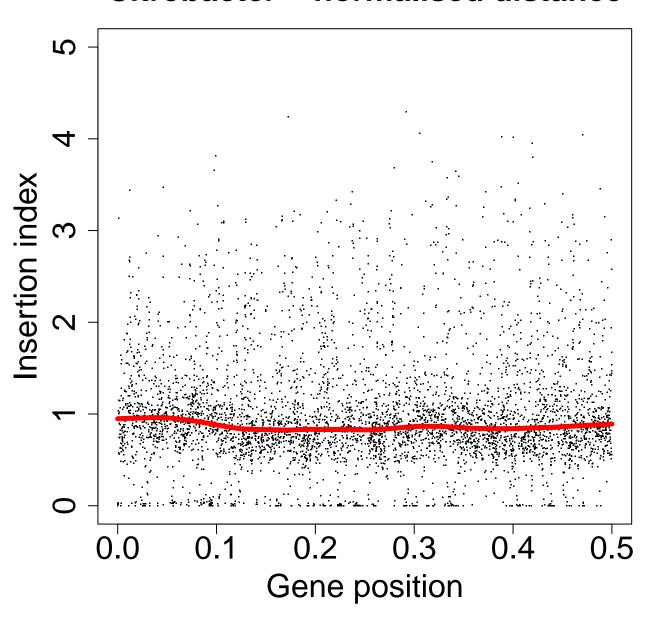
Distance bias - Citrobacter



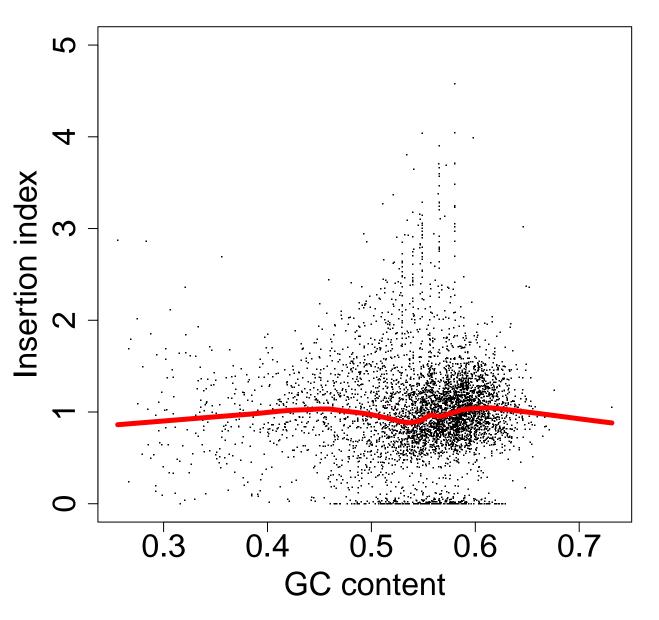


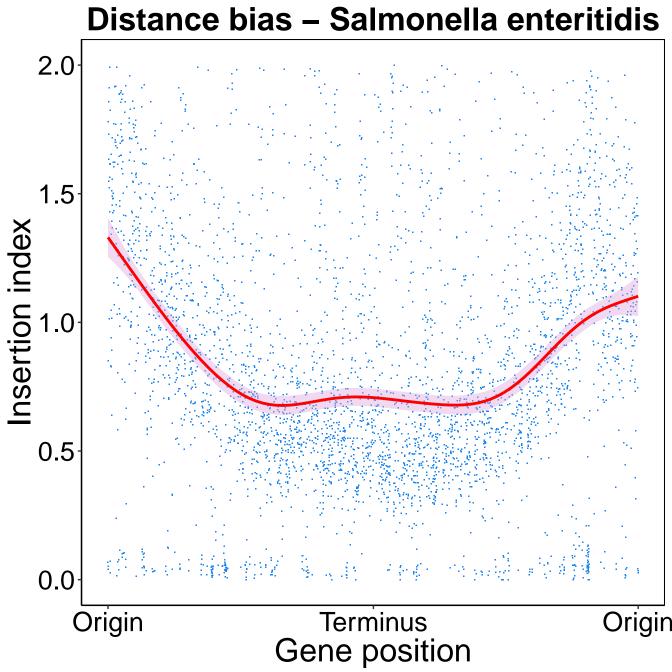


Citrobacter - normalised distance

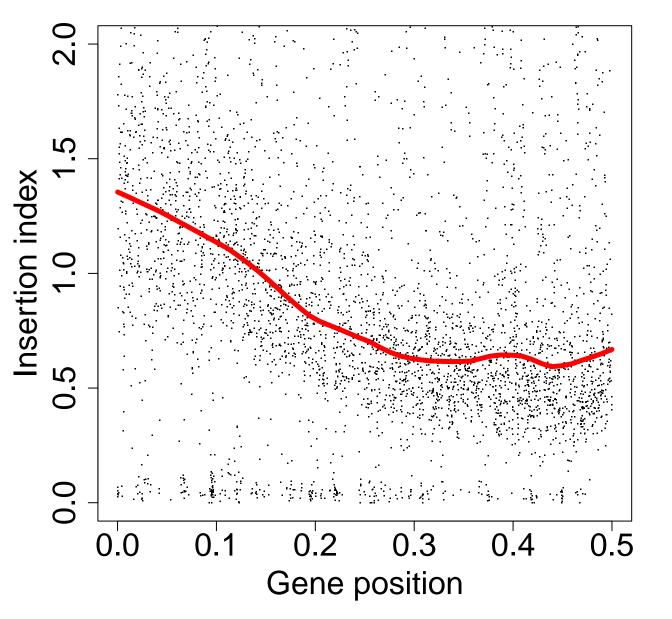


