

**Supplementary Figure S1**. MPNSTs are sensitive to combined HDAC and mTOR inhibition. **A)** p53 WT MPNST cell line T265 was treated with vehicle (veh), sapanisertib (200nM, sap), vorinostat ( $2\mu$ M, vor), or vorinostat and sapanisertib (200nM) for 3 days. Left y-axis indicates log 2 of fold change in cell number at 3 days relative to day 0. Right y-axis indicates percent change in cell number at 3 days on a log 2 scale. Error bars ±SD from technical triplicates. Two other p53 WT MPNST cell lines (88-14 and SNF96.2) were also sensitive to the this combination (data not shown). **B)** Vorinostat and sapanisertib are synergistic according to Gaddum's non-interaction model, which is the most appropriate model for assessing synergy on cytotoxicity when at least one agent (sapanisertib) is purely cytostatic. 90-8TLs were treated with indicated drug concentrations for three days and viability was assessed by cell titer glow. The percent decrease in viability realtive to day 0 was calculated as a proxy for cell death. The excess over the highest single agent was calculated and is shown in the matrix, with 100 indicating complete synergy, and -100 indicating complete antagonism. Boxes are shaded according to the color scale shown. \*p=0.000006, \*\*p<0.00001

## Supplementary Figure S2



Supplementary Figure S2. Change in animal weight while on combiniation treatment. A) Weight of same animals in Figure 1G treated with rapamycin and vorinostat. Y-axis indicates relative weight (normalized to day 0 weight of individual animal). X-axis indicates days on treatment. Legend at right shows mouse number of each mouse. B) Weight of same animals in Figure 1J treated with sapanisertib and panobinostat. Y-axis indicates relative weight (normalized to day 0 weight of individual animal). X-axis indicates to day 0 weight of individual animal). X-axis indicates and panobinostat. Y-axis indicates relative weight (normalized to day 0 weight of individual animal). X-axis indicates days on treatment. Legend at right shows mouse number of each mouse.

## Supplementary Figure S3



**Supplementary Figure S3**. **A)** 90-8TL cells were treated with Nexturastat A at concentration indicated, with (black) or without (white) 100nM sapanisertib. Graph depicts the mean log2 fold change of cell number after 72 hours, relative to day 0 ±SD. At right, immunoblots depict acetylated lysine 9 on histone H3 (Ac. H3K9) and phosphorylated S6 (pS6). Total S6 and vinculin serve as a loading controls. **B)** 90-8TLs were treated with entinostat at indicated concentrations, with (black) or without (white) 100nM sapanisertib. At right, immunoblots depict acetylated lysine 9 on histone H3 (Ac. H3K9) and phosphorylated S6 (pS6). Total S6 and vinculin serve as a loading controls. **B)** 90-8TLs were treated with entinostat at indicated concentrations, with (black) or without (white) 100nM sapanisertib. At right, immunoblots depict acetylated lysine 9 on histone H3 (Ac. H3K9) and phosphorylated S6 (pS6). Actin serves as a loading control.



**Supplementary Figure S4**. Combined HDAC and mTOR inhibition increases reactive oxygen species in MPNSTs. **A)** Graph depicts relative mean fluorescence intensity of S462 cells stained with dichlorofluorescin diacetate (DCFDA), a dye that measures reactive oxygen species (ROS) and treated with vehicle (veh), sapanisertib (sap), vorinostat (vor), or vorinostat and sapanisertib (sap + vor) for 24 hours. Error bars indicate SD from three technical triplicates. \*p=0.007277, \*\*p=0.000441

## Supplementary Figure S5



Supplementary Figure S5. TXNIP is required for combination induced MPNST cell death. A) 90-8TLs were infected with lentiCRISPRv2 expressing guides targeting LACZ or TXNIP as indicated. Immunoblot of TXNIP levels after 16 hours of sapanisertib and vorinostat treatment. Vinculin serves as a loading control. B) As in A, 90-8TLs were infected with lentiCRISPRv2 expressing guides against LACZ or TXNIP as indicated and treated with sapanisertib and vorinostat. Left y-axis indicates the log2 of fold change in cell number after 3 days. The right y-axis indicates the percent change in cell number after 72 hours relative to day 0 on a log2 scale. Error bars indicate SD of technical triplicates. \*p=0.016. \*\*p=0.00413 C) 90-8TLs were transfected with pooled siRNAs targeting MAP3K5 (ASK1) or non-targeting (CTRL) and treated with vehicle, sapanisertib (100nM), vorinostat (2µM), or sapanisertib and vorinostat. Left y-axis indicates the log2 of fold change in cell number after 3 days. The right y-axis indicates the percent change in cell number after 72 hours relative to day 0 on a log2 scale. Error bars indicate SD of technical triplicates. Vehicle and combination data are reprinted from Fig. 3K for clarity. Immunoblot showing ASK1 in 90-8TLs 72 hours after transfection with indicated siRNA (MAP3K5 is the gene name for ASK1). Tubulin serves as a control. This immunoblot is a control for the experiment shown in Fig. 3K. D) Immunoblot showing GLUT1 in 90-8TLs 72 hours after transfection with indicated siRNA (SLC2A1 is the gene name for GLUT1). Tubulin serves as a control. This immunoblot is a control for the experiment shown in Fig. 3L.

