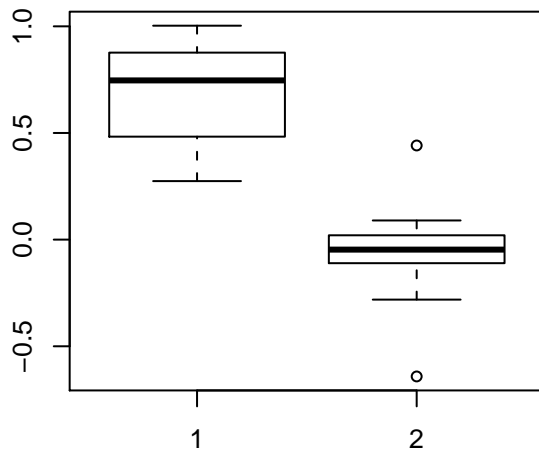
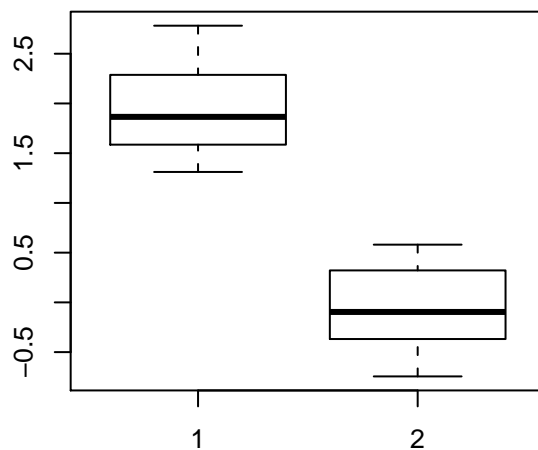