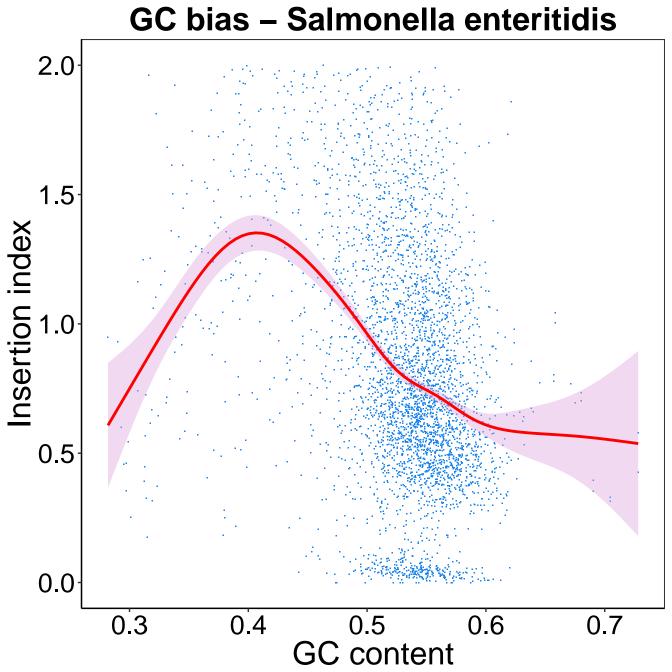
Citrobacter - normalised GC

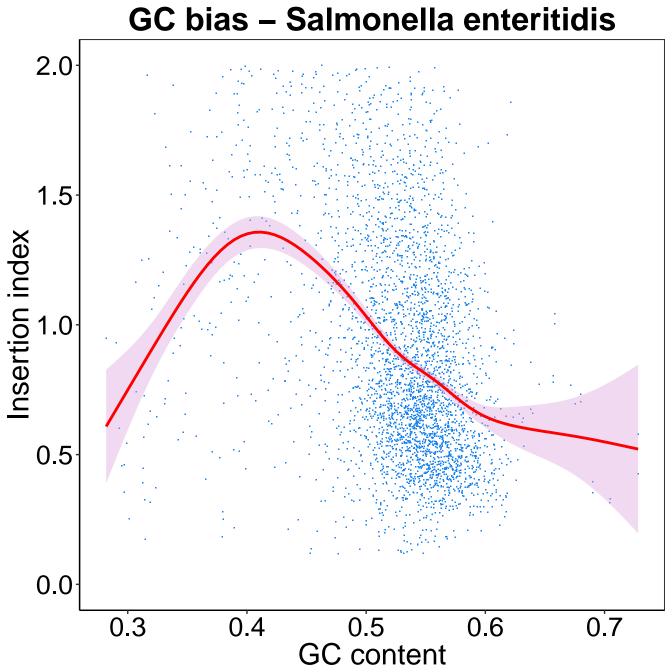




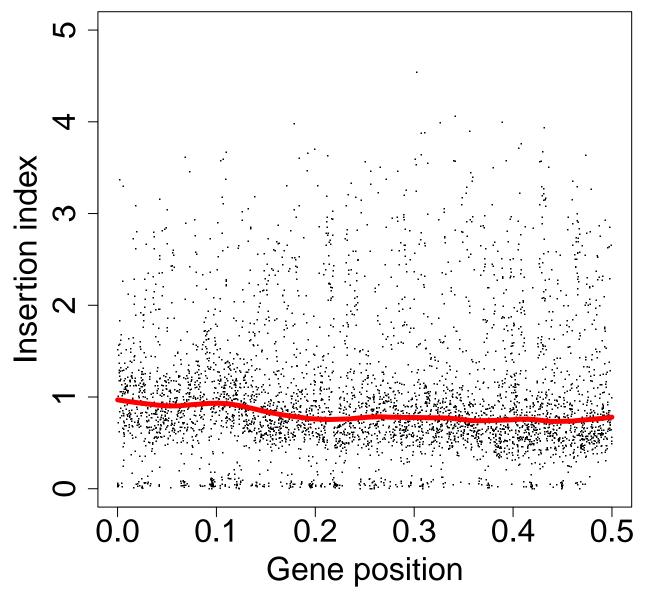
Distance bias - Salmonella enteritidis



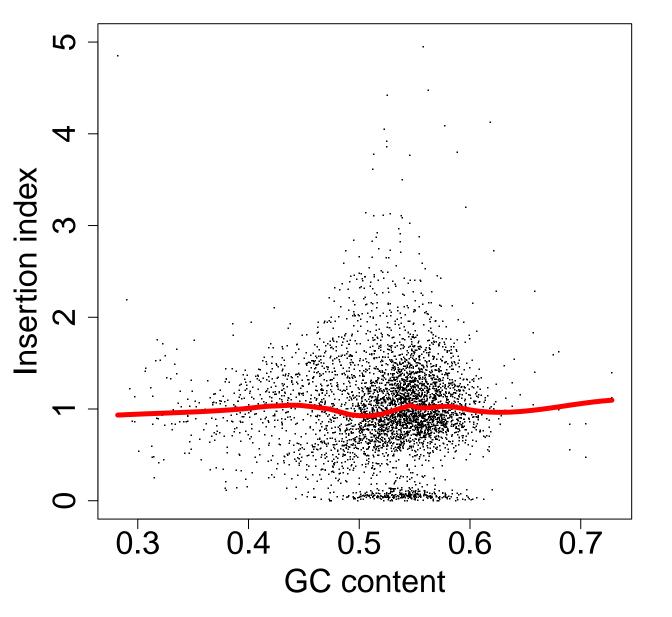


