

# The Intracellular Delivery Of Anti-HPV16 E7 scFvs Through Engineered Extracellular Vesicles Inhibits The Proliferation Of HPV-Infected Cells

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**Purpose:** Single-chain variable fragments (scFvs) are one of the smallest antigen-binding units having the invaluable advantage to be expressed by a unique short open reading frame (ORF). Despite their reduced size, spontaneous cell entry of scFvs remains inefficient, hence precluding the possibility to target intracellular antigens. Here, we describe an original strategy to deliver scFvs inside target cells through engineered extracellular vesicles (EVs). This approach relies on the properties of a Human Immunodeficiency Virus (HIV)-1 Nef mutant protein referred to as Nef<sup>mut</sup>. It is a previously characterized Nef allele lacking basically all functions of wt Nef, yet strongly accumulating in the EV lumen also when fused at its C-terminus with a foreign protein. To gain the proof-of-principle for the efficacy of the proposed strategy, the tumor-promoting Human Papilloma Virus (HPV)16-E7 protein was considered as a scFv-specific intracellular target. The oncogenic effect of HPV16-E7 relies on its binding to the tumor suppressor pRb protein leading to a dysregulated cell duplication. Interfering with this interaction means impairing the HPV16-E7-induced cell proliferation.

**Methods:** The Nef<sup>mut</sup> gene was fused in frame at its 3'-terminus with the ORF coding for a previously characterized anti-HPV16-E7 scFv. Interaction between the Nef<sup>mut</sup>-fused anti-HPV16-E7 scFv and the HPV16-E7 protein was tested by both confocal microscope and co-immunoprecipitation analyses on co-transfected cells. The in cis anti-proliferative effect of the Nef<sup>mut</sup>/anti-HPV16-E7 scFv was assayed by transfecting HPV16-infected cells. The anti-proliferative effect of EVs engineered with Nef<sup>mut</sup>/anti-HPV16-E7 scFv on HPV16-E7-expressing cells was evaluated in two ways: i) through challenge with purified EVs by a Real-Time Cell Analysis system and ii) in transwell co-cultures by an MTS-based assay.

**Results:** The Nef<sup>mut</sup>/anti-HPV16-E7 scFv chimeric product is efficiently uploaded in EVs, binds HPV16-E7, and inhibits the proliferation of HPV16-E7-expressing cells. Most important, challenge with cell-free EVs incorporating the Nef<sup>mut</sup>/anti-HPV16-E7 scFv led to the inhibition of proliferation of HPV16-E7-expressing cells. The proliferation of these cells was hindered also when they were co-cultured in transwells with cells producing EVs uploading Nef<sup>mut</sup>/anti-HPV16-E7 scFv.

**Conclusion:** Our data represent the proof-of-concept for the possibility to target intracellular antigens through EV-mediated delivery of scFvs. This finding could be relevant to design novel methods of intracellular therapeutic interventions.

**Keywords:** single-chain variable fragments, Human Papilloma Virus E7 protein, exosomes, human immunodeficiency virus Nef protein, scFv intracellular delivery

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## Introduction

Monoclonal antibodies (mAbs) are largely used in therapy, diagnostics, and biotechnology. However, in view of their large heterotetrameric composition, the

efficacy of whole Ab molecules is limited for a number of applications. The complex Ab structure can be reduced in several formats without affecting the binding efficacy. Single-chain variable fragment (scFv) is one of the smallest molecule retaining antigen-binding activity.<sup>1,2</sup> This molecule consists of Ab variable regions of both heavy and light chains joined by a peptide linker. Although scFvs have the invaluable advantage of being expressed by a single open reading frame (ORF), their access to intracellular targets is severely impaired by plasma membrane. The most popular way to provide target cells with scFvs relies on the use of gene therapy. In this manner, transduced cells express the specific scFv (in this case referred to as intrabody), which can bind the intracellular target.<sup>3</sup> The effects of gene therapy-driven scFv delivery are operative in actually transduced cells, with little or no benefit for neighboring ones. Our ambition was developing a method to extend the beneficial effects induced by vector-expressed scFvs from the producer to bystander cells. To this aim, we envisioned a strategy relying on *in vivo* engineering of extracellular vesicles (EVs) through a method we recently described.<sup>4</sup>

EVs comprise a heterogeneous population of membrane vesicles whose size ranges from 50 to 500 nm. They are classified into two major categories: exosomes (50–200 nm) and microvesicles (150–500 nm).<sup>5</sup> Exosomes originate intracellularly upon inward invagination of endosome membranes<sup>6</sup> and formation of intraluminal vesicles (ILVs), which then become part of multivesicular bodies (MVBs). They can traffic to either lysosomes for degradation or plasma membrane. In the latter case, MVBs release their contents in the extracellular space upon fusion with plasma membrane. EVs formed by direct budding of plasma membrane are referred to as microvesicles.<sup>7</sup>

We have previously identified a Human Immunodeficiency Virus (HIV)-1 Nef mutant protein (i.e., Nef<sup>mut</sup>) acting as an exosome-anchoring protein.<sup>8</sup> Nef<sup>mut</sup> is a functionally defective protein mutant lacking the Nef functions typically associated with HIV pathogenesis.<sup>9</sup> It exhibits extraordinary levels of incorporation into both exosomes and microvesicles, ranging from 50 to 100 fold those of the wild-type isoform.<sup>8</sup> Nef<sup>mut</sup> can be fused to proteins of choice while retaining its EV-anchoring properties.<sup>10–12</sup> In this way, Nef<sup>mut</sup> allows incorporation into EVs of large amounts of the products fused to it, which thus remain protected from external neutralization/degradation factors.

In an effort to develop a novel approach to deliver scFv into target cells, we planned to exploit the strategy based

on EV engineering we previously described.<sup>4</sup> It relies on the intracellular expression of a DNA vector expressing the product of fusion between Nef<sup>mut</sup> and the ORF of choice. This event leads to the release of EVs engineered with the fusion proteins whose entry into target cells allows intracellular delivery of the EV cargo. In the case Nef<sup>mut</sup> is fused with an ORF coding for an scFv, this method has the potential to target and inhibit intracellular antigens in DNA vector expressing cells as well as in bystander ones through the cell-to-cell transfer of scFv-engineered EVs.

Here, we demonstrate the possibility to engineer EVs with scFvs and provide experimental evidence supporting the idea that EV-incorporated scFvs can be functional in bystander cells.

## Materials And Methods

### DNA Constructs

The previously described ORF coding for the anti-HPV16-E7 43M2 scFv<sup>13,14</sup> was PCR-amplified using primers including *Apa* I (forward: 5' GGCCGGGCCCATGGCCGAGGTGC AGCTGGTGG 3') and *Acc* I (reverse: 5' CCGGGTCTACCT ACTTGTCATCGTCGCCTTGTAG 3') restriction sites. The PCR product was then inserted in *Apa* I/*Acc* I digested pTarget-Nef<sup>mut</sup> expression vector<sup>10</sup> to generate an in frame Nef<sup>mut</sup>-43M2 scFv ORF. The ORF coding for the anti-glucose oxidase from *Aspergillus niger*<sup>15</sup> was PCR-amplified from the previously described expression vector using primers including *Apa* I (forward: 5' ATTGGGCCCGCCATGGCCGAG 3') and *Sal* I (reverse: 5' ATTGTCGACCTACTAATGGTGATG 3') restriction sites, and cloned in *Apa* I/*Sal* I restriction sites of pTarget-Nef<sup>mut</sup> to generate an in frame Nef<sup>mut</sup>/GO scFv ORF. All restriction enzymes were from New England Biolabs. A flag tag was inserted at the C-terminus of both Nef<sup>mut</sup>-based scFvs. The DNA vector expressing HPV16-E7 fused to the red fluorescent protein was kindly provided by David Pim, ICGEB, Trieste. The pcDNA3 vector expressing a HPV16-E7 coded by a nucleotide sequence optimized as previously described<sup>16</sup> and 6×His tag flagged at its C-terminus was obtained as a synthesis product from Eurofins. The DNA vectors expressing Nef<sup>mut</sup> and sg25GFP have been already described.<sup>8,9</sup>

### Cell Cultures, Co-Cultures, And Transfections

HEK293T (ATCC, CRL-11268), SiHa (ATCC HTB-35), HeLa (ATCC, CCL-2), and TC-1 (a generous gift of prof.