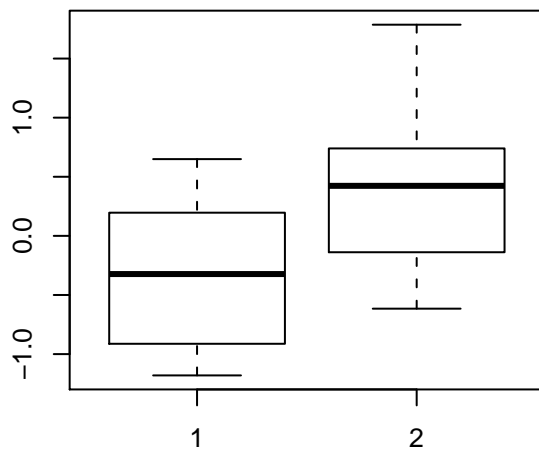
**Gene 1**



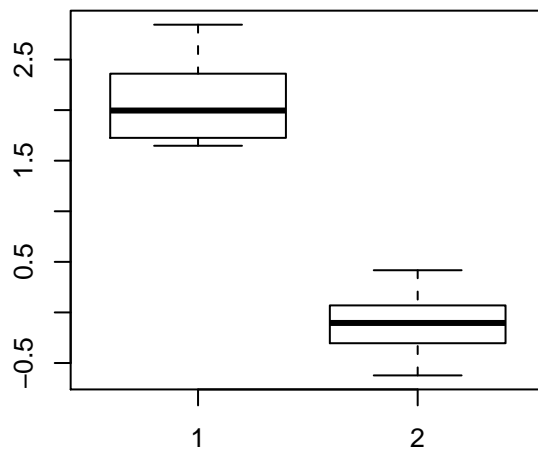
**Gene 2**

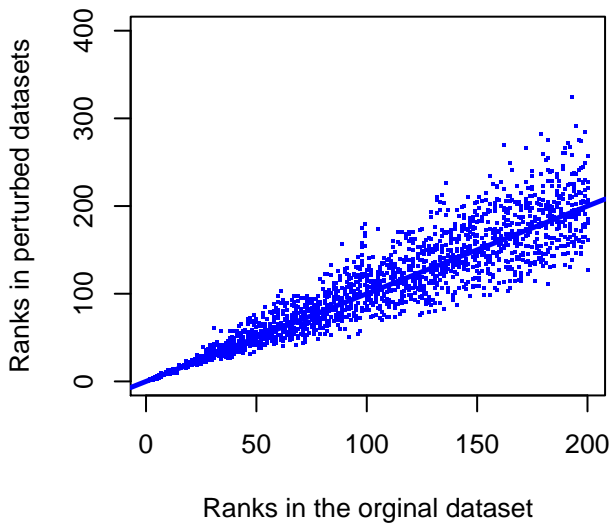
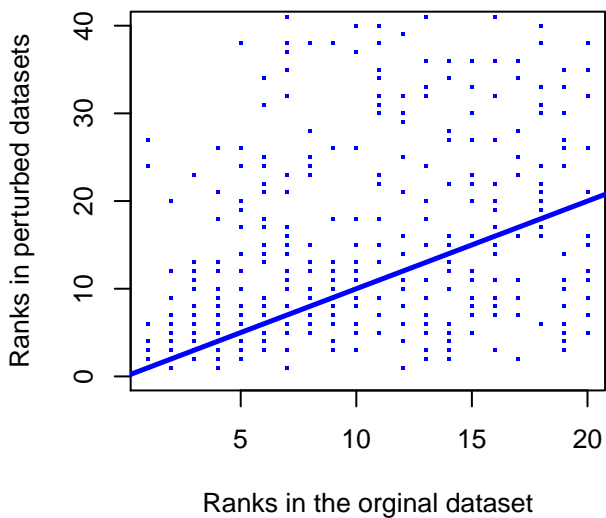