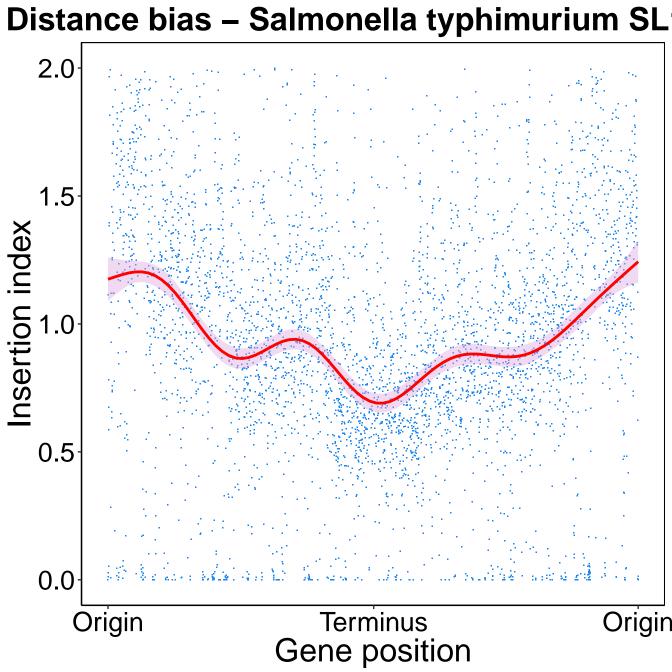


Salmonella enteritidis – normalised distan

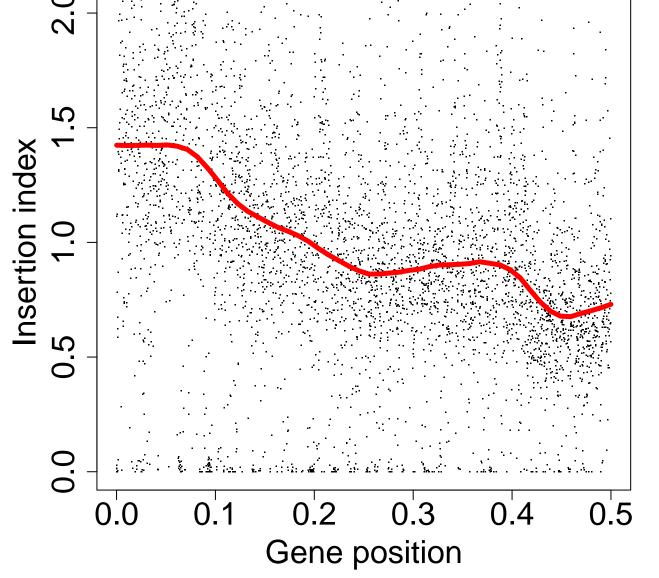


Salmonella enteritidis – normalised GC





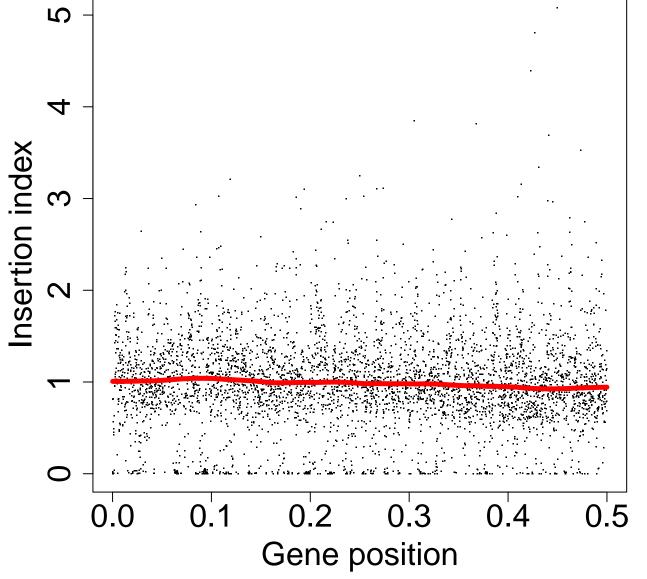
Distance bias – Salmonella typhimurium SL1