Wu, Johns Hopkins Medical Institutes, Baltimore, MD) cells were grown in 10% heat-inactivated fetal calf serum (FCS)-supplemented Dulbecco's modified Eagle's (DMEM, Sigma). Transwell co-cultures were carried out in 6-well plates using Cell Culture Insert Falcon Membrane (25 mm diameter, 0.4  $\mu$ m pore size, Becton Dickinson). Transfection assays were carried out by the Lipofectamine 2000-based method (Invitrogen, Thermo Fisher Scientific), which, with the exception of HEK293T cells, was performed by adding liposomes to freshly trypsinized cells. In detail, for a 10 cm diameter dish,  $5 \times 10^6$  cells were seeded the day before transfection in medium without antibiotics. The day of transfection, the medium volume was brought to 9 mL, and 1 mL of transfection mix (i.e., 20  $\mu$ L of Lipofectamine plus 30  $\mu$ g of DNA in DMEM) was added after 20 mins incubation at room temperature. After additional 24 hrs, the medium was replaced.

## Exosome Isolation And Titration

Twenty-four hours after HEK293T cell transfection, cultures switched to fresh complete medium containing 5% exosome-depleted FCS, which was obtained after ultracentrifugation at  $70,000 \times g$ , 3 hrs at 4°C. Cell supernatants were harvested from 48 to 72 hrs after transfection and centrifuged at  $500 \times g$  for 10 mins. Clarified supernatants were therefore processed for exosome purification by differential centrifugations as previously described.<sup>17</sup> Briefly, supernatants underwent a first ultracentrifugation at  $10,000 \times g$  for 30 mins at 4°C, followed by 0.22  $\mu$ m pore-size filtration. Supernatants were then ultracentrifuged at  $70,000 \times g$  for 2.5 hrs at 4°C. Pelleted vesicles were washed with phosphate-buffered saline (PBS), and ultracentrifuged again at  $70,000 \times g$  for 1 hr at 4°C. Pellets were finally resuspended in either DMEM (1:200 of the initial supernatant volume) or 0.1% Triton-X100 TNE (1:300 of the initial volume).

Exosome preparations were quantified by measuring the activity of acetylcholinesterase (AChE), i.e., a typical exosome marker,<sup>18</sup> through the Amplex Red kit (Molecular Probes) following the manufacturer's recommendations. The AChE activity was measured as mU/mL, where 1 mU is defined as the amount of enzyme which hydrolyzes 1 pmole of acetylcholine to choline and acetate per minute at pH 8.0 at 37°C.

## Western Blot

Western blot analysis on lysates of up to  $5 \times 10^6$  cells was performed by washing cells twice with  $1 \times$  PBS (pH 7.4) and lysing them for 20 mins on ice with lysis buffer (20 mM HEPES pH 7.9, 50 mM NaCl, 10 mM EDTA, 2 mM

EGTA, 0.5% nonionic detergent IGEPAL CA-630, 0.5 mM dithiothreitol, 20 mM sodium molybdate, 10 mM sodium orthovanadate, 100 mM sodium fluoride, 10  $\mu$ g/mL leupeptin, 0.5 mM phenylmethylsulfonyl fluoride). Whole cell lysates were centrifuged at  $6000 \times g$  for 10 mins at 4°C. The protein concentration of cell extracts was determined by the Lowry protein quantitation assay. Aliquots of cell extracts containing 30 to 50  $\mu$ g of total proteins were resolved by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred by electroblotting on a 0.45- $\mu$ m pore size nitrocellulose membrane (Amersham) overnight using a Bio-Rad Trans-Blot. Exosomes were lysed and analyzed as described for cell lysates. For immunoassays, membranes were blocked with 5% non-fat dry milk in PBS containing 0.1% Triton X-100 for 1 hr at room temperature, then incubated overnight at 4°C with specific antibodies diluted in PBS containing 0.1% Triton X-100. The antibodies used in immunoblots were: 1:2000 diluted sheep polyclonal anti-Nef ARP444 (a generous gift from M. Harris, Leeds, UK); 1:500 diluted rabbit horseradish peroxidase (HRP)-conjugated-anti- $\beta$ -Actin (13E5, cat. 5125) (Cell Signaling Technology); 1:500 diluted H-270 rabbit polyclonal anti-Alix (Santa Cruz, cat. sc99010). When appropriate, immune complexes were detected by 1:2000 diluted HRP-conjugated secondary antibodies (Merck, cat. AP132P and AP147P). Finally, signals were revealed by enhanced chemiluminescence reaction (ECL, Millipore).

## Co-Immunoprecipitation Assay

Cells expressing 6 $\times$ His-tagged HPV16-E7 and/or Nef<sup>mut</sup>/43M2 were lysed in  $1 \times$  RIPA lysis buffer (cat. 20188, Millipore) in the presence of protease inhibitors. The cell lysis was carried out also in the presence of 10  $\mu$ M of the HPV16-E7<sub>10-21</sub> peptide previously identified as the major binding domain of 43M2<sup>19</sup> to avoid the HPV16-E7-Nef<sup>mut</sup>/43M2 binding in cell lysates. A total of 500  $\mu$ g of the cell lysates was incubated o.n. at 4°C with 1  $\mu$ g of anti-6 $\times$ His mAb (cat. 13964A, Bio-Rad), and then for 4 hrs at 4°C with 20  $\mu$ L of Dynabeads conjugated with goat anti-mouse IgGs (Fisher Scientific), previously saturated in FCS. Co-immunoprecipitates were then magnetically separated, washed 5 times in NaCl 100 mM, Tris-Hcl 30 mM, Tween 0.5%, pH 7.4, and finally revealed by Western blot analysis using both ARP444 sheep anti-Nef and rabbit anti-6 $\times$ His Abs (Bethyl, cat. A190-114A).

## Confocal Microscope Analysis

A total of  $4 \times 10^4$  HEK293T cells were seeded on chamber glass slides (BD Biosciences), and co-transfected with vectors expressing HPV16-E7/RFP and either Nef<sup>mut</sup>/GO or Nef<sup>mut</sup>/43M2. Twenty-four hours later, cells were labeled with 1:2,000 diluted anti-Nef mAb MATG020 (kindly provided by O. Schwartz, Paris, France), followed by incubation with 1:2500 diluted Alexa 488-conjugated goat anti-mouse (Invitrogen, cat. A28175). Images were acquired using a Leica TCS SP5 confocal microscope and analyzed by the LAS AF version 1.6.3 software (Leica microsystems).

## Cytofluorimetric Analysis

Cells transfected with a GFP-expressing DNA vector were trypsinized, fixed with a  $1 \times$  PBS 2% v/v paraformaldehyde (PFA), and the fluorescence analyzed in a BD FACScalibur flow cytometer after labeling with  $1 \mu\text{g/mL}$  of 7-aminoactinomycin D (AAD) (Sigma) to exclude died cells. For Ki-67 detection, cells were fixed and permeabilized with Cytofix/Cytoperm solutions (BD Pharmingen) according to the manufacturer's instructions, and then labeled with 1:1000 diluted Alexa 488-conjugated anti-Ki-67 mAb (BioLegend, cat. 350508) for 30 mins at room temperature. Finally, cells were analyzed in a BD FACScalibur cytometer.

Fluorescent exosomes isolated from supernatants of transfected cell cultures were analyzed upon binding with aldehyde/sulfate latex beads (Molecular Probes, Thermo Fisher). To this end, 2 mU equivalent of exosomes were incubated with  $5 \mu\text{L}$  of beads overnight at room temperature on a rotating plate, washed, and incubated for 1 hr at  $4^\circ\text{C}$  with either 1:1000 diluted phycoerythrin (PE)-conjugated anti-CD63 (clone H5C6, BD Pharmingen, cat. 556020) or the same concentration of a control isotype. Finally, beads were resuspended and fixed in  $1 \times$  PBS-2% v/v paraformaldehyde, and FACS analyzed.

For the detection of exosome internalization into SiHa cells,  $5 \times 10^4$  cells were seeded in microwells and challenged with different amounts of fluorescent exosomes by spinoculation at 1200 g for 30 mins at  $4^\circ\text{C}$ . Then, the cells were incubated either at  $37^\circ\text{C}$  or  $4^\circ\text{C}$ . At different times, i.e., from 1 to 5 hrs, the cells were treated for 10 mins with cold trypsin, harvested, washed, fixed, and analyzed by FACS.

## Anti-HPV-E7 ELISA

Methods to produce, purify, and characterize recombinant HPV16-E7, HPV18-E7, and the 43M2 scFv have been already described.<sup>14</sup> ELISA was carried out as published

with minor modifications. Briefly, recombinant HPV16-E7 and HPV18-E7 proteins were adsorbed onto Maxisorp microtiter plates (NUNC, Thermofisher Scientific) overnight at  $4^\circ\text{C}$ . Afterwards, wells were saturated with 2% non-fat dry milk (Bio-Rad) at  $37^\circ\text{C}$  for 2 hrs, and then incubated at  $37^\circ\text{C}$  overnight with  $3 \mu\text{g/mL}$  of purified scFv43M2 followed by 1:3000 dilution of anti-flag M2 mAb (Sigma) for 1 hr. Wells were then washed, and a 1:20,000 dilution of anti-mouse HRP-conjugated polyclonal IgG (SBA Inc.) was added for 40 mins. After washing, the antigen-antibody complexes were detected by tetramethylbenzidine (BM blue, POD substrate, Roche). After 30 mins, the enzymatic reaction was stopped, and optical density (OD) measured at 450 nm.

