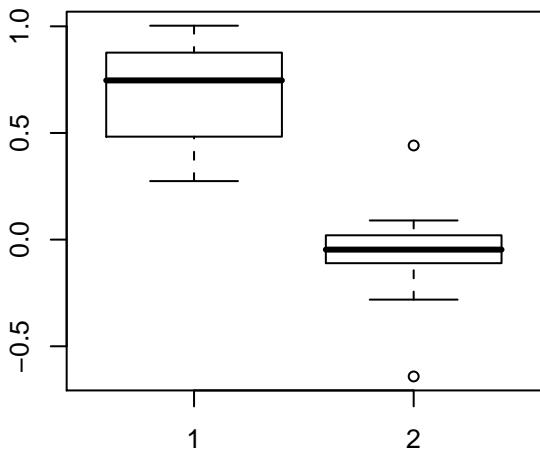
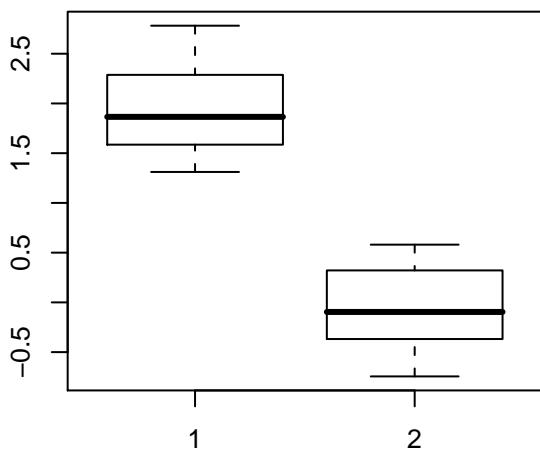
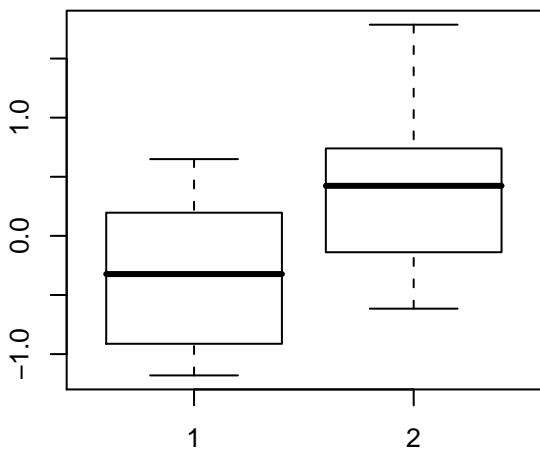
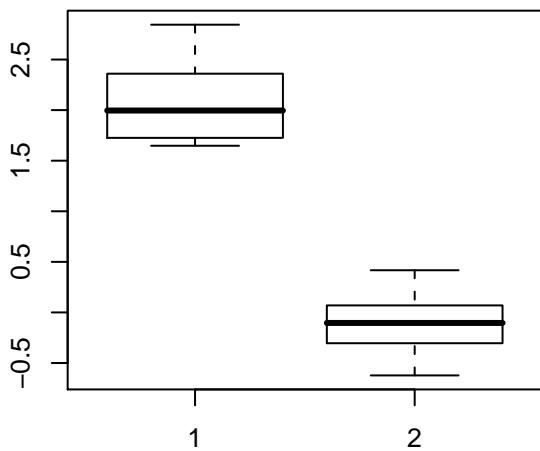
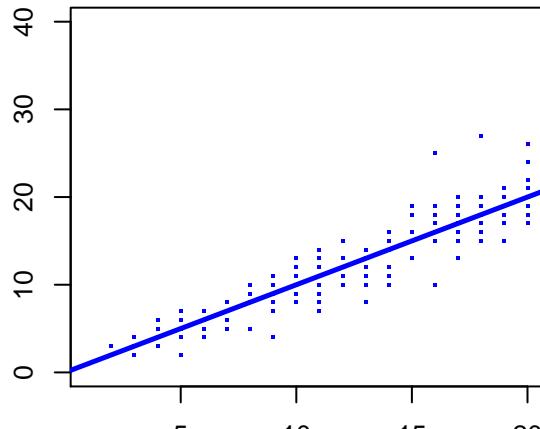


