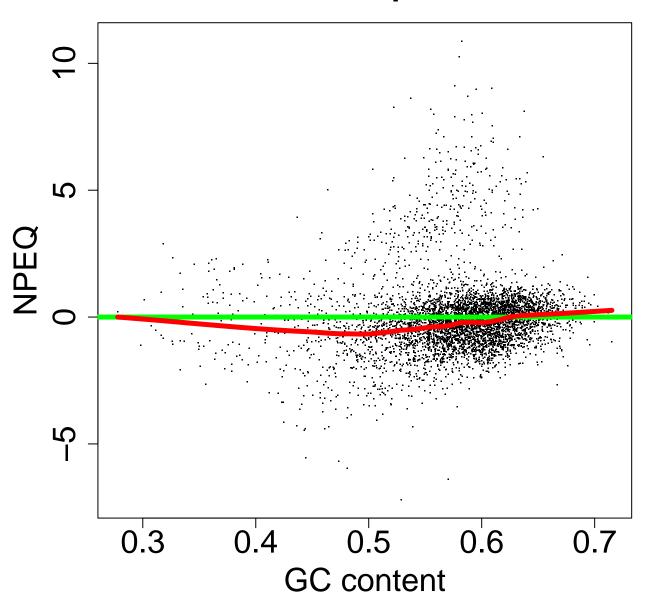
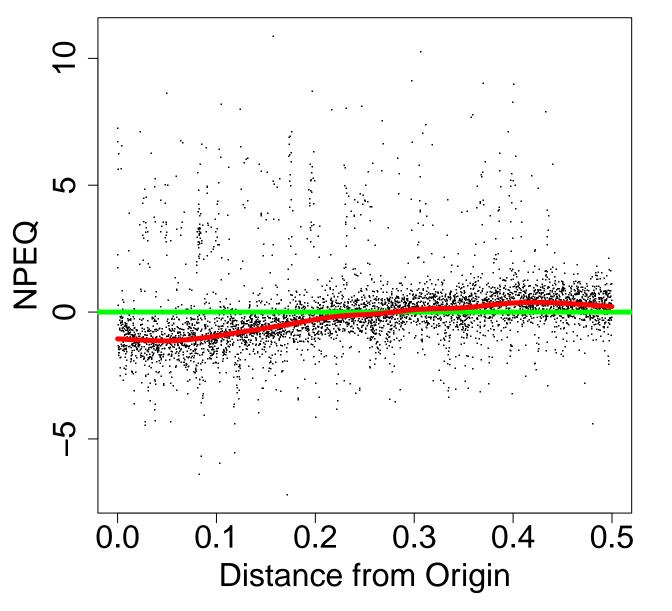
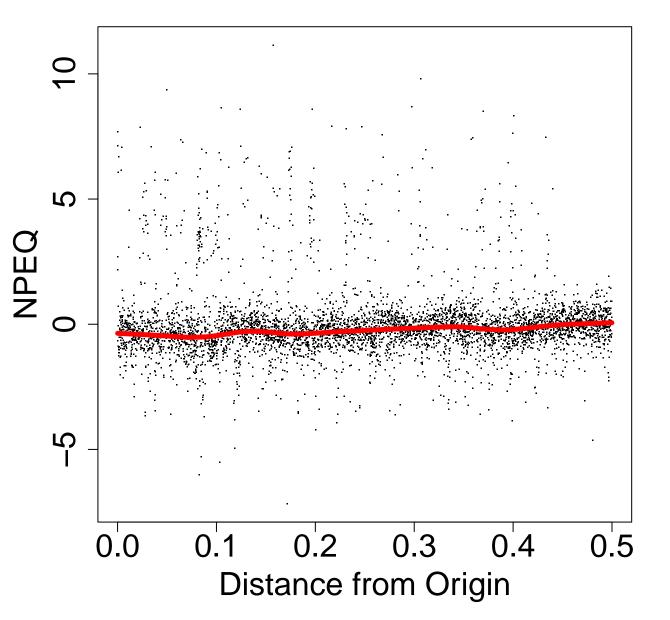
# GC bias - Klebsiella pneumoniae Ecl8



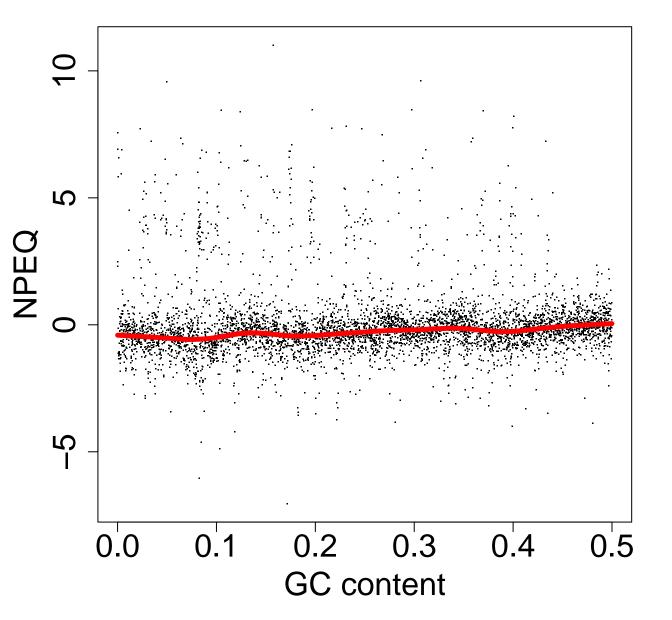
# Distance bias – Klebsiella pneumoniae Ecl8



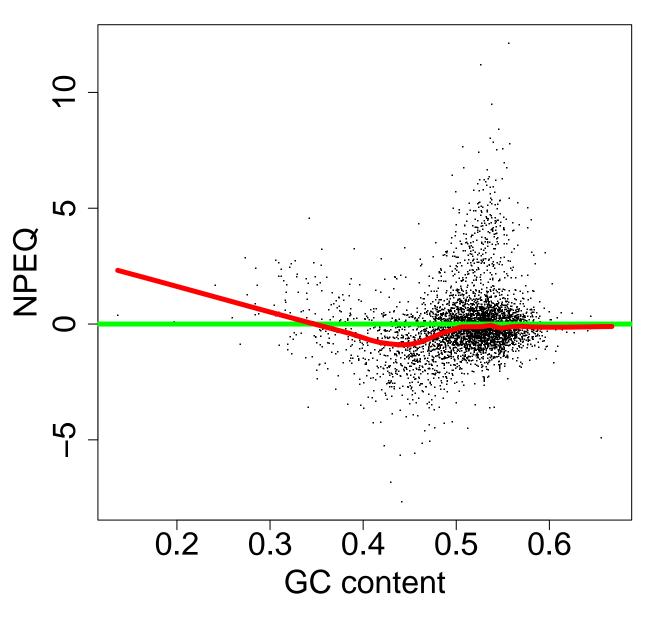
#### **BN373 – normalised distance**



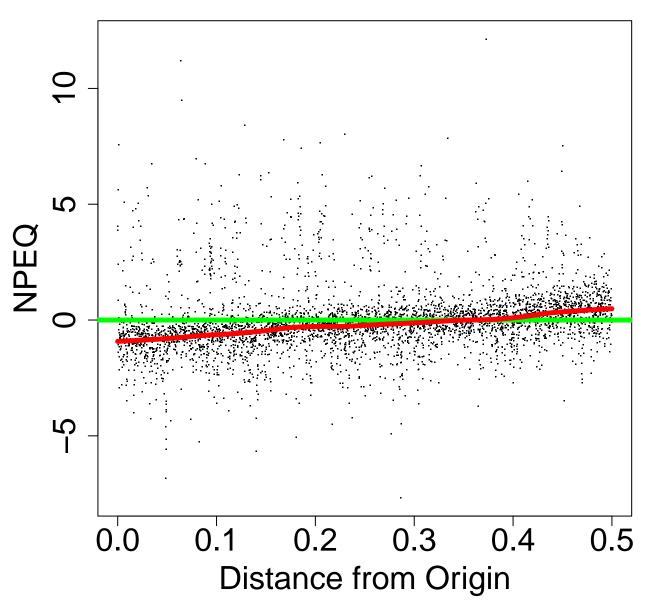
#### **BN373 - normalised GC bias**



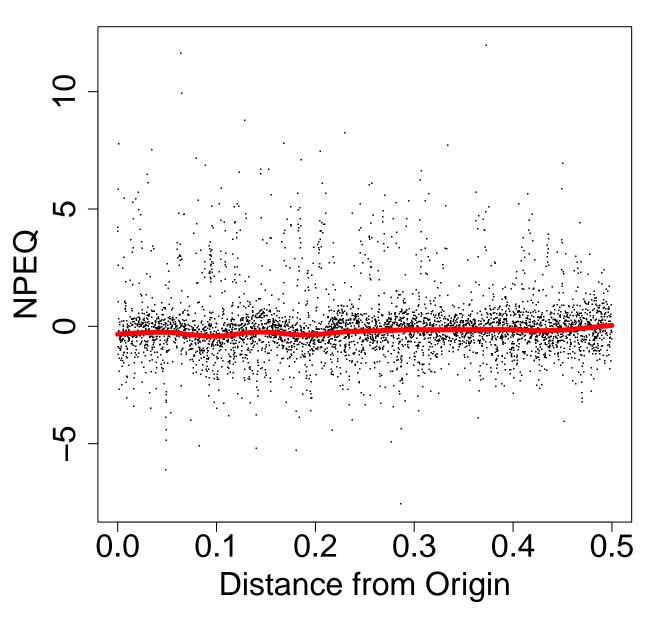
# GC bias - Escherichia coli BW25113



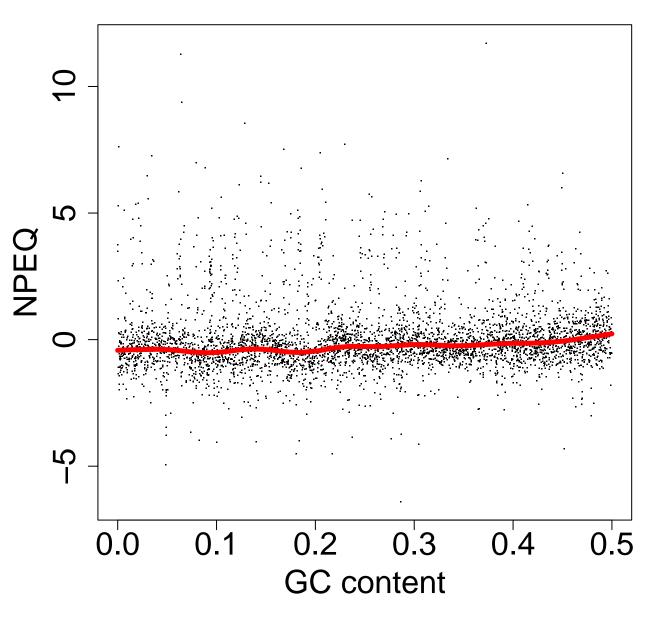
#### Distance bias - Escherichia coli BW25113



