



Extracellular Vesicle-Serpine-1 Affects Neural Progenitor Cell Mitochondrial Networks and Synaptic Density: Modulation by Amyloid Beta and HIV-1

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Abstract

Brain endothelial extracellular vesicles carrying amyloid beta (EV-A β) can be transferred to neural progenitor cells (NPCs) leading to NPC dysfunction. However, the events involved in this EV-mediated A β pathology are unclear. EV-proteomics studies identified Serpine-1 (plasminogen activator inhibitor 1, PAI-1) as a major connecting “hub” on several protein–protein interaction maps. Serpine-1 was described as a key player in A β pathology and was linked to HIV-1 infection as well. Therefore, the aim of this work was to address the hypothesis that Serpine-1 can be transferred via EVs from brain endothelial cells (HBMEC) to NPCs and contribute to NPC dysfunction. HBMEC concentrated and released Serpine-1 via EVs, the effect that was potentiated by HIV-1 and A β . EVs loaded with Serpine-1 were readily taken up by NPCs, and HIV-1 enhanced this event. Interestingly, a highly specific Serpine-1 inhibitor PAI039 increased EV-A β transfer to NPCs in the presence of HIV-1. PAI039 also partially blocked mitochondrial network morphology alterations in the recipient NPCs, which developed mainly after HIV + A β -EV transfer. PAI039 partly attenuated HIV-EV-mediated decreased synaptic protein levels in NPCs, while increased synaptic protein levels in NPC projections. These findings contribute to a better understanding of the complex mechanisms underlying EV-Serpine-1 related A β pathology in the context of HIV infection. They are relevant to HIV-1 associated neurocognitive disorders (HAND) in an effort to elucidate the mechanisms of neuropathology in HIV infection.

Keywords Serpine-1 · Extracellular vesicles · Blood–brain barrier · HIV-1 · Amyloid beta · Neural progenitor cells

Abbreviations

A β Amyloid beta
AD Alzheimer's disease
BBB Blood-brain barrier

bFGF Basic fibroblast growth factor
DAPI 4',6-Diamidino-2-phenylindole
ECAR Extracellular acidification rate
EVs Extracellular vesicles
EGF Epidermal growth factor
ELISA Enzyme linked immunosorbent assay
HAND HIV-associated neurocognitive disorders
GFP Green fluorescent protein
HBMEC Human brain microvascular endothelial cells
HIV-1 Human immunodeficiency virus type 1
LRP1 Lipoprotein receptor related protein 1
MiNA Mitochondrial network analysis
NPCs Neural progenitor cells
NTA Nanoparticle tracking analysis
OCR Oxygen consumption rate
PAI-1 Plasminogen activator inhibitor 1
PBS Phosphate buffered saline
RAGE Receptor for advanced glycation end products
RFP Red fluorescent protein
Serpine-1 Plasminogen activator Inhibitor Type 1

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PSD95 Postsynaptic density protein 95
tPA Tissue plasminogen activator

Introduction

Extensive evidence indicates amyloid pathology in HIV-1 infection. Studies reported that HIV-infected brains have increased amyloid beta (A β) deposition mostly in the perivascular space [1–4] when compared to age-matched controls [1, 5–9]. These earlier findings have recently been reevaluated and expanded, as new evidence from autopsy of HIV-1 brains did not confirm an overall enhanced A β deposition. Instead, duration of infection was correlated with A β accumulation independent of age. These findings suggest that the length of HIV infection, and not the age of patients, predicts elevated brain A β levels, confirming an accelerated brain senescence in people living with HIV [10]. For example, methylome-wide analysis of chronic HIV-1 infection revealed five-year increase in biological age of infected individuals [11]. HIV-related age acceleration has been shown to be associated with reductions in total gray matter using epigenetic age as a biomarker for age acceleration [12]. Evidence indicates that increased and persistent senescence during HIV-1 infection contributes to chronic inflammation, immune failure and mitochondrial dysfunction [13–15].

It was demonstrated that the blood–brain barrier (BBB) is critical for A β homeostasis and contributes to elevated A β deposition in the brain [16, 17]. A β transport into the brain across the BBB involves BBB transfer mechanisms such as the receptor for advanced glycation end products (RAGE) [18]. Moreover, we found that HIV-1 contributes to A β accumulation in brain endothelial cells via upregulation of RAGE [19]. Extracellular vesicles (EVs) also appear to be important players in A β pathology by carrying biologically active proteins and genetic material. We demonstrated that exposure to HIV-1 increases EV release from human brain microvascular endothelial cells (HBMEC) and alter their A β cargo [20]. EVs having diverse origin, size, and cargo can be taken up by the target cells via several routes [21]. Importantly, HBMEC-derived EVs (HBMEC-EVs) carrying A β can be transferred to other cells of the neurovascular unit, including neural progenitor cells (NPCs), causing NPC dysfunction and their aberrant neurogenesis [22]. The mechanisms of this EV-mediated A β pathology are not clear; however, proteomics of HBMEC-EVs revealed a complex protein cargo with elaborate functional interactions as mapped in STRING [23]. These maps had several “hubs” with more connections to other proteins. One of the main proteins at such crossroad interactions was Serpine-1, known also as plasminogen activator inhibitor 1, PAI-1.

Serpine-1 is crucial for the coagulation/fibrinolysis homeostasis. By inhibiting tissue plasminogen activator (tPA) and

urokinase type activator (uPA), Serpine-1 promotes coagulation [24]. These processes are important in both HIV-1 infection and Alzheimer’s amyloidopathy. For instance, endothelial dysfunction in HIV-1 infection is associated with leucocyte recruitment, platelet adhesion and aggregation, blood clotting activation and fibrinolysis derangement that may be responsible for occlusive thrombotic events [25]. In addition, HIV-1 replication alters the composition of extrinsic pathway coagulation factors and increases thrombin generation, leading to a procoagulant state [26]. HIV-1 infection constitutes a higher risk of cardiovascular events, including stroke and blood–brain barrier (BBB) disruption, due to coagulation abnormalities and dyslipidemia [27]. Regarding A β pathology, activation of plasmin via the coagulation cascade results not only in fibrinolysis but also in degradation of APP and A β . Consequently, Serpine-1, by decreasing tPA-plasminogen-plasmin-mediated APP/A β degradation, can contribute to an increase in A β levels [28].

Because both HIV-1 infection [29] and A β pathology [30, 31] were linked to elevated levels of Serpine-1, the aim of the present study was to address the hypothesis that Serpine-1 can be transferred via EVs from brain endothelial cells to NPCs affecting their functions. Mechanistically, we focused on the role of EV-associated Serpine-1 (EV-Serpine-1) in the NPC mitochondrial changes in the context of HIV-1. We also evaluated the involvement of Serpine-1 in NPC synaptic protein alterations. The obtained results indicate that EV-Serpine-1 may be a crucial player in vascular A β pathology in HIV-infection.

Materials and Methods

Cell Cultures

Human Brain Microvascular Endothelial Cells (HBMEC)

HBMEC used in the present study represent a stable, well characterized, and differentiated human brain endothelial cell line [32]. Briefly, normal human brain endothelial cells were transduced by lentiviral vectors incorporating human telomerase or SV40T antigen. Among several stable immortalized clones obtained by sequential limiting dilution cloning of the transduced cells, the hCMEC/D3 cell line (referred here as HBMEC) was selected as expressing normal endothelial markers and demonstrating blood–brain barrier characteristics. HBMEC for the present study were supplied by Dr. Couraud (Institut Cochin, Paris, France). HBMEC were cultured on collagen type I (BD Biosciences Pharmingen, San Jose, CA) coated dishes in EBM-2 medium (Clonetics, East Rutherford, NJ) supplemented with VEGF, IGF-1, EGF, basic FGF, hydrocortisone, ascorbate,

gentamycin and 0.5% exosome depleted fetal bovine serum (Exo-FBS; System Biosciences, Mountain View, CA).

Human Neural Progenitor Cells (NPCs)

An immortalized NPC line ReNcell VM, derived from 10-week human ventral mesencephalon, was obtained from Millipore and cultured according to the manufacturer's protocols. The cells were validated for high expression of Sox2 and nestin as well as for their self-renewal and differentiation capacity. Cells were grown on laminin coated tissue culture dishes in a maintenance medium (Millipore), containing 20 ng/ml FGF-2 and 20 ng/ml of rhEGF. Cells were used for experiments at <80% confluence, three days after plating.

HIV-1 Infection and Treatment Factors

HIV-1 stock was generated using human embryonic kidney (HEK) 293 T cells (ATCC) transfected with pYK-JRCSF plasmid containing full-length proviral DNA. Throughout the study, HBMEC were exposed to HIV-1 particles at the p24 level of 30 ng/ml as previously reported [22]. Treatment was terminated by removing cell culture media containing HIV-1, followed by washing the cells with PBS.

A β (1–40) and A β (1–40) HiLyte 647 were purchased from Anaspec (San Jose, CA) and dissolved in PBS. Freshly solubilized A β solutions without pre-aggregation were used for experiments as such a form of A β was demonstrated to induce proinflammatory reactions [33]. A β (1–40) HiLyte was dissolved first in a basic buffer (0.1 M NH₄OH) and then diluted further in PBS as suggested by the manufacturer. Cells were treated with A β (1–40) or A β (1–40) HiLyte at the concentration of 100 nM in complete medium.

PAI039 (Tiplaxtinin, Catalog # PZ0295) was purchased from Millipore Sigma, Burlington, MA, USA). PAI039 is a potent and selective Serpine-1 inhibitor [34] and demonstrated efficacy *in vivo* in multiple models of acute arterial thrombosis. A 20 mM stock solution was prepared in DMSO. In a typical experiment, NPCs were cotreated with isolated EVs and/or 2 μ M PAI039 for 24 h. Literature indicates that 1 μ M PAI-039 can effectively inhibit Serpine-1 activity *in vitro* [35]. PAI039 exerts its activity by binding close to the vitronectin binding site [36].

Treatment of Brain Endothelial Cells and EV Isolation

Confluent HBMEC were exposed to HIV-1, and/or A β (1–40)/A β (1–40) HiLyte for 48 h. EVs were isolated from the media using ExoQuick-TC exosome precipitation solution (System Biosciences, Mountain View, CA) according to the manufacturer's specifications. Briefly, 10 ml culture medium from 1.7×10^7 cells at confluency cultured in a P100

dish was centrifuged at 3000 g for 15 min to remove cells and debris. Then, the samples were mixed thoroughly with 2 ml of Exo-Quick exosome precipitation solution and incubated overnight at 4 °C. The next day, the samples were centrifuged at 1500 g for 30 min; the supernatants were removed and centrifuged again at 1500 g for 5 min. The EV pellets were resuspended in 400 μ l PBS and used for further studies. The aliquots of 20 μ l of EV suspension for every 100 μ l of cell culture media was used for NPC treatment.