Supplementary Table S1. Differentially expressed gene sets in MPNSTs treated with HDAC and mTOR inhibitors. To identify the pathways that were deregulated in response to vorinostat and rapamycin 90-8TLs were treated with vehicle, sapanisertib (100nM), vorinostat (2µM), or combined sapanisertib + vorinostat for 24 hours and a microarray analysis was performed. Recurrent pathway-related gene sets that were differentially expressed in the combination treated cells relative to the other treatment groups (vehicle and monotherapies) are shown. Gene sets were considered significant if they reached a nominal p-value of <0.005 in at least one of the Efron-Tibshirani GSA test, LS permutation test, or KS permutation test as in Figure 2A. Gene sets were considered recurrent if at least three related gene sets reached significance.

Oxidative Stress
CHUANG_OXIDATIVE_STRESS_RESPONSE_UP
CHUANG_OXIDATIVE_STRESS_RESPONSE_DN
WEIGEL_OXIDATIVE_STRESS_BY_TBH_AND_H2O2
GARGALOVIC RESPONSE TO OXIDIZED PHOSPHOLIPIDS TURQUOISE DN
KYNG_RESPONSE_TO_H202
GARGALOVIC RESPONSE TO OXIDIZED PHOSPHOLIPIDS TURQUOISE UP
MOOTHA_ROS
MOOTHA FFA OXYDATION
ER stress
REACTOME UNFOLDED PROTEIN RESPONSE
REACTOME_ACTIVATION_OF_CHAPERONES_BY_IRE1_ALPHA
REACTOME ACTIVATION OF CHAPERONE GENES BY ATF6 ALPHA
REACTOME_ACTIVATION_OF_CHAPERONES_BY_ATF6_ALPHA
Senescence and aging
FRIDMAN_SENESCENCE_DN
TANG_SENESCENCE_TP53_TARGETS_DN
LY_AGING_MIDDLE_DN
LY_AGING_OLD_DN
LY_AGING_PREMATURE_DN
KING NORMAL AGING UP
Нурохіа
HARRIS_HYPOXIA
WEINMANN_ADAPTATION_TO_HYPOXIA_DN
MENSE HYPOXIA UP
KRIEG HYPOXIA VIA DOMBA
WEIMMANN, ADAPTATION_IO_HYPOXIA_UP
IFN DEACOME ANTINUDAL MECHANION DV IEN CTIMULATED, CENEC
DER IEN GAMMA BESPONSE UP
REACTOME INTERFERON GAMMA SIGNALING
REACTOME_INTERFERON_ALPHA_BETA_SIGNALING
TNF
SANA TNF SIGNALING UP
PID TNFPATHWAY
PHONG_TNF_TARGETS_DN
PHONG TNF TARGETS UP
TIAN_TNF_SIGNALING_NOT_VIA_NFKB
VEGF
WESTON_VEGFA_TARGETS
ABE_VEGFA_TARGETS_2HR
WESTON_VEGFA_TARGETS_12HR
HELLEBREKERS_SILENCED_DURING_TUMOR_ANGIOGENESIS
Viral infection
BROWNE_HCMV_INFECTION_20HR_DN
ZHU CMV 24 HR UP
Radiation
AMUNDSON GAMMA RADIATION RESPONSE