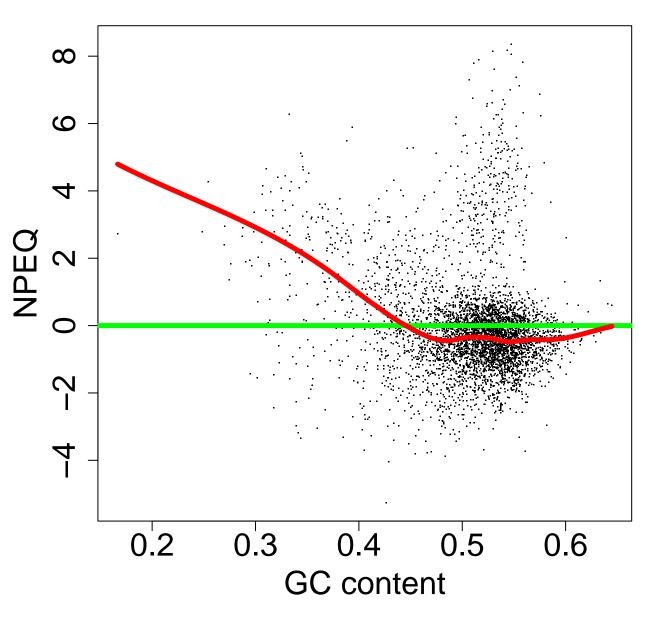
#### **BW25113 – normalised distance**



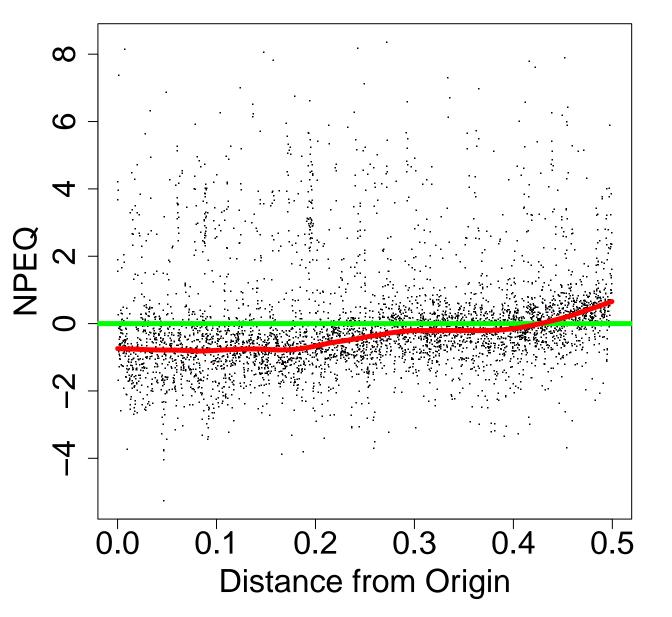
#### **BW25113 - normalised GC bias**



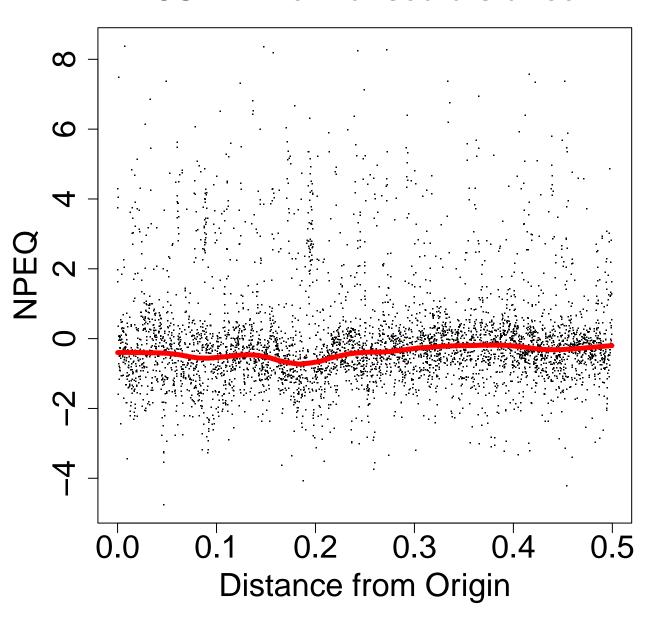
# GC bias - Escherichia coli ETEC CS17



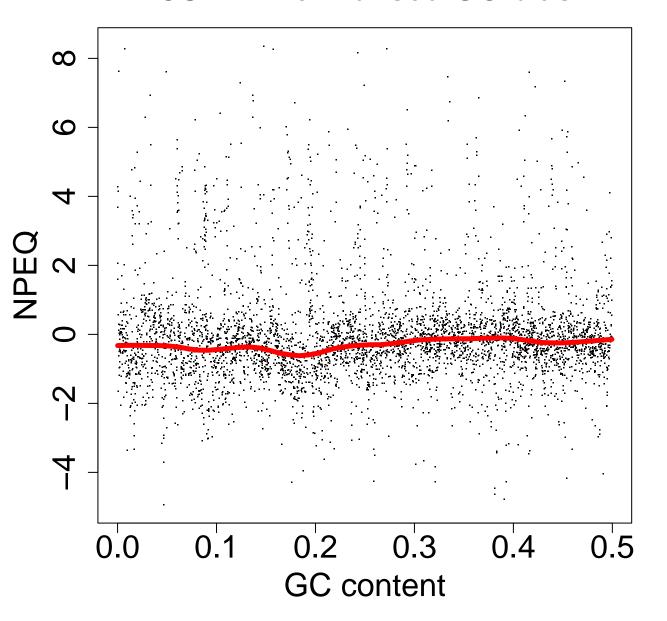
# Distance bias - Escherichia coli ETEC CS17



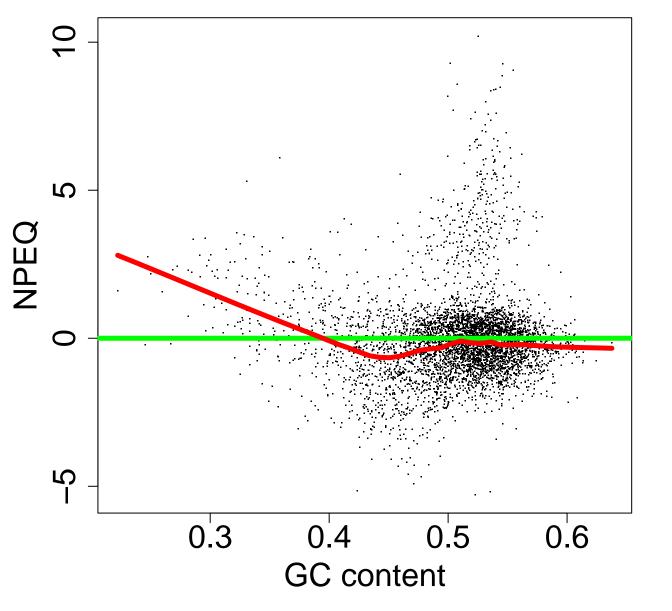
#### **CS17 – normalised distance**



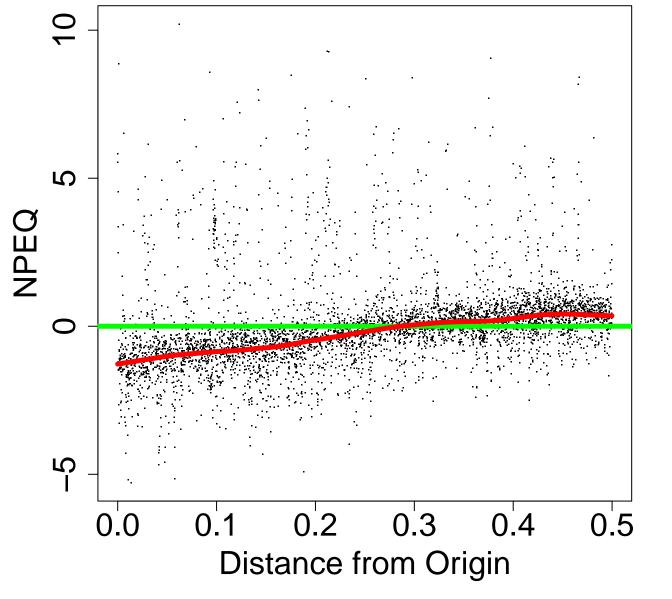
#### CS17 - normalised GC bias



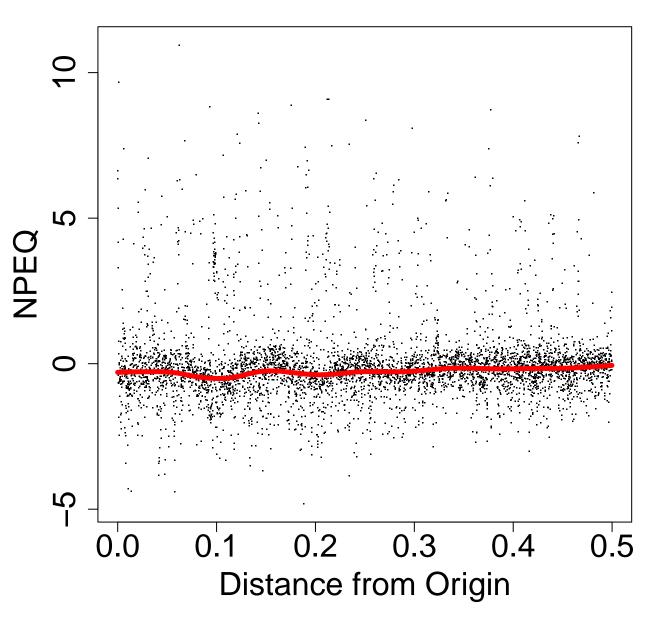
# GC bias – Escherichia coli ST131 EC958



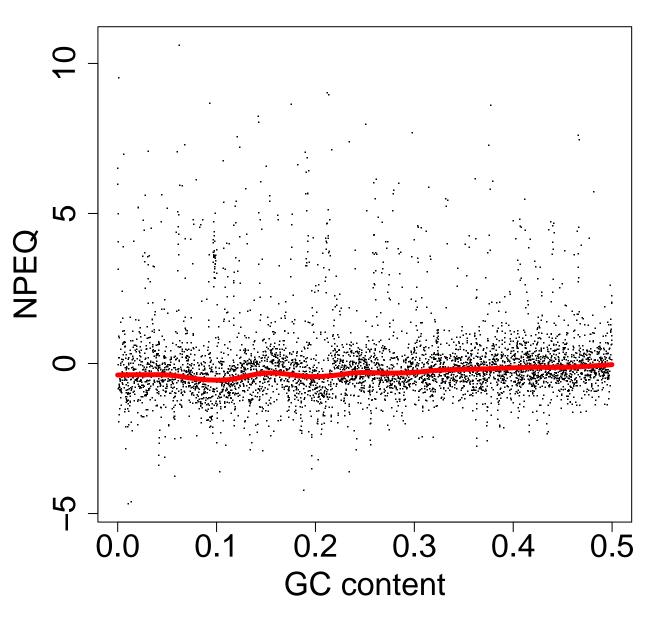
# Distance bias – Escherichia coli ST131 EC958