Transfection of Brain Endothelial Cells

HBMEC were transfected with the CD63 RFP and Serpine-1 GFP constructs (Vectorbuilder, Chicago, IL, USA) using Purefection Transfection Reagent (System Biosciences) following the manufacturer's protocol. We measured transfection efficiency from confocal images in cells 72 h post transfection and found that 90–95% of the cells were transfected with both constructs and expressed Serpine-1 GFP and CD63 RFP. Twenty-four hours post transfection, cells were exposed to HIV-1 or/and A β (1–40) HiLyte for 48 h followed by EV isolation from the media. We have observed that even minor changes of HIV-1 stock preparation can lead to differences in EV release from HBMEC exposed to HIV-1. If HEK 293 T cells were cultured in Exo-FBS, their morphology changed, and the resulting viral stock had vastly different effects on EV release and EV isolation. Therefore, HEK 293 T cells were cultured in media containing regular FBS consistent with our previous reports [20, 22]. As a control for HIV-1 exposure, isolates from mock-transfected HEK 293 T cells were used.

Nanoparticle Tracking Analysis (NTA)

EVs were analyzed by NanoSight model NS300 (Malvern Instruments Company, NanoSight, Malvern, United Kingdom) as described earlier [20]. Briefly, EV pellet samples obtained in the process of EV isolation were resuspended in 4% paraformaldehyde-PBS and further diluted 50-fold in PBS for analysis. During analysis, five 15 s videos were recorded for each sample. The obtained data were analyzed using Nanosight NTA Analytical Software (Malvern Instruments Company) with the detection threshold optimized for each sample and screen gain at 10 to track the maximal number of particles with minimal background.

EV-A β HiLyte Transfer Assay

EV-derived A β HiLyte fluorescence was quantified in the recipient NPCs using a plate reader. Briefly, NPCs were plated in 96-well black plates (9,000 cells/well) and differentiated for 3 days with the last 24 h in the presence of the isolated EVs. To block Serpine-1 activity, selected

NPC cultures were cotreated with 2 μ M PAI039 and brain endothelial EVs for 24 h. After treatment, cells were washed with PBS, fixed with ethanol for 30 min at 4 °C, washed again with PBS and A β HiLyte fluorescence was measured with a plate reader (Molecular Devices, SpectraMax iD3) as suggested by the manufacturer (Anaspec, Ex/Em 503/528 nm). Cell nuclei were stained with DRAQ5 (Cell Signaling, Catalog #4084L, dilution 1:1000) for 5 min, washed with PBS and DRAQ5 fluorescence was measured with the same plate reader (Ex/Em 647/681). A β HiLyte fluorescence was then normalized to nuclear DRAQ5 fluorescence.

Enzyme Linked Immunosorbent Assay (ELISA) and Serpine-1 Activity Assay

ELISA was used to determine levels of human Serpine-1 (PAI-1) (Catalog # DSE100, R&D Systems, Minneapolis, MN, USA) in the parent cells, isolated brain endothelial EVs, NPC cell culture media, and NPCs.

Serpine-1 (PAI-1) activity was determined using the Plasminogen Activator Inhibitor Type 1 Human Chromogenic Activity Assay Kit (Abcam, #ab108894) according to the manufacturer's instructions. Briefly, a fixed amount of tissue-type plasminogen activator (tPA) was added in excess to undiluted sample, which allowed Serpine-1 and tPA to form an inactive complex. The assay measures plasminogen activation by residual tPA in coupled assays that contain tPA, plasminogen, and a plasmin-specific synthetic substrate. The amount of plasmin produced was quantitated using a highly specific plasmin substrate releasing a yellow para-nitroaniline chromophore. The absorbance of the chromophore at 405 nm was inversely proportional to the Serpine-1 enzymatic activity. One arbitrary unit (AU) of inhibition was defined as the amount of Serpine-1 that can inhibit one IU of tPA/ml under the testing conditions. EV-Serpine-1 activity was also measured in the presence of Tx-100 1%, which was used to lyse EVs.

Tissue Plasminogen Activator (tPA) Assay

tPA activity was measured with the Tissue type Plasminogen Activator Activity Assay Kit (Abcam, #ab108905) following the manufacturer's instructions. Briefly, the tPA assay protocol measures the ability of tPA to activate the plasminogen to plasmin in coupled or indirect assays that contain tPA, plasminogen, and a plasmin-specific synthetic substrate. The amount of plasmin produced was quantified using a highly specific plasmin substrate releasing a yellow para-nitroaniline chromophore. The change in absorbance of the chromophore in the reaction solution at 405 nm was directly proportional to the tPA enzymatic activity.

Confocal Microscopy

HBMEC cultured on collagen I coated chambered glass slides (Becton Dickinson Biosciences, San Jose, CA) were transfected with the CD63 RFP and Serpine-1 GFP constructs as described above. The parent cells expressing CD63 RFP and Serpine-1 GFP were imaged live after 24 h using an inverted confocal laser-scanning microscope (Olympus Fluoview 3000). Twenty-four hours post transfection, cells were exposed to HIV-1 or/and A β (1–40) HiLyte 647 for 48 h. At the end of treatment, the culture media was removed for fluorescent EV isolation; cells were washed with PBS and fixed with ethanol for 30 min at 4 °C. CD63 RFP, Serpine-1 GFP and A β (1–40) HiLyte 647 fluorescence was imaged with an inverted confocal laser-scanning microscope (Olympus Fluoview 3000, objective lens UPLXAPO100XO 100X oil, numerical aperture 1.45) and analyzed using CellSens software.

After isolation from the media, fluorescent EVs were pipetted onto cleaned glass slides, heat fixed for 10 min at 95 °C, and then fixed with ethanol for 30 min at 4 °C followed by PBS wash. Slides were mounted using ProLong Gold Antifade reagent with or without 4',6-diamidino-2-phenylindole (DAPI, Invitrogen, Carlsbad, CA, USA) to visualize the nucleic material in EVs. Specimens were covered with coverslips and the fluorescent images were evaluated and captured under confocal microscopy. Red fluorescence originating from EV-CD63 RFP, blue fluorescence from DAPI, green fluorescence from EV-Serpine-1 GFP and far-red fluorescence from EV-A β HiLyte-Alexa Fluor 647 was acquired directly using confocal microscopy (Olympus, Fluoview 3000, 100 \times oil immersion lens, room temperature). Serpine-1 GFP positive, CD63 RFP positive EVs were counted with the CellSens software and expressed as percentage of the total DAPI positive EVs. Serpine-1 GFP and CD63 RFP double positive EVs were counted and expressed as percentage of the total Serpine-1 GFP positive EV number.

NPCs were seeded on laminin coated 8-well chambered glass slides (15,000 cells/well) and incubated overnight at 37 °C in maintenance culture medium (Millipore), containing 20 ng/ml FGF-2 and 20 ng/ml of rhEGF. The following day, the medium was changed to maintenance medium without growth factors to induce differentiation. Cells were allowed to differentiate for a total of 3 days, with the last 24 h in the presence of the isolated EVs containing Serpine-1 GFP, CD63 RFP \pm A β (1–40) HiLyte 647. After 1 h of EV exposure, CD63 RFP, Serpine-1 GFP and A β (1–40) HiLyte 647 fluorescence in the acceptor non-fluorescent NPCs was imaged live with an inverted confocal laser-scanning microscope. After 24 h of EV treatment, NPC media was removed, cells were washed with PBS and fixed with ethanol for 30 min at 4 °C. CD63 RFP, Serpine-1 GFP

and A β (1–40) HiLyte 647 fluorescence in the acceptor non-fluorescent NPCs was imaged randomly as above.

HBMEC were exposed to HIV-1 or/and A β (1–40) HiLyte 488 for 48 h followed by EV isolation from the conditioned media. NPCs cultured on chambered slides and differentiated as above, were exposed to EV-A β HiLyte 488 for 24 h. Then, the media was removed, cells were washed with PBS and fixed with ethanol for 30 min at 4 °C. Cells were washed again with PBS and transferred A β HiLyte 488 fluorescence was assessed by confocal microscopy.

On some confocal images we have increased brightness or contrast for better visibility of the fluorescence signal. These changes were consistent across all treatment groups to preserve the integrity of the data.

Mitochondrial Network Analysis (MiNA)

Mitochondria were visualized with Mitotracker Deep Red (Invitrogen, Catalog #M22426). The total area of fluorescence intensity on the acquired images was normalized to the number of NPC nuclei. Confocal microscopy images were analyzed using FIJI software (v. 2.3.0, NIH). Images were pre-processed using Bio-Formats (v.6.9.0, <https://github.com/ome/bioformats>). Mitochondrial network parameters were analyzed using the Stuart Lab Mitochondrial Network Analysis (MiNA) (v. 3.0.1, <https://github.com/StuarLab>) plugin [37].

Mitochondrial Stress Seahorse Assay

The Seahorse XFe24 Analyzer was used to calculate oxygen consumption rate (OCR; a measure of mitochondrial respiration) and extracellular acidification rate (ECAR; a measure of glycolysis) in NPCs using the Agilent Seahorse XF Cell Mito Stress Test Kit (Agilent Technologies, Santa Clara, CA, USA). Cells were seeded in a 24XF cell culture microplate at 30,000 cells/well differentiated and treated with brain endothelial EVs as described above. After EV treatment for 24 h, the cell culture media were replaced with 500 μ l of the prepared Seahorse medium (containing Seahorse XF DMEM media, 2 mM L-glutamine, 10 mM glucose, and 1 mM pyruvate) and incubated at 37 °C for 1 h. Measurements of mitochondrial respiration and glycolysis were carried out as previously described [38]. In brief, cells were treated with 1.5 μ M concentrations of oligomycin (ATP synthase inhibitor of complex V), 1 μ M carbonyl cyanide-p-trifluoromethoxyphenylhydrazone (FCCP, electron transport chain (ETC) uncoupler), and 0.5 μ M rotenone with antimycin A (both ETC inhibitors of complex I and III, respectively) throughout the analysis. These treatments, which were added to the cells at specified time points, allowed for calculations of mitochondrial respiratory parameters, such as baseline OCR, ATP production, maximal respiration,

proton leak, and non-mitochondrial oxygen consumption. After three basal measurements of OCR and ECAR were recorded, oligomycin was injected to inhibit ATP synthase and two more measurements were recorded to assess proton leak. Next, FCCP was injected to uncouple respiration and measure maximal respiration. Finally, antimycin A was injected to measure non-mitochondrial respiration. All OCR measurements were normalized to non-mitochondrial respiration and the final values normalized to NPC protein concentration in each well. Reserve capacity is the difference between maximal respiration and basal respiration, while ATP-linked OCR is the difference between basal and proton leak. The data were analyzed using the Wave Software (Agilent Technologies). All conditions were measured in 4–6 samples/group, and three repeats were performed for this experiment.

