

Correlations between the stationary GC-content and the current GC-content, the crossover rate and distance to telomeres.

Data set: 1 Mb genome tiling

GC*: stationary GC-content

GC: current GC-content

CO: crossover rate (HapMap data)

LDT: Log distance to telomere in bp

GC* vs (GC) :	$R^2 = 0.25$
GC* vs (LDT) :	$R^2 = 0.35$
GC* vs (CO) :	$R^2 = 0.36$
GC* vs (GC, LDT) :	$R^2 = 0.42$
GC* vs (GC, CO) :	$R^2 = 0.44$
GC* vs (CO, LDT) :	$R^2 = 0.47$
GC* vs (GC, CO, LDT) :	$R^2 = 0.51$