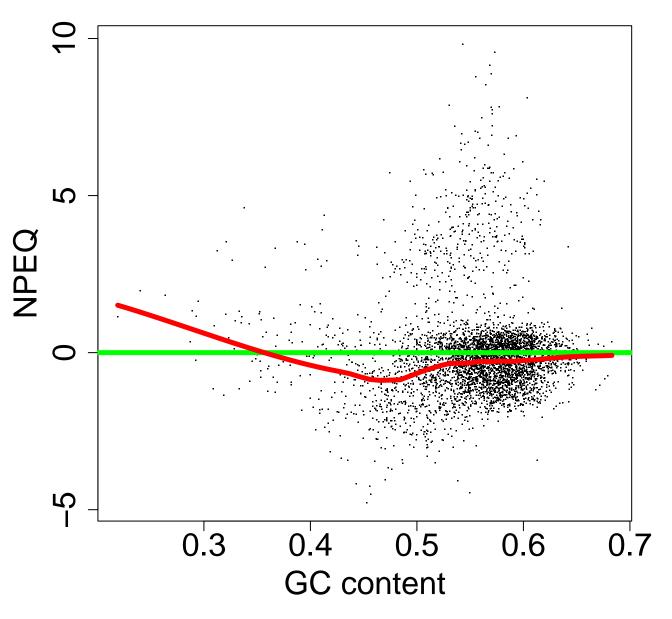
#### EC958 - normalised distance



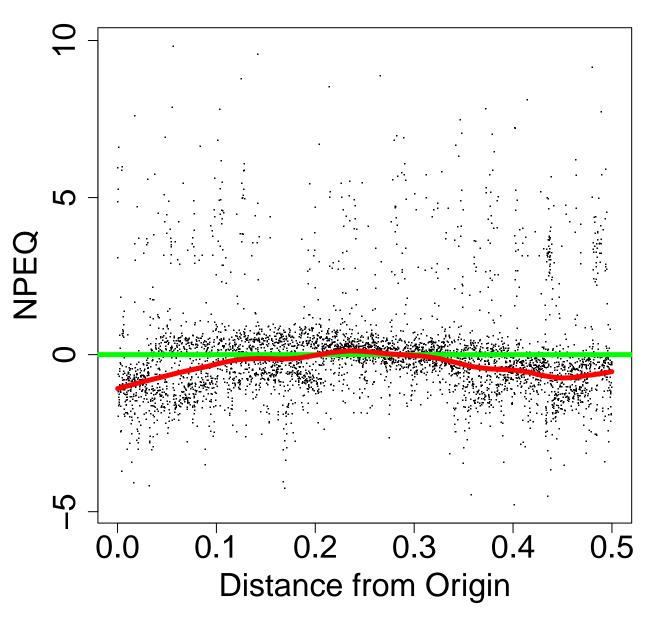
# EC958 - normalised GC bias



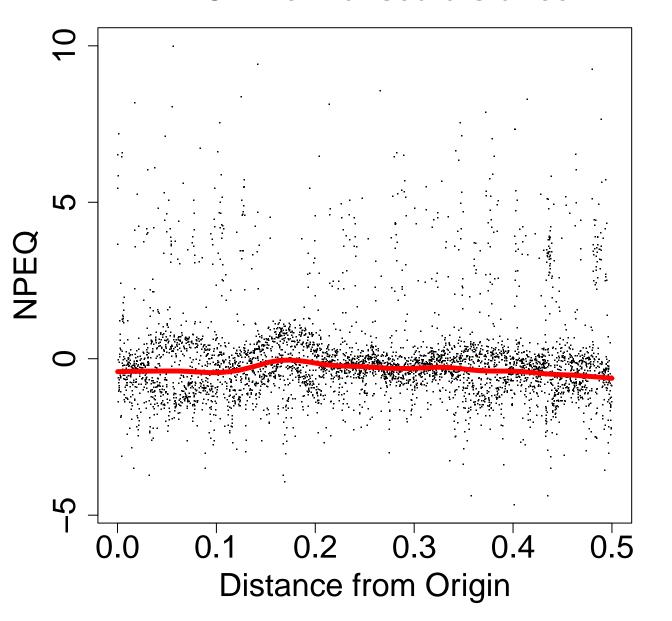
## GC bias - Enterobacter



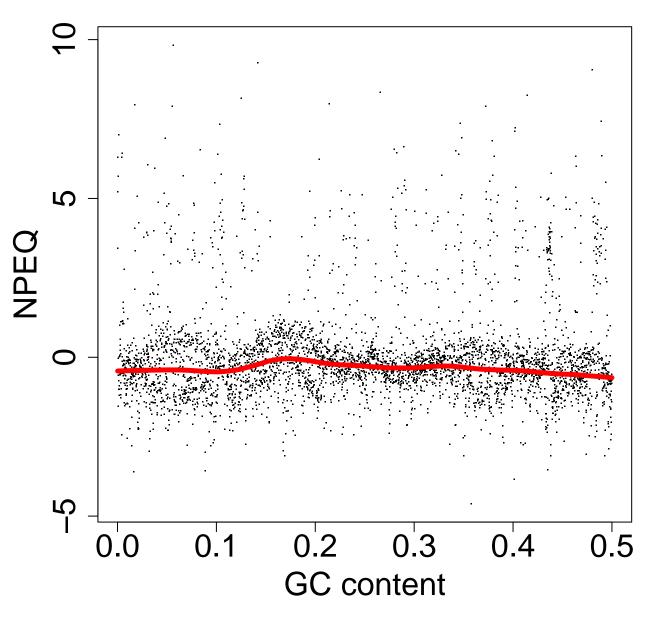
#### **Distance bias – Enterobacter**



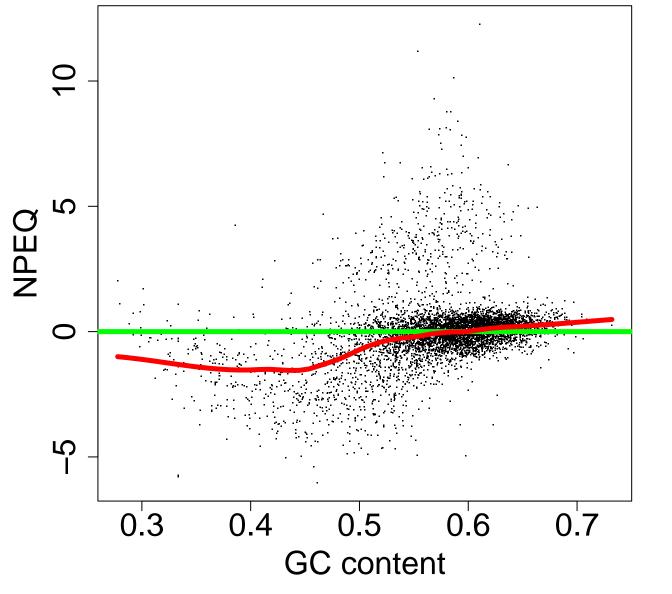
## **ENC** – normalised distance



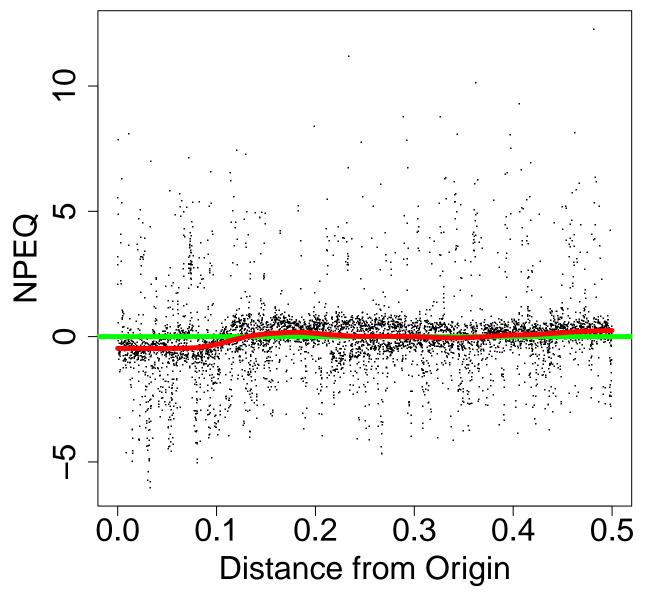
#### **ENC – normalised GC bias**



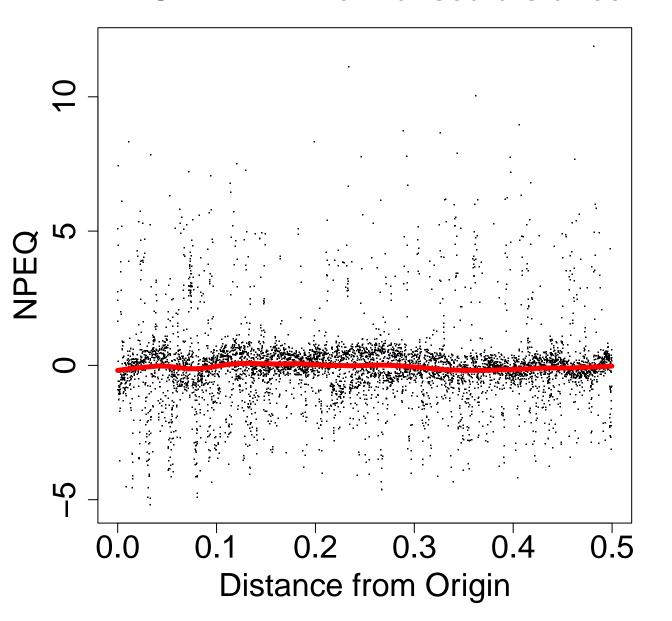
# GC bias – Klebsiella pneumoniae RH20120