Expression of Synaptic Proteins

EV-treated NPCs cultured on laminin coated chambered glass slides (ibidi USA, Madison, WI, USA) were fixed with ethanol for 30 min at 4 °C. After washing with PBS and blocking with 3% bovine serum albumin in PBS for 30 min at room temperature, samples were incubated overnight at 4 °C with the primary antibody: mouse anti-PSD95 monoclonal antibody (Abcam, Waltham, MA, USA, Catalog #192,757, 1:1000) or rabbit anti-synaptophysin polyclonal antibody (Abcam, 1:1000). Then, the excess of primary antibody was removed, slides were washed with PBS, and incubated with Alexa Fluor 488/594-conjugated secondary antibodies (1:1000, Invitrogen) for 2 h at room temperature. Nuclei were stained with Hoechst 33342 (Invitrogen, Catalog #H3570). The immunofluorescent images were evaluated and captured under confocal microscopy. Red fluorescence originating from PSD95, blue fluorescence from Hoechst 33,342, and green fluorescence from synaptophysin were acquired directly using confocal microscopy (Olympus, Fluoview 3000, 100 \times oil immersion lens, room temperature).

Quantitative analysis of synaptic protein expression was performed similarly to a previously published method, with modifications [39]. After immunofluorescence staining for PSD95 and synaptophysin, fields were selected randomly using the guidance of DAPI nuclear staining (three images/treatment group, total 9 images from three independent experiments), then confocal images were taken. On each confocal image (brightfield), 20 identical rectangular areas were randomly superimposed on different segments of NPC projections. The selected areas were analyzed for green (synaptophysin) and red (PSD95) fluorescence intensity using the NIH Image J software (Bethesda, MD, USA). Mean Fluorescence Intensity (MFI) for each area was normalized by subtracting the background fluorescence intensity for that

image. Synaptic protein densities were expressed relative to those in control NPCs. For total synaptophysin and PSD95 assessment, mean fluorescence intensity on the acquired images was normalized to the number of NPC nuclei.

Statistical Analysis

Data were analyzed using GraphPad Prism 9.0 (Graphpad Software, San Diego, CA). ANOVA was used to compare responses among treatments. Treatment means were compared using All Pairwise Multiple Comparison Procedures and $p < 0.05$ was considered significant.

Results

Serpine-1 is Concentrated in EVs Released from Control and HIV-1 Exposed Brain Endothelial Cells

Secretion of Serpine-1 in EVs was traced by cotransfection of HBMEC with the Serpine-1 GFP and CD63 RFP constructs. The tetraspanin CD63 is a membrane protein, which is predominantly localized to the vesicles, and; therefore, commonly used as a biomarker for EVs. As demonstrated by live fluorescence microscopy 24 h post transfection, transfected cells appeared to concentrate and secrete Serpine-1 GFP in CD63 RFP-positive EVs. Indeed, green fluorescence, corresponding to Serpine-1, noticeably overlapped with CD63 RFP-positive red fluorescent EVs budding off from the parent cells (arrow heads for single markers, arrows for overlapping fluorescence, Fig. 1A). We also measured the total number of EVs in the media originating from non-transfected HBMEC. EVs released from control HBMEC had a total EV concentration of $43.05 \pm 6.65 \times 10^8$ particles/ml, while EVs released from HIV-1 treated HBMEC had a total EV concentration of $83.40 \pm 34.60 \times 10^8$ particles/ml. This HIV-related increase in EVs release is consistent with our previous report [20].

Because both EVs and Serpine-1 were shown to play a role in A β pathology [20, 30, 31], we next evaluated the impact of HIV-1 on Serpine-1 and A β release via EVs. For these experiments, 24 h after cotransfection with Serpine-1 GFP and CD63 RFP, HBMEC were exposed to HIV-1 (30 ng p24/ml) and/or 100 nM A β (1–40) HiLyte 647 for 48 h. The treatment was terminated by removing the cell culture media for EV isolation, followed by washing with PBS and fixing the parent cells. As illustrated in Figs. 1B and C, Serpine-1 GFP green fluorescence and CD63 RFP red fluorescence partially overlapped with the fluorescent A β HiLyte 647 (yellow) taken up by the parent cells (arrow heads for single-, arrows for overlapping fluorescence). The images were quantified for Serpine-1, CD63, or A β HiLyte

colocalization (Fig. 1D), indicating that Serpine-1 GFP and A β HiLyte 647 colocalization significantly increased in the HIV + A β group when compared to control (Fig. 1D, lower right graph).

Next, Serpine-1 GFP, CD63 RFP and A β (1–40) HiLyte 647 fluorescence was visualized in EVs isolated from cell culture media. Serpine-1 GFP was detected in EVs of different sizes in control, A β and/or HIV-1 treated samples (Fig. 2A) indicating that the parent cells secrete Serpine-1 via EVs. Interestingly, overall fewer CD63-RFP positive EVs were present in all groups and A β HiLyte 647 occasionally colocalized with both Serpine-1 GFP and CD63-RFP in the secreted EVs. In the A β groups we frequently observed aggregates of A β HiLyte 647 with associated EVs (Fig. 2A, arrow heads for single markers, arrows for overlapping fluorescence). We also stained the EV genetic material with DAPI (blue fluorescence). Most of the EVs showed DAPI fluorescence indicating DNA/RNA cargo.

To further characterize the isolated EVs, we counted the number of Serpine-1 GFP-, CD63 RFP- and DAPI positive EVs from the confocal microscopy images. As quantified on the graphs from Fig. 2B, most of the DAPI containing EVs were also positive for Serpine-1 GFP. This percentage did not change significantly in any treatment group. In contrast, the percentage of CD63 RFP positive EVs was much lower and significantly decreased in the HIV-EV and the HIV + A β -EV groups as compared to the A β -EV group. Similarly, the number of Serpine-1 GFP and CD63 RFP double positive EVs changed in the same way (Fig. 2B).

HIV-1 Impacts Serpine-1 Levels and Activity in the Released EVs

Serpine-1 levels in the EV lysates were next assessed by ELISA in non-transfected HBMEC and normalized either to cell culture media volume or to EV protein content (Figs. 3A). EVs isolated from control, A β , and/or HIV-treated HBMEC cultures contained Serpine-1. Importantly, Serpine-1 levels were significantly higher in the HIV-1 group as compared to the control or A β groups when normalizing to cell culture volume (Fig. 3A, left graph). Serpine-1 levels in the HIV + A β group were not significantly different from other groups, when statistical analysis was performed on all treatment groups at the same time. However, the changes in Serpine-1 levels between the HIV + A β group vs the Control group gained significance when comparing only these two groups (Fig. 3A, middle graph). In the light of the entire study, we propose that these differences may have important biological implications and may be relevant in prolonged HIV-1 infection or in older HIV-1 patients where both HIV-1 and A β may play a role in neuropathogenesis. Nevertheless, EV Serpine-1 concentration was similar in all treatment groups after normalization to EV

Fig. 1 Serpine-1 is concentrated in vesicular structures in brain endothelial cells. HBMEC were transfected with the Serpine-1 GFP and CD63 RFP plasmids (**A**) and 24 h after transfection, cells were exposed to HIV-1 (30 ng p24/ml) and/or 100 nM A β (1–40) HiLyte 647 for 48 h (**B**). **A** The images represent live imaging of Serpine-1 GFP and CD63 RFP (arrow heads) in the parent HBMEC 24 h after transfection. Scale bar: 10 μ m. **B** Serpine-1 GFP, CD63 RFP, or A β HiLyte 647-positive fluorescence (arrow heads) in the fixed parent cells 48 h after vehicle (control), HIV-1, and/or A β exposure. Arrows indicate overlapping positive fluorescence of Serpine-1 GFP, CD63 RFP, and/or A β HiLyte 647. Scale bar: 10 μ m. Representative images from four experiments. **C** Confocal z-stack imaging of Serpine-1 GFP, CD63 RFP, and A β HiLyte 647 colocalization (arrows) in the fixed parent cells 48 h after HIV-1 and A β exposure. Scale bar: 5 μ m. **D** Quantification of colocalization of Serpine-1 GFP and CD63 RFP in the live parent cells after transfection (upper graph), unpaired t-test; Colocalization of Serpine-1 GFP and CD63 RFP (lower left graph) and Serpine-1 GFP and A β HiLyte 647 (lower right graph) in the fixed parent cells 48 h after HIV-1 and/or A β exposure. Values are mean \pm SEM, $n = 3–9$. One- and two-way ANOVA with Tukey's multiple comparisons test. *Statistically significant at $p < 0.05$