## Cell Viability

Cell viability was assessed by both trypan blue exclusion method and MTS CellTiter 96® AQueous One Solution Cell Proliferation Assay (Promega) according to manufacturer's instructions. It is based on the use of the tetrazolium inner salt which in metabolically active cells is reduced by NAD(P)H-dependent dehydrogenase enzymes to a colored formazan product that is soluble in cell culture media.

Alternatively, cell viability was monitored after seeding cells in E-plate of xCELLigence RTCA DP system (ACEA Biosciences Inc., San Diego, CA, USA). In detail, 24 hrs after transfection,  $4 \times 10^4$  cells were seeded in each well, and cell proliferation monitored in real time for 80 to 96 hrs. The impedance (referred to as Cell Index) of each well of RTCA device was monitored at 15-min to 3-hr intervals in the first 24 hrs, and at 15-min intervals in the remaining time frames.

In the case of exosome-challenged cells,  $4 \times 10^4$  cells were resuspended in culture medium in the presence of exosome-deprived FCS and then seeded in duplicate in microwells of xCELLigence RTCA DP system. After 24 hrs, cells were challenged with 2 mU of each exosome preparation in a total of  $10 \mu\text{L}$  of medium. Cell Index of each well of RTCA device was monitored at 15-min intervals for the 96-hr observation.

## Statistical Analysis

Statistical tests were performed using Graph Pad InStat3 software. Either nonparametric Mann-Whitney *U*-Test or parametric Student *T* Test was used to determine *p* values, as appropriate. All statistical analyses were carried out at a two-sided 5% significance level.

## Results

### Recovery Of scFv-Engineered Exosomes

43M2 is a previously characterized anti-HPV16-E7 scFv.<sup>13,14</sup> In the form of intrabody, it efficiently counteracts the proliferation of HPV16-E7 expressing cells by means of the inhibition of HPV16-E7-pRb interaction.<sup>19,20</sup>

We generated DNA vectors expressing the product of fusion of Nef<sup>mut</sup> with this scFv as well as an scFv binding the glucose oxidase (GO) from *Aspergillus niger*<sup>15</sup> as negative control (Figure 1A). Western blot analysis of HEK293T cells transfected with these vectors indicated that both fusion products are produced in the absence of apparent truncated forms. Most important, exosomes isolated from supernatants of transfected cultures were found to upload well-detectable amounts of the Nef<sup>mut</sup>/scFv fusion product (Figure 1B and C). Conversely, 43M2 does not associate with exosomes when expressed in the absence of the Nef<sup>mut</sup> fusion (Figure 1D).

We concluded that, as already reported for a large number of both cellular and viral products,<sup>8,10–12</sup> scFvs also can be efficiently incorporated into exosomes when fused with Nef<sup>mut</sup>.

### Intracellular Interaction Between HPV16-E7 And Nef<sup>mut</sup>/43M2

Next, we investigated whether the 43M2 scFv retained the already described HPV16-E7 binding activity when expressed in the form of product of fusion with Nef<sup>mut</sup>. It is known that HPV-E7 prevalently accumulates in cell nucleus, where it binds and inactivates the tumor-suppressor protein pRb.<sup>21</sup> For its part, Nef<sup>mut</sup> prevalently localizes both in cytoplasm and, in view of its N-terminal myristoylated and palmitoylated ends, at the intracellular membranes.<sup>4</sup>

With these premises, we tried to establish whether co-expression of Nef<sup>mut</sup>/43M2 and HPV16-E7 would result in changes in HPV16-E7 intracellular localization. HEK293T cells were co-transfected with DNA vectors expressing HPV16-E7/RFP together with either Nef<sup>mut</sup>/43M2 or Nef<sup>mut</sup>/GO. After 48 hrs, the cells were labeled with an anti-Nef mAb and analyzed by confocal microscope. We observed that the co-expression of Nef<sup>mut</sup>/43M2, but not Nef<sup>mut</sup>/GO, led to a quite impressive HPV16-E7 delocalization from nucleus to cytoplasm (Figure 2A). Consistently, regions of scFv co-localization with HPV16-E7 can be detected in cells expressing Nef<sup>mut</sup>/43M2, but not Nef<sup>mut</sup>/GO.

Next, to strengthen the idea that Nef<sup>mut</sup>/43M2 indeed binds HPV16-E7 intracellularly, co-immunoprecipitation assays were carried out. To this end, 293T cells were co-transfected

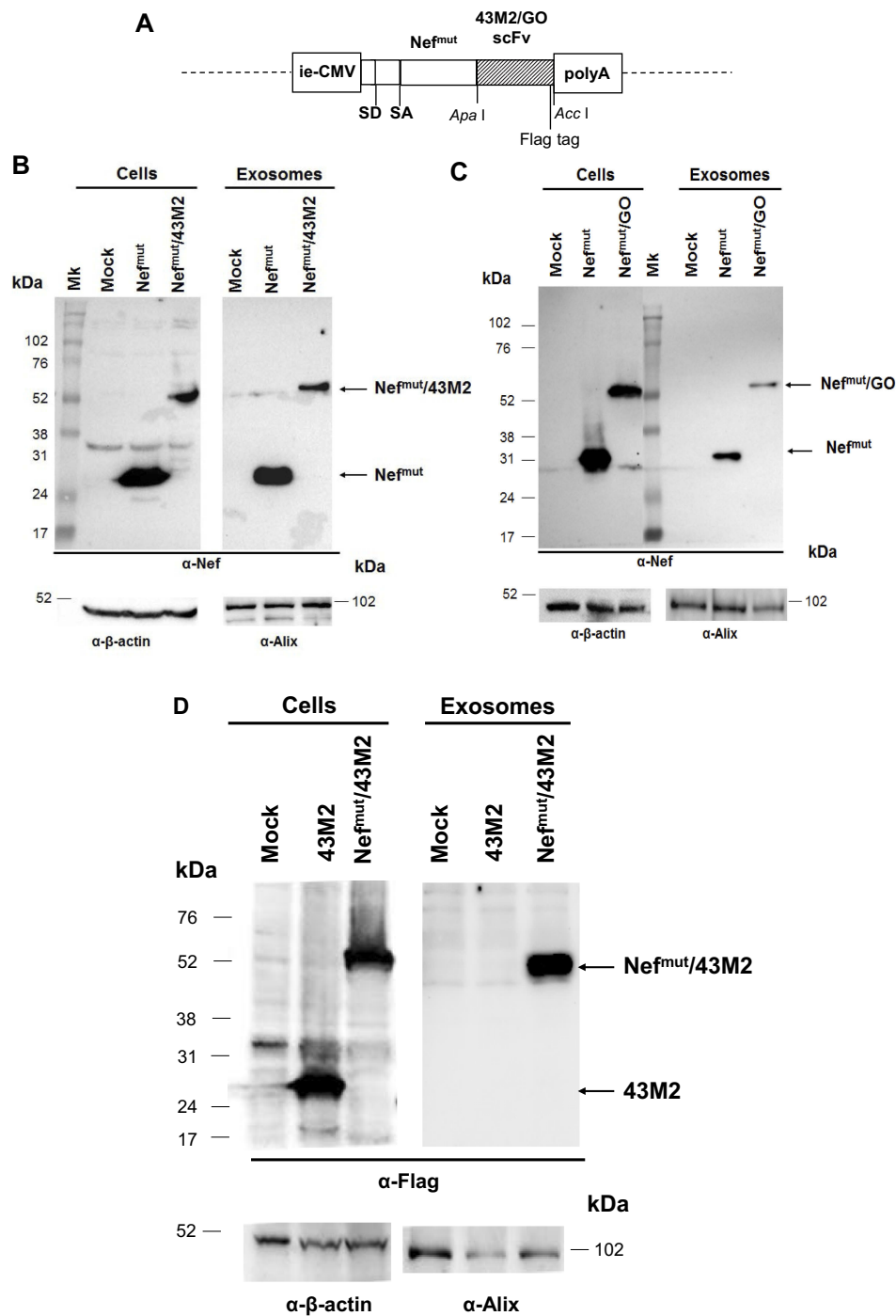
with DNA vectors expressing Nef<sup>mut</sup>/43M2 and an optimized HPV16-E7 sequence<sup>16</sup> tagged at its C-terminus with 6×His. As control, cells were either transfected with the DNA vector expressing Nef<sup>mut</sup>/43M2 alone or mock-transfected. After two days, the cells were harvested, lysed, and co-immunoprecipitation assay was performed using an anti-6×His mAb. Immunocomplexes were revealed with anti-Nef Abs and, as control, rabbit polyclonal anti 6×His Abs. As shown in Figure 2B, Nef<sup>mut</sup>/43M2 was detectable only in immunocomplexes precipitated from lysates of cells co-expressing HPV16-E7. This result confirmed that Nef<sup>mut</sup>/43M2 binds HPV16-E7 inside the cells.

