



	GS follow link to MSigDB	GS DETAILS	SIZE	ES	NES	NOM p-val	FDR q-val	FWER p-val	RANK AT MAX	LEADING EDGE
1	GO_DNA_REPAIR	<a href="#">Details ...</a>	2	0.50	0.79	0.741	0.769	0.809	9	tags=50%, list=16%, signal=57%
1	REACTOME_DNA_REPAIR	<a href="#">Details ...</a>	1	-0.76	-1.01	0.461	0.936	0.500	15	tags=100%, list=27%, signal=134%
2	KAUFFMANN_DNA_REPAIR_GENES	<a href="#">Details ...</a>	2	-0.48	-0.88	0.596	0.648	0.640	31	tags=100%, list=55%, signal=216%

Supplementary Figure 4