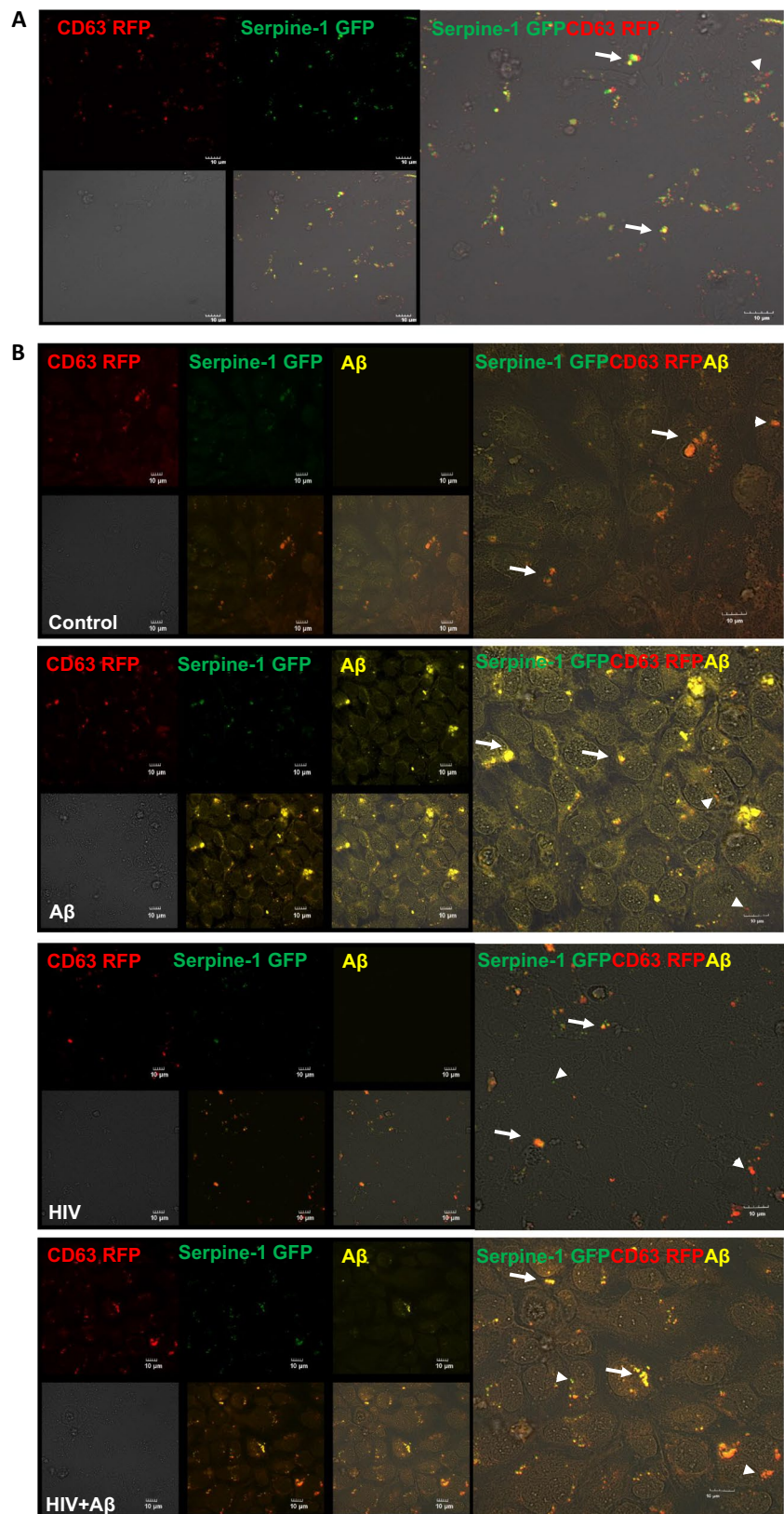
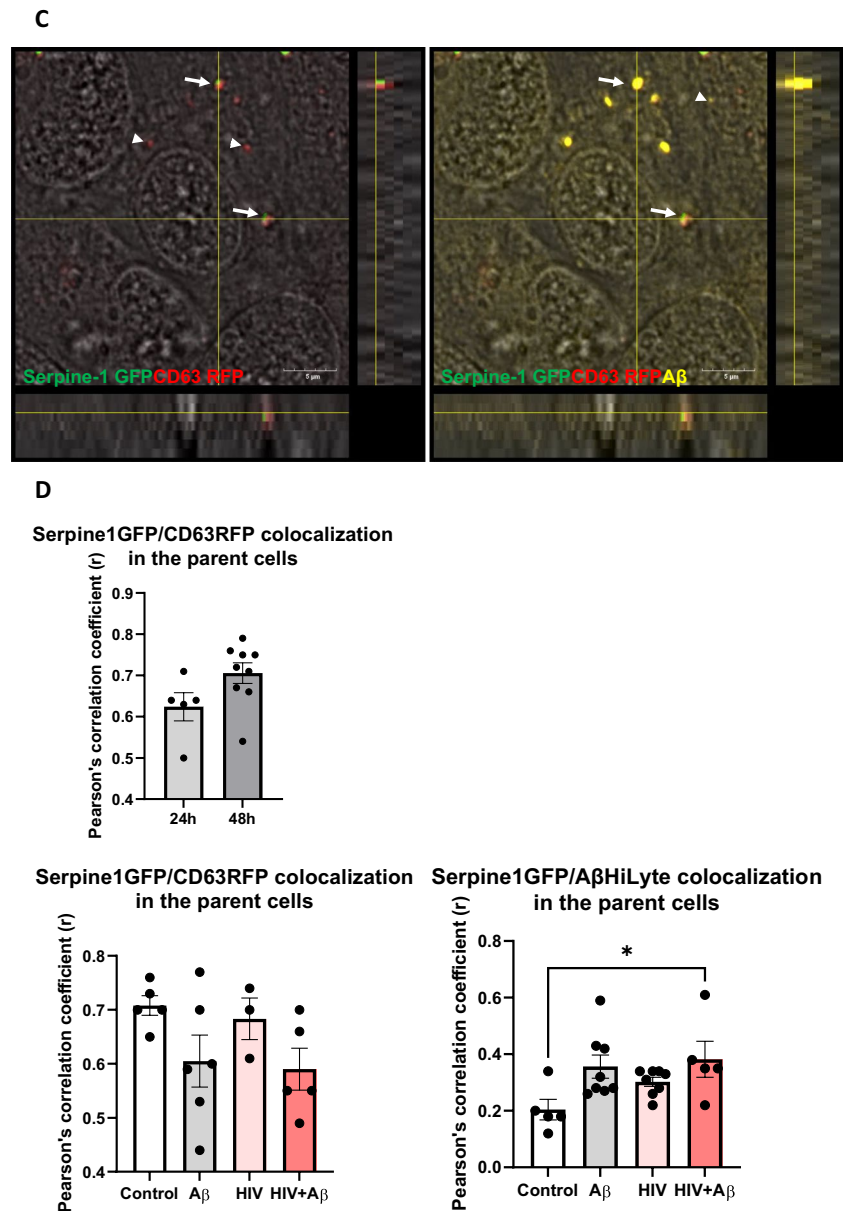


Fig. 1 (continued)



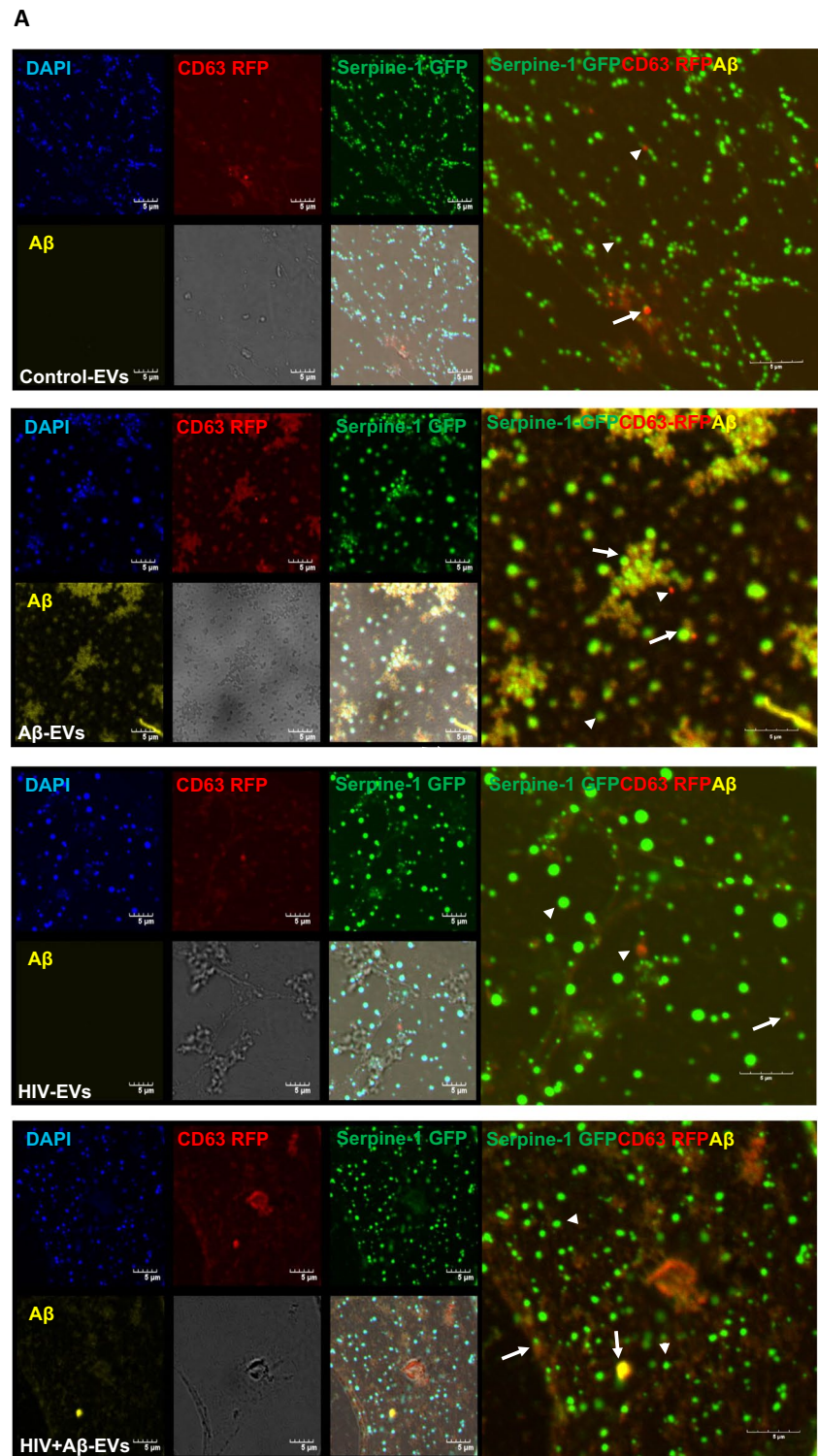
protein levels (Fig. 3A, right graph), presumably because exposure to HIV-1 increases the overall EV number produced as reported before [20]. Interestingly, Serpine-1 levels in the parent cells were in the picogram range and thus much lower as compared to the nanogram range of Serpine-1 in EVs. Moreover, Serpine-1 levels in the parent cells did not change after HIV-1 and/or A β treatment (Fig. 3B).

Serpine-1 is an enzyme, therefore, its activity was also measured from the isolated EVs lysed with 1% Tx100 (see Methods). HIV + A β -EVs had the highest Serpine-1 activity, which was statistically significant when compared to the control and A β groups and remained significant even after normalizing the results to EV protein content (Fig. 3C). Interestingly, this increase in EV Serpine-1 activity occurred even though Serpine-1 protein levels did not

change (Fig. 3A). While the mechanism of this apparent discrepancy is not clear, A β in combination with HIV-1 may change the EV microenvironment that promotes Serpine-1 activity without changes in its protein levels.

Because Serpine-1 is known to inhibit tPA activity [24]; therefore, we also measured tPA activity in the isolated EVs. As illustrated on Fig. 3D, tPA activity trends were the opposite of those of Serpine-1 activity. Treatment with A β , HIV-1 and HIV + A β of the parent cells had a significant impact on tPA activity in EVs, which significantly decreased in these groups as compared to control, when normalizing to cell culture volume (Fig. 3D, left graph). A decreasing trend was also observed when normalizing the results to EV protein content (Fig. 3D, right graph) but the changes were not statistically significant.

