

G9A AND GLP METHYLATE LYSINE 373 IN THE TUMOR SUPPRESSOR P53

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Supplementary Experimental Procedures

siRNA sequences:

G9a_siRNA#1: sense, UGAGAGAGGAUGAUUCUUAUU; anti-sense, 5'-
pUAAGAAUCAUCCUCUCUCAUU

G9a_siRNA#2: sense, GAACAUCGAUCGCAACAUCUU; anti-sense, 5'-
pGAUGUUGCGAUCGAUGUUCUU

Glp_siRNA#1: sense, CAAACAGCGUGGUCAAGUAUU; anti-sense, 5'-
pUACUUGACCACGCUGUUUGUU

Glp_siRNA#2: sense, CAAGAAAGGCCACUACGAAUU; anti-sense, 5'-
pUUCGUAGUGGCCUUUCUUGUU

Realtime PCR primer sequences:

G9a mRNA: forward, 5'-TCATGCTGCACCAAGACCTG-3'; reverse, 5'-
GGGAACTGAAGAAGGCGATG-3'

Glp mRNA: forward, 5'-TCCAAGGCCAAAGAGGTGAC-3'; reverse, 5'-
CACGGTCGAGGTGGTGTCT-3'

GAPDH mRNA: forward, 5'-TGGGCTACACTGAGCACCAG-3'; reverse, 5'-
GGGTGTCGCTGTTGAAGTCA-3'

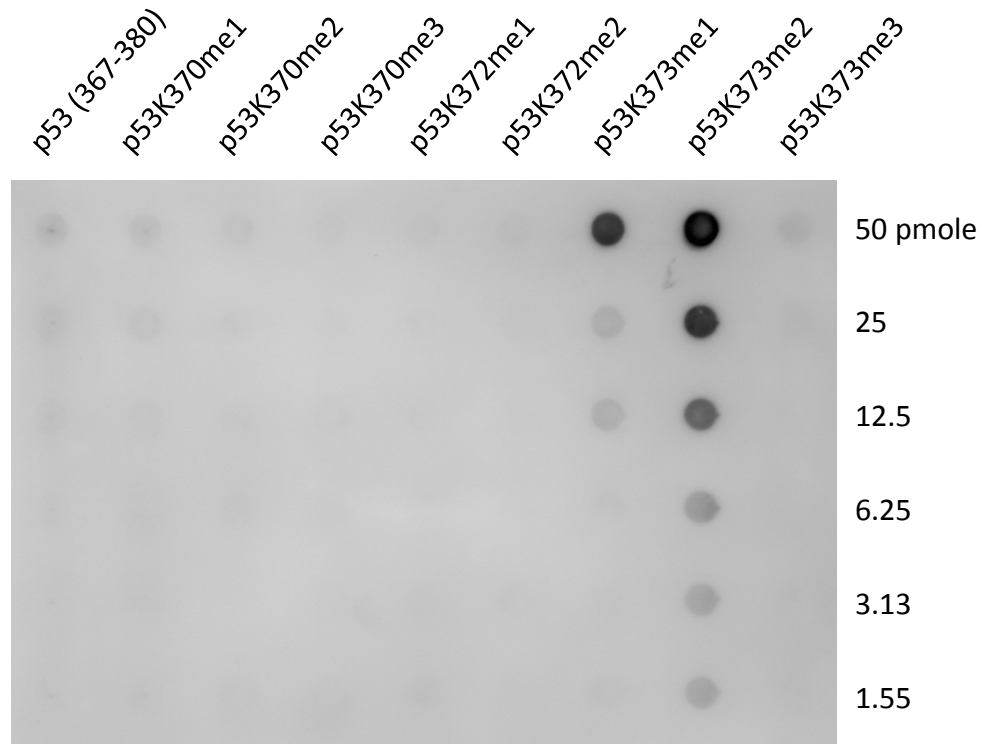


Figure S1. Dotblot analysis to characterize the p53K373me2 antibody. Serial dilution of p53 peptides (as indicated) are dotted on the PVDF membrane followed by blotting with 1:1,000 dilution of p53K373me2 antibody.

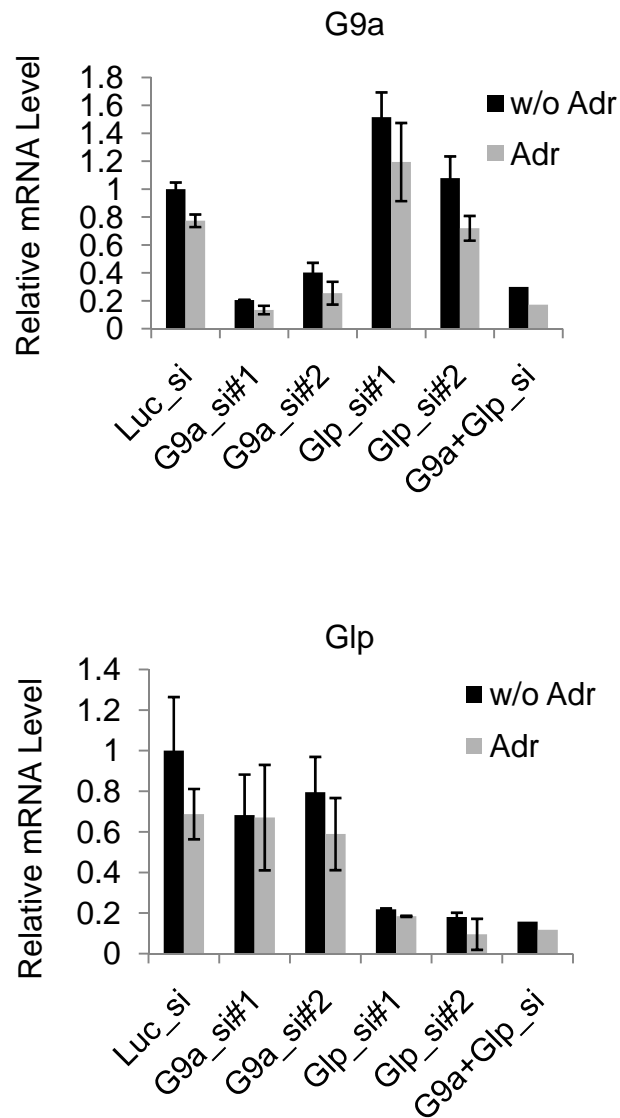


Figure S2. Realtime PCR analysis to assess the knockdown efficiency of siRNAs for luciferase (Luc_si), G9a (G9a_si#1 or #2), Glp (Glp_si#1 or #2) and both (G9a+Glp_si). Shown are normalized values of G9a or Glp mRNA levels to those of GAPDH (internal controls). The normalized value for Luci_si without adriamycin treatment was set as 1. Shown are the relative mRNA level from the averages of two independent repeats.