## **Description of Additional Supplementary Information**

File Name: Supplementary Data 1

Description: : Annotation of 1,034 human cancer cell lines used in our study and the source of RNA-seq data. All cell lines are part of the GDSC cancer cell line project (see Cell Model Passport and

COSMIC IDs).

File Name: Supplementary Data 2

Description: List and annotation of 10,514 fusion transcripts identified in 1,011 cell lines.

File Name: Supplementary Data 3

Description: Significant results from differential gene expression for recurrent fusions.

File Name: Supplementary Data 4

Description: Annotation of gene fusions for aberrant expression of the 3-prime end gene.

File Name: Supplementary Data 5

Description: High-throughput cell line drug sensitivity data (IC50's) used in this study together with

compound annotation. Column names are COSMIC id's of cell lines.

File Name: Supplementary Data 6

Description: Significant ANOVA results from the drug-association analysis using cancer functional

events.

File Name: Supplementary Data 7

Description: Significant ANOVA results from the drug-association analysis using gene fusions.

File Name: Supplementary Data 8

Description: Fusion essentiality score and significance calculation for all 2,821 fusion transcripts with mapping guides. For 525 fusion transcripts where multiple data sets contained mapping guides, both are reported.

File Name: Supplementary Data 9

Description: List of primers sequences used to validate gene fusions.