Fig. 2 Serpine-1 is released in EVs from control, A β , and/or HIV-1 exposed brain endothelial cells (**A**) Visualization by confocal microscopy of Serpine-1 GFP (green), CD63 RFP (red), and A β (1–40) HiLyte (yellow) (arrow heads) in EVs isolated from media of transfected and treated HBMEC as in Fig. 1. Examples of overlapping fluorescence of Serpine-1 GFP, CD63 RFP, and/or A β (1–40) HiLyte associated with EVs are indicated by arrows. DAPI stains the genetic material in EVs. Representative images from three experiments. Scale bar: 5 μ m. **B** Quantification of Serpine-1 GFP-positive, CD63 RFP-positive, and Serpine-1 GFP/CD63 RFP-double positive EVs. Values are mean \pm SEM, $n=5-7$. One- and two-way ANOVA with Šídák's and Tukey's multiple comparisons tests. *Statistically significant at $p < 0.05$

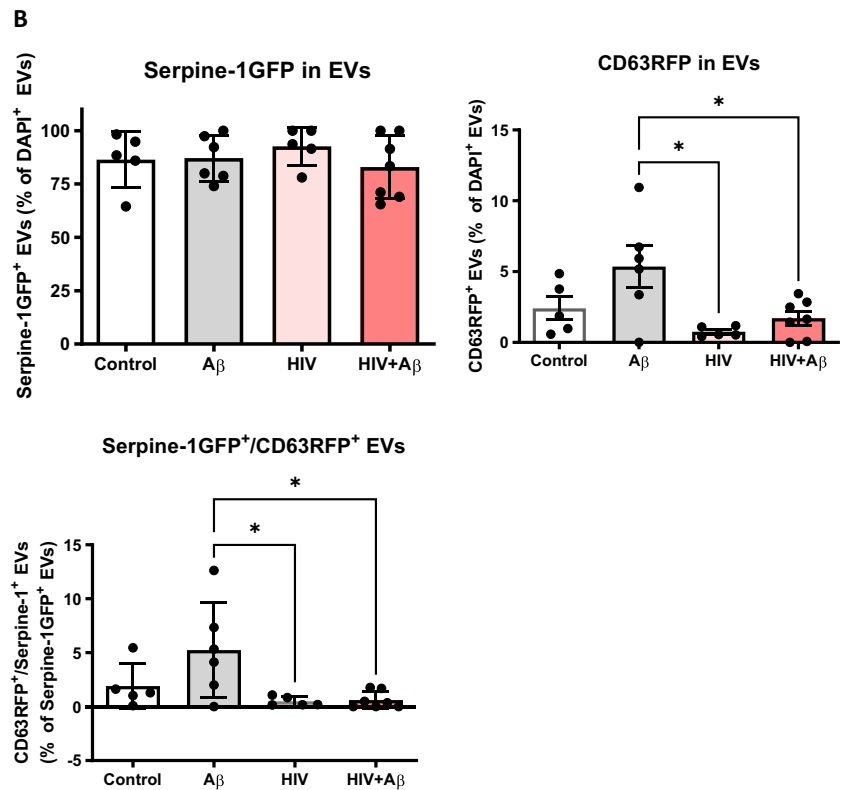


HBMEC-derived EVs Transfer Serpine-1 Cargo to Recipient Neural Progenitor Cells

HBMEC are part of a functional unit at the BBB that consists of pericytes, perivascular astrocytes, microglia, and neurons, called the neurovascular unit [40]. Therefore, we

hypothesized that HBMEC-derived EVs can transfer Serpine-1 to neighboring cells of the neurovascular unit, including neural progenitor cells (NPCs). It was shown that ~47% of dividing progenitor and 46% of transit amplifying cells (precursors of neuroblasts) are located in close proximity to the brain endothelium [41, 42].

Fig. 2 (continued)



In order to assess Serpine-1 transfer to NPCs, HBMEC transiently transfected with Serpine-1 GFP and CD63-RFP were exposed to 100 nM A β HiLyte 647 and/or HIV-1 for 48 h, resulting in secretion of Serpine-1 GFP positive and CD63 RFP positive EVs, with some of them containing fluorescent A β cargo. EVs were isolated from the cell culture media and then employed to differentiating NPCs for 24 h. Green fluorescence signals (corresponding to EV-derived Serpine-1 GFP), red fluorescent signals (indicating EV-CD63 RFP) and yellow A β HiLyte 647 fluorescence (indicating EV-A β cargo) in the acceptor non-fluorescent NPCs was assessed by confocal microscopy.

Representative images of NPC cultures exposed for 1 h to EVs derived from control, HIV plus/or A β -treated HBMEC are illustrated in Fig. 4A, with a variety of vesicular and non-vesicular structures. Some EVs show red fluorescence due to the presence of the EV marker CD63 RFP. EVs also exhibit yellow fluorescence indicating A β transfer via EVs derived from A β -treated HBMEC. Some of the EVs with fluorescent A β cargo show an overlapping red or green fluorescence indicating colocalization with Serpine-1 and/or CD63 (Fig. 4A, arrow heads for individual-, arrows for overlapping fluorescence). After 24 h EV exposure, NPC media was removed, and the acceptor NPCs were fixed and imaged again. Figure 4B visualizes Serpine-1 GFP, CD63 RFP and A β HiLyte transfer to NPCs by EVs derived from HIV and/or A β HiLyte-exposed HBMEC. Some of the transferred

fluorescent A β appeared to be concentrated in large yellow aggregates (arrows), particularly in the HIV + A β group (Fig. 4B).

We also quantified HBMEC-derived EV-Serpine-1 cargo transfer to NPCs. For these experiments, non-transfected HBMEC were exposed to 100 nM non-fluorescent A β and/or HIV-1 for 48 h. EVs were isolated from the cell culture media and used for the subsequent NPC exposure for 24 h, followed by PBS wash. First, Serpine-1 levels in the NPC culture media were assessed by ELISA 1 h and 24 h after EV exposure (Fig. 4C). As illustrated, 1 h after EV exposure, Serpine-1 levels were significantly higher in the HIV-EV and HIV + A β -EV treated groups as compared to the Control-EV group. In addition, Serpine-1 levels in all EV-treated groups were significantly higher when compared to the No EV group, indicating that Serpine-1 in the NPC culture media, indeed, originated from the HBMEC-derived EVs and was not secreted by the recipient NPCs. This trend was maintained at 24 h after EVs exposure as well, indicating that most of the Serpine-1 was still present in the culture media and originated from the employed EVs (Fig. 4C, right graph).

Next, NPC culture media samples were used to assess Serpine-1 activity 1 h and 24 h after EVs treatment (Fig. 4D). Although Serpine-1 activity measurements were low, they showed a statistically significant increase in the HIV + A β -EV group when compared to the control-EV

Fig. 3 Serpine-1 levels and activity in the released EVs. Non-transfected HBMEC were exposed to HIV-1 (30 ng p24/ml) and/or 100 nM A β (1–40) for 48 h, followed by isolation of released EVs from the culture media. Serpine-1 levels in EVs and the parent cells were measured by ELISA. Serpine-1 activity and tPA activity were determined by specific activity assays. **A** Serpine-1 levels in isolated EVs normalized to cell culture media volume (left and middle graph) or to EV protein content (right graph). **B** Serpine-1 levels in the parent cells. Values are mean \pm SEM, $n=4-8$. **C** Serpine-1 activity in the isolated EVs normalized to cell culture media volume (left graph) or to EV protein content (right graph). **D** tPA activity in the isolated EVs normalized to cell culture media volume (left graph) or to EV protein content (right graph). Values are mean \pm SEM, $n=4-5$. One- and two-way ANOVA with Tukey's multiple comparisons test. *Statistically significant at $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, or **** $p < 0.0001$