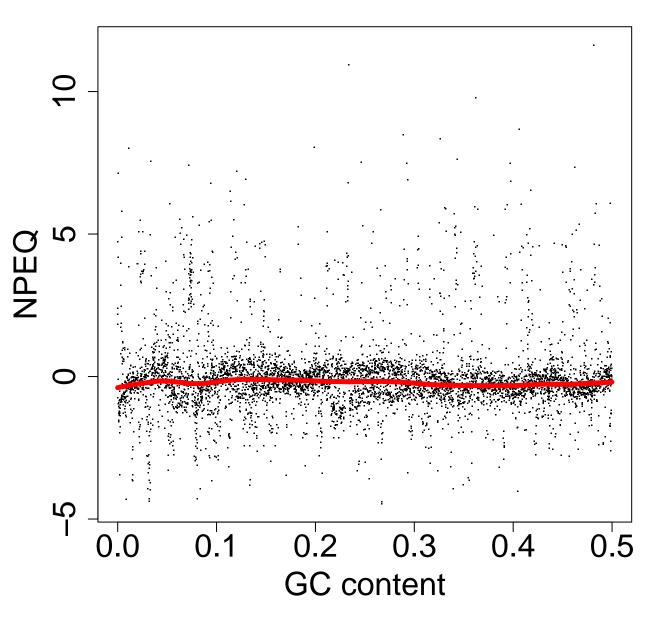
# Distance bias – Klebsiella pneumoniae RH2012



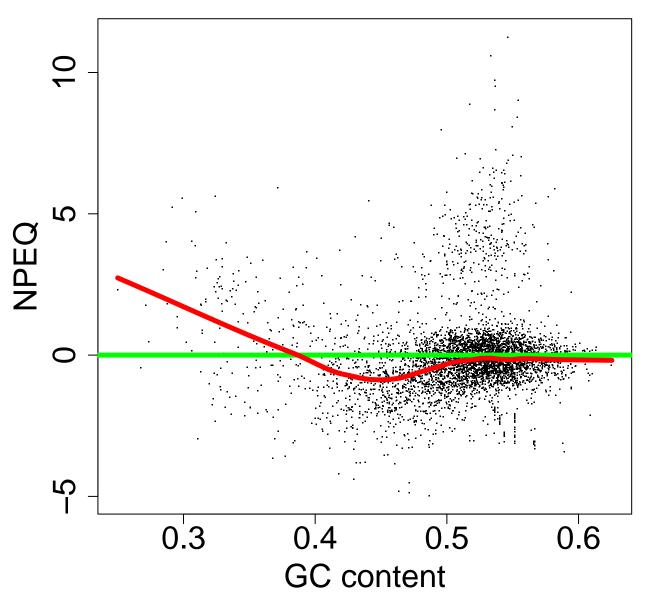
#### ERS227112 - normalised distance



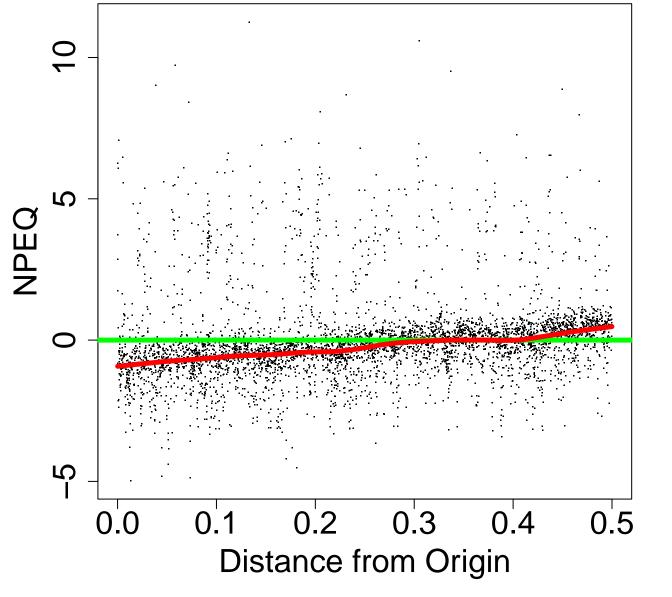
## **ERS227112 – normalised GC bias**



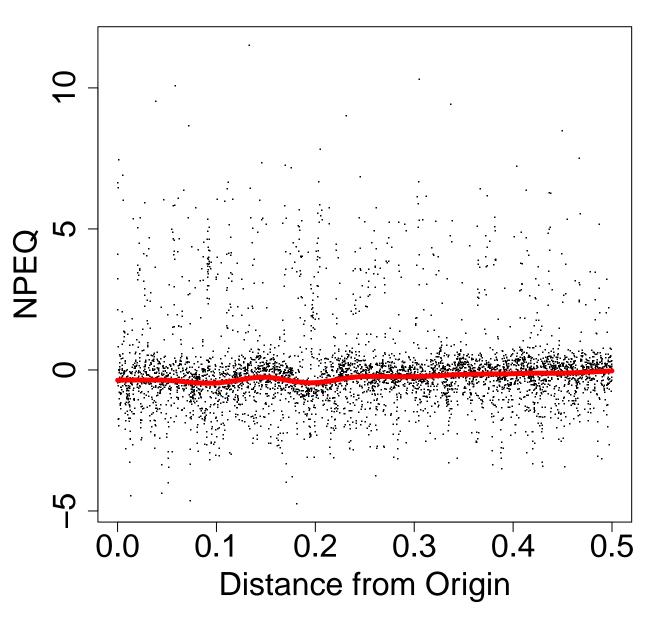
# GC bias – Escherichia coli ETEC H10407



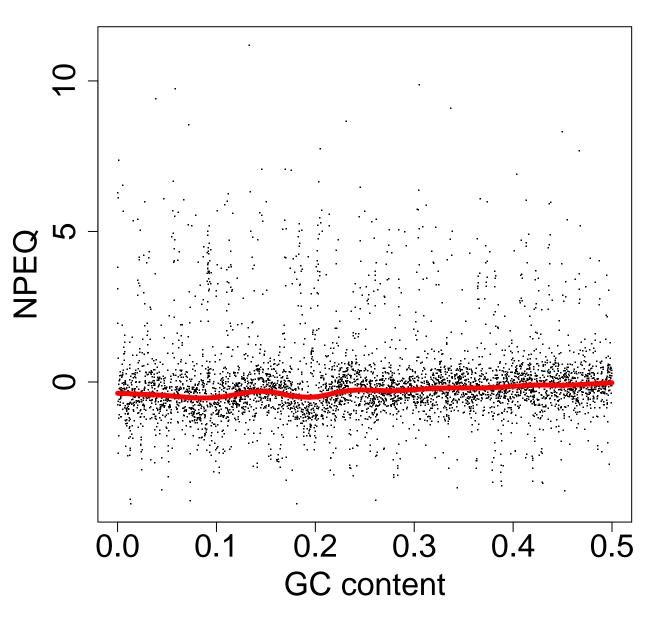
# Distance bias – Escherichia coli ETEC H1040



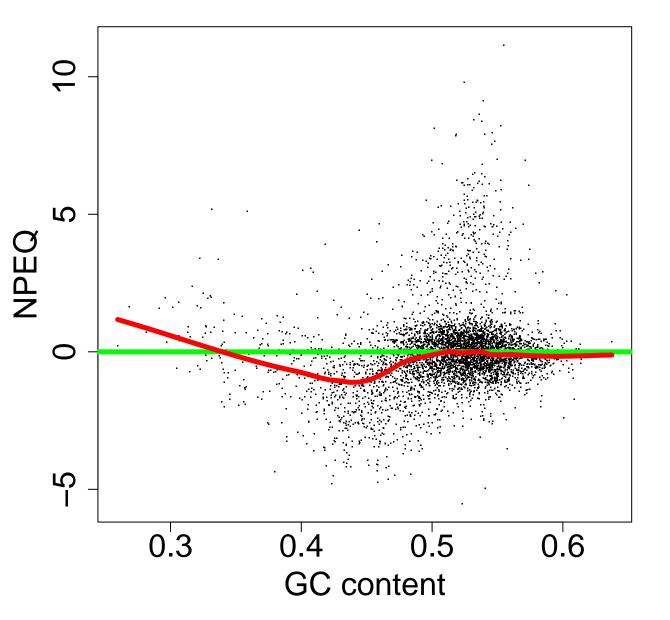
#### **ETEC – normalised distance**



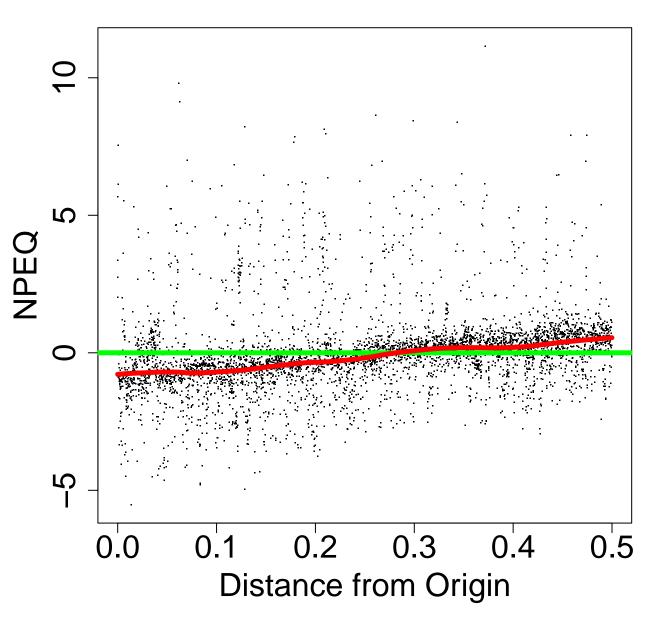
#### **ETEC – normalised GC bias**



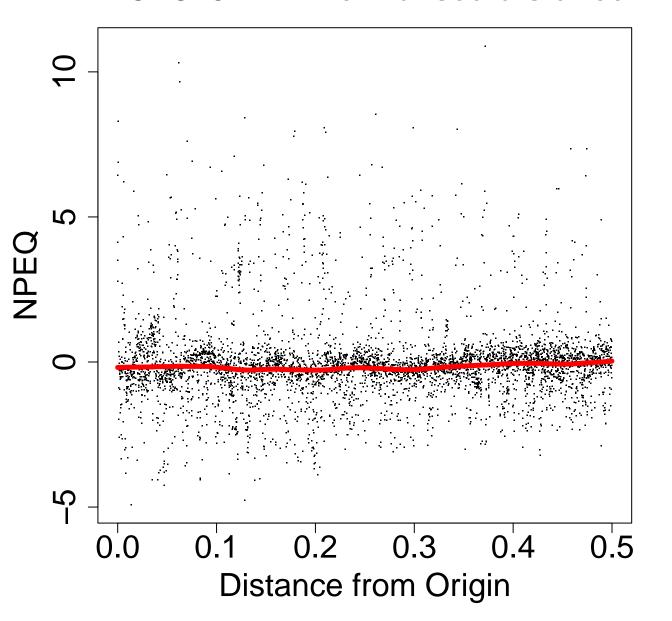
## GC bias - Escherichia coli UPEC



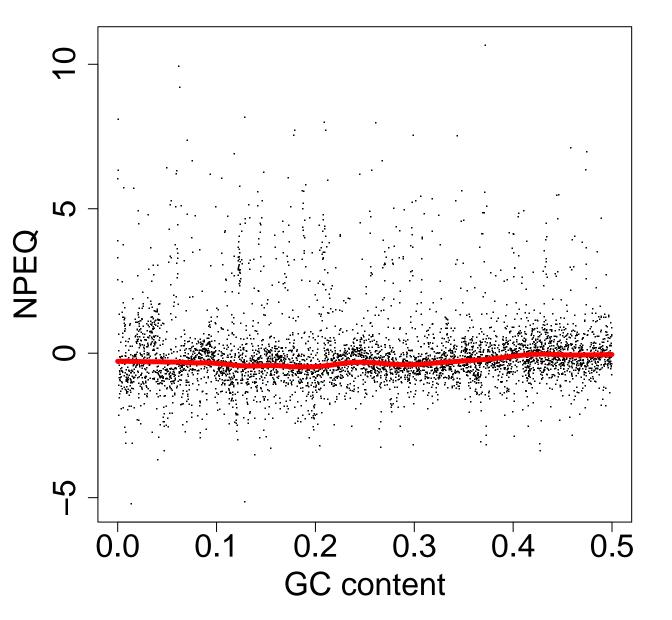
#### Distance bias - Escherichia coli UPEC



