

Anti-oxidative stress response genes: bioinformatic analysis of their expression and relevance in multiple cancers - Rotblat et al

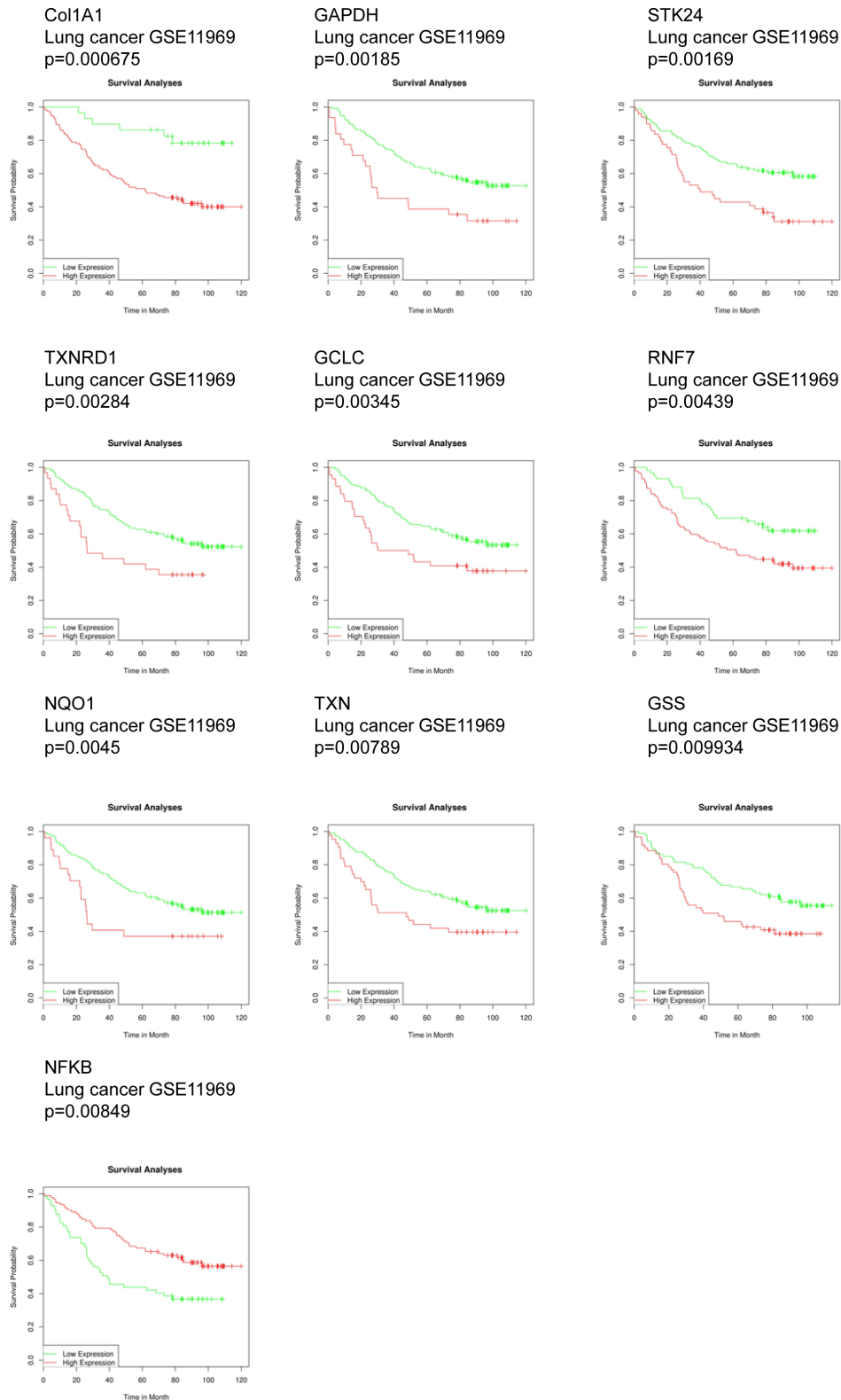
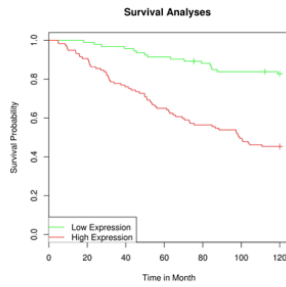
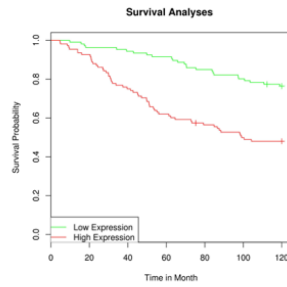


