

Supplemental Fig. S1. Human papillomavirus-transformed cervical carcinoma cell-lines and E6 expressing transfected cells contain increased levels of HIF-1 α and HIF-2 α . (A) The hypoxic markers HIF- 1α and HIF-2 α were detected in HPV18+ (HeLa and C-4 I) and HPV16+ (SiHa) cervical carcinoma celllines by immunoblotting with densitometric quantification. (B) HT-1080 cells were transfected with expression constructs for HPV18 or HPV16 E6 (HA-tagged), or an empty CBS vector as negative control and the expression of HIF-1 α , HIF-2 α , and the viral E6 (HA) oncoprotein was detected by immunoblotting. Actin is shown as a protein loading control. N-value $= 3$. The asterisks denote statistical significance as determined using Student's t-test (**P* < 0.0332, ***P* < 0.0021).

Cervical cancer cell-lines

Supplemental Fig. S2. Hypoxia-independent mitochondrial targeting of the TIGAR protein in hrHPVtransformed cervical carcinoma cells. (A) Mitochondrial localization of the TIGAR protein in HPV16 transformed (SiHa and Ca Ski) and HPV18-transformed (MS751 and C-4 I) cervical carcinoma cell-lines was visualized by labeling the samples with MitoTracker Orange (red signal) and then performing immunofluorescence-confocal microscopy using an Anti-TIGAR primary antibody (green signal; *the MS751 cell-line also contains HPV45 sequences). Immortalized human HFL1 fibroblasts (HPV-negative) are shown for comparison and exhibited reduced, broadly diffuse TIGAR expression. DAPI nuclearstaining (blue signal) is provided for reference in the merged images. Scale bar, 20 μ m. The enlarged inset areas (bottom panels) depict colocalization between TIGAR and the MitoTracker Orange probe in hrHPVtransformed cells. (B) The hypoxia-independent mitochondrial targeting of TIGAR in transfected cells expressing the HPV18 E6 oncoprotein was further verified by cotransfecting the cells with CMV-TIGAR (FLAG-tagged) and labeling the cells with MitoTracker Orange, followed by immunofluorescenceconfocal imaging analysis. Scale bar, 20 μ m. The TIGAR (FLAG) protein in cells containing HPV18 E6 notably exhibited punctate colocalization with the MitoTracker Orange probe (see inset). (C) The expression of the FLAG-tagged TIGAR protein was detected by immunoblotting extracts from HPV18+ HeLa cells transfected with CMV-TIGAR (FLAG). An empty CBS vector was included as a negative control. Actin is shown as a protein loading control. N-value $= 3$.

Supplemental Fig. S3. The hrHPV E6-induced mitochondrial targeting of TIGAR requires the activation of cellular kinases. (A) HPV18-transformed HeLa cervical carcinoma cells were treated with various chemical kinase inhibitors (i.e., staurosporine, wortmannin, or SB203580; Millipore-Sigma) or a DMSO solvent control and then the cells were labeled with MitoTracker Orange (red signal) and immunofluorescence-confocal microscopy was performed using an Anti-TIGAR primary antibody (green signal) to visualize mitochondrial targeting of the TIGAR protein. UT, untreated cells. DAPI nuclear-

staining (blue signal) is provided for reference. Scale bar, 20 μ m. The enlarged inset areas show the subcellular distribution of the TIGAR protein. The inhibitors, wortmannin (inhibits PI3K-related kinases) and SB203580 (inhibits p38MAPK), but neither staurosporine nor the DMSO solvent control, markedly blocked the localization of TIGAR to mitochondrial membranes (see arrows) consistent with their ability to inhibit serine-phosphorylation of the TIGAR protein as shown in Fig. 2B. The relative fluorescenceintensities and overlap between the TIGAR and MitoTracker Orange signals are graphically represented in the 2.5D Merged images (lower right panels). (B) The expression of the constitutively-active mutants of PI3K (HA), PI5P4K (FLAG), and AKT (HA) as well as the TIGAR (FLAG-tagged) protein used in Fig. 2C was detected by immunoblotting. "ns" denotes a nonspecific band. (C) The involvement of PI3K/PI5P4K and MAPK in the E6-induced mitochondrial targeting of TIGAR suggests that Ras-signaling could activate these Ras effector-loop binding factors. Indeed, the HPV18+ cervical carcinoma cell-lines, HeLa, MS751, and C-4 I, were found to contain elevated levels of H-Ras as compared to (HPV-negative) HFL1 fibroblasts. The H-Ras protein was detected by immunoblotting and quantified by densitometry. Nvalue = 3. The asterisks denote statistical significance as determined using Student's t-test (**P* < 0.0332).