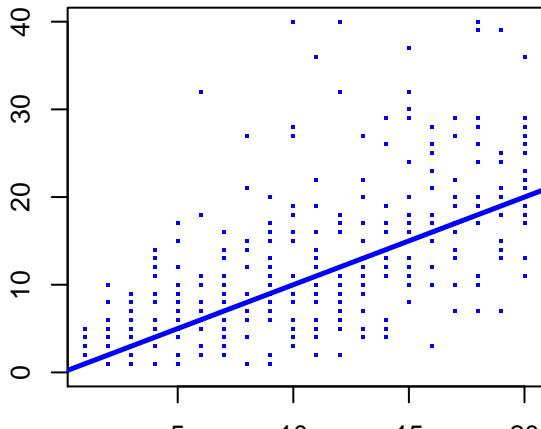
**Gene 1****Gene 2****Gene 3****Gene 4**

Ranks in perturbed datasets



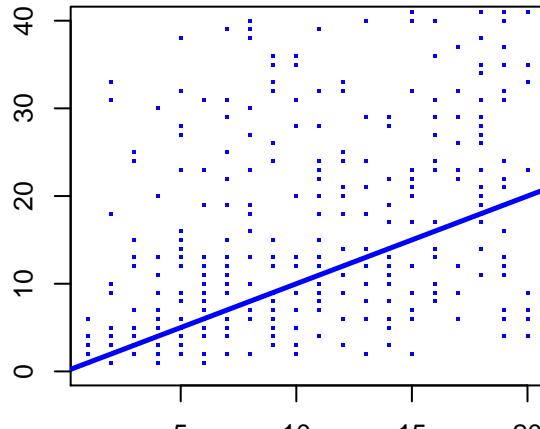
Ranks in the orginal dataset

Ranks in perturbed datasets



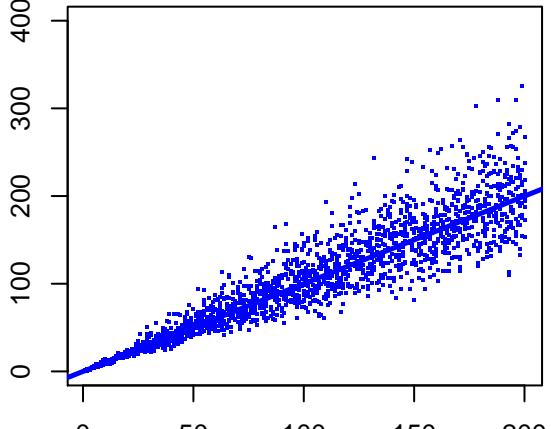
Ranks in the orginal dataset

Ranks in perturbed datasets

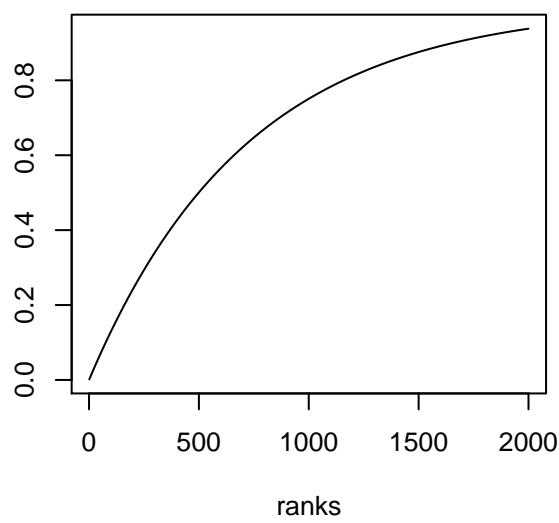
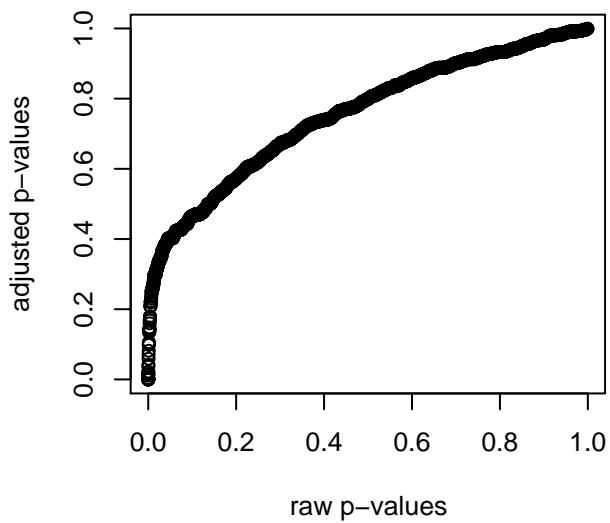