Figure S1: Kaplan-Meier plots for all oxidative stress response genes found to be highly expressed in cancers whose expression significantly correlates with survival in lung cancer.

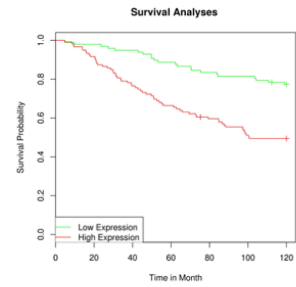
FOXM1
Breast cancer GSE22220
 $p < 10^{-7}$



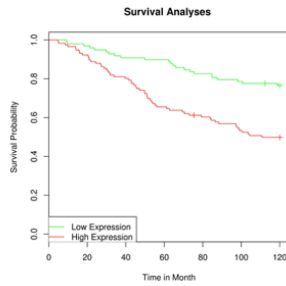
TXNRD1
Breast cancer GSE22220
 $p < 10^{-5}$



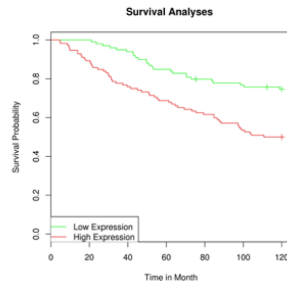
CDC2
Breast cancer GSE22220
 $p < 10^{-4}$



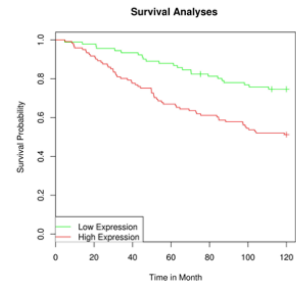
CASP3
Breast cancer GSE22220
 $p < 10^{-4}$



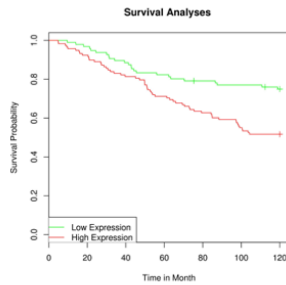
ECT2
Breast cancer GSE22220
 $p = 0.00012$



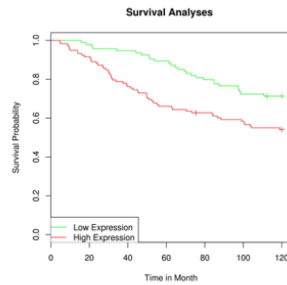
HMOX1
Breast cancer GSE22220
 $p = 0.00012$



NUDT1
Breast cancer GSE22220
 $p = 0.0016$



SERPINE1
Breast cancer GSE22220
 $p = 0.00167$



PRDX4
Breast cancer GSE22220
 $p = 0.00276$

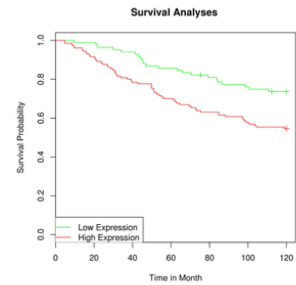


Figure S2-S3: Kaplan-Meier plots for all oxidative stress response genes found to be highly expressed in cancers whose expression significantly correlates with survival in breast cancer.

