| Gene  | TR−DDR Score |
|-------|--------------|
| BRCA  | Melanoma Cohort1 |
| COAD  | Melanoma Cohort2 |
| HNSC  | ccRCC |
| KIRC  | Urothelial cancer cohort1 |
| KIRP  | Urothelial cancer cohort2 |
| LIHC  | BRCA |
| LUAD  | COAD |
| LUSC  | HNSC |
| PRAD  | KIRC |
| STAD  | KIRP |
| THCA  | LIHC |

| Gene  | TR−DDR Score |
|-------|--------------|
| UBE2T | 0            |
| EXO1  | 0.2          |
| GEN1  | 0.4          |
| BARD1 | 0.6          |
| FANCD2| 0.8          |
| POLQ  | TR−DDR Score |
| DLX5  |             |
| XRC2  |             |
| FANC2 |             |
| FANCB |             |
| FEN1  |             |
| PRKDC |             |
| BRCA2 |             |
| FANC3 |             |
| RAD1  |             |
| FANC1 |             |
| FANCA |             |
| BRCA1 |             |
| EME1  |             |
| BRIP1 |             |
| TR−DDR Score | 0.8 |
FCRL4.txt

Melanoma Cohort1
Melanoma Cohort2
cCRCC
Urothelial cancer cohort1
Urothelial cancer cohort2
BRCA
COAD
HNSC
KIRC
KIRP
LIHC
LUAD
LUSC
PRAD
STAD
THCA

TR-DDR Score

0
0.2
0.4
0.6
0.8
MAGEA4.txt

TR-DDR Score

Melanoma Cohort1
Melanoma Cohort2
cCRCC
Urothelial cancer cohort1
Urothelial cancer cohort2
BRCA
COAD
HNSC
KIRC
KIRP
LIHC
LUAD
LUSC
PRAD
STAD
THCA

UBE2T    EXO1    GEN1    BARD1    FANCD2    POLO    DLX5    XRCC2    FANCB    PRKDC    FEN1    BRC2    XRC3    RAD51    FANCI    BLM    FANCA    BRCA1    EME1    BRIP1

0        0.2      0.4      0.6      0.8