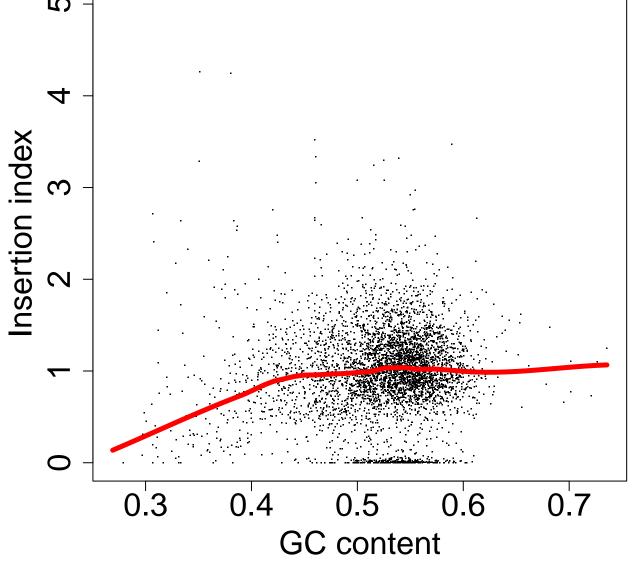
GC bias - Salmonella typhimurium SL134 2.0 1.5 Insertion index 0.5 0.0 0.4 0.3 0.5 0.6 0.7 GC content

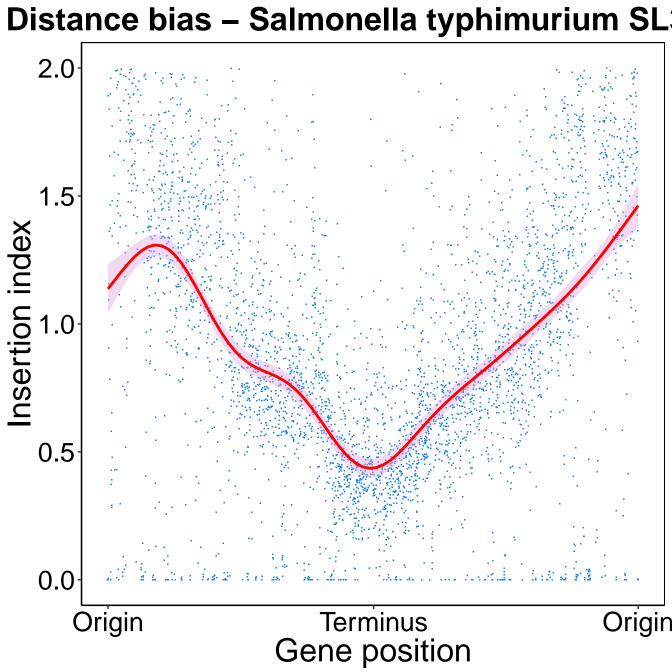
GC bias - Salmonella typhimurium SL134 2.0 1.5 Insertion index 1.00 0.5 0.0 0.4 0.6 0.7 0.3 0.5 GC content

monella typhimurium SL1344 – normalised d

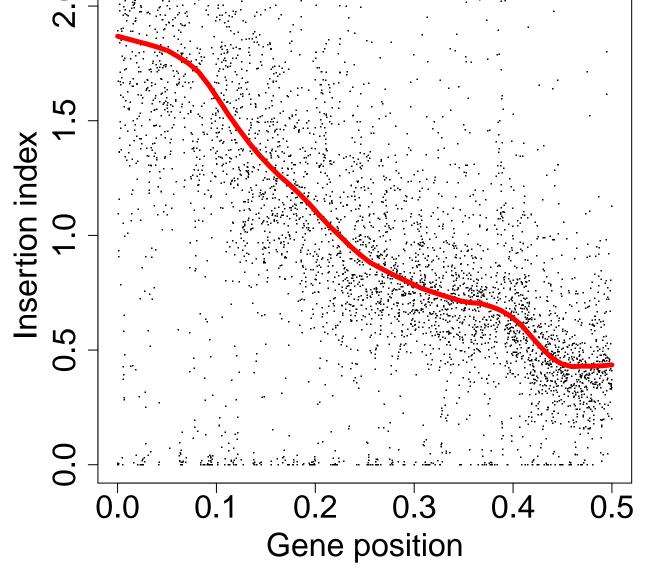


Salmonella typhimurium SL1344 – normalised ഹ –





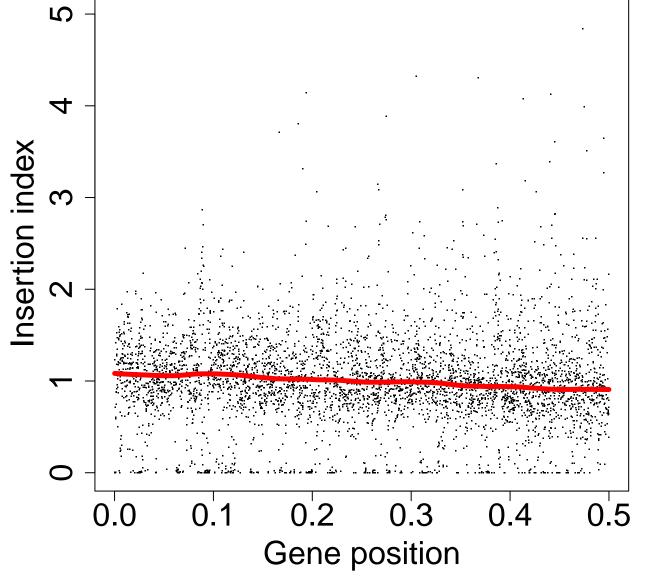
Distance bias – Salmonella typhimurium SL3