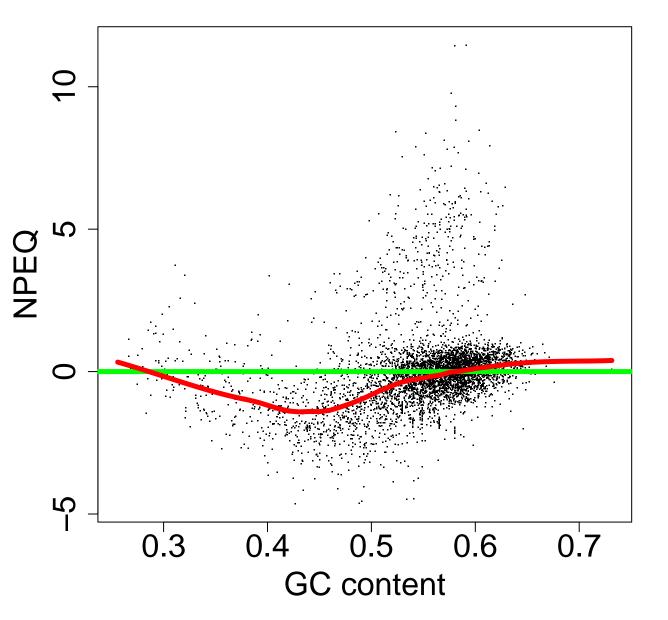
#### NCTC13441 - normalised distance



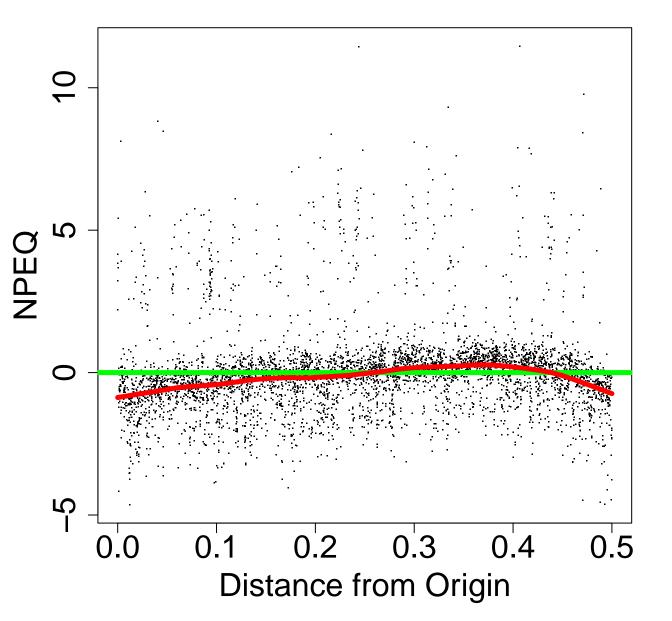
# NCTC13441 - normalised GC bias



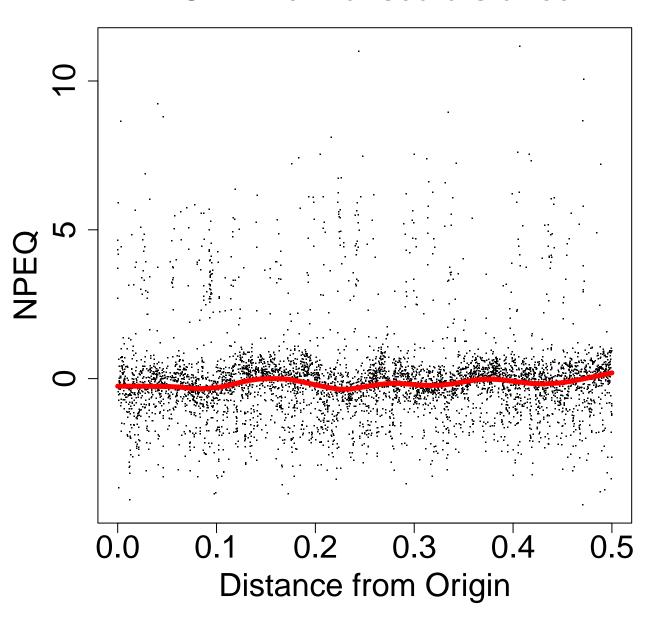
## GC bias - Citrobacter



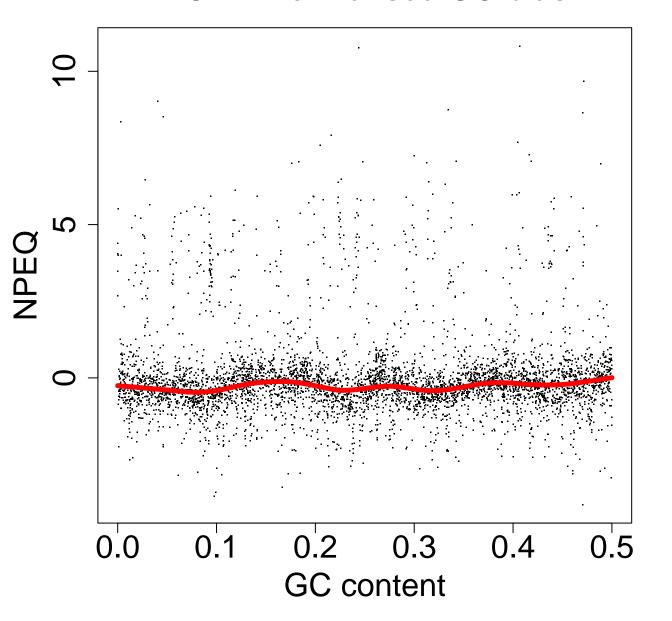
#### **Distance bias – Citrobacter**



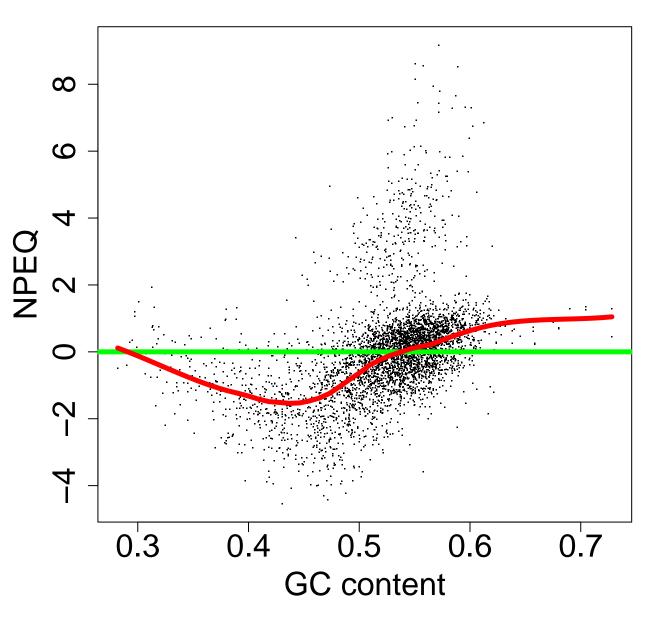
#### **ROD – normalised distance**



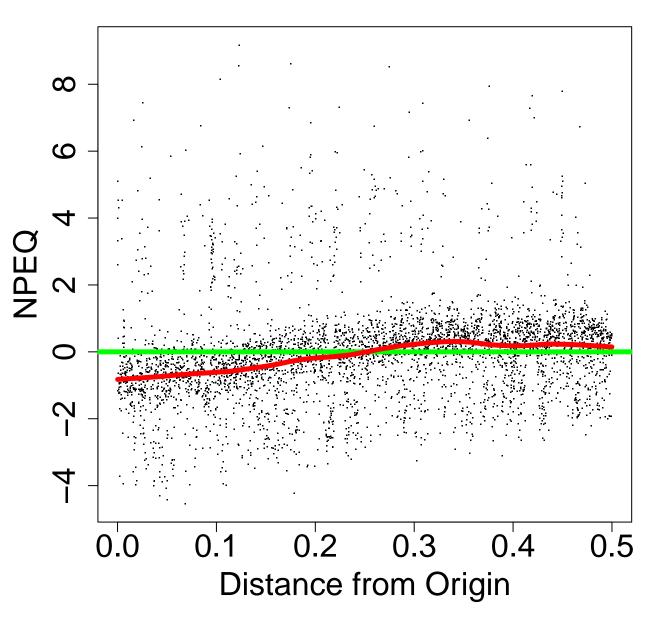
#### **ROD – normalised GC bias**



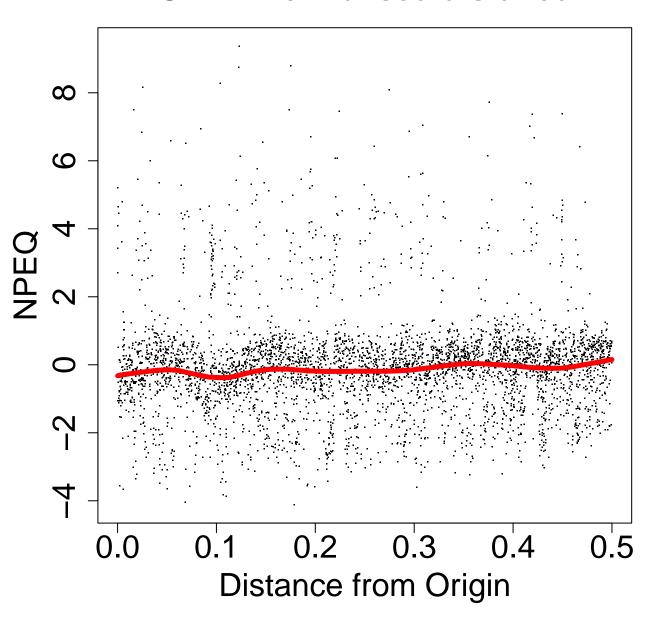
#### GC bias - Salmonella enteritidis



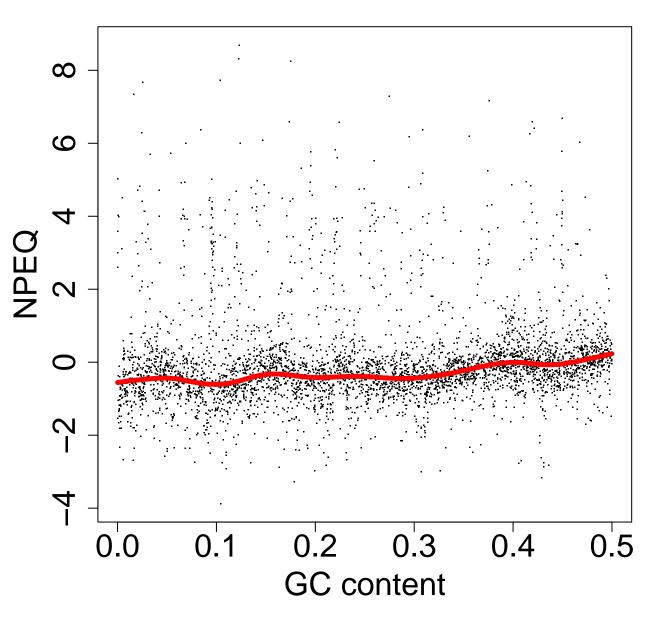