Ranks in the orginal dataset

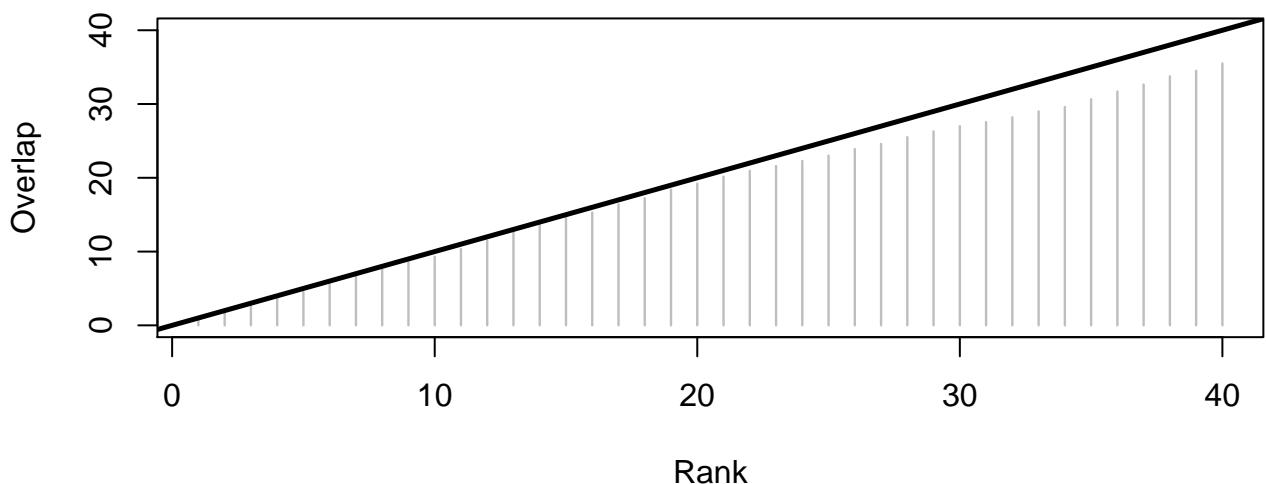
Ranks in perturbed datasets



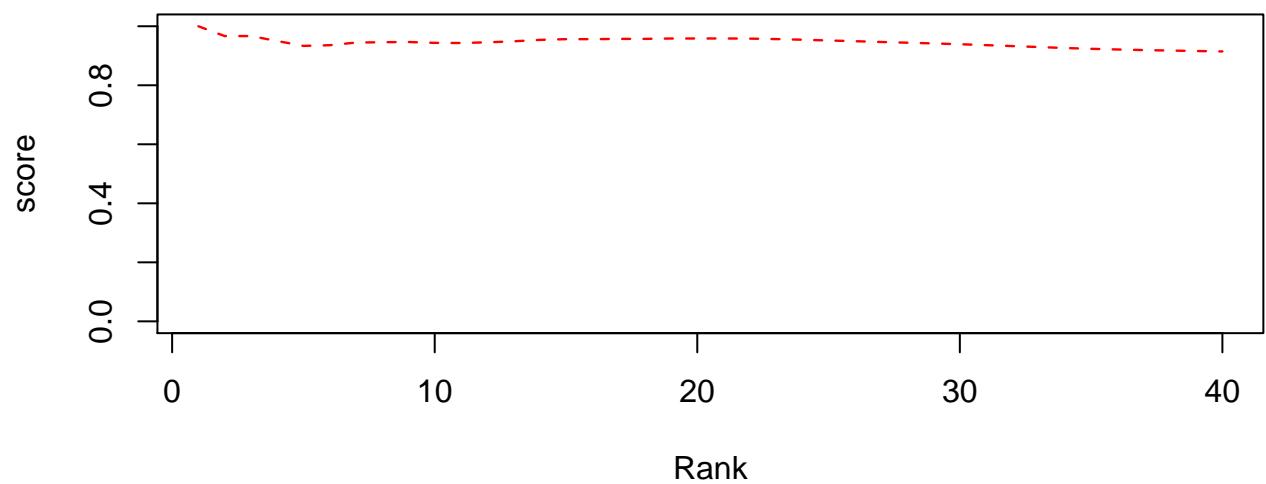