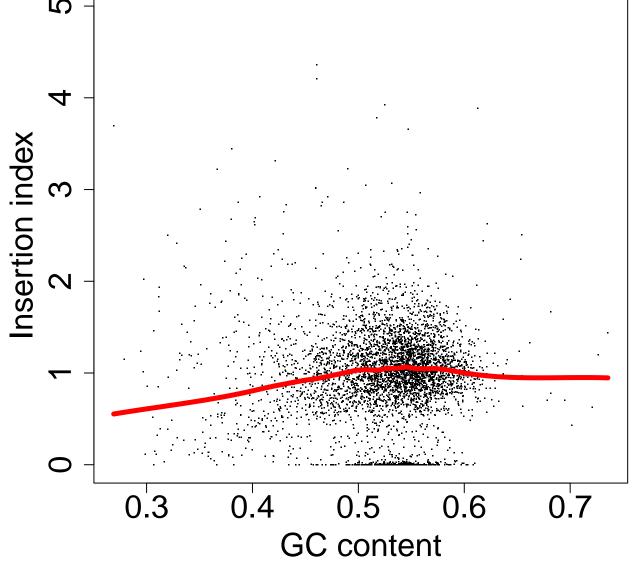
GC bias - Salmonella typhimurium SL326 2.0 1.5 Insertion index 0.5 0.0 0.3 0.4 0.5 0.6 0.7 GC content

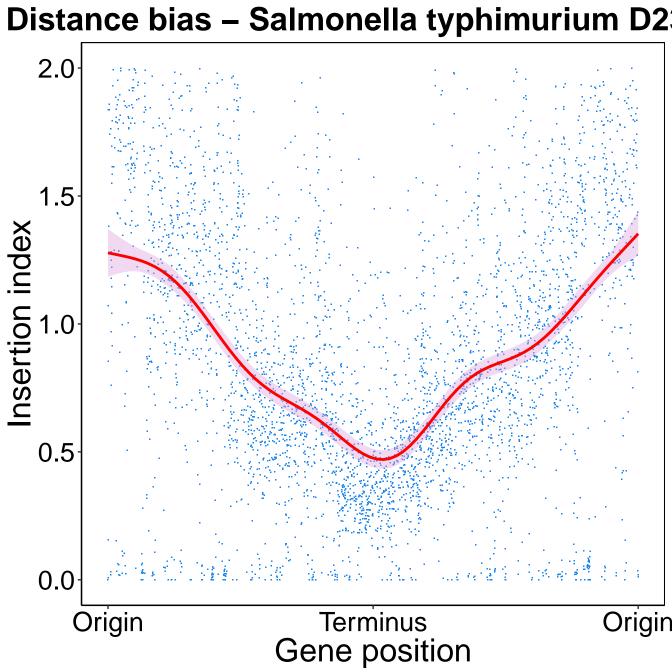
GC bias - Salmonella typhimurium SL326 2.0 1.5 Insertion index 1.00 0.5 0.0 0.5 0.4 0.6 0.7 0.3 GC content

monella typhimurium SL3261 – normalised d

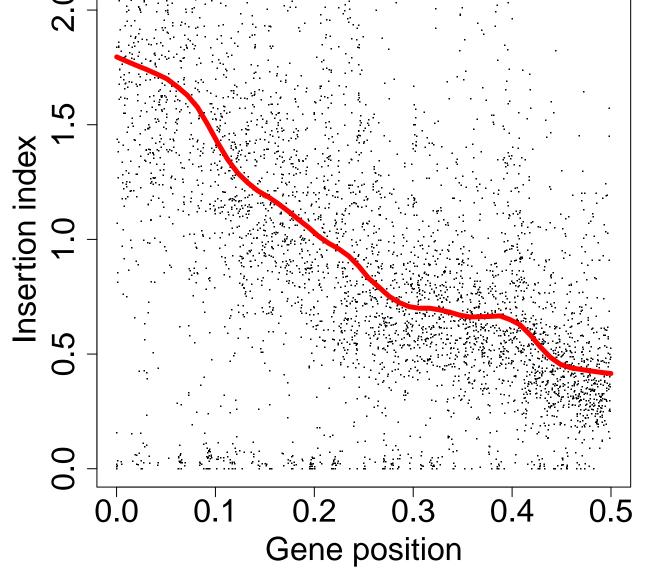


Salmonella typhimurium SL3261 – normalised ഹ –





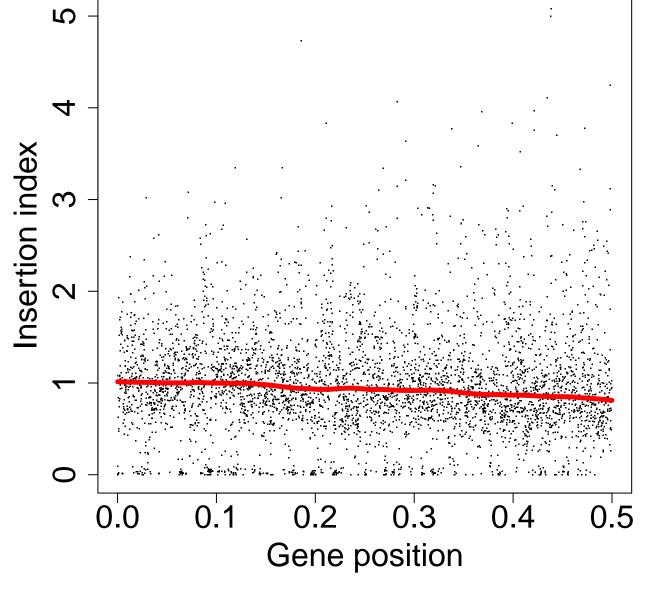
Distance bias – Salmonella typhimurium D23