Together, these data strongly support the idea that Nef<sup>mut</sup>/43M2 actually recognizes intracellular HPV16-E7, thus indicating that the fusion with Nef<sup>mut</sup> does not affect the scFv binding activity.

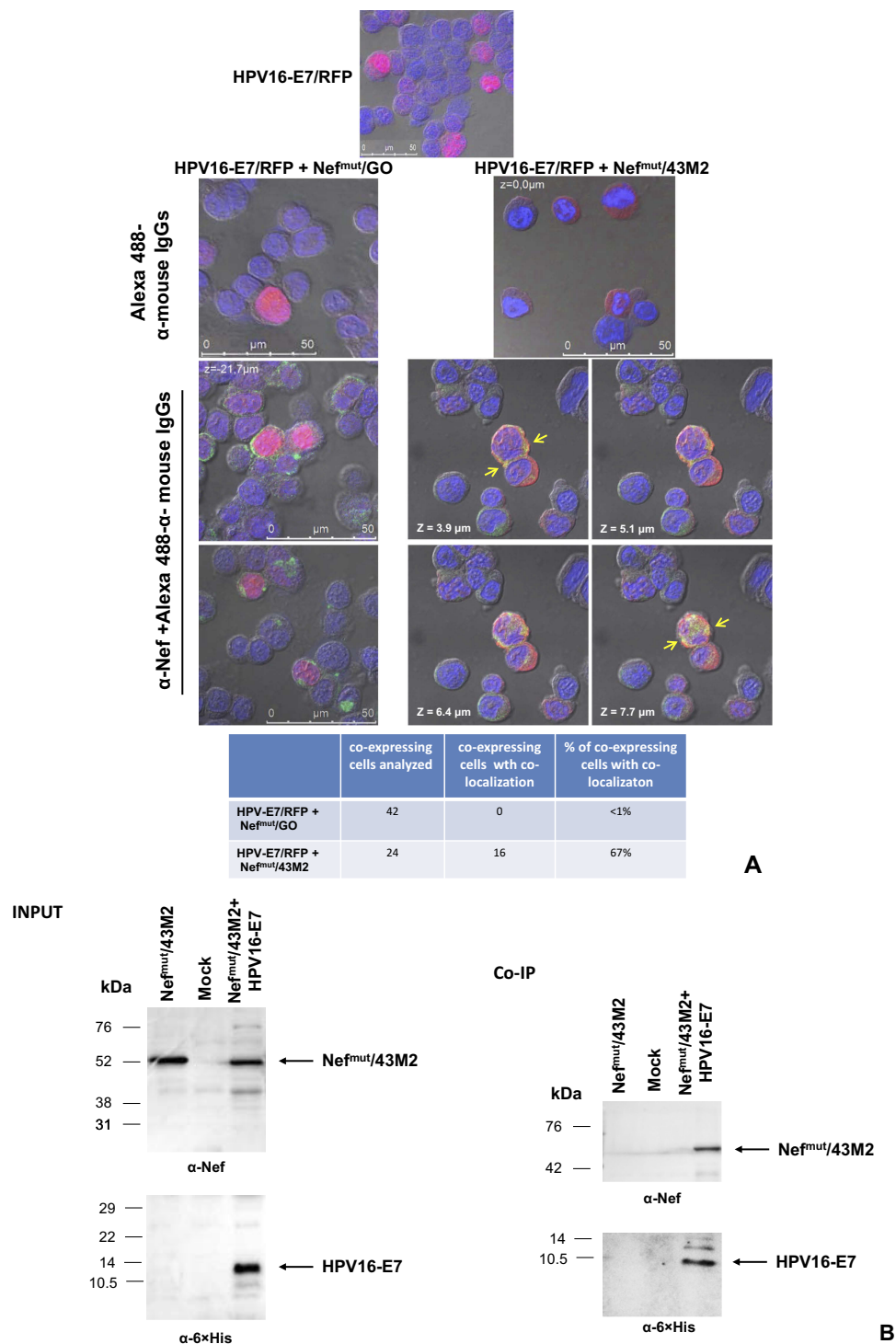
### The Expression Of Nef<sup>mut</sup>/43M2 Generates A Block Of Proliferation In HPV-Infected Cells

Next, we investigated whether the Nef<sup>mut</sup>/43M2 intracellular binding to HPV16-E7 resulted in a decrease of cell proliferation. To this end, SiHa cells, i.e., a human cell line expressing the whole HPV16 genome,<sup>22</sup> were transfected with the DNA vectors expressing either Nef<sup>mut</sup>/43M2 or Nef<sup>mut</sup>/GO, and cell viability scored 48 hrs after transfection by the trypan blue exclusion assay. As controls, SiHa cells were also transfected with vectors expressing either GFP, Nef<sup>mut</sup>, or 43M2 alone. Of note, accordingly with previously published data,<sup>19</sup> the intracellular expression of 43M2 alone was expected to be ineffective in the absence of tag sequences disposing the scFv either into nucleus or at secretory compartments.

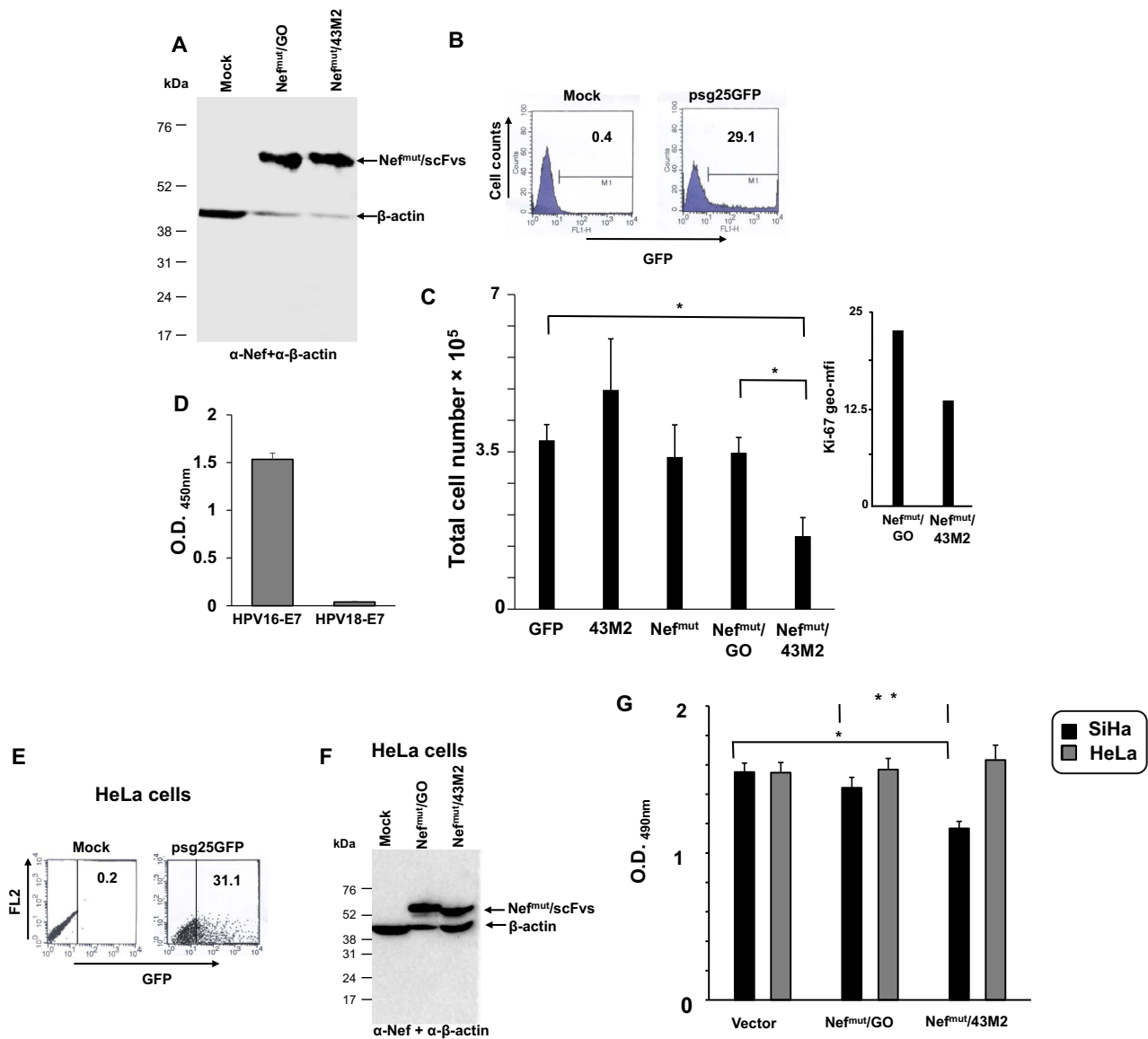
Western blot analysis served to verify that both Nef<sup>mut</sup>-fused scFvs were expressed in SiHa transfected cells (Figure 3A). In the presence of a transfection efficiency ranging from 25% to 35%, as assessed by FACS analysis of cells transfected with a GFP-expressing vector (Figure 3B), we noticed a statistically significant decrease in cell proliferation of Nef<sup>mut</sup>/43M2-expressing cells compared to control GFP-transfected cells, i.e, a mean of  $1.5 \times 10^5$  versus  $4 \times 10^5$  of total cells, respectively (Figure 3C). The inhibition of cell proliferation associated with a reduction of the expression of the cell proliferation marker Ki-67 (Figure 3C inset). As expected, the expression of neither Nef<sup>mut</sup> nor 43M2 alone inhibited the cell proliferation, and the same was for cells expressing