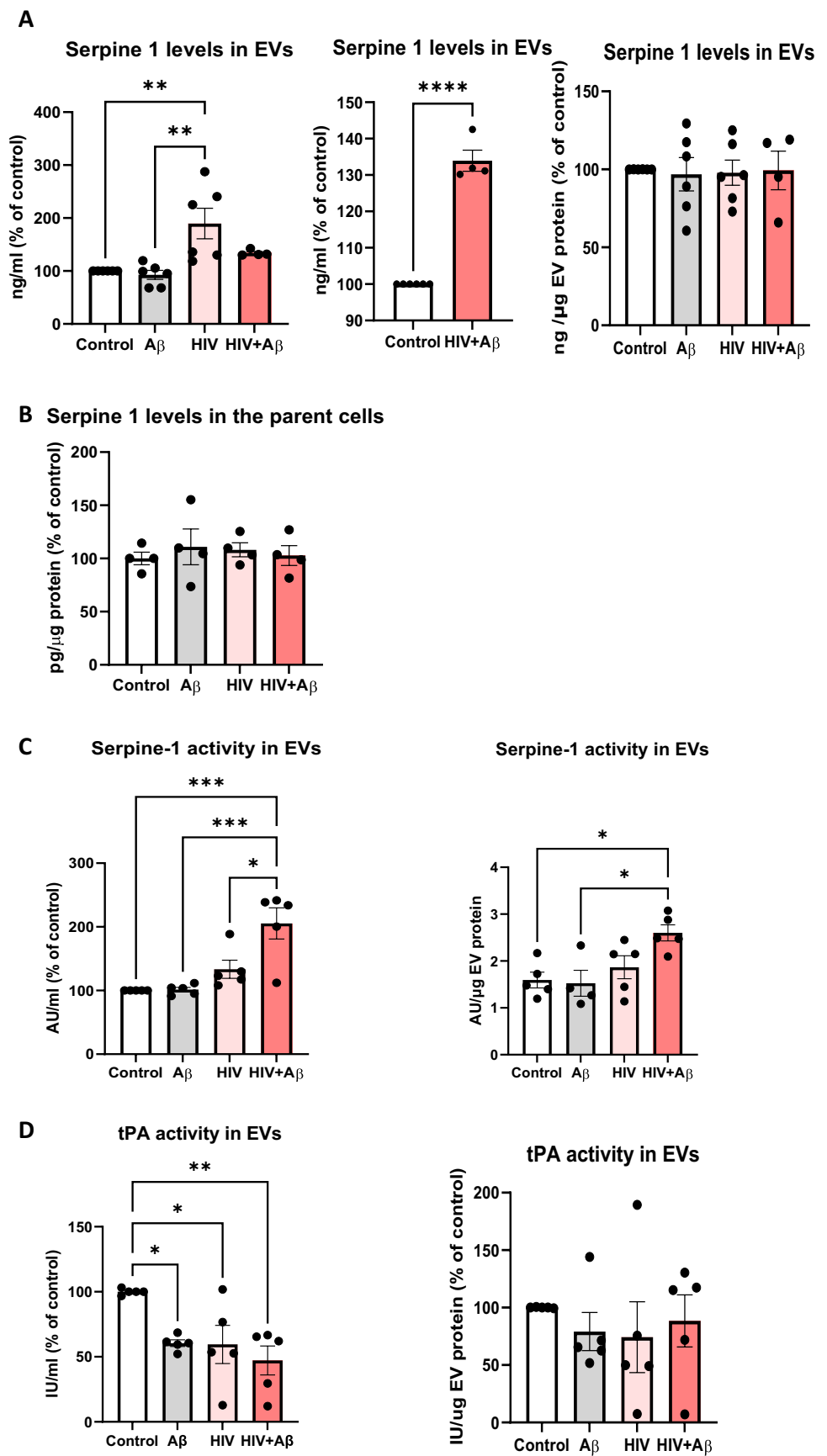
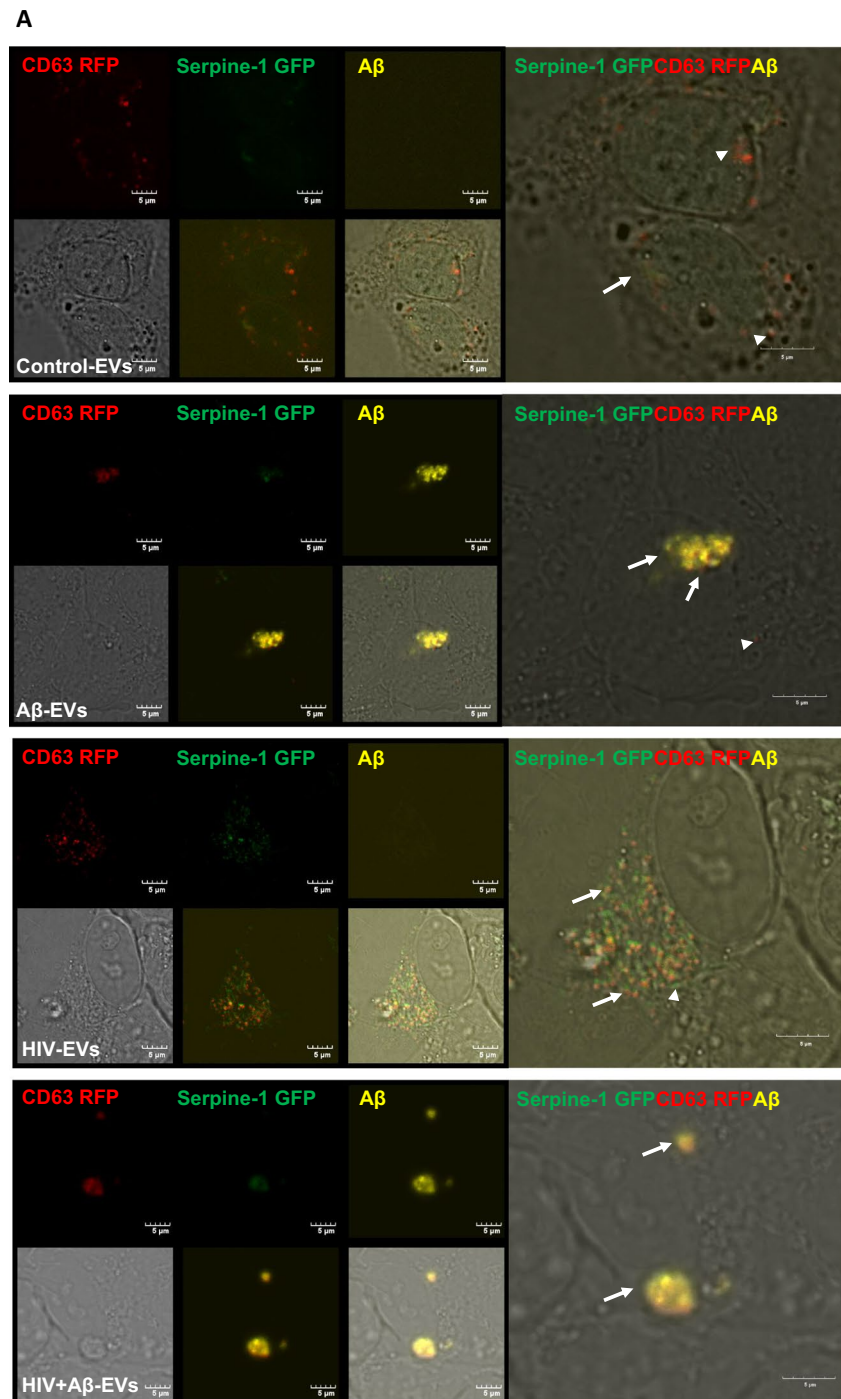


Fig. 4 Transfer of Serpine-1 from donor HBMEC-derived EVs to recipient neural progenitor cells (NPCs). HBMEC were transfected and treated as in Fig. 1, followed by isolation of EVs from the cell culture media and treatment of NPCs for up to 24 h. All images were performed by confocal microscopy. **A** Live imaging of Serpine-1 GFP (green), CD63 RFP (red) and A β (1–40) HiLyte (yellow) in non-fluorescent NPC cultures (arrow heads) after 1 h exposure to EVs from Control HBMEC, HIV-1 and/or A β -treated HBMEC. Scale bar: 5 μ m. **B** Visualization of Serpine-1 GFP (green), CD63 RFP (red) and A β (1–40) HiLyte (yellow) in fixed non-fluorescent NPC cultures (arrow heads) after 24 h of EVs exposure. Examples of overlapping fluorescence of Serpine-1 GFP, CD63 RFP and A β (1–40) HiLyte in (A) and (B) are indicated by arrows. Scale bar: 10 μ m. Representative images from three experiments. **C–F** Non-transfected HBMEC were treated as in Fig. 3. EVs were isolated from the culture media and employed for NPC treatment for 1 (left panels) or 24 h (right panels). Serpine-1 levels (C), Serpine-1 activity (D), and tPA activity (E) were determined in NPC media as in Fig. 3. **F** Serpine-1 levels in the recipient NPCs 1 h (left graph) and 24 h (right graph) after EVs exposure. Values are mean \pm SEM, $n=4-7$. One- and two-way ANOVA with Tukey's multiple comparisons test. *Statistically significant at $p<0.05$, ** $p<0.01$, *** $p<0.001$, and **** $p<0.0001$

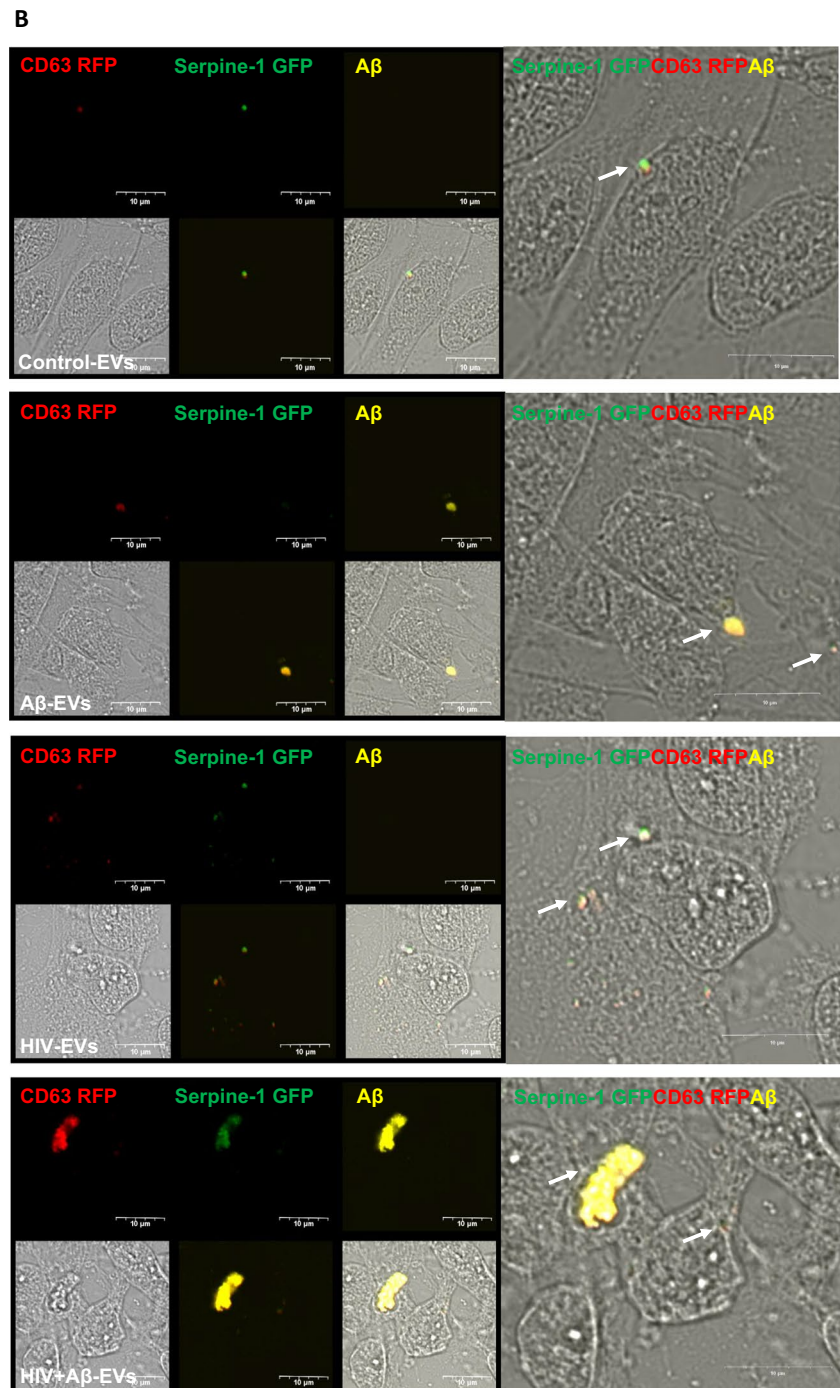


group, corresponding to Serpine-1 levels measured by ELISA. An increasing trend in the HIV-EV group was also observed although values in the No EV group were scattered (Fig. 4D).

In the following experiments, we explored whether Serpine-1 in the NPC media can inhibit tPA activity. As illustrated on Fig. 4E (left graph), tPA activity after 1 h was the highest in the control-EV group and showed a decreasing trend in the other groups. tPA activity in the Control-EV

and A β -EV groups was significantly higher as compared to the No EV group (Fig. 4E). Overall, tPA activity change trends were the opposite of the Serpine-1 activity trends, confirming that the transferred EV-Serpine-1 can inhibit the transferred EV-tPA activity. Interestingly, 24 h after EV exposure, tPA activity levels were similar in all EV-treated groups and most of them were significantly higher as compared to the No EV group (Fig. 4E, right graph), verifying that tPA activity in NPCs originated mainly from the

Fig. 4 (continued)



employed EV exposure and that NPCs have no or minimal endogenous tPA activity.