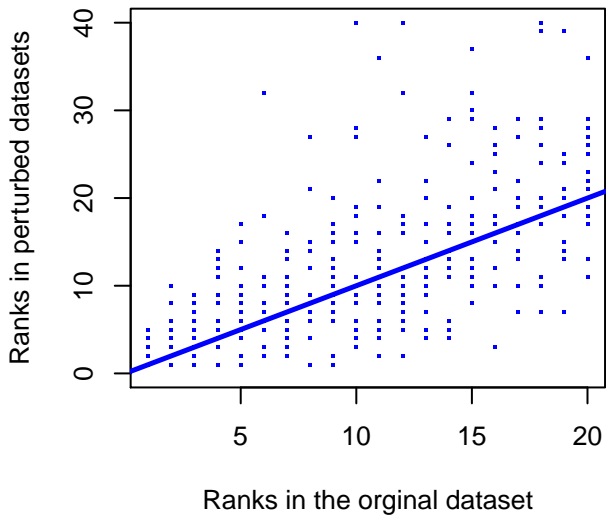
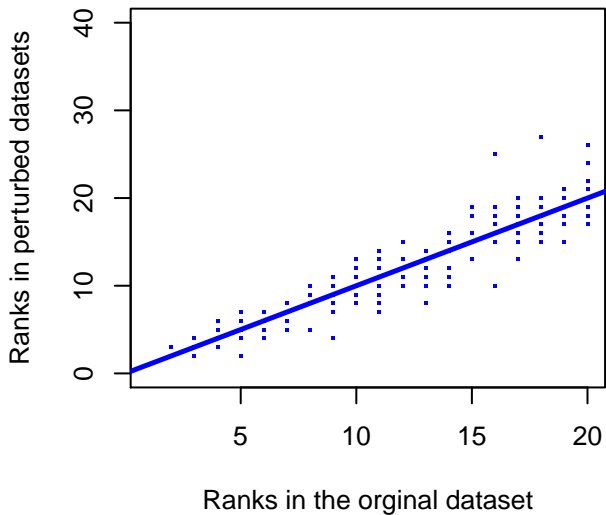


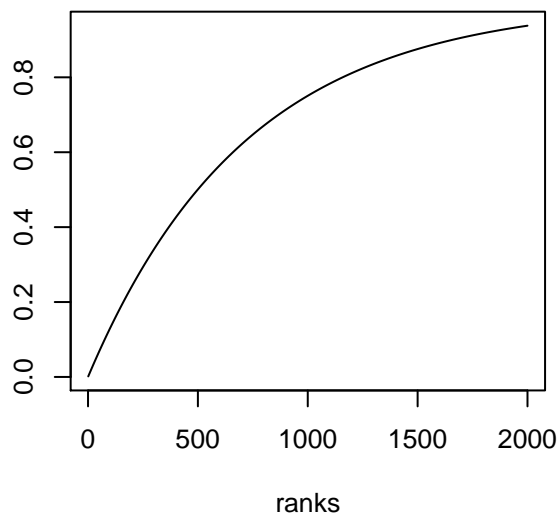
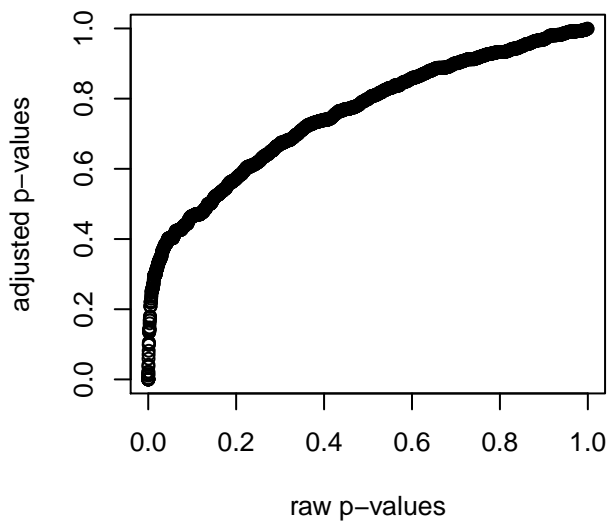
**Gene 3**