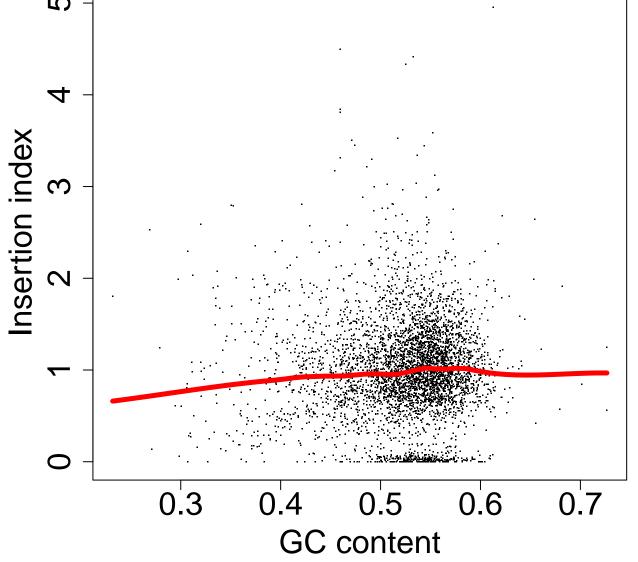
GC bias - Salmonella typhimurium D2358 2.0 1.5 Insertion index 0.5 0.0 0.3 0.4 0.5 0.6 GC content

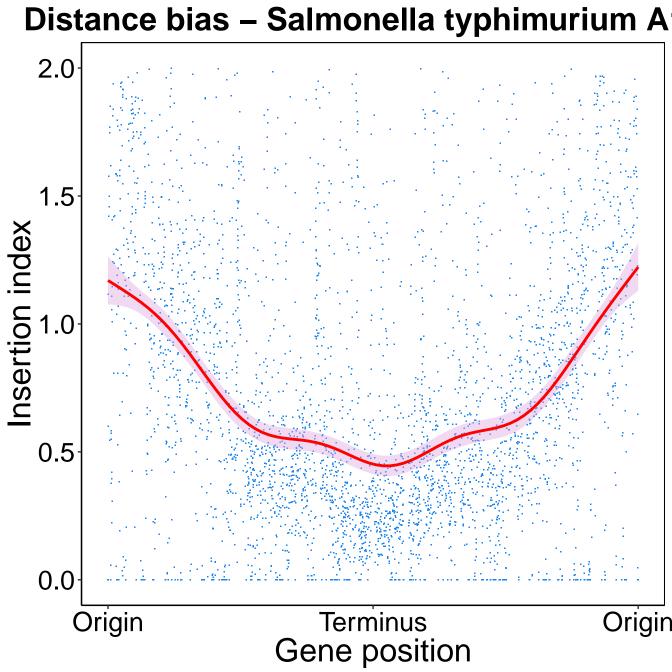
GC bias - Salmonella typhimurium D2358 2.0 1.5 Insertion index 1.00 0.5 0.0 0.3 0.4 0.5 0.6 0.7 GC content

monella typhimurium D23580 – normalised d

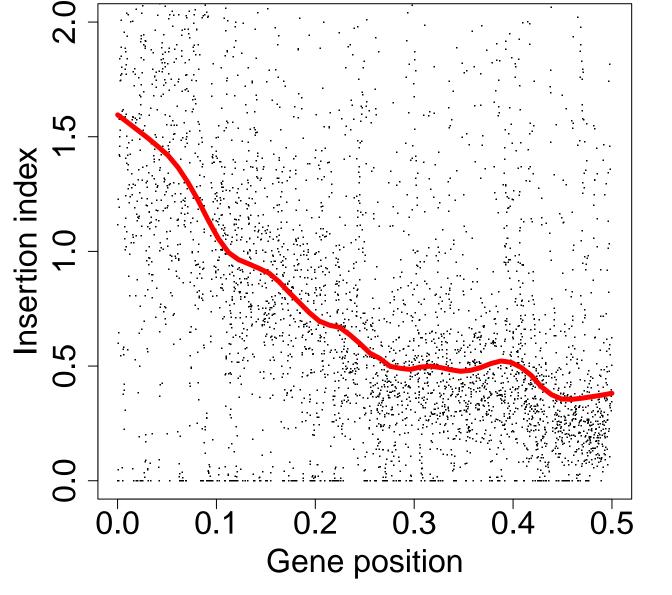


Salmonella typhimurium D23580 – normalised ഹ –





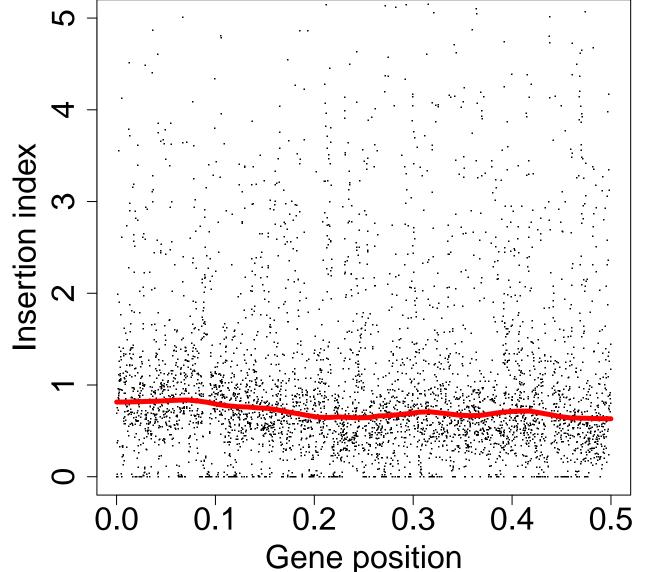
Distance bias – Salmonella typhimurium A1