Serpine-1 levels and activity were also measured in the recipient NPCs. As shown on Fig. 4F, Serpine-1 levels were much lower (picogram range) in the recipient NPCs as compared to the media (nanogram range). These levels were significantly higher in the HIV + A β -EV treated group as compared to the Control-EV group and to the No EV group, again indicating that Serpine-1 in the NPCs

originated from the HBMEC-derived EVs. This trend was similar at 1 and 24 h after EVs exposure (Fig. 4F). Finally, we measured Serpine-1 activity and tPA activity in the recipient NPCs 1 h and 24 h after EVs exposure; however, the levels were mostly very low or undetectable (data not shown). These results are consistent with Serpine-1 and tPA being secreted enzymes; therefore, they may primarily exert their effects on the recipient NPCs from the extracellular side.

Fig. 4 (continued)

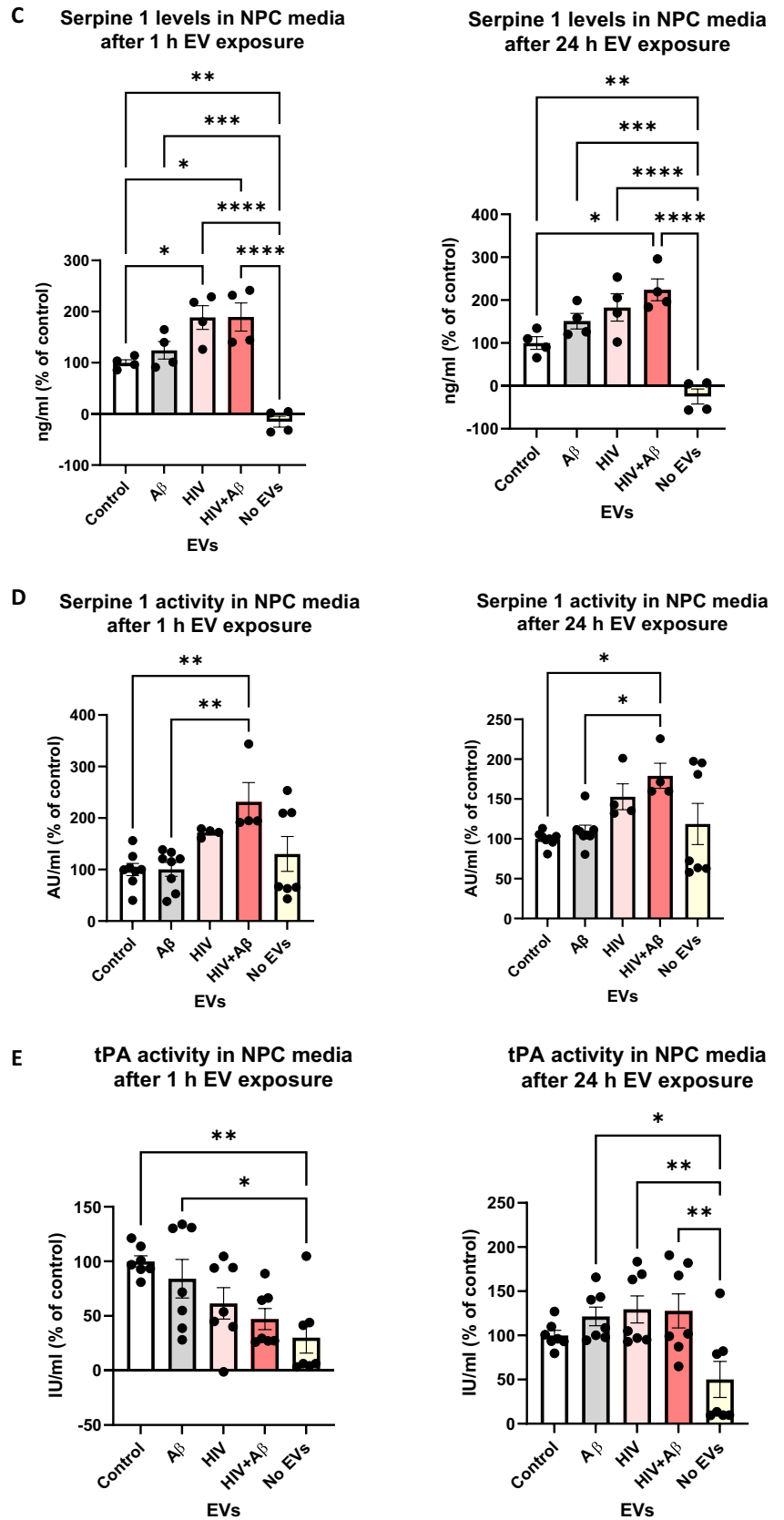
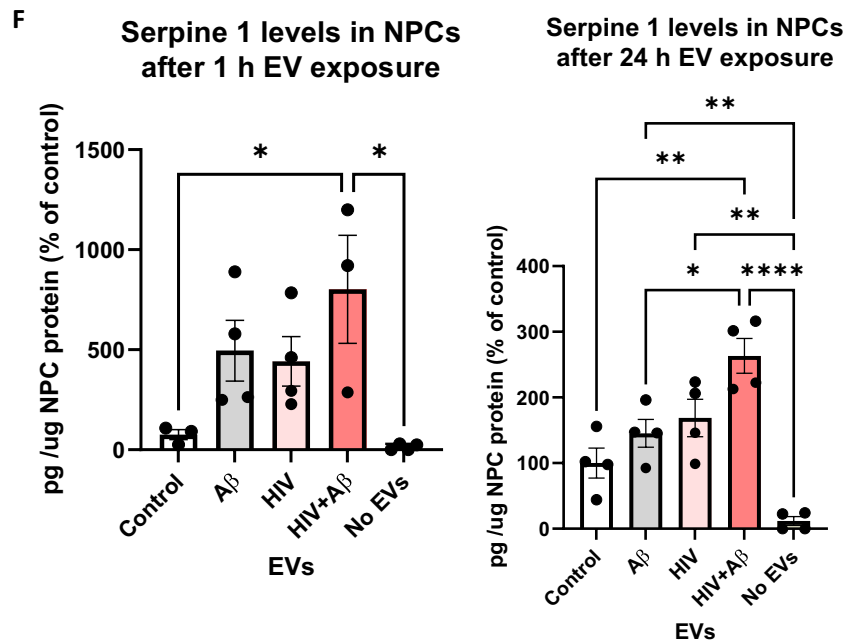


Fig. 4 (continued)



Serpine-1 is Involved in EV-mediated Transfer of A β Cargo to Recipient NPCs

After establishing that HBMEC-derived EVs can transfer both Serpine-1 and A β , we explored if Serpine-1 can be involved in A β transfer and/or uptake by NPCs. Non-transfected HBMEC were exposed to 100 nM A β HiLyte 488 and/or HIV-1 for 48 h, followed by isolation of EVs, which were then employed for NPC exposure for 24 h. Representative images of NPCs exposed to fluorescent EVs for 24 h visualize the transferred A β HiLyte (green fluorescence) and the NPC mitochondria traced with Mitotracker (red fluorescence) (Fig. 5A). To assess the involvement of Serpine-1 in this process, NPCs were cotreated with the Serpine-1 inhibitor PAI039 for 24 h. PAI039 was used at low concentration of 2 μ M, which did not affect NPC viability (Fig. 5B).

EV-derived A β HiLyte fluorescence was next quantified in the recipient NPCs using a plate reader and normalized to nuclear DRAQ5 fluorescence (Fig. 5C). A significant fluorescence increase was observed upon treatment with EVs derived from A β HiLyte-treated HBMEC and HIV-1 plus A β HiLyte-treated HBMEC as compared to treatment with EVs from control HBMEC. PAI039 affected the EV-A β HiLyte transfer only in NPCs exposed to EVs derived from HIV-1 plus A β HiLyte HBMEC. Specifically, cotreatment with PAI039 significantly increased A β transfer in this group as compared to the HIV + A β -EV group without inhibitor (Fig. 5C). These results indicated that transfer of EV-derived A β cargo could be modulated by Serpine-1 only in specific conditions, such as HIV-1 exposure.

When we incubated only the isolated EVs with the inhibitor, we have observed that PAI039 paradoxically increased

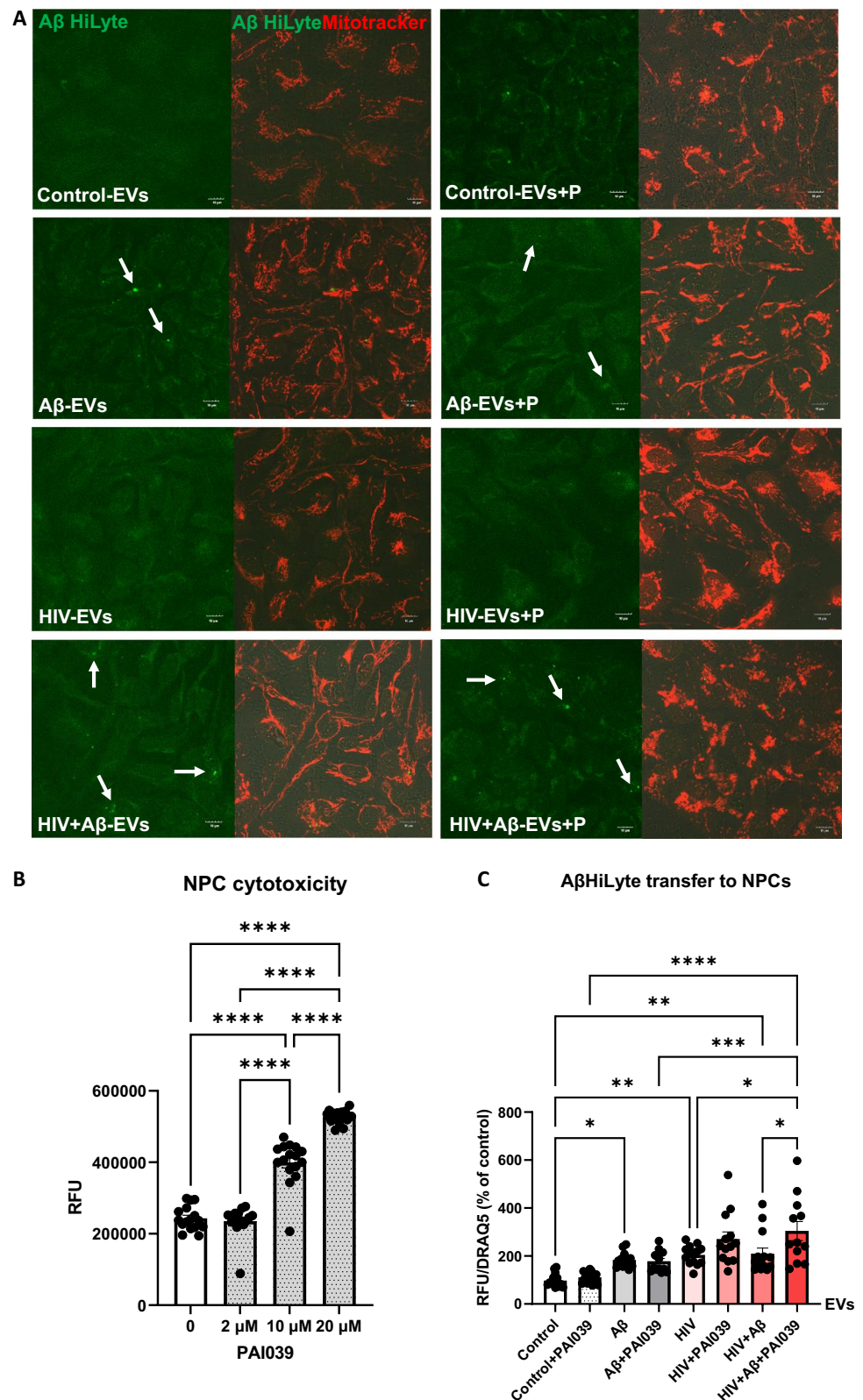
EV-Serpine-1 activity in the HIV groups and caused an increasing trend in the non-HIV groups when compared to the respective groups without PAI039 (Supplementary Fig. 1A). Although observed in a different experimental setup, these unexpected effects may have contributed to the impact of PAI039 on EV-A β transfer to NPCs in the context of HIV-1. On the other hand, PAI039 did not change tPA activity in the isolated EVs (Supplementary Fig. 1B). It is possible that A β in the presence of HIV-1 creates a unique cellular environment that may be responsible for an unexpected increase in EV-A β transfer to NPCs in the presence of PAI039.

