Interaction between the human papillomavirus 16 E7 oncoprotein and gelsolin ignites cancer cell motility and invasiveness

Supplementary Materials



Supplementary Figure S1: E7 expression in CC cell lines. (A) *Left panel*. Cytofluorimetric analysis after single cell staining with anti-E7 MAb. Numbers represent the median fluorescence intensity obtained in a representative experiment. *Micrographs*. IVM analysis after anti-E7 MAb/Hoechst double cell staining. Magnification, $1,300\times$. (B) Western blot analysis using an anti-E7 MAb. E7 positive control cells displayed a protein with a higher apparent molecular mass due to the addition of a T7 tag sequence. Loading control was evaluated using anti- α -tubulin MAb. Density of each band in the same gel was analyzed and the densitometric E7/ α -tubulin ratios are shown in the bar graph in the right panel. (*) indicates P < 0.01 vs. C-33A.



Cells transfected with a construct emitting in FL1 channel to restrict the analysis to the transfected cells only В



Cells transfected with a construct emitting in FL1 channel stained or not with an anti-BrdU MAb

Positive sample: MAb α-BrdU+α-mouse-Cy5+PI

Supplementary Figure S2: Experimental set up for differential analyses of transfected and non-transfected cells. Exemplification of a flow cytometry analysis to differentiate transfected and non-transfected cells. (A) Left panel. Non-transfected cells were selected on the basis of their physic parameters (R1 region). Right panel. Analysis of FL1 fluorescence emission in R1 cells subpopulation to define FL1-negative and FL1-positive (R2) regions. (B) Left panel. Selection of a population of transfected cells on the bases of its physic parameters (R1). Right panel. FL1 fluorescence emission analysis of cells included in R1 region to quantify the percentage of actually transfected cells on the basis of FL1 negative and positive regions defined in non-transfected cells sample. (C) Seventy-two hours after transfection, cells were incubated with BrdU and then fixed and permeabilized as reported in Material and Methods. After this step, each experimental sample was divided into two: a negative control (IgG1 + anti-mouse Cy5-conjugated + PI) and a positive sample (MAb anti-BrdU + anti-mouse Cy5-conjugated + PI). Left panel. Multiparametric analyses of negative control (FL1, FL2 and FL4 fluorescence channels) to distinguish a FL4-negative and a FL4-positive region (R3). Right panel. Multiparametric analyses of a positive sample (FL1, FL2 and FL4 fluorescence channels) to evaluate the percentage of cells in FL4- positive region (R3, cells positive to anti-BrdU). Analysis was restricted to FL1 positive (transfected) cells only.

Negative control: Mouse IgG+a-mouse-Cy5+PI

Control samples of siRNA experiments

Supplementary Figure S3: siRNA experiment control samples. (A) Cytofluorimetric analysis of fluorescence emission in cells transfected with FITC-siRNA. The percentage of FITC-positive cells (corresponding to transfected cells) was considered indicative of the transfection efficiency. (B) Cytofluorimetric evaluation of E7 expression level 48 h after siRNA transfection. The numbers represent the median fluorescence intensity. A representative experiment among three is shown.

Fibronectin determination

Supplementary Figure S4: Fibronectin determination in C-33A, SiHa and CaSki cells. Western blot analysis of fibronectin (FN) on C-33A, SiHa and CaSki cells transfected with control siRNA and E7 siRNA. FN appears as multiple bands, due to transcriptional and post-translational modifications (alternative splicing and glycosylation) of the protein. Loading control was evaluated using an anti-α-tubulin MAb.