#### Distance bias - Salmonella enteritidis



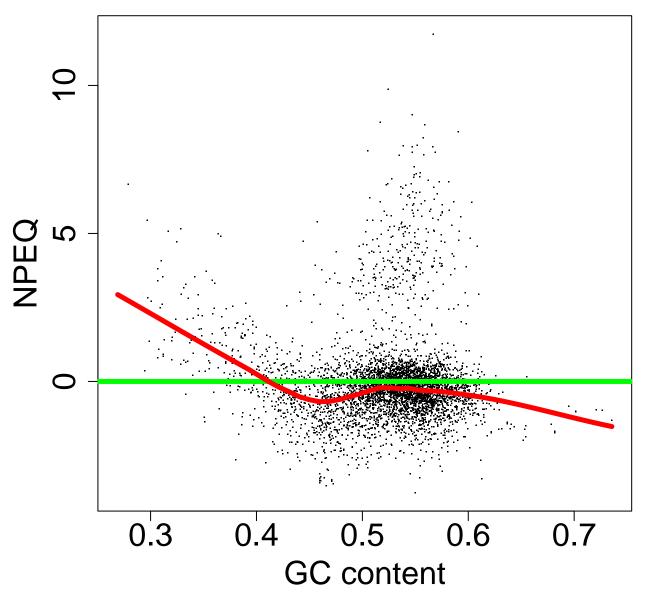
#### **SEN** – normalised distance



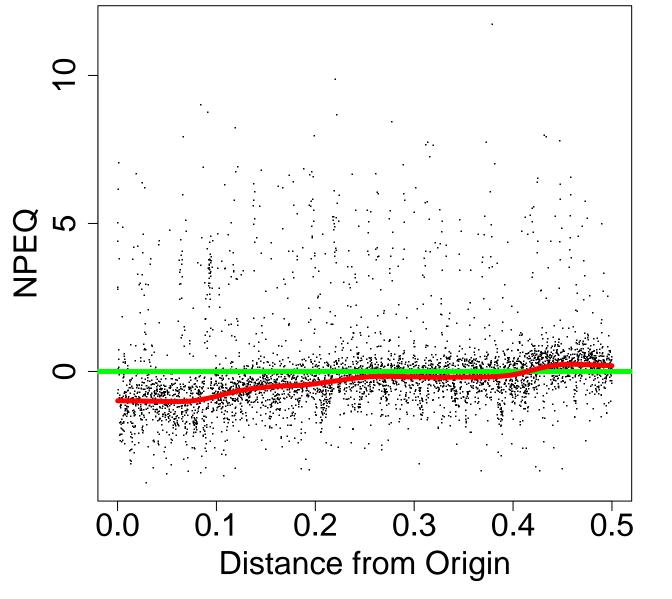
#### **SEN – normalised GC bias**



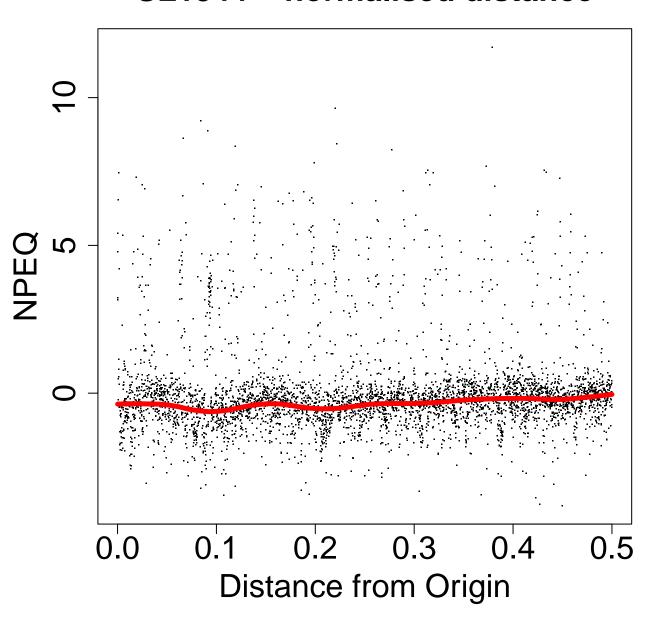
## GC bias – Salmonella typhimurium SL134



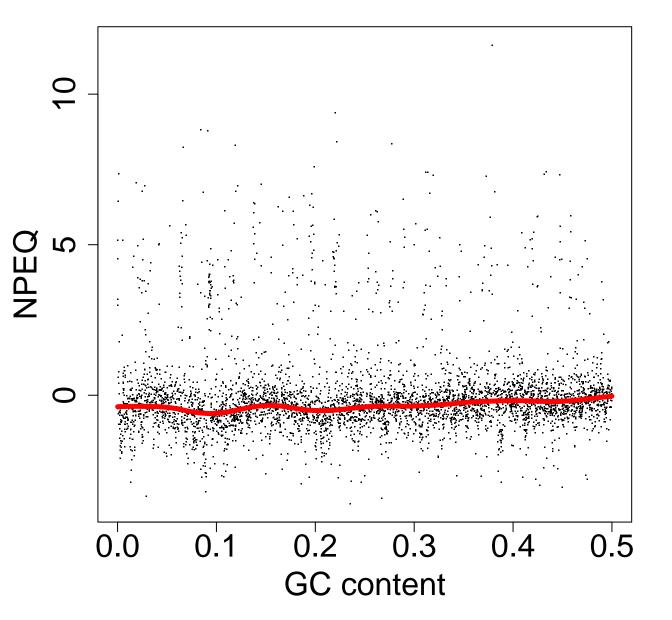
# Distance bias – Salmonella typhimurium SL13



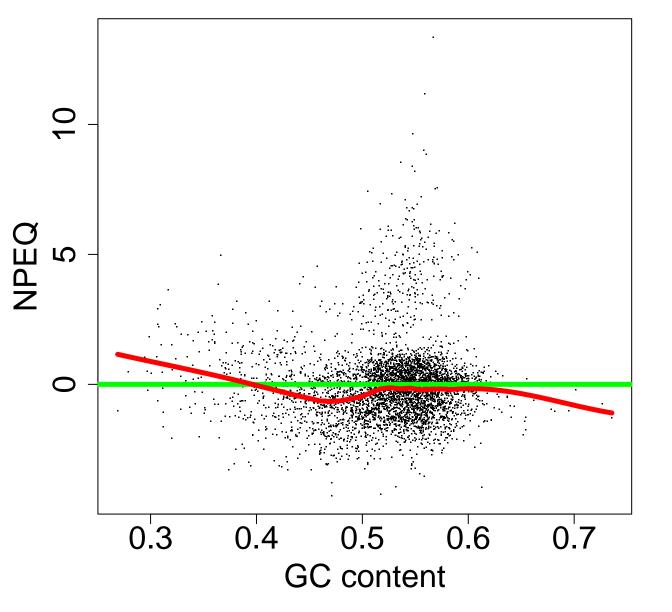
#### SL1344 - normalised distance



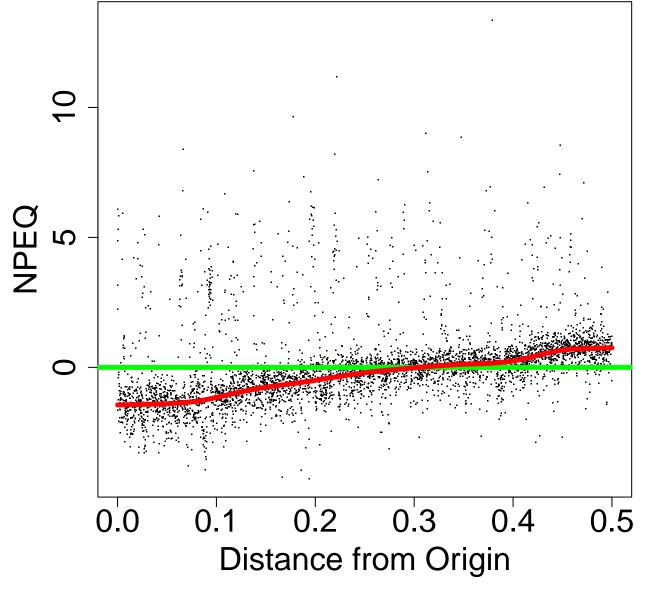
#### **SL1344 – normalised GC bias**



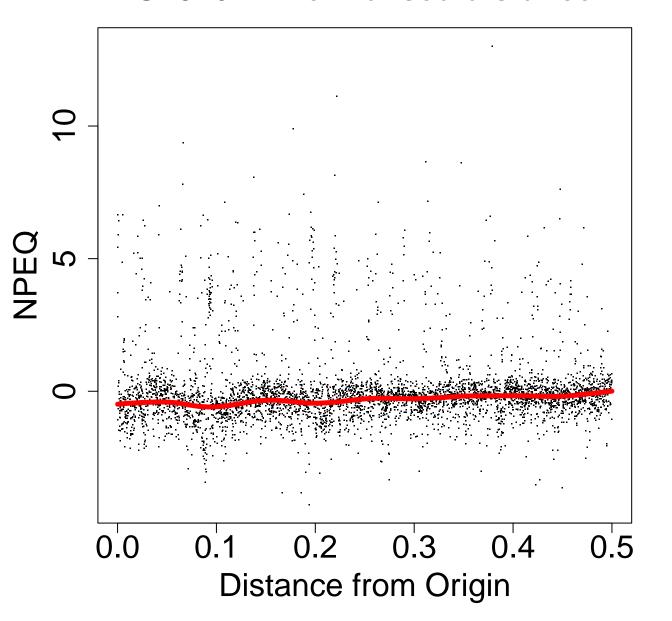
## GC bias – Salmonella typhimurium SL326