**Figure 1** Design and expression of vectors coding for Nef<sup>mut</sup>-fused scFvs. **(A)** Linear map of the DNA vectors expressing either Nef<sup>mut</sup>/43M2 or Nef<sup>mut</sup>/GO. Shown are the relevant restriction sites of the pTarget-Nef<sup>mut</sup>- based expression vector, as well as the flag tag position. ie-CMV, immediate-early CMV promoter; SD, major splice donor site; SA, major splice acceptor site; polyA, polyadenylation site. **(B, C)** Detection of Nef<sup>mut</sup>-based fusion products in both transfected cells and exosomes. Western blot analysis of both 30 µg of cell lysates from HEK293T cells transfected with DNA vectors expressing either 43M2 **(B)** or GO **(C)** scFvs fused with Nef<sup>mut</sup>, and 2 mU of exosomes purified from the respective supernatants. As control, samples from mock-transfected cells as well as cells transfected with the vector expressing Nef<sup>mut</sup> alone were included. Polyclonal anti-Nef Abs served to detect Nef<sup>mut</sup>-based products, while β-actin and Alix were revealed as cell and exosome markers, respectively. The relevant protein products are highlighted. Molecular weight markers are given in kilodaltons (kDa). The results are representative of three independent experiments. **(D)** Lack of 43M2 scFv incorporation in exosomes. Western blot analysis of 30 µg of cell lysates from HEK293T cells transfected with DNA vectors expressing 43M2 scFv either alone or fused with Nef<sup>mut</sup>, and of 2 mU of exosomes purified from the respective supernatants. As control, samples from mock-transfected cells were included. Anti-flag tag M2 mAb served to detect 43M2-based products, while β-actin and Alix were revealed as cell and exosome markers, respectively. Molecular weight markers are given in kilodaltons (kDa). The results are representative of two independent experiments.



**Figure 2** Intracellular interaction between HPV16-E7 and Nef<sup>mut</sup>/43M2. **(A)** HPV16-E7 disposition in Nef<sup>mut</sup>/43M2 expressing cells. Confocal microscope analysis of HEK293T cells co-transfected with HPV16-E7/RFP and either Nef<sup>mut</sup>/43M2 or Nef<sup>mut</sup>/GO expressing vectors. Top panels show representative fields from cells either transfected with E7-RFP alone or co-transfected as indicated and labeled with Alexa 488-conjugated anti-mouse IgGs. Below, shown are representative fields from co-transfected cell cultures labeled first with an anti-Nef mAb, and then with Alexa 488-conjugated anti-mouse IgGs. In the case of cells co-transfected with HPV16-E7/RFP and Nef<sup>mut</sup>/43M2 expressing vectors, four Z-stack sections are shown. Here, major green to red overlapping areas are indicated by arrows. DAPI (blue fluorescence) was used to highlight cell nuclei. The images were produced by differential interference contrast. Scale bars are reported. The results are representative of two independent experiments. At the bottom, both numbers and percentages of cells showing fluorescence co-localization are reported. **(B)** Co-immunoprecipitation assay. 293T cells were co-transfected either with vectors expressing Nef<sup>mut</sup>/43M2 and 6×His-tagged HPV16-E7, or with Nef<sup>mut</sup>/43M2 alone. As control, mock-transfected cells were also included. Forty-eight hours later, cell extracts were incubated with an anti-6×His mAb, and then incubated with anti-mouse IgG-coupled Dynabeads. Immunocomplexes were finally revealed by anti-Nef and, as control, anti-6×His Abs. On the top, both anti-Nef and anti-6×His tag Western blot analysis on 30 μg of each cell lysate are shown. At the bottom, shown are both anti-Nef and anti-6×His tag Western blot analysis on immunocomplexes isolated from 500 μg of each cell lysate. Molecular weight markers are given in kilodaltons (kDa). The results are representative of three independent experiments.



**Figure 3** Analysis of the anti-proliferative effect of the expression of Nef<sup>mut</sup>/43M2 vector in HPV16-infected cells. **(A)** Western blot analysis of the expression of Nef<sup>mut</sup>-derived products in transfected SiHa cells. Filter was revealed by both anti-Nef and HRP-conjugated anti-β-actin Abs. Molecular mass markers in kDa are reported on the left. Results are representative of three independent experiments. **(B)** FACS analysis of SiHa cells after transfection with a GFP-expressing vector. As control, mock-transfected cells also were analyzed. The analysis was carried out on 7-AAD negative cells. Percentages of positive cells are indicated. Shown are results representative of six independent experiments. **(C)** Cell viability of SiHa cells 48 hrs after transfection with the indicated vectors. A total of 2 × 10<sup>5</sup> cells was transfected with 1 μg of each DNA vector, and cell viability was scored 48 hrs later by the trypan blue exclusion method. Whatever the DNA vector used, an average of 10–20% of the dead cells were detected 24 hrs after transfection. Shown are mean values + standard errors of data from four independent experiments carried out in duplicates. \*p<0.05 by two-tailed Mann-Whitney U-Test. Inset, geometric mean-fluorescence intensity as calculated by FACS analysis conducted on SiHa cells transfected with the indicated vectors and labeled with anti-Ki67 mAb. Shown are the mean values calculated from an experiment performed with duplicate conditions. **(D-G)** Effects induced by the expression of the Nef<sup>mut</sup>/43M2 vector on the proliferation of cells expressing HPV18-E7. **(D)** ELISA for the binding of 43M2 to recombinant E7 from either HPV16 or HPV18 strains. The results are representative of three independent experiments carried out in quadruplicate conditions. **(E)** FACS analysis of HeLa cells 48 hrs after transfection with a GFP-expressing vector. Percentages of DNA vector-expressing cells are reported. The analysis was carried out on 7-AAD negative cells. Results are representative of two independent experiments. **(F)** Western blot analysis of the expression of Nef<sup>mut</sup>-derived products in transfected HeLa cells. Filter was revealed by both anti-Nef and HRP-conjugated anti-β-actin Abs. Molecular mass markers in kDa are reported on the left. Results are representative of three independent experiments. **(G)** Cell viability of both SiHa and HeLa cells 48 hrs after transfection as assessed by MTS-based cell proliferation assay. Transfection procedures were as described for panel C, except that, to perform the MTS assay, the cells from each well of the 24-well plates were split in four microwells of a 96-well plate 48 hrs after transfection. Shown are the mean values of absorbance at 490<sub>nm</sub> (OD) + standard errors of the results obtained from four independent experiments performed in quadruplicate. \*p: 0.0002, \*\*p: 0.0054 by two-tailed Student T Test.

Nef<sup>mut</sup>/GO. The anti-proliferative effect was observed also in TC-1 cells, i.e., murine tumor cells constitutively expressing HPV16-E7<sup>23</sup> (not shown).

