

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: qPCR Primers used in this study.

File Name: Supplementary Data 2

Description: shRNA used in this study.

File Name: Supplementary Data 3

Description: CRISPR/Cas9 screen data processed by MAGeCK algorithm.

File Name: Supplementary Data 4

Description: IP-MS data analyzed using MaxQuant version 1.6.0.1 search against Mouse Fasta database, with label free quantification and match between runs functions enabled.