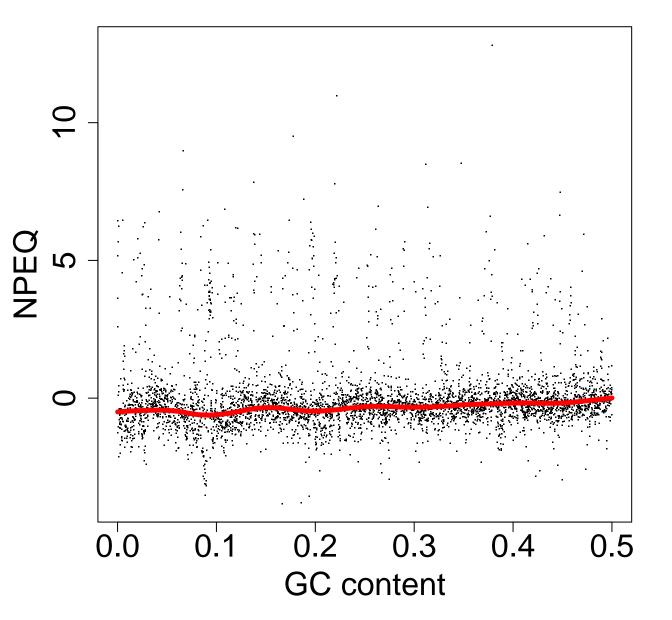
# Distance bias – Salmonella typhimurium SL32



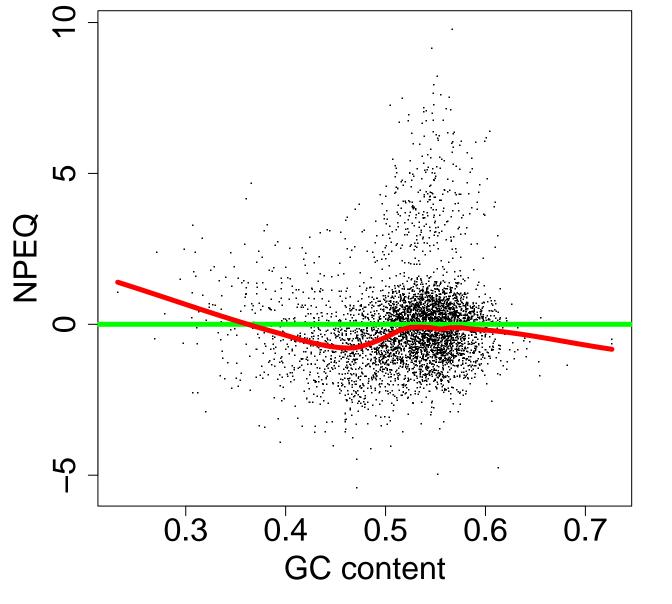
#### SL3261 - normalised distance



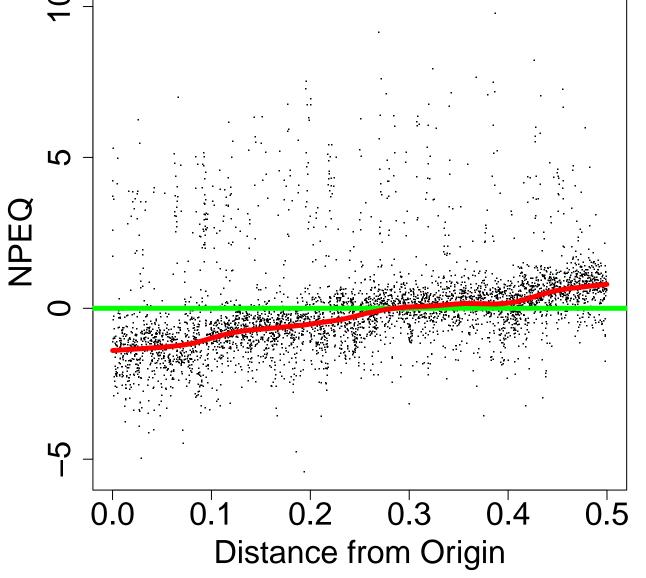
#### **SL3261 – normalised GC bias**



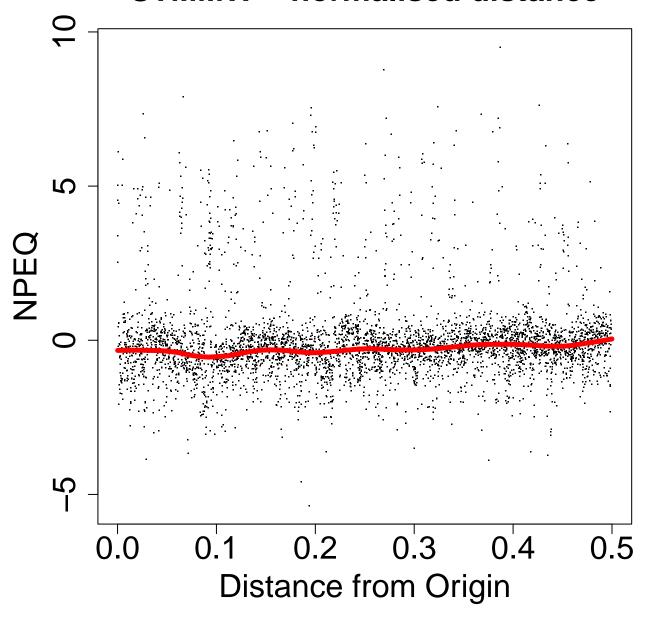
# GC bias – Salmonella typhimurium D2358



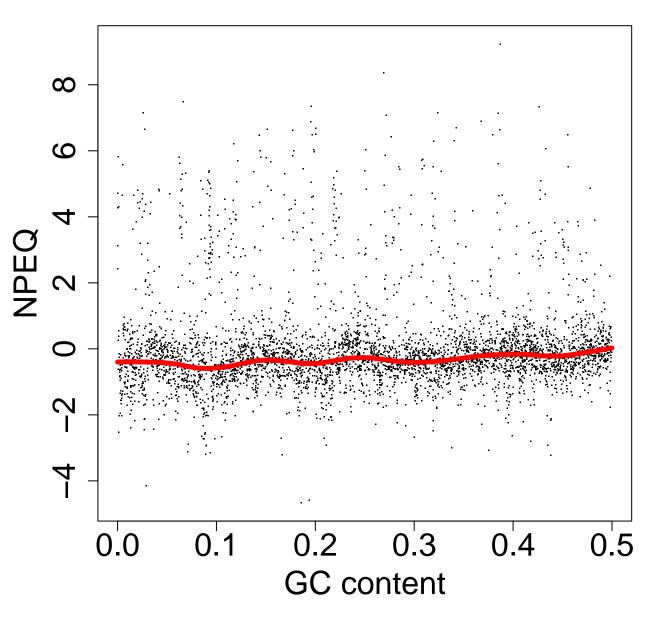
# Distance bias – Salmonella typhimurium D235