To further check the specificity of the Nef<sup>mut</sup>/43M2-induced effects, we reproduced these experiments using HeLa cells. This cell line expresses the HPV18-E7



isoform, which is not recognized by the 43M2 scFv as sensed by the ELISA assay whose results are reported in [Figure 3D](#). Cell proliferation was evaluated by the MTS-based assay. In detail, both SiHa and HeLa cells were transfected with either Nef<sup>mut</sup>/43M2 or Nef<sup>mut</sup>/scFvGO expressing vectors. As detected after 48 hrs, the transfection efficiency on HeLa cells ([Figure 3E](#)) appeared comparable to what we observed with SiHa cells. Western blot analysis demonstrated that both Nef<sup>mut</sup>-fused scFvs were stable in HeLa cells too ([Figure 3F](#)). Nonetheless, no reduction of cell proliferation was detectable in Nef<sup>mut</sup>/43M2-expressing HeLa cells compared to cells transfected with the control vectors, in the presence, however, of the expected inhibition in SiHa cell cultures ([Figure 3G](#)).

Taken together, these data indicated that Nef<sup>mut</sup>/43M2 binds intracellular HPV16-E7, thereby inducing a specific anti-proliferative effect.

## Proliferation Of HPV16-Infected Cells Is Impaired By Challenge With Nef<sup>mut</sup>/43M2-Loaded Exosomes

We next analyzed the activity of exosome-associated Nef<sup>mut</sup>/43M2 in challenged HPV-positive cells. In a first instance, we sought to establish whether the cargo of Nef<sup>mut</sup>-engineered exosomes could be delivered inside HPV16-expressing cells at detectable extents. To this aim, HEK293T cells were transiently transfected with a Nef<sup>mut</sup>/GFP expressing vector<sup>4</sup> ([Figure 4A](#)), and culture supernatants collected from 48 to 72 hrs thereafter. Fluorescent exosomes were then isolated and analyzed by flow cytometry for the presence of CD63, i.e., a tetraspanin typically expressed on the exosome cell membrane<sup>24,25</sup> ([Figure 4B](#)), and by Western blot for the detection of the exosome marker Alix<sup>5</sup> ([Figure 4C](#)). Then, SiHa cells were challenged with different amounts of the GFP-labeled exosomes and, at different times, FACS analyzed for the intracellular presence of exosome-associated fluorescence. We detected cell-associated fluorescence until 5 hrs after the exosome challenge performed at 37°C but not at 4°C. Furthermore, the percentage of fluorescent cells increased with the doses of challenging exosomes ([Figure 4D](#)). This result supported the idea that the cargo of engineered exosomes can be delivered into SiHa cells preserving an apparently unaltered conformation, considering that the GFP fluorescence relies on its intact secondary and tertiary structures.<sup>26</sup>

Next, we set up experiments to determine the effects of Nef<sup>mut</sup>/43M2-engineered exosomes on SiHa cell proliferation.

For obvious quantitative reasons, we assumed that the exosome treatment would result in an antiproliferative effect weaker than that observed in DNA-transfected cells. To optimize the readout sensitivity, we monitored cell proliferation through the sensitive, automated, and label-free xCELLigence Real-Time Cell Analysis (RTCA) system.<sup>27</sup> In the absence of variation in the extra-cellular matrix composition and overall cell morphology, this assay provides a powerful means to monitor cell viability and proliferation in real-time and in a non-invasive way through the measurement of cell culture impedance.

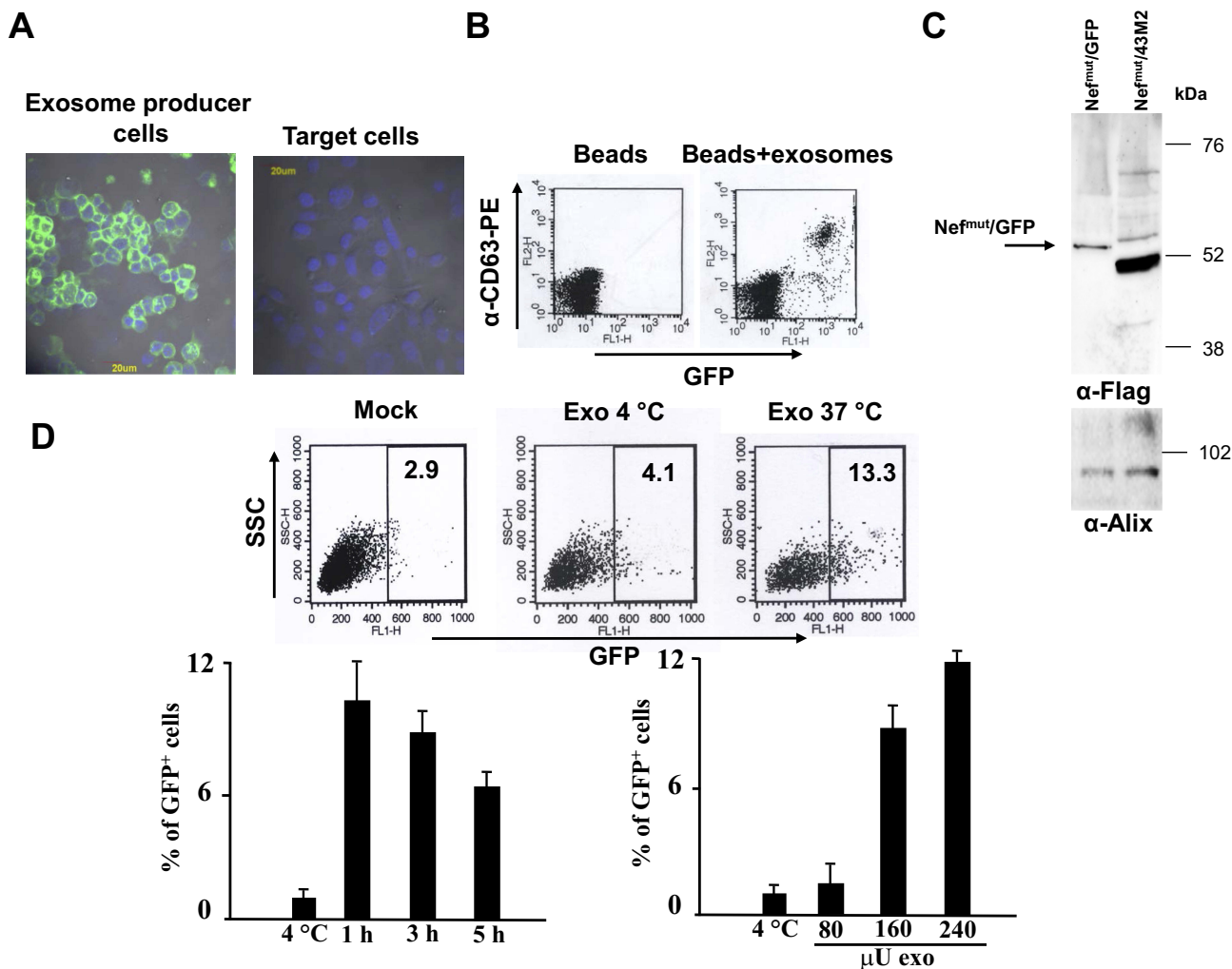
To establish whether the use of the RTCA assay was a suitable tool to evaluate the anti-proliferative effect of Nef<sup>mut</sup>/43M2, both SiHa and HeLa cells were transfected with either Nef<sup>mut</sup>/GO- or Nef<sup>mut</sup>/43M2-expressing vectors. After 24 hrs, the cells were seeded in RTCA micro-wells, and then cell proliferation was followed in real time for 80 hrs. Except for a first time lag required for cell adhesion, we noticed both consistent and reproducible lower values of the cell index (CI, directly proportional to cell proliferation) in SiHa cultures expressing Nef<sup>mut</sup>/43M2 compared to cells expressing Nef<sup>mut</sup>/GO. Conversely, such a decrease was not detectable in transfected HeLa cells ([Figure 5A](#)). We concluded that the RTCA represented a suitable way to evaluate the Nef<sup>mut</sup>/43M2-induced anti-proliferative effect.

Next, we tried to assess the effects of the treatment with Nef<sup>mut</sup>/43M2-engineered exosomes on cell proliferation. To this end, 24 hrs after seeding in the RTCA microwells, both SiHa and HeLa cells were challenged with equal amounts of exosomes incorporating either Nef<sup>mut</sup>/43M2, Nef<sup>mut</sup>/GO, or, in the case of SiHa cells only, with void exosomes. Starting from 48 hrs after exosome challenge, the real-time monitoring of the cell cultures revealed a specific reduction in the proliferation of SiHa cells challenged with Nef<sup>mut</sup>/43M2-engineered exosomes ([Figure 5B](#)).

These results represented a first indication that scFvs delivered in target cells by means of engineered exosomes can preserve their functional activity.