Supplemental Fig. S4. The HPV16+ cervical cancer cell-line, SiHa, exhibits hypoxia-independent phosphorylation of the TIGAR protein that is inhibited by the chemical kinase-inhibitors wortmannin and SB203580, but not staurosporine. SiHa cells were treated with various chemical kinase inhibitors as described and then immunoprecipitations were performed using phospho-specific monoclonal antibodies (Anti-Phospho-Serine and Anti-Phospho-Threonine) and Protein-G agarose. As a control for the phosphospecific antibodies, the SiHa cell extract was treated for 1 hr at 37°C with a recombinant Serine/threonineprotein phosphatase (2,000 U; Abcam) prior to immunoprecipitation (UT + Phos lanes). The input levels of TIGAR and Actin were detected by immunoblotting and are shown in the left panels. The immunoprecipitated phosphorylated TIGAR protein was detected by immunoblotting and quantified by densitometry (right panels). N-value = 3. The asterisks denote statistical significance as determined using Student's t-test (**P* < 0.0332, ***P* < 0.0021).

Supplemental Fig. S5. Human papillomavirus-transformed cells are generally more resistant to oxidative stress-induced cytotoxicity than HPV-negative cells. (A and B). 293 HEK (HPV-neg) cells and HPV18+ HeLa cells were treated with hydrogen peroxide (H₂O₂; 2 μ M) for 2 hrs and then the samples were stained with Annexin V-FITC and PI and the relative percentages of apoptotic cells were quantified by confocal microscopy and counting at 20x magnification. A DIC filter was included in the merged images to visualize all the cells (in B). Scale bar, 20 μ m. N-value = 3. The asterisks denote statistical significance as determined using unpaired two-tailed Student's t-tests (**P* < 0.0332).

Supplemental Fig. S6. High-risk HPV E6 oncoproteins prevent c-Myc-induced apoptosis in a TIGARdependent manner. (A and B) The ability of the viral E6 oncoprotein to prevent c-Myc-induced cellular apoptosis was evaluated by cotransfecting HT-1080 cells with CF-c-Myc and expression constructs for HPV18 E6 or HPV16 E6. Certain samples were also transduced with lentiviral-siRNA-*tigar* to inhibit endogenous TIGAR expression or an empty pLenti vector as negative control. The cells were cotransfected with C β F-c-Myc and CMV-TIGAR for comparison. As a positive apoptosis control, one sample was treated with CCCP (50 μ M). The cells were then stained with Annexin V-FITC (green signal) and PI (red signal) and confocal microscopy was performed to quantify the relative percentages of apoptotic cells (Annexin V-FITC and/or PI-positive cells) per field by counting triplicate visual fields at 200x magnification (n-value $=$ 3). Error bars represent mean \pm SD. N-value $=$ 3. Scale bar, 20 μ m. The asterisks denote statistical significance as determined using unpaired two-tailed Student's t-tests (**P* < 0.0332, ***P* < 0.0021).

Supplemental Fig. S7. The ability of the HPV18 E6 oncoprotein to protect against c-Myc-induced oxidative damage and genotoxicity is dependent upon TIGAR functions. (A) Intracellular ROS were detected using the fluorescent probe, CellROX-Deep Red, and the cells were visualized by confocal microscopy with the inclusion of a DIC filter in merged images. Representative micrographs are shown for HT-1080 cells that were cotransfected with CBF-c-Myc and an expression construct for HPV18 E6 and then transduced with either lentiviral-siRNA-*tigar* or an empty pLenti vector as negative control. The data were quantified and graphically depicted in Fig. 3B. (B) Mitochondrial damage/membrane depolarization (green signal) and normal polarized mt membranes (red signal) were visualized in cells expressing HPV18 E6/c-Myc and lentiviral-siRNA-*tigar* or an empty pLenti vector by staining the cells with the fluorescent JC-1 probe for analysis by confocal microscopy. Representative merged images are shown. The quantified data is graphically presented in Fig. 3C. (C) Oxidative DNA-damage was visualized in cotransfected cells expressing HPV18 E6/c-Myc and either lentiviral-siRNA-*tigar* or an empty pLenti vector by staining the cells using a 594 nm fluorescent Click-iT Plus TUNEL kit (Invitrogen) and then performing confocal microscopy. Hoechst 33342 nuclear-staining is provided in the merged images. The quantified data is shown in Fig. 4A. (D) Incorporation of the phosphorylated histone variant, γ H2A.X, at sites of damaged chromatin (red foci) in cells expressing HPV18E6/c-Myc and lentiviral-siRNA-*tigar* or an empty pLenti vector was visualized by immunofluorescence-confocal microscopy. DAPI nuclear-staining is provided for reference in the merged images. Enlarged inset areas are shown in the lower panels. The data was quantified and is graphically presented in Fig. 4B. Scale bars, 20 μ m. N-value = 3.

