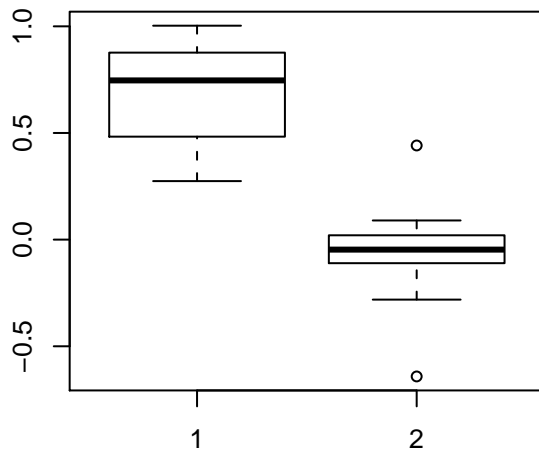
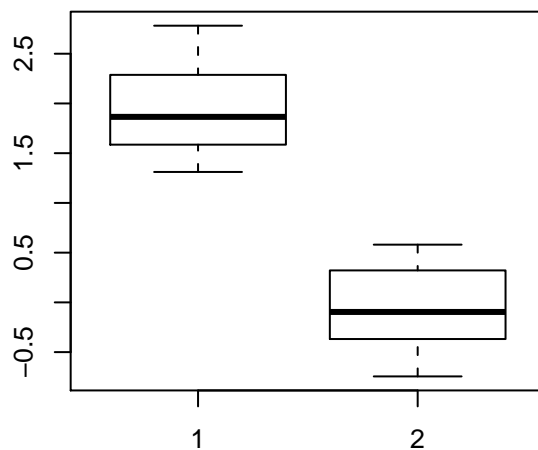