## Decreased Proliferation Of HPV16 Infected Cells Co-Cultured In Transwell Chambers With Cells Producing Nef<sup>mut</sup>/43M2 Engineered EVs

To strengthen the significance of the data we obtained through exosome challenge, we designed experiments where the anti-proliferative effect was evaluated in cells

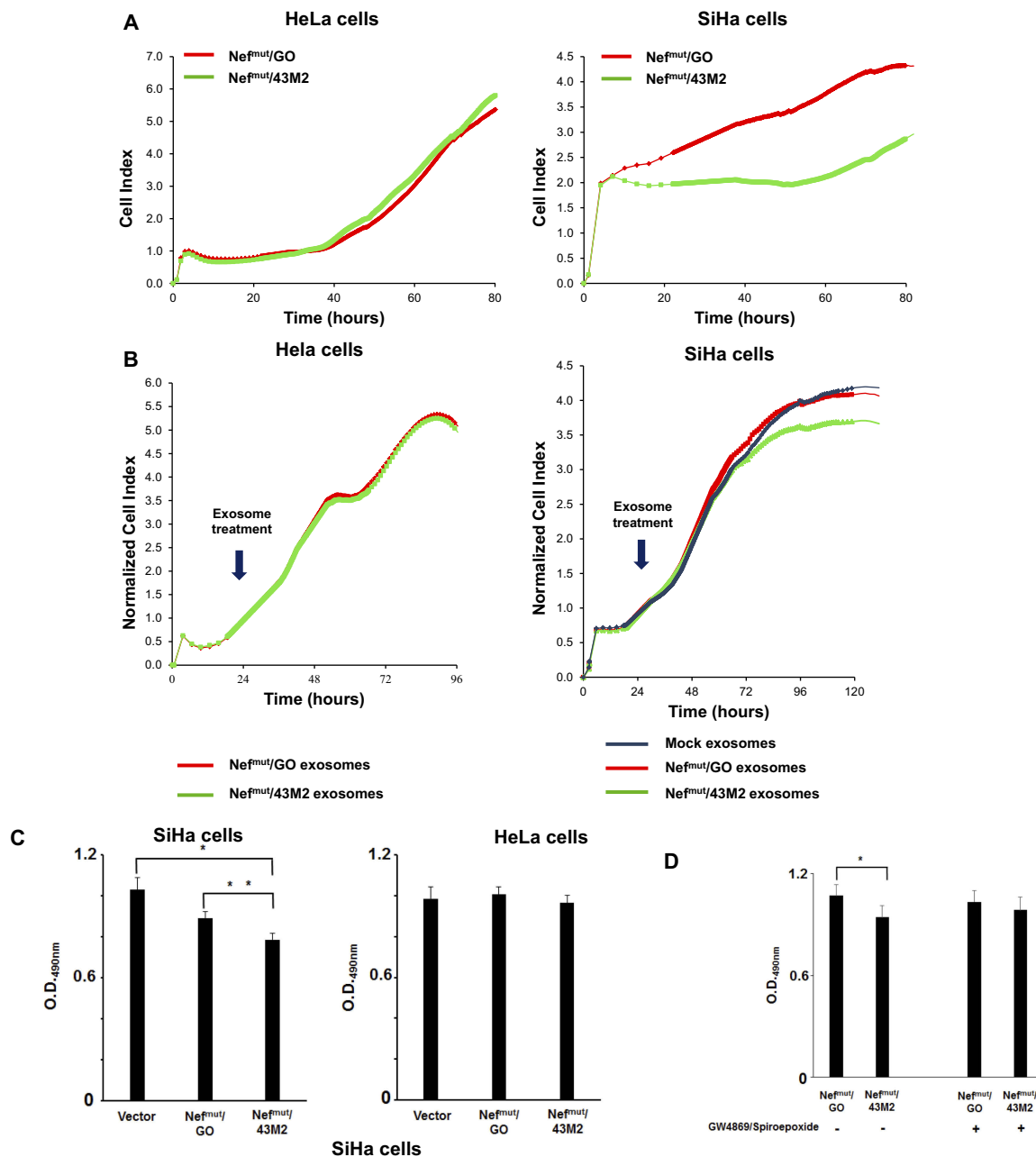


**Figure 4** Detection of fluorescent exosomes in challenged SiHa cells. **(A)** On the left: fluorescence microscope analysis of HEK293T cells transfected with Nef<sup>mut</sup>/GFP vector. On the right, the same analysis was carried out on mock-transfected SiHa cells. The cell nuclei were stained with DAPI (blue fluorescence). **(B)** FACS analysis of fluorescent exosomes produced by Nef<sup>mut</sup>/GFP expressing cells. Exosomes were bound to surfactant-free white aldehyde/sulfate latex beads and then labeled with PE-conjugated anti-CD63 mAbs. The analysis included singularly dispersed beads only, as identified through the FSC/SSC plots. Shown are the results representative of six independent experiments. **(C)** Western blot analysis of 2 μU of exosomes purified from the supernatants of Nef<sup>mut</sup>/GFP expressing cells. As control, exosomes from cells transfected with the vector expressing Nef<sup>mut</sup>/43M2 were included. Polyclonal anti-Nef Abs served to detect Nef<sup>mut</sup>-based products, while Alix was revealed as exosome marker. Molecular weight markers are given in kilodaltons (kDa). The results are representative of two independent experiments. **(D)** FACS analysis on SiHa cells challenged with GFP-fluorescent exosomes. A total of 2×10<sup>4</sup> SiHa cells was challenged with different amounts of fluorescent exosomes (i.e., from 80 to 240 μU). After spinoculation, cells were incubated either at either 4°C or 37°C. At different times, cells were detached with cold trypsin, washed, fixed, and FACS analyzed. On the top, shown are the raw data from cells challenged or not (Mock) with 160 μU of exosomes, and analyzed 3 hrs later. As control, the results from exosome-challenged cells incubated at 4°C are shown. On the bottom left, shown are the results from FACS analysis of cells challenged with 160 μU of exosomes and harvested at different times. As control, conditions where exosome-challenged cells were incubated at 4°C and harvested at the latest time point were included. On the bottom right, shown are the results from FACS analysis of cells challenged with the indicated amounts of fluorescent exosomes, and harvested 3 hrs after spinoculation. As control, data from the analysis of cells challenged with the highest amounts of exosomes and incubated at 4°C were included. The results are presented as mean values +SD of triplicates after subtraction of background values (i.e., the fluorescence levels of mock-challenged cells), and are representative of two independent experiments.

growing in the continuous presence of Nef<sup>mut</sup>/43M2-engineered EVs. To this end, we set up transwell co-cultures comprising donor cells, i.e., HEK293T cells transiently transfected with Nef<sup>mut</sup>-based DNA vectors which were seeded in the upper chamber, and either SiHa or HeLa target cells placed in the lower one. After four days, the proliferation of both SiHa and HeLa cells was assessed by the MTS-based colorimetric assay. A significant reduction

of cell proliferation was detected in SiHa cells co-cultured with the donor Nef<sup>mut</sup>/43M2-expressing HEK293T cells, whereas no differences were observed using HeLa as target cells (Figure 5C).

To assess the role of Nef<sup>mut</sup>/43M2 engineered exosomes in the inhibition of cell proliferation, we repeated the transwell experiments in the presence or absence of 1 μM of both GW4869 and spiroepoxide, i.e., two inhibitors of exosome



**Figure 5** Functional analysis of Nef<sup>mut</sup>/43M2 engineered exosomes. **(A)** Viability of SiHa cells after challenge with Nef<sup>mut</sup>/43M2-uploading exosomes. RTCA proliferation assay in both HPV18-positive HeLa and HPV16-positive SiHa cells after transfection with either Nef<sup>mut</sup>/GO or Nef<sup>mut</sup>/43M2 expressing vectors. Results shown are representative of five independent experiments. **(B)** RTCA proliferation assay in both HeLa and SiHa cells challenged with exosomes uploading either Nef<sup>mut</sup>/GO or Nef<sup>mut</sup>/43M2. As an additional control, SiHa cells were challenged with equivalent amounts of exosomes isolated from the supernatants of mock-transfected cells. Arrows indicate the time of exosome treatment, i.e., 24 hrs after seeding. Results are expressed as normalized cell index, i.e., the ratio between cell index at a given time and cell index at the time of exosome treatment, and are representative of two independent experiments. **(C)** Viability of SiHa cells co-cultured in transwell chambers with HEK293T cells producing Nef<sup>mut</sup>/43M2 engineered EVs. HEK293T cells were transfected with vectors expressing either Nef<sup>mut</sup>/43M2, Nef<sup>mut</sup>/GO, or empty vector. After 24 hrs, transfected cells were seeded in the upper transwell chamber while either SiHa or HeLa cells were plated in the lower one. Four days later, cell viability was assessed by MTS-based cell proliferation assay. At this time, cells from each well were detached, resuspended in 0.4 mL of complete medium, and four microwells of a 96-well plate were seeded with 100  $\mu$ L of each cell culture. Shown are mean OD values + standard errors from the results of six complete experiments performed in quadruplicates. \**p*: 0.0094; \*\**p*: 0.0274 by two-tailed Mann-Whitney *U*-Test. **(D)** Cell viability as assessed by MTS-based cell proliferation assay 4 days after setting the co-culture of SiHa cells with HEK293T cells previously transfected with the indicated vectors, and carried out in the presence or absence of 1  $\mu$ M GW4869 and spiroepoxide. SiHa cell cultures were processed for the MTS assay as described for panel A. Shown are mean OD values + standard errors from the results of four independent experiments conducted in quadruplicates. \**p*: 0.0396 by two-tailed Student *T* Test. Differences in cell viability between SiHa cells co-cultivated with donor cells transfected with the two different vectors in the presence of the inhibitors were not statistically significant (*p*: 0.5471 by two-tailed Student *T* Test).