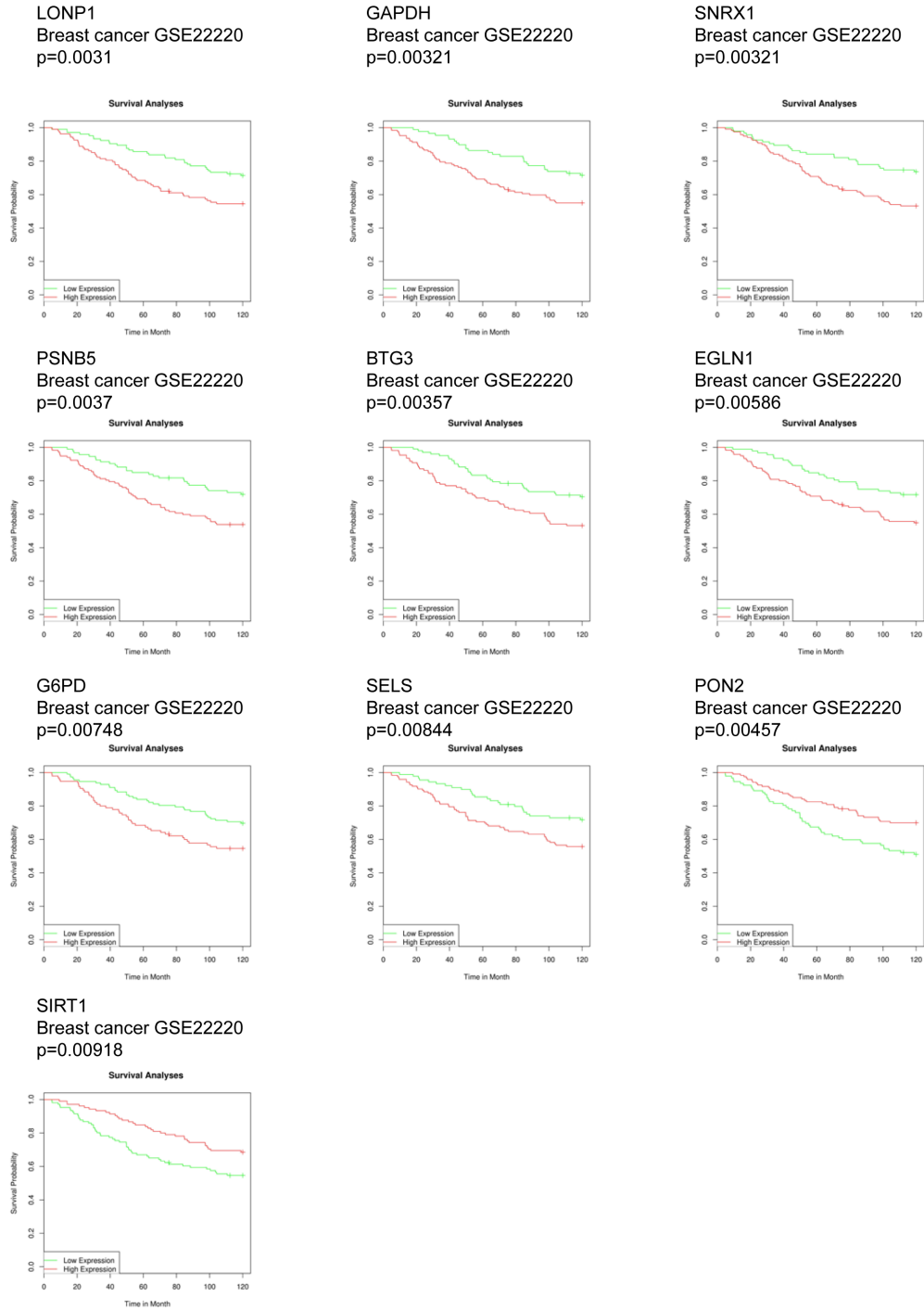


Figure S2-S3: Kaplan-Meier plots for all oxidative stress response genes found to be highly expressed in cancers whose expression significantly correlates with survival in breast cancer.