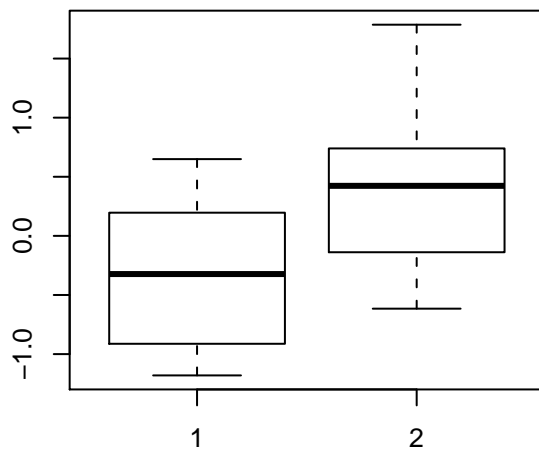
**Gene 1**



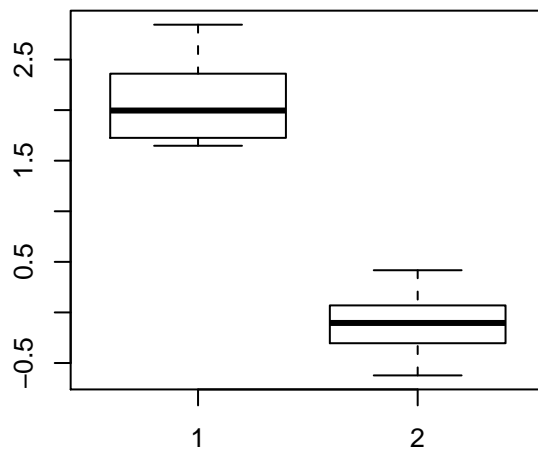
**Gene 2**



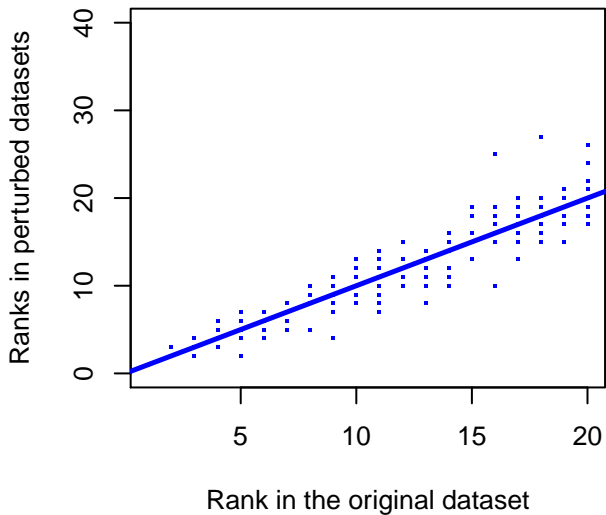
**Gene 3**



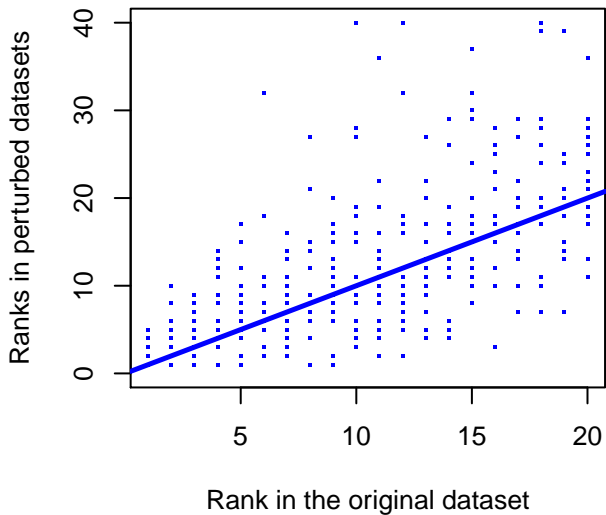
**Gene 4**



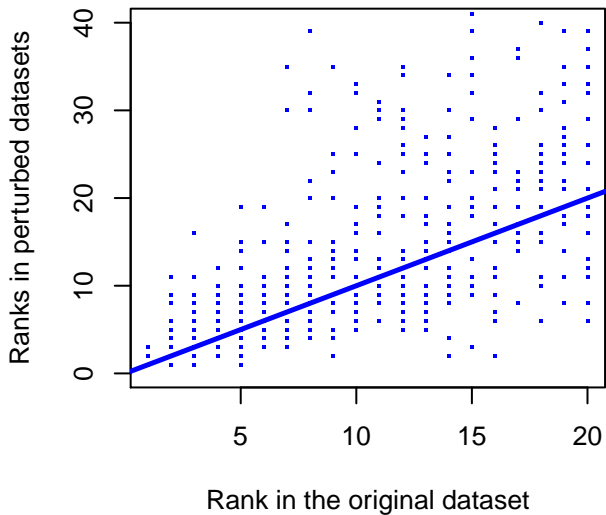