biosynthesis.<sup>28–33</sup> In our hands, the actual inhibition of exosome release from HEK293T cells treated with 1  $\mu$ M

GW4869 and spiroepoxide ranged from 60% to 80%, as assessed through the acetylcholinesterase assay (not

shown). Interestingly, treatment with the inhibitors restored the cell proliferation levels of SiHa cells co-cultured with Nef<sup>mut</sup>/43M2 expressing donor cells (Figure 5D). These results strongly suggested that the proliferation of target cells was indeed affected by the release of Nef<sup>mut</sup>/43M2-engineered exosomes from donor cells.

Together with what we observed by challenging SiHa cells with cell-free exosomes, these results support the idea that Nef<sup>mut</sup>-based engineered EVs can represent an efficient tool to deliver scFvs inside target cells.

## Discussion

ScFvs, in their different configurations, can represent effective surrogates of tetrameric Abs, since they were shown to reproduce both agonist and antagonist activities. As with classic Abs, however, plasma membrane hinders their access to the intracellular milieu. Developing strategies aimed at favoring scFv intracellular delivery have been a matter of major interest in the last decade. Previously described methods are based on coupling scFvs with a number of polymers/nanoparticles, e.g., cationic liposomes,<sup>34,35</sup> or cell-penetrating peptides.<sup>36,37</sup> Although in both cases intracellular antigens have been successfully targeted, major limitations in possible therapeutic applications still remained. The aim of our investigations was achieving the proof-of-concept for a novel strategy of intracellular scFv delivery potentially suitable for clinical development. The idea was based on our recent finding concerning the possibility to generate engineered EVs through a DNA vector expressing the Nef<sup>mut</sup> exosome-anchoring protein fused with polypeptides of interest.<sup>11</sup> When the system is translated *in vivo*, engineered EVs protect their cargo from degradation factors present in both extracellular milieu and serum, and can safely deliver it upon entry into target cells. In the case scFvs are uploaded into EVs, intracellular therapeutic targets can be inactivated.

To evaluate the effectiveness of our strategy in an *in vitro* system, we considered the anti-HPV16-E7 43M2 scFv.<sup>13,14,19</sup> In the form of intrabody, this scFv has been previously characterized for its ability to counteract the proliferative effect of HPV16-E7 both *in vitro* and *in vivo*.<sup>13,20</sup> Here, we show that both 43M2 and GO scFvs are readily uploaded into EVs when fused to the Nef<sup>mut</sup> C-terminus. In this form, 43M2 strongly reduces the nuclear accumulation of HPV16-E7 as assessed by confocal microscope analysis. In addition, the well-detectable areas of Nef<sup>mut</sup>/43M2-HPV16-E7 co-localization were consistent with the idea that the HPV16-E7 translocation was consequence of the interaction with 43M2. This is a

relevant issue since it was previously shown that 43M2 counteracts the HPV16-E7 activity exclusively when provided with signals for intracellular localization at either nucleus or secretory compartment.<sup>19</sup> Hence, fusion with Nef<sup>mut</sup> overcomes the need to modify 43M2 to optimize its effect against the target protein. It is conceivable that the strong affinity of Nef<sup>mut</sup> for intracellular membranes favors both the stability of Nef<sup>mut</sup>/43M2 and its interaction with neo-synthesized HPV16-E7.

Nef<sup>mut</sup>/43M2 expressed in transfected cells was shown to specifically inhibit cell proliferation of HPV16-E7 expressing cells. This conclusion was drawn through trypan blue exclusion test, Ki-67 detection, MTS, and RTCA assays. Hence, 43M2 preserves its anti-proliferative activity even when fused with Nef<sup>mut</sup>. The overall potency of this effect appeared comparable to what previously described when 43M2 fused to a signal for retention in the endoplasmic reticulum was expressed intracellularly.<sup>13</sup>

As for the mechanism underlying the anti-proliferative effect in transfected cell cultures, it should be considered that at least part of the Nef<sup>mut</sup>/43M2 molecules could associate with exosomes released by DNA-vector expressing cells, thus contributing to HPV16-E7 neutralization in bystander cells. This mechanism may account for the extents of cell proliferation inhibition sometime exceeding the percentages of DNA transfected cells as evaluated by FACS analysis. This hypothesis is consistent with the results we obtained from the experiments performed by challenging SiHa cells with engineered exosomes, as well as those from co-cultures of SiHa cells with HEK293T cells producing Nef<sup>mut</sup>/43M2-uploaded EVs. In the latter instance, the fact that treatment of transwell co-cultures with inhibitors of exosome biosynthesis strongly reduced the anti-proliferative effect highlighted the role of engineered exosomes in the inhibitory effect.

The impairment of cell proliferation consequent of the treatment with cell-free engineered exosomes appeared somewhat reduced and delayed compared to what we detected in Nef<sup>mut</sup>/43M2 transfected cells. Likely, quantitative differences in the actual intracellular Nef<sup>mut</sup>/43M2 availability may account for such apparent discrepancy. In fact, it is conceivable that, whereas the expression of transfected DNA leads to a both continuous and massive production of Nef<sup>mut</sup>/43M2, conversely providing cells with a single pulse of engineered exosomes likely results in a less abundant and, most important, temporarily limited availability of the anti- HPV16-E7 scFv. In addition, efficiency of both nanovesicle cell entry and escape from

intracellular degradative pathways may influence the overall scFv availability in EV-challenged cells.

## Conclusion

Engineering EVs with scFvs has the potential to be exploited as a novel therapeutic approach to target intracellular pathogenic factors. In vivo, this strategy could be applied by engineering endogenous EVs through the intramuscular injection of Nef<sup>mut</sup>-related DNA vectors, as we already described.<sup>4,10</sup> The delivery of DNA expressing Nef<sup>mut</sup>-fused scFvs, possibly also by viral vectors, would result in a continuous source of engineered EVs. These nanovesicles can safely deliver therapeutic scFvs into bystander cells, where the binding to specific intracellular pathogenic factors can generate the desired effect.

Previous literature data report protein engineering of EVs essentially through manipulation of membrane-associated proteins. For instance, tetraspanins (e.g., CD9, CD63, CD81),<sup>38,39</sup> spontaneously associating with exosome membranes at high levels, as well as proteins of the endosome/lysosome apparatus, e.g., Rab5a<sup>40</sup> Lamp2b<sup>41</sup> were used as scaffolds to engineer exosomes with fluorescent proteins. Similarly, the transmembrane domain of platelet-derived growth factor receptor<sup>42</sup> and the C1C2 domains of lactadherin<sup>43,44</sup> represented the platform of chimeric proteins expressed in exosome membranes. In addition, specific cell targeting of EVs was achieved by building chimeric proteins recognizing specific cell membrane receptors, for instance through the fusion with receptor-specific scFvs. As an example, exosomes displaying anti-HER2 scFv to target cancer mammary cells were recently described by two independent groups.<sup>45,46</sup> Similarly, exosomes engineered with anti-CD19 scFv fused with CD63 were shown to efficiently target lymphoma cells.<sup>47</sup>

With these premises, vectors co-expressing Nef<sup>mut</sup> fused with therapeutic scFvs and membrane molecules engineered with receptor-specific scFvs have the potential to generate EVs efficiently expanding the scFv-induced therapeutic effect to appropriate bystander cells.

## Ethics

The use of all cell lines had Istituto Superiore di Sanità (ISS) review board approval.

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## Disclosure

The authors report no conflicts of interest in this work.

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