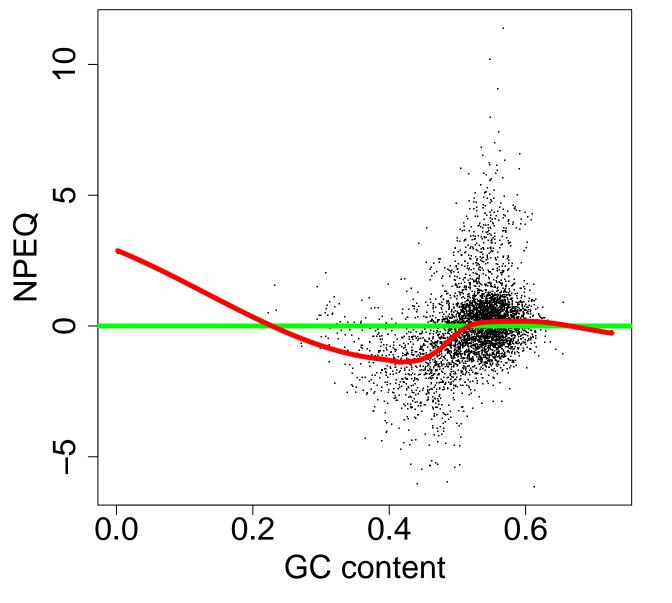
#### STMMW - normalised distance



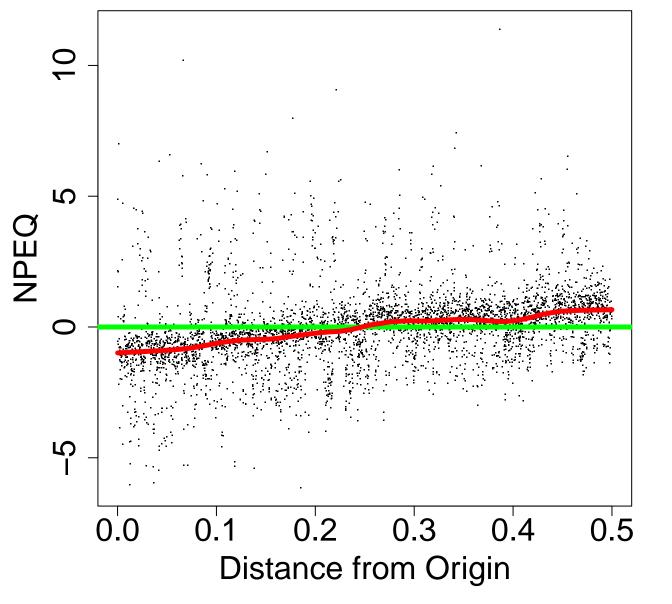
#### **STMMW – normalised GC bias**



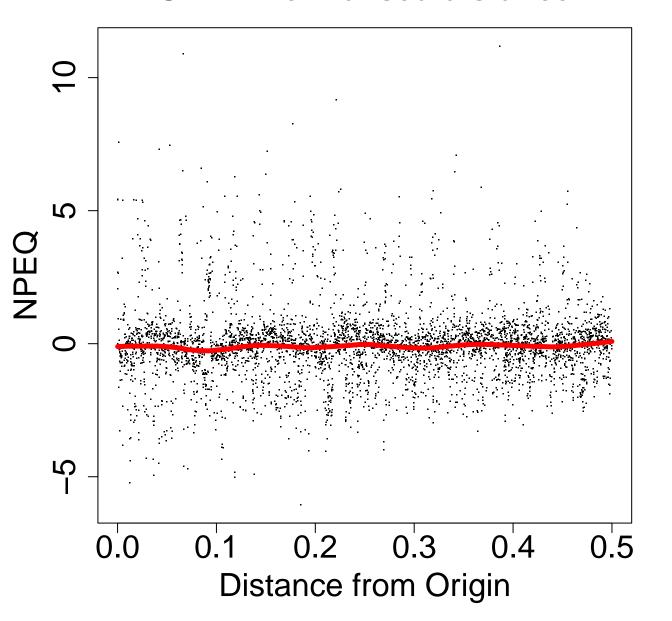
# GC bias – Salmonella typhimurium A130



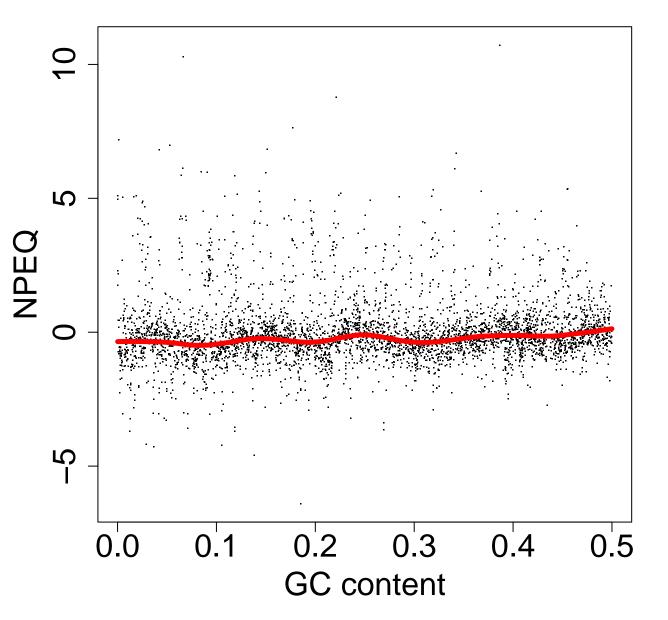
## Distance bias – Salmonella typhimurium A13



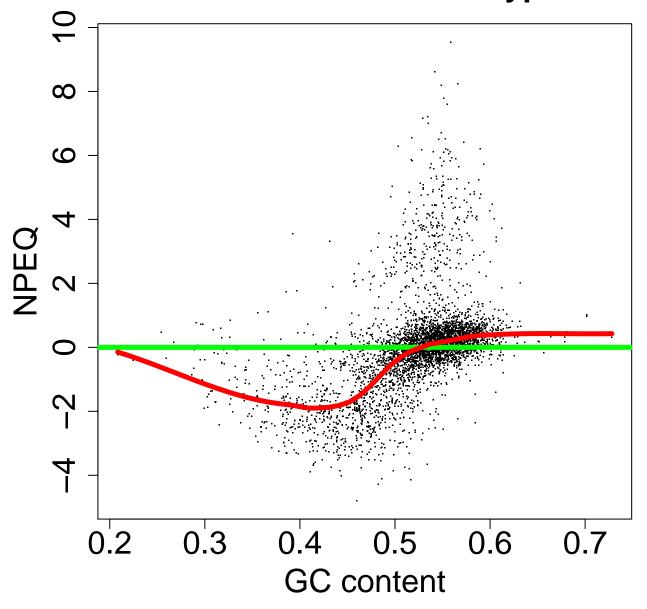
#### **STM** – normalised distance



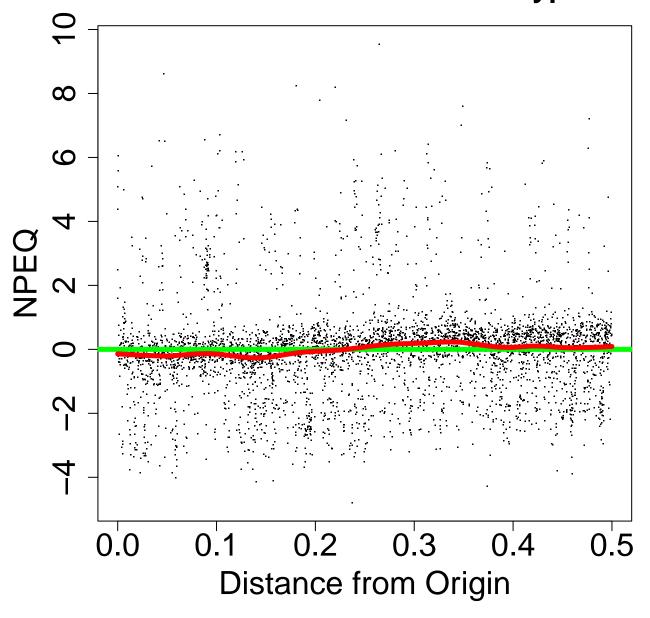
#### STM - normalised GC bias



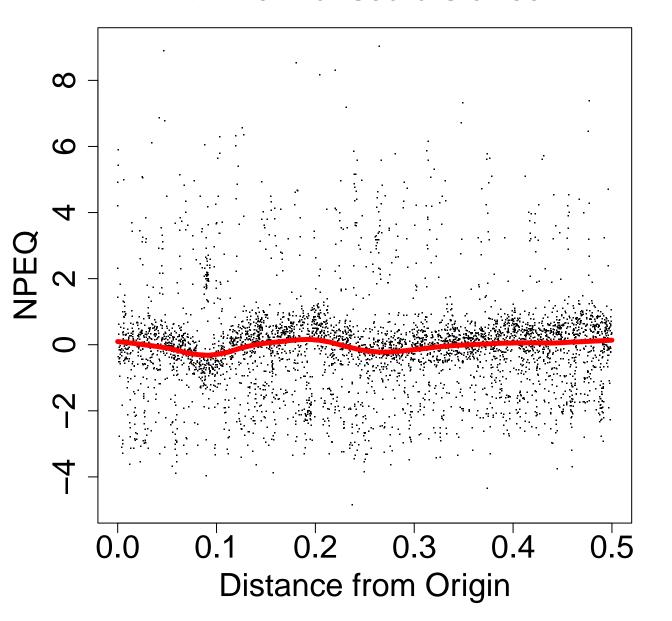
## GC bias - Salmonella typhi



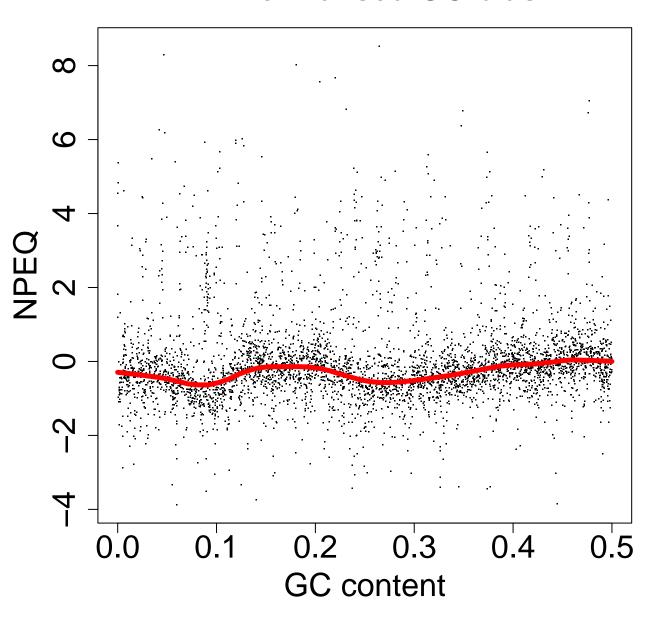
#### Distance bias - Salmonella typhi



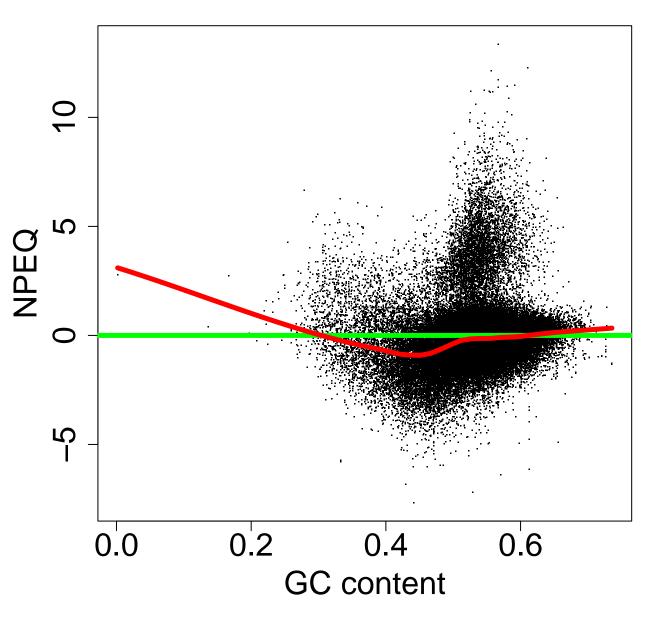
#### t - normalised distance



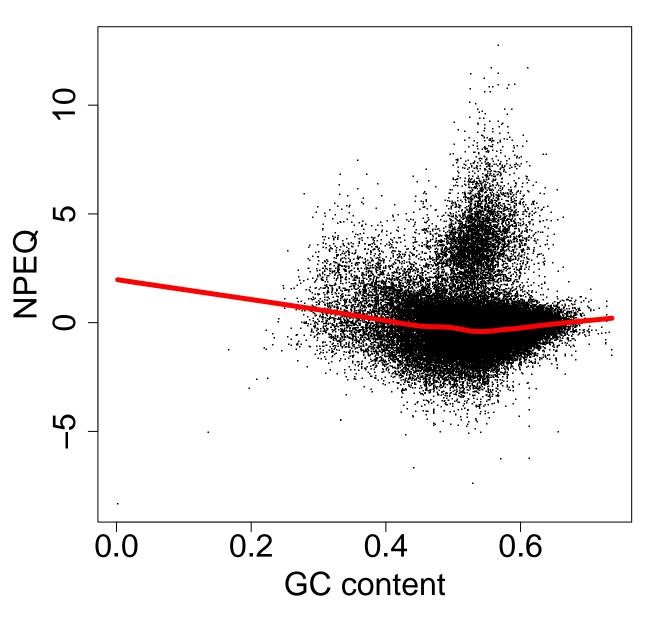
#### t - normalised GC bias



## **GC** bias



#### GC bias - normalised



## Length bias