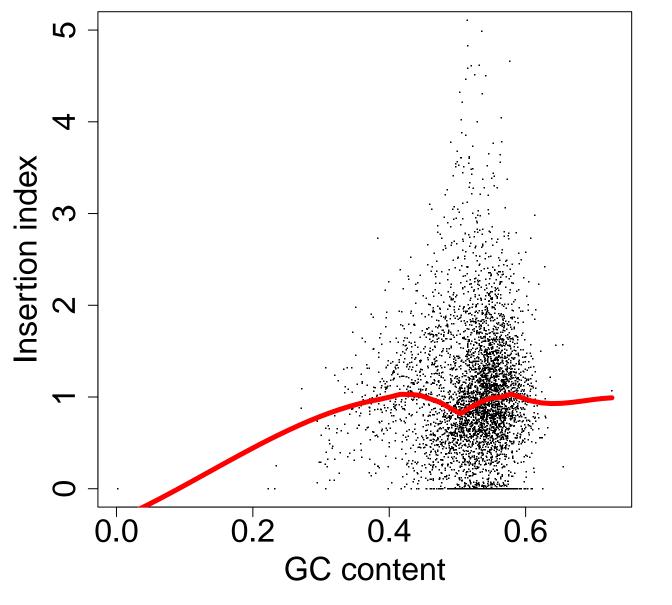
GC bias - Salmonella typhimurium A130 2.0 1.5 Insertion index 0.5 0.0 0.2 0.0 0.4 0.6 GC content

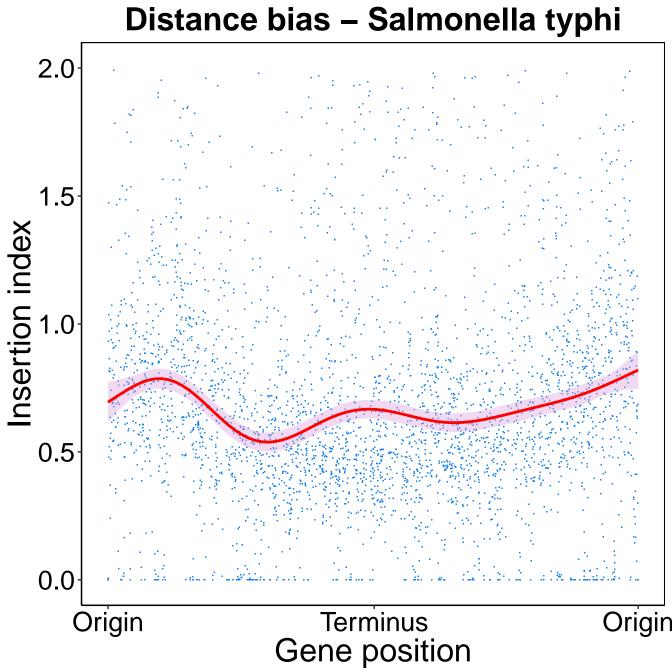
GC bias - Salmonella typhimurium A130 2.0 1.5 Insertion index 1.00 0.5 0.0 0.3 0.4 0.5 0.6 0.7 GC content