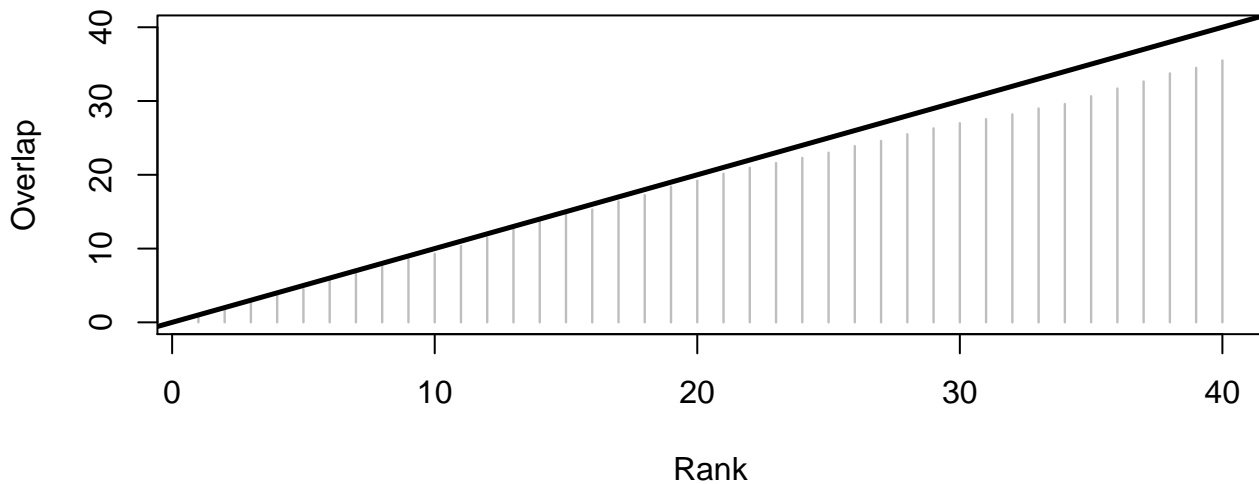
**Gene 4**



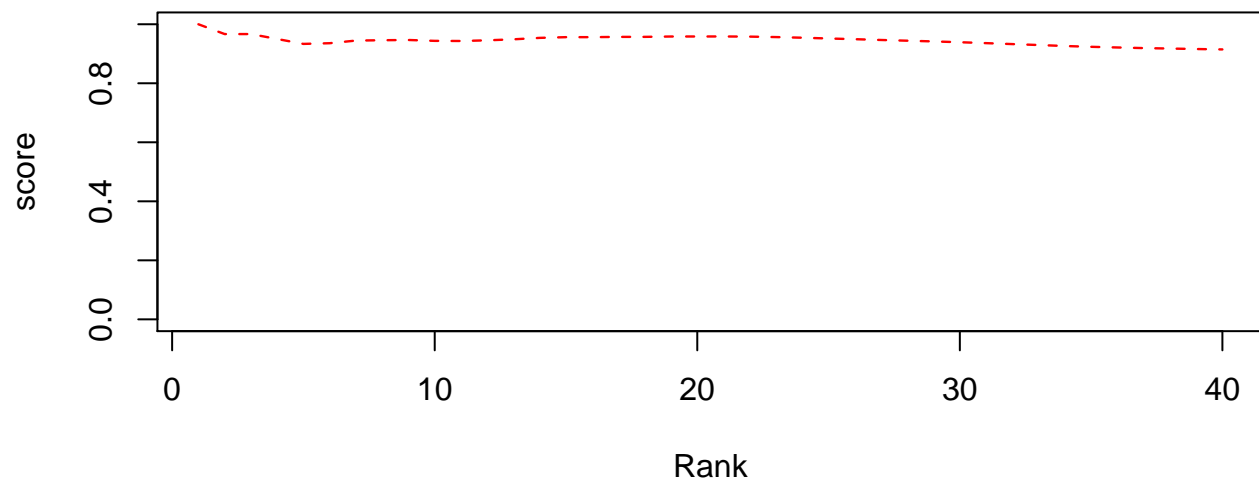


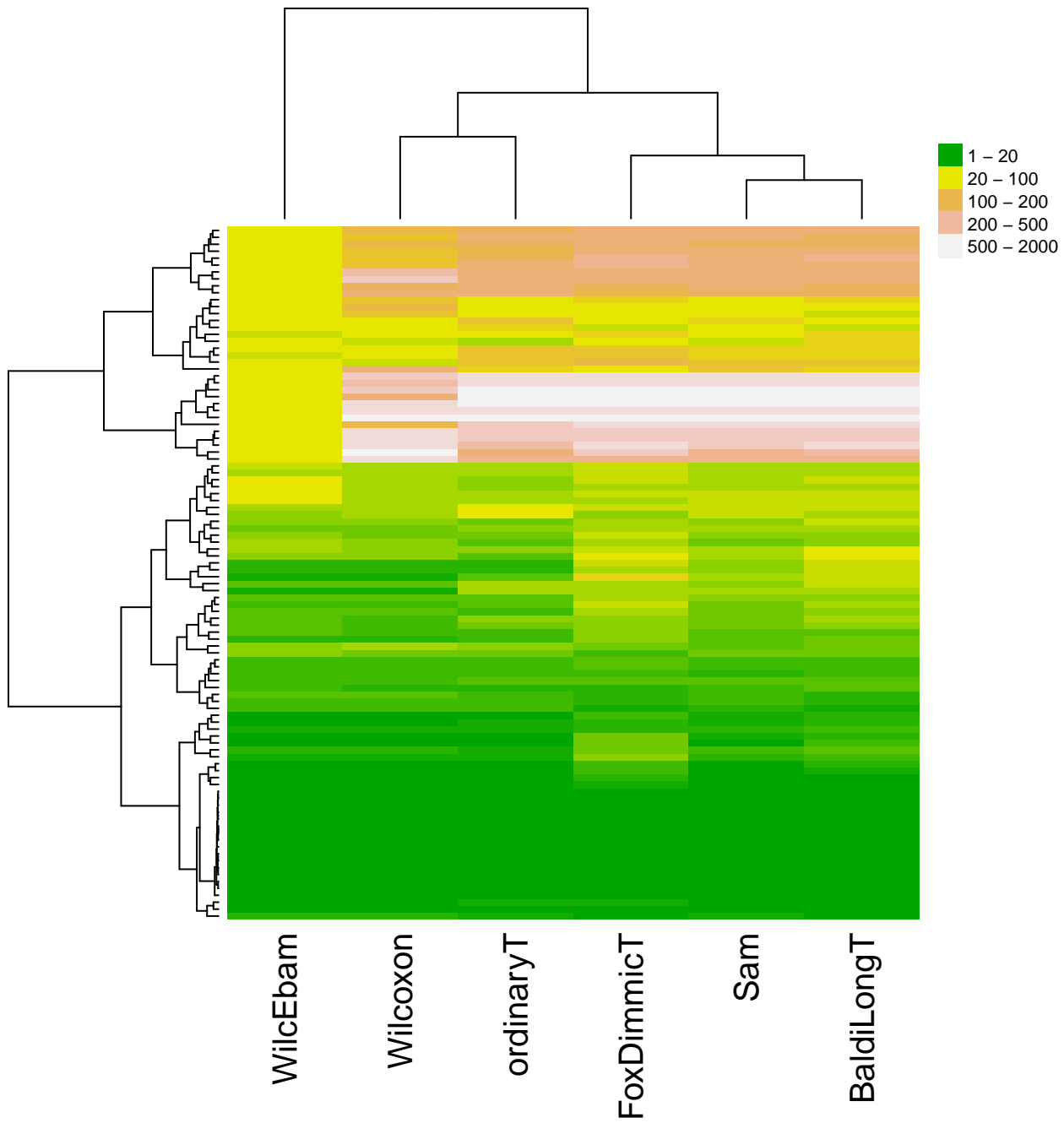


### Mean Overlap

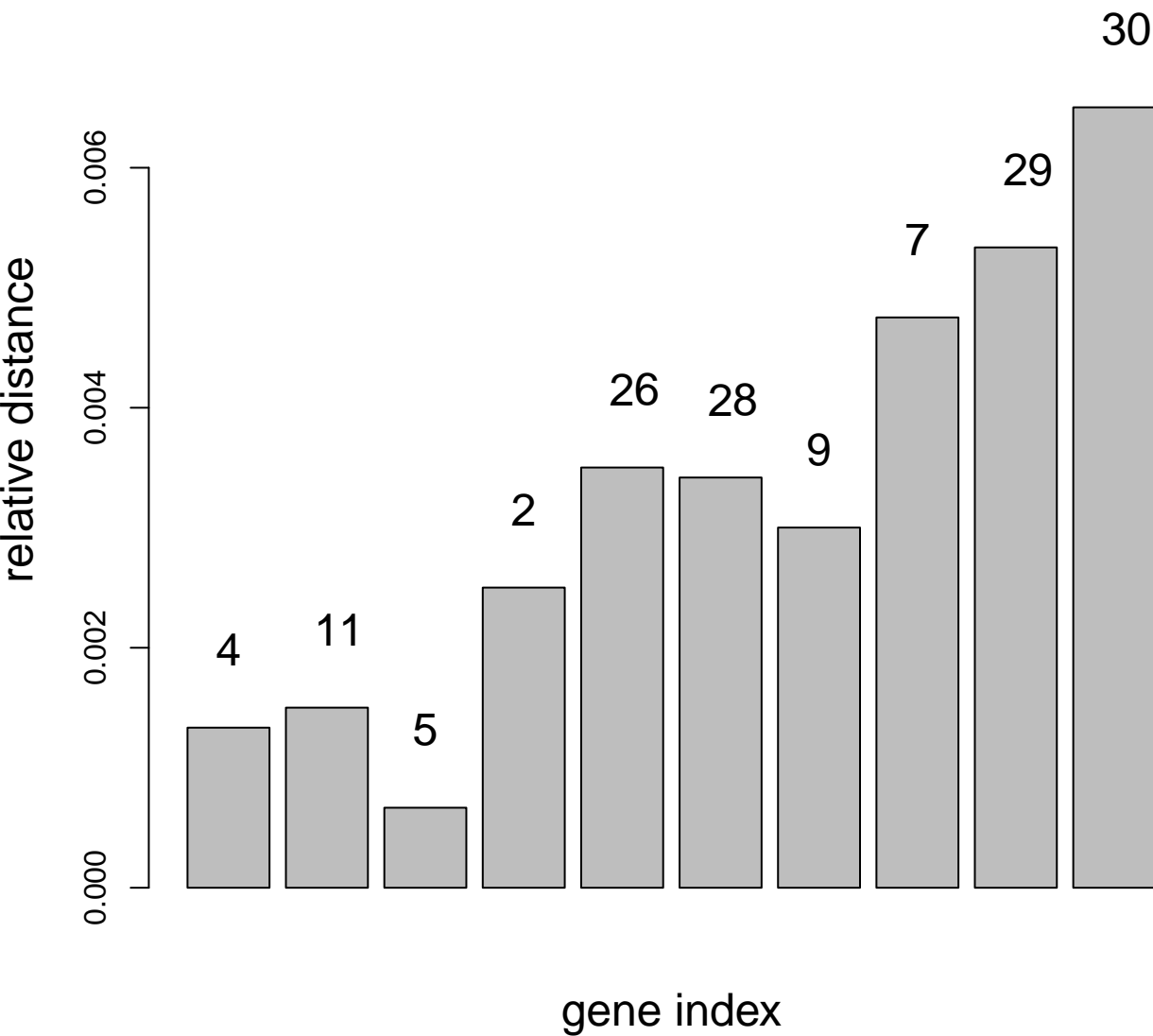


### Mean Score





**GeneSelector relative distance plot**



GeneInfoScreen for gene 30

selected ?	statistic	rank
+	Sam	15
+	Wilcoxon	15
+	BaldiLongT	17
+	FoxDimmicT	7
+	ordinaryT	15
+	WilcEbam	15

1 – 2020 – 100100 – 200200 – 500500 – 2000