Ranks in the orginal dataset

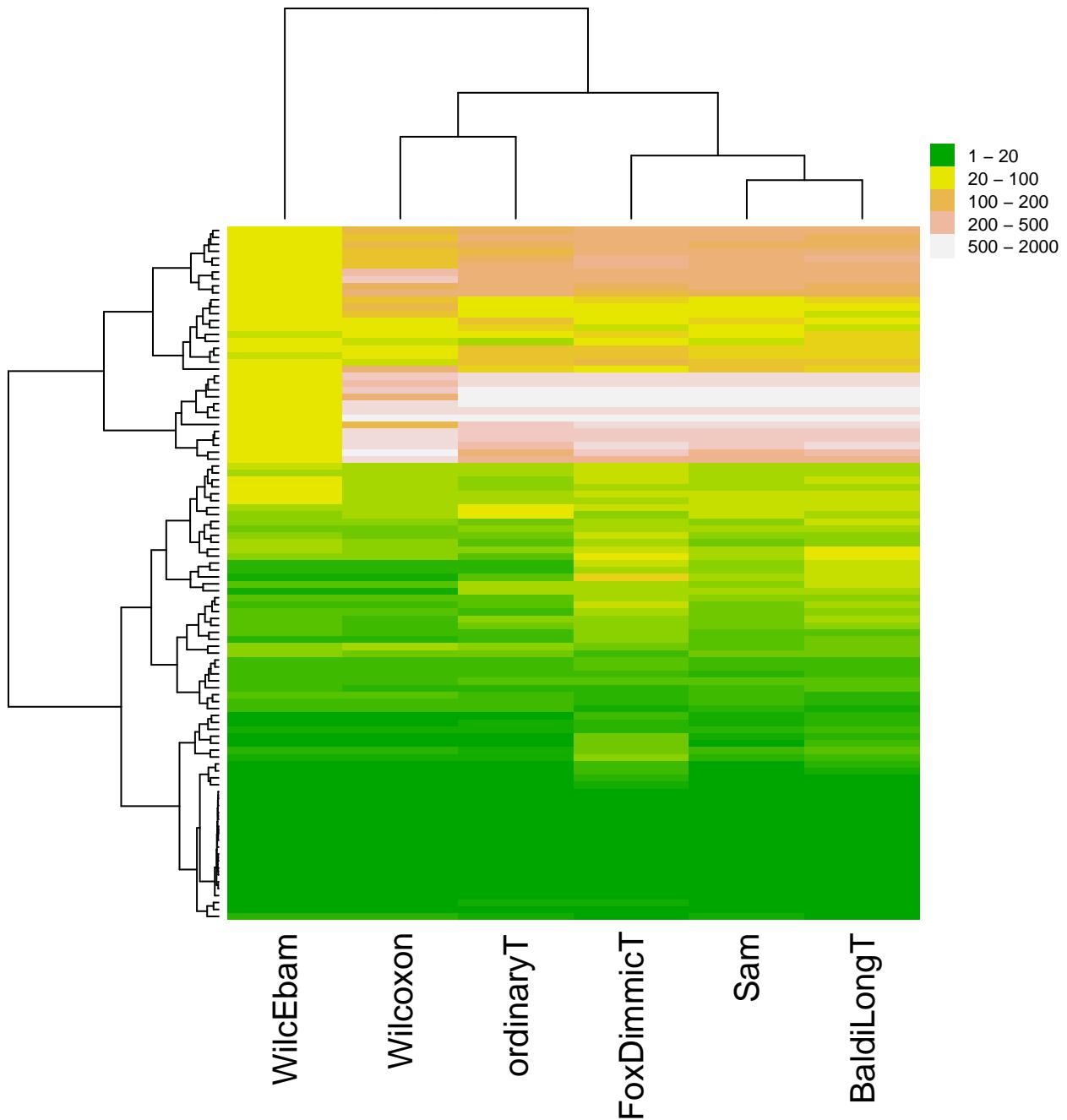


## Mean Overlap

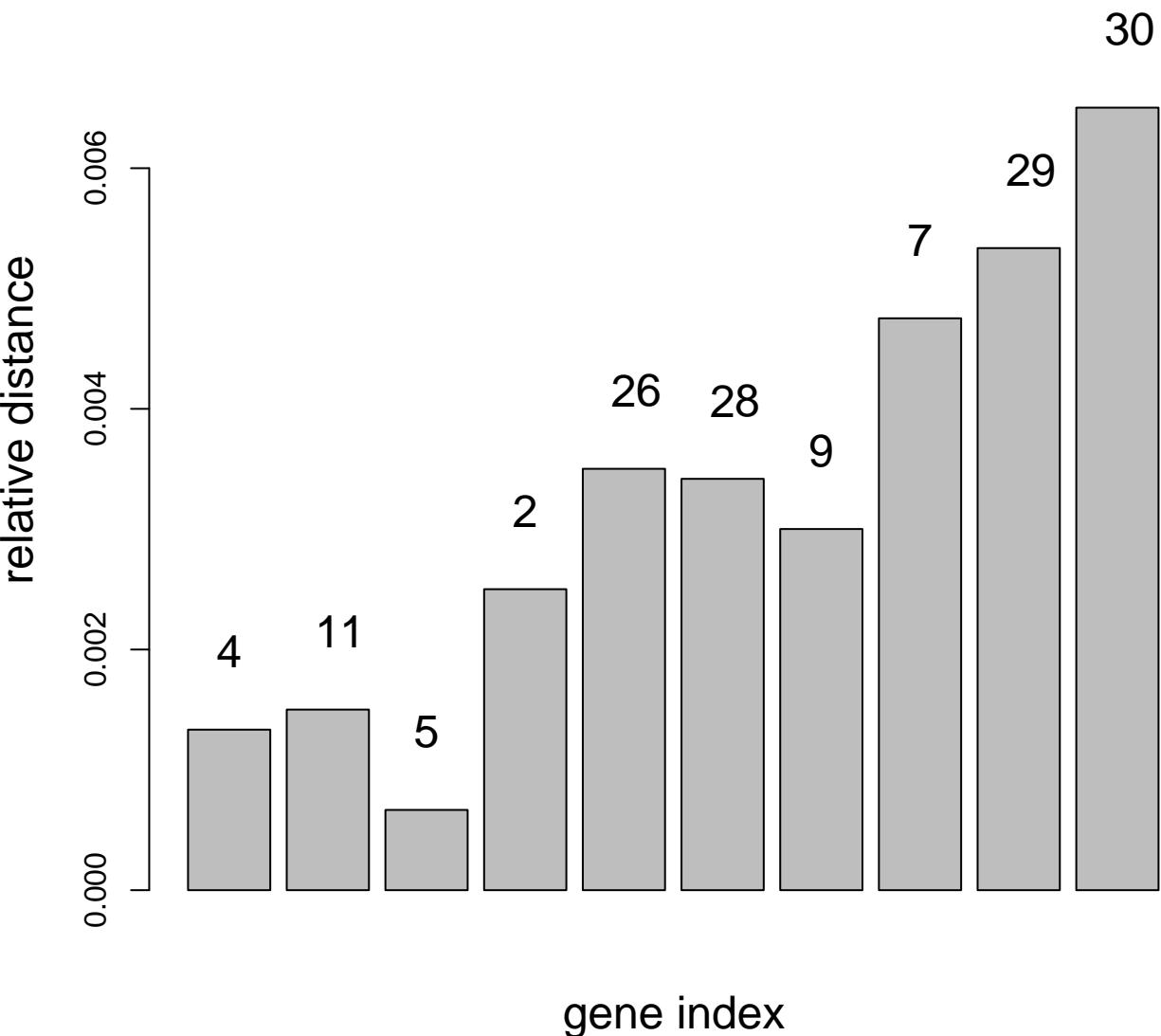


## Mean Score





## GeneSelector relative distance plot



## GeneInfoScreen for gene 30

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