

Supplementary Materials

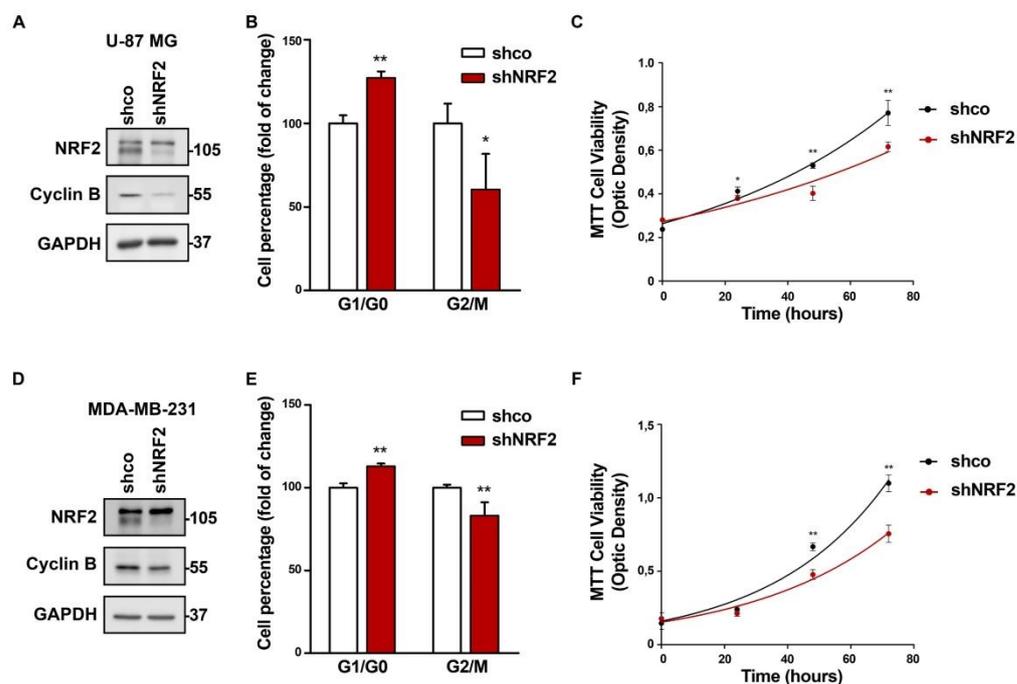


Figure S1. NRF2 increases G1/G0 cell fraction and reduces cell proliferation rate. (A–C) U-87 MG glioblastoma cells. (D–F) MDA-MB-231 breast adenocarcinoma cells. Both cell lines were transduced with lentiviral vectors containing control (shco) or shNRF2-1 (shNRF2). (A,D) Representative immunoblots of NRF2, Cyclin B, and GAPDH as a loading control. (B,E) Flow cytometry analysis of changes in cell cycle distribution in Hoechst-stained cells. Data are presented as mean \pm S.D. ** $p \leq 0.01$; * $p \leq 0.05$ according to a Student's *t*-test ($n = 3$). (C,F) Proliferation rate in shNRF2 cells based on MTT viability assay. The curve was fitted according to an exponential growth model with the least square fit. Data are presented as mean \pm S.D. ** $p \leq 0.01$; * $p \leq 0.05$ vs. shco according to a two-way ANOVA test ($n = 4$).

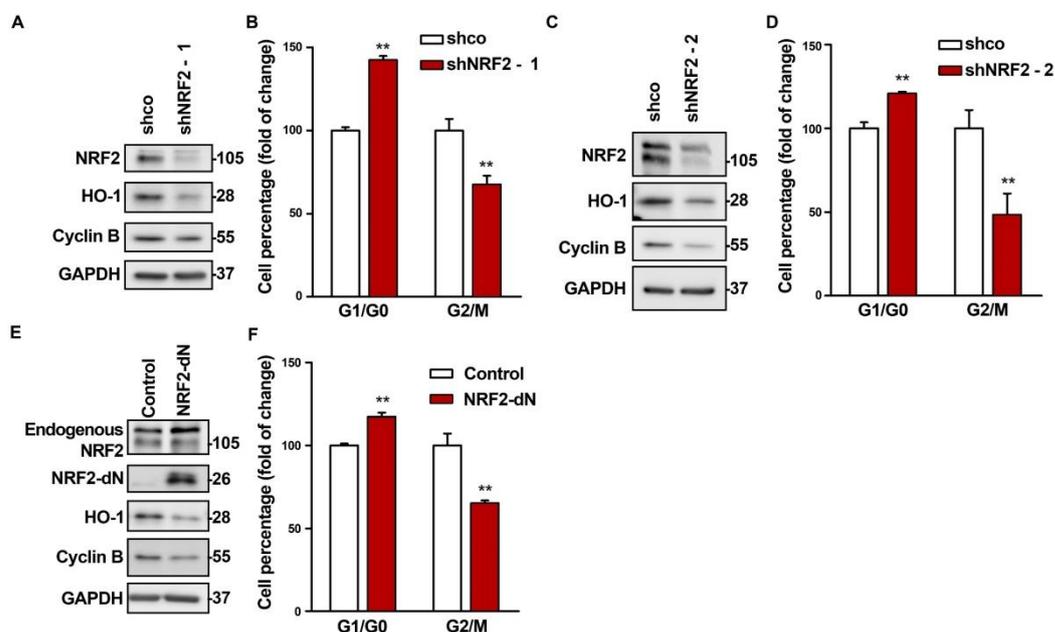


Figure S2. NRF2 downregulation increases G1 cell numbers. U-373 MG cells were transduced with different lentiviral vectors (A,B) Knockdown of NRF2 with lentiviral vectors containing control (shco) or shNRF2-1. (C,D) Knockdown of NRF2 with lentiviral vector containing control (shco) or

shNRF2-2. For A and C, representative immunoblots of NRF2, HO-1, Cyclin B, and GAPDH as a loading control. For B and D, flow cytometry analysis of G1 and G2/M phases in Hoechst-stained cells. Data are presented as mean \pm S.D. ** $p \leq 0.01$ according to a Student's t-test ($n = 3$). **(E,F)** U-373 MG glioblastomas cells were transduced with lentiviral vectors containing NRF2 dominant-negative mutant (NRF2-dN) or an empty vector (Control). **(E)** Representative immunoblots of NRF2, HO-1, Cyclin B, and GAPDH as a loading control. **(F)** Flow cytometry analysis of G1 and G2/M phases in Hoechst-stained cells. Data are presented as mean \pm S.D. ** $p \leq 0.01$ according to a Student's t-test ($n = 3$).