SESTO RESPONSE TO UV C8
GENTILE UV RESPONSE CLUSTER D8
GENTILE_UV_RESPONSE_CLUSTER_D9
GENTILE UV LOW DOSE UP
GENTILE OV RESPONSE CLOSTER DS
GENTILE_UV_RESPONSE_CLUSTER_D7
DAZABD LIV BESPONSE CLUSTER G1
SMIRNOV_RESPONSE_TO_IR_6HR_DN
ZHOU CELL CYCLE GENES IN IR RESPONSE 6HR
SMIBNOV BESPONSE TO IB 2HB DN
SMIRNOV RESPONSE TO IR 2HR UP
ZHOU CELL CYCLE GENES IN IR RESPONSE 2HR
AMUNDSON DNA DAMAGE RESPONSE TP53
AMBROSINI FLAVOPIRIDOL TREATMENT 1P53
KEGG_P53_SIGNALING_PATHWAY
SCIAN CELL CYCLE TABGETS OF TP53 AND TP73 DN
WU_APOPTOSIS_BY_CDKN1A_VIA_TP53
KUMAMOTO RESPONSE TO NUTLIN 3A DN
KUMAMOTO RESPONSE TO NUTUR 24 UR
BRCA and DNA repair
HUNHADU_DHEASI_CANCER_DHCAI_VS_DHCA2
PUJANA BREAST CANCER WITH BRCA1 MUTATED UP
BAE BRCAL TARGETS LIP
MACLACHLAN_BRCA1_TARGETS_UP
PID ATR PATHWAY
OLSSON F2F3 TABGETS DN
CROSBY F2F4 TABGETS
REN_BOUND_BY_E2F
EGUCHI CELL CYCLE RB1 TARGETS
PEACTOME RECILIATION OF MITOTIC CELL CYCLE
MOLENAAR_TARGETS_OF_CCND1_AND_CDK4_DN
VERNELL BETINOBLASTOMA PATHWAY UP
Myc
SCHUHMACHER MYC TARGETS UP
MENSSEN_MYC_TARGETS
MENSSEN_MYC_TARGETS
COLLER MYC_LARGETS
MENSSEN_MYC_TARGETS           COLLER_MYC_TARGETS_UP           SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY
MENSSEN_MYC_IARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN
MENSSEN_MYC_IARGETS_UP         COLLER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         COLLER_MYC_TARGETS_DN
MENSSEN_MYC_IARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN
MENSSEN_MYC_IARGETS_UP         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN
MENSSEN_MYC_IARGETS_UP         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TFS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN
MENSSEN_MYC_IARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN
MENSSEN_MYC_IARGETS_UP         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED BY_MYC_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_ON         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF
MERSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGE_RESPONSE_120_HELA
MENSSEN_MYC_IARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_DESPONSE_120_HELA
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_ON         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_480_MCF10A
MENSSEN_MYC_IARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_20_MCF10A         AMIT_EGF_RESPONSE_20_HELA
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_ON         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_480_MCF10A         AMIT_EGF_RESPONSE_20_HELA         ZWANCE_GE_INTERPONSE_DN
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS         VIDENSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS OF MYC AND TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS TARGETS OF TP53 AND MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS OF MYC AND TFRC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS OF MYC AND TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_20_NEF10A         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_ON         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_480_MCF10A         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP         AMAGASHIMA_EGF_SIGNALING_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_ON         COLLER_MYC_TARGETS_ON         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_400_MCF10A         AMIT_EGF_RESPONSE_400_MCF10A         AMIT_EGF_RESPONSE_20 HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_ON         KIM_MYC_AMPLIFICATION_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         CEBALLOS_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP         AGASHIMA_EGF_SIGNALING_UP         ALURORA_B_PATHWAY
MENSSEN_MYC_IARGETS       UP         COLLER_MYC_TARGETS_UP       ODONNELL_TARGETS_OF_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_ON       COLLER_MYC_TARGETS_DN         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN       CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_SERUM_RESPONSE_DN       ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         DANG_REGULATED_BY_MYC_UP       EGF         AMIT_EGF_RESPONSE_120_HELA       AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP       NAGASHIMA_EGF_SIGNALING_UP         NAGASHIMA_EGF_SIGNALING_UP       PID_AURORA_B_PATHWAY         PID_AURORA_B_PATHWAY       PID_AURORA_A_PATHWAY
MENSSEN_MYC_IARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TFRC_UP         COLLER_MYC_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_AND_SERUM_RESPONSE_DN         ODONNELL_TARGETS_OF_MYC_AND_TFRC_DN         CEBALLOS_TARGETS_OF_MYC_AND_TERC_DN         ODONNELL_TARGETS_OF_MYC_AND_TERC_DN         DONNELL_TARGETS_OF_MYC_AND_TERC_DN         ODONNELL_TARGETS_OF_MYC_AND_TERC_DN         DANG_REGULATED_BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP         AMIT_EGF_RESPONSE_SIGNALING_UP         NAGASHIMA_EGF_SIGNALING_UP         NAGASHIMA_EGF_SIGNALING_UP         PID_AURORA_B_PATHWAY         PID_AURORA_A_PATHWAY         OUMONNEL_AURUMA_APATHWAY
MENSSEN MYC_IARGETS         COLLER MYC_TARGETS UP         SCHLOSSER MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS OF         COLLER_MYC_TARGETS_ON         COLLER_MYC_TARGETS_ON         CEBALLOS_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TP53_AND_TERC_DN         ODONNELL_TARGETS_OF_TP53_AND_TERC_DN         ODONNELL_TARGETS_OF_MYC_AND_TERC_DN         DANG REGULATED BY_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_40_MCF10A         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP         Aurora         PID_AURORA_B_PATHWAY         PID_AURORA_B_PATHWAY         PID_AURORA_A_TARGETS
Menssen MYC_TARGETS         COLLER_MYC_TARGETS_UP         SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY         ODONNELL_TARGETS_OF_MYC_AND_TERC_UP         COLLER_MYC_TARGETS_DN         CEBALLOS_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_TPS3_AND_MYC_DN         SCHLOSSER_MYC_TARGETS_OF_MYC_AND_TERC_DN         ODONNELL_TARGETS_OF_MYC_AND_TERC_DN         ODONNELL_TARGETS_OF_MYC_UP         EGF         AMIT_EGF_RESPONSE_120_HELA         AMIT_EGF_RESPONSE_480_MCF10A         AMIT_EGF_RESPONSE_20_HELA         ZWANG_EGF_INTERVAL_UP         NAGASHIMA_EGF_SIGNALING_UP         AMIT_EGF_RESPONSE_20_HELA         PID_AURORA_B_PATHWAY         PID_AURORA_A_PATHWAY         PID_AURORA_A_PATHWAY         OHASHI AURKA_TARGETS
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