Imonella typhimurium A130 – normalised dis

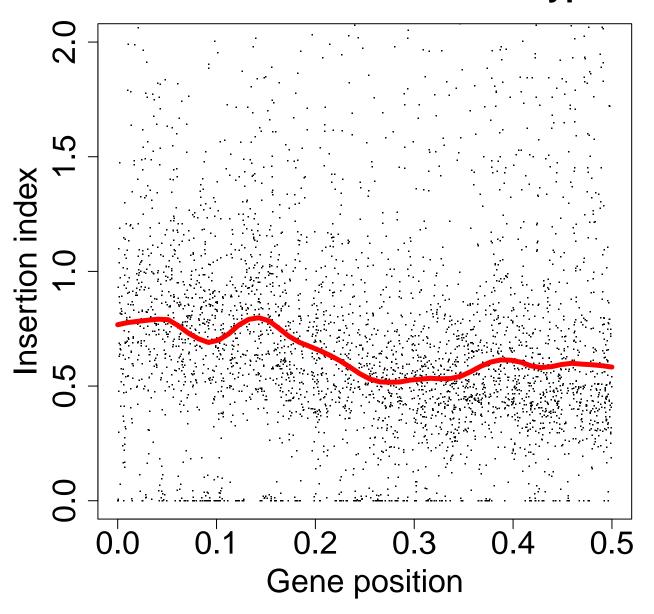


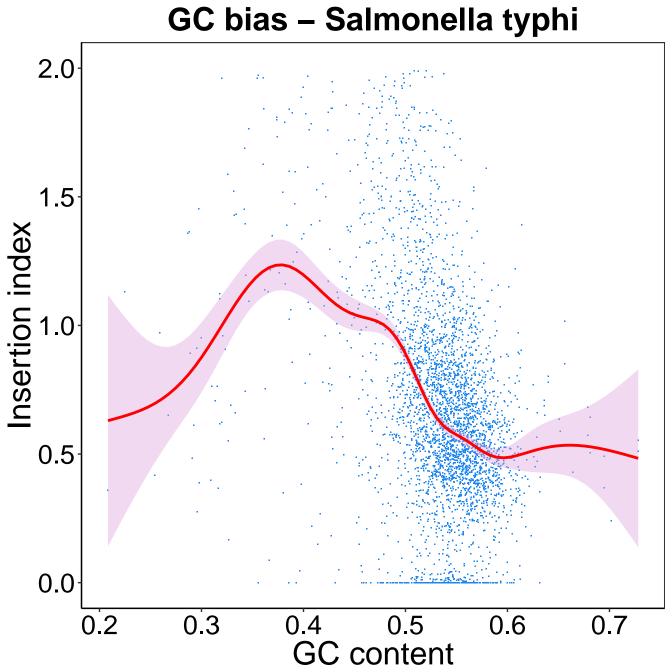
Salmonella typhimurium A130 – normalised

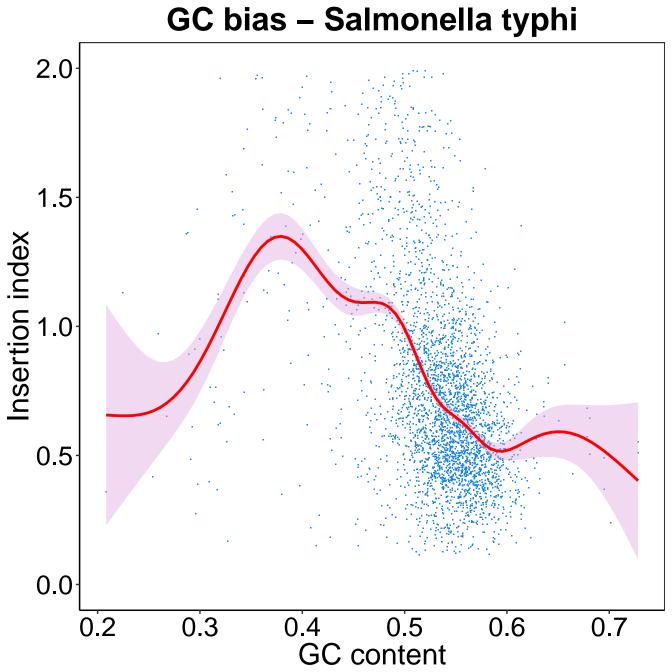




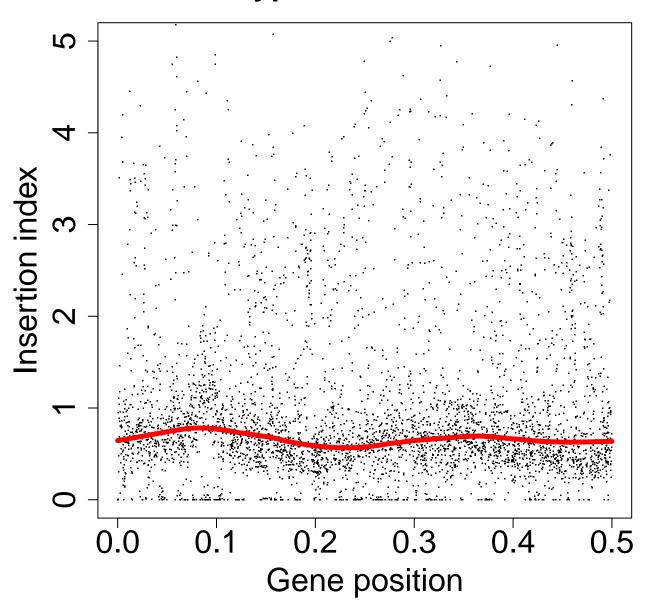
Distance bias - Salmonella typhi



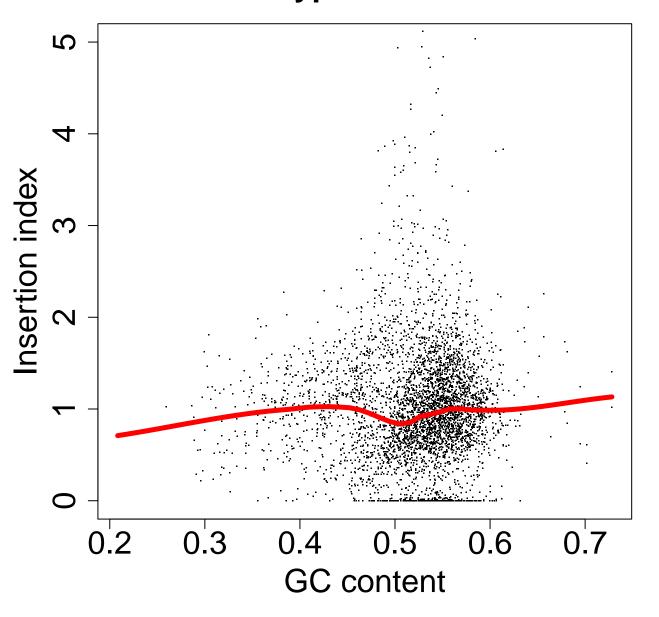




Salmonella typhi – normalised distance



Salmonella typhi – normalised GC



GC bias