**jackknife**



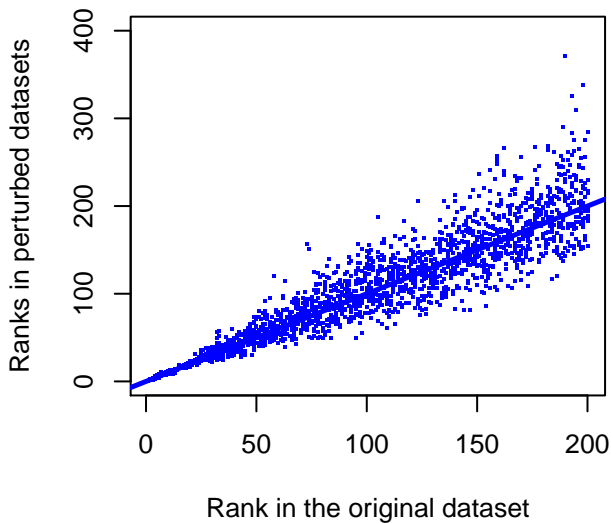
**label exchange**



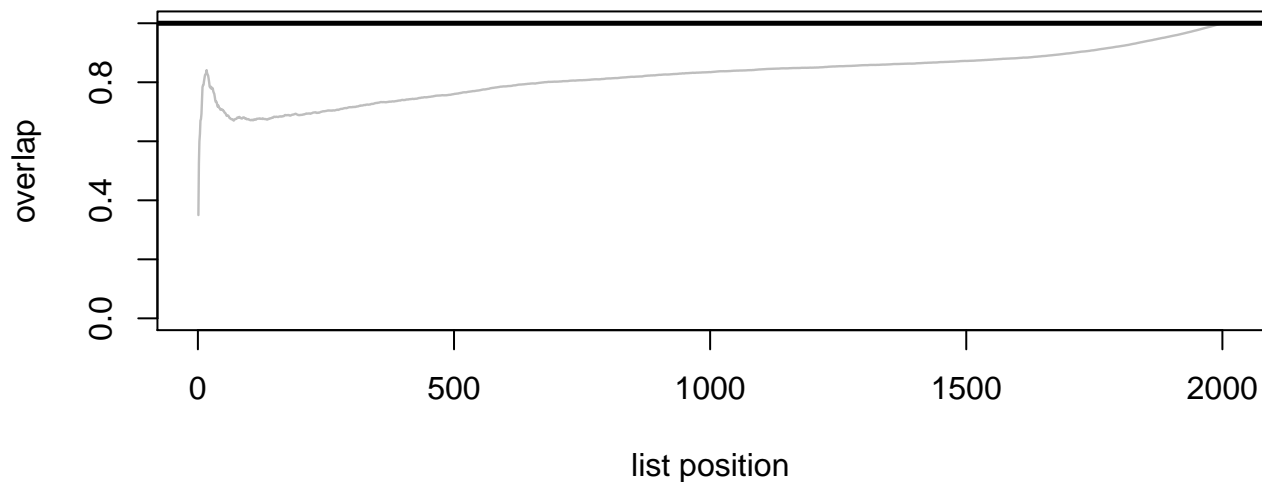
**bootstrap**



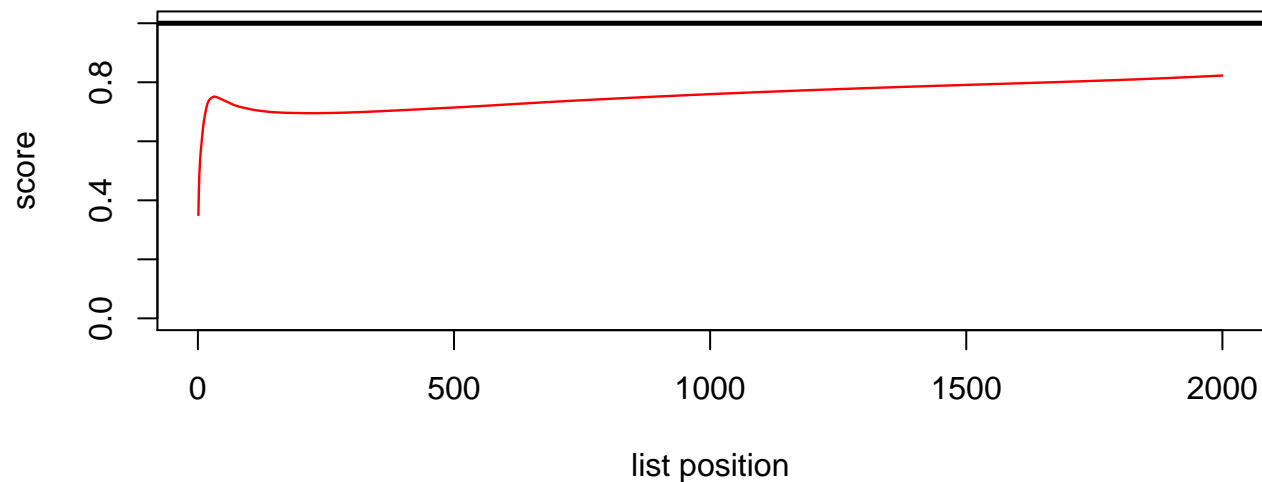
**noise**

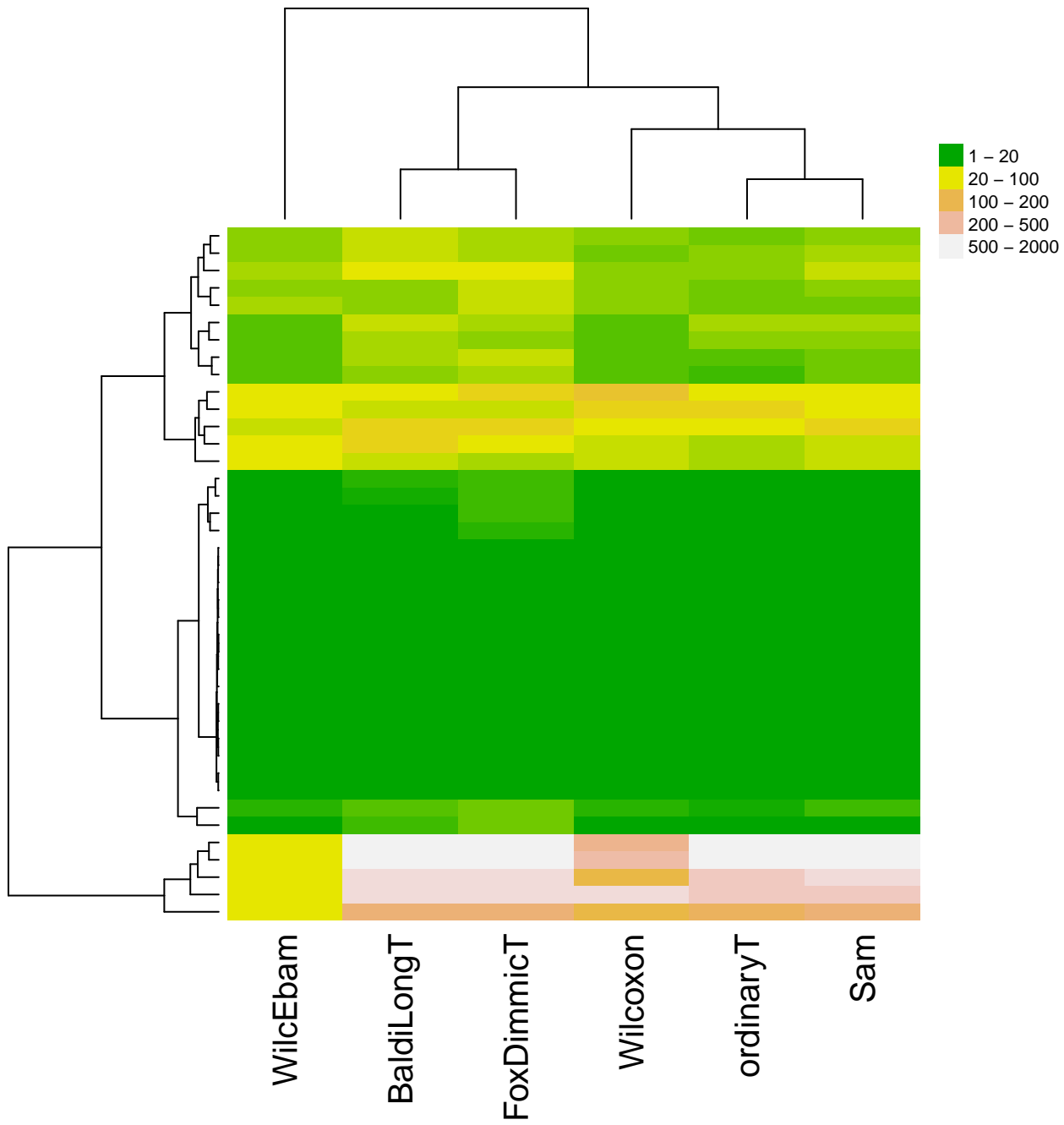


**percentage of overlap**



**average overlap score**





GeneInfoScreen for gene 1

selected ?	criterion	rank
+	ordinaryT	11
−	BaldiLongT	357
−	FoxDimmicT	544
−	Sam	53
+	Wilcoxon	11
+	WilcEbam	14

1 – 20

20 – 100

100 – 200

200 – 500

500 – 2000