 $\mathbf C$ A HCT116 p53-null (CRISPR KO)

Supplemental Fig. S8. Expression of the p53 and TIP60 proteins in transfected cells. (A) The p53 protein in HCT116 *p53^{-/-}* (homozygous *p53* CRISPR KO) carcinoma cells that were transfected with expression constructs for HPV18 E6, c-Myc, and either wild-type p53 or an empty CS vector control, as shown in Figs. 5C and 5D, was detected by immunoblotting. (B) The ectopically expressed TIP60 (HA) protein was detected in transfected HPV18+ C-4 I cervical carcinoma cells by immunoblotting using a rabbit polyclonal Anti-HA (Y-11) primary antibody. "ns" denotes nonspecific bands. (C) The HPV18-transformed cervical carcinoma cell-lines, HeLa and C-4 I, were transfected with a pOZ-TIP60 (HA-tagged) expression construct or an empty C β S vector as shown in Figs. 5A and 5B, and the HA-tagged TIP60 (green signal) and p53 (red signal) proteins were detected by immunofluorescence-microscopy. DAPI nuclear-staining (blue signal) and DIC phase-contrast merged images are provided for reference. Scale bar, 20 μ m. N-value = 3.

Supplemental Fig. S9. The siRNA-inhibition of TIGAR expression inhibits in vivo tumorigenesis in a xenograft model of hrHPV-induced carcinoma. (A) To determine if TIGAR functions are required for HPVinduced tumor formation and disease progression in vivo, athymic NIH III-nude mice were subcutaneously engrafted over each hind flank with HPV18+ HeLa adenocarcinoma cells that had been transduced with either lentiviral-siRNA-*tigar* or an empty lentiviral (pLenti) vector as a negative control. Another group of animals was injected with the Vehicle alone for comparison (n-value = 12). The experimental animals were closely monitored over a period of 8 weeks for the development and growth of primary tumors at the injection sites. A Kaplan-Meier plot of event-free survival demonstrates that the HeLa/siRNA-*tigar* engrafted animals exhibited delayed tumor formation as compared to the HeLa/pLenti vector sample group.

(B and C) Although a similar number of animals (58% versus 66%) developed tumors in both the HeLa/siRNA-*tigar* and HeLa/pLenti vector experimental groups (in B), fewer HeLa/siRNA-*tigar* engrafted animals exhibited bilateral tumor-growth as compared to the HeLa/pLenti vector group (in C). (D) The HPV18+ HeLa/pLenti vector xenograft tumors express H-Ras, p53, and the angiogenic markers, HIF-1 α and VEGF. The primary tumor masses were harvested from experimental NIH III-nude animals that were engrafted with HPV18+ HeLa cells transduced with the empty pLenti vector (n-value = 12) and then fixed and immunostained using Anti-human Ki67 (huKi67), Anti-HIF-1 α , Anti-VEGF, Anti-H-Ras, Anti-p53, and Anti-HPV18 E6 primary antibodies. DAPI nuclear-staining is included for reference in the merged images. Immunofluorescence-confocal microscopy was performed to visualize the expression of HIF-1 α . VEGF, and p53 (green signals) in the huKi67-positive (red signal) engrafted HeLa tumor cells, as well as H-Ras expression (red signal) in the HPV18 E6-positive (green signal) HeLa tumor cells. The data shown is representative of triplicate sections analyzed for each primary tumor.

Supplemental Fig. S10. Primary HPV16+ cervical cancer clinical isolates contain detectable TIGAR, c-Myc, H-Ras, and p53 protein expression. (A) To determine how TIGAR, c-Myc, and p53 contribute to hrHPV-induced carcinogenesis, biopsied HPV16+ cervical carcinoma tissue samples (n-value = 9) were

kindly provided by the Pathology Shared Resource of the University of Hawaii Cancer Center and analyzed by immunofluorescence-confocal microscopy to quantify the relative percentages of HPV16 E6-positive cells (green signal) that contain TIGAR, c-Myc, or p53 protein expression (red signal) as shown in Fig. 6G. DAPI nuclear-staining (blue signal) and DIC phase-contrast are provided in the merged images. Human HFL1 fibroblasts were immunostained as an HPV-negative antibody (Ab) control (right panels). (B) The expression of H-Ras (red signal) was visualized within the HPV16 E6-positive tumor cells using immunofluorescence-confocal microscopy. Scale bar, 20 μ m. The data in A and B is representative of triplicate fields for each tissue section.

Supplemental Table S1. Antibodies, Chemicals and reagents, Cell-lines, Clinical samples, Animal strains, Lentiviruses, and DNA plasmids.

Antibodies

Chemicals and reagents

Cell-lines

Primary HPV16+ cervical carcinoma clinical samples

Animal strains

Lentiviruses

DNA plasmids