Serpine-1 Impacts Mitochondrial Networks and Functions in NPCs Exposed to HBMEC-derived EVs

In the next series of experiments, we investigated the implications of Serpine-1 transfer to NPCs via EVs by evaluating the mitochondrial networks and functions. In support of this line of investigation, there is evidence that proper mitochondrial functions of NPCs are essential for correct neurogenesis [43].

To assess mitochondrial morphology changes, we employed Mitotracker Deep Red to stain mitochondrial networks in the NPCs treated with EVs isolated from control, A β , and/or HIV-exposed HBMEC (Fig. 6A). The experiments also involved treatment with PAI039. Morphological analysis was performed using mitochondrial network analysis (MiNA), an Image J plug-in, as previously published [37] to quantify the networks pre-processed and skeletonized by the software (skeletonized images on Fig. 6A

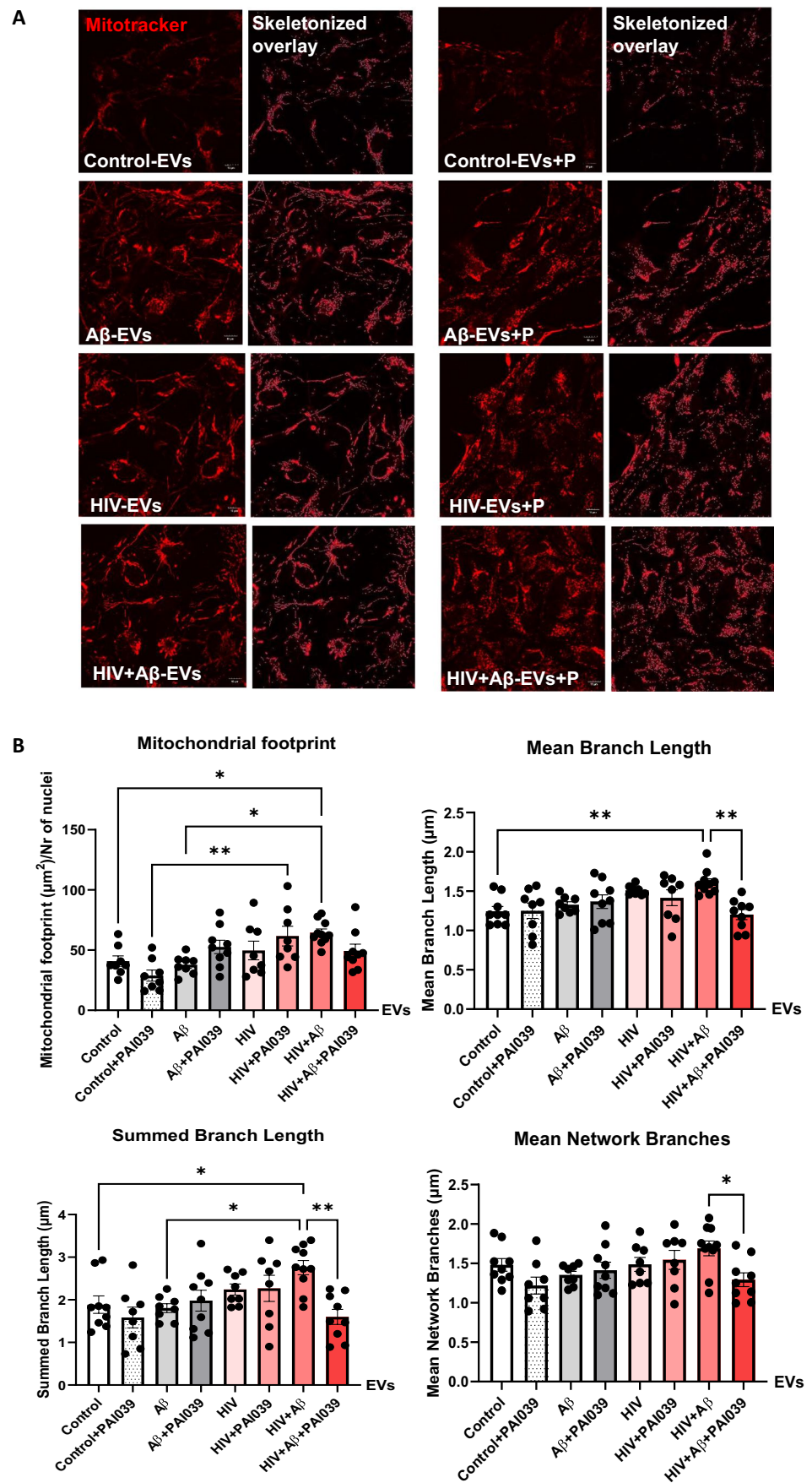
Fig. 5 Involvement of Serpine-1 in the transfer of A β cargo from HBMEC-derived EVs to recipient NPCs. HBMEC were exposed to HIV (30 ng/ml) and/or 100 nM A β (1–40) HiLyte 488 for 48 h, followed by isolation of EVs from the cell culture media and treatment of NPCs for 24 h in the presence or absence of PAI039. **A** Confocal images of recipient NPCs with A β HiLyte fluorescence (green) and Mitotracker (red). **B** Dose-dependent PAI039 cytotoxicity in NPCs. **C** Quantification of A β HiLyte fluorescence in recipient NPCs. NPCs grown on 96-well plates were exposed to HBMEC-derived fluorescent EVs for 24 h. Controls were exposed to non-fluorescent EVs from HBMEC. Selected NPCs were cotreated with the Serpine-1 inhibitor PAI039 (2 μ M) and EVs for 24 h. After washing with PBS, A β HiLyte fluorescence was measured (Abs/Em 503/528 nm) in a plate reader. The values were normalized to nuclear DRAQ5 fluorescence. Values are mean \pm SEM, $n = 12$ –14. One-, two- and three-way ANOVA with Tukey's multiple comparisons test. *Statistically significant at $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$



and graphs on Fig. 6B). The mitochondrial footprint (the total area of the image with the Mitotracker fluorescence signal, normalized to the number of nuclei), was significantly

increased in the HIV + A β -EV group when compared to the A β -EV and control-EV groups. In addition, mitochondrial footprint was significantly higher in the HIV-EV + PAI039

Fig. 6 Mitochondrial network analysis (MiNA) of mitochondrial morphology in EV-exposed NPCs. Non-transfected HBMEC were treated as in Fig. 3. EVs were isolated from the culture media and employed for NPC treatment for 24 h. Selected NPCs were cotreated with PAI039 (2 μ M) and EVs for 24 h. **A** Confocal images of NPCs stained with Mitotracker Deep Red (red) for tracking the mitochondria. MiNA plugin on ImageJ was used to skeletonize the mitochondria. Scale bar: 10 μ m. **B** Quantification of the mitochondrial footprint, mean branch length, total branch length and mean network branches. Values are mean \pm SEM, $n = 8-10$. One-, two- and three-way ANOVA with Tukey’s multiple comparisons test. *Statistically significant $p < 0.05$, ** $p < 0.01$



group as compared to the control-EV + PAI039 group, indicating HIV-1-mediated impact (Fig. 6B, upper left graph). The mean branch length was also significantly increased in the HIV + A β -EV group when compared to the control-EV group, and this effect was significantly blocked by the PAI039 exposure (Fig. 6B, upper right graph). The presence of long branches may indicate hyperfusion, leading to an elongated branched network. The summed or total branch length showed a similar change with a significant increase in the HIV + A β -EV group when compared to the control-EV or A β -EV only groups and PAI039 abolished this effect (Fig. 6B, lower left graph). Consistent with these results, mean network branches were also significantly decreased in the HIV + A β -EV + PAI039 group when compared to the HIV + A β -EV group (Fig. 6B, lower right graph). Taken together, these results indicate that in the HIV + A β -EV group the mitochondrial branches are longer; however, PAI039 can reverse these effects.

Next, we assessed whether mitochondrial network changes were accompanied by functional changes as determined using the Seahorse Mito Stress assay. For these experiments, HBMEC were treated with A β and/or HIV, followed by isolation of EVs as in Fig. 3. Then, human NPCs seeded on 24-well Seahorse plates (30,000/well) were differentiated for 3 days with exposure to these isolated EVs for the last 24 h. Selected NPC cultures were also cotreated with PAI039 (2 μ M) and EVs for 24 h. Mitochondrial oxidative phosphorylation was measured by the oxygen consumption rate (OCR) and glycolysis by analyzing the extracellular acidification rate (ECAR) in real-time in live NPCs (Supplementary Figs. 1C–L). Data from the Seahorse were normalized to NPC protein concentration per well.

Exposure to PAI039 alone did not induce any changes in OCR in NPCs. While ECAR was slightly lower in the PAI039 group as compared to control, these changes did not reach statistical significance (Supplementary Fig. 1C). In NPCs exposed to EVs, the obtained results on Supplementary Fig. 1D–L indicate no or modest impact of Serpine-1 inhibition on bioenergetics of mitochondria. While the changes are only moderate, increases in basal and maximal respiration, elevated proton leak and ATP production were all observed in the HIV + A β + PAI039-EV group. A decrease in spare respiratory capacity in this group suggests that Serpine-1 inhibition may impair the ability of cells to respond to stress or metabolic demand (Supplementary Fig. 1D–L).

Serpine-1 Affects Synaptic Protein Expression in the Developing NPCs

Literature reports on potential neurotoxicity of Serpine-1 are conflicting. There are indicators that Serpine-1 can block the damage of neuronal networks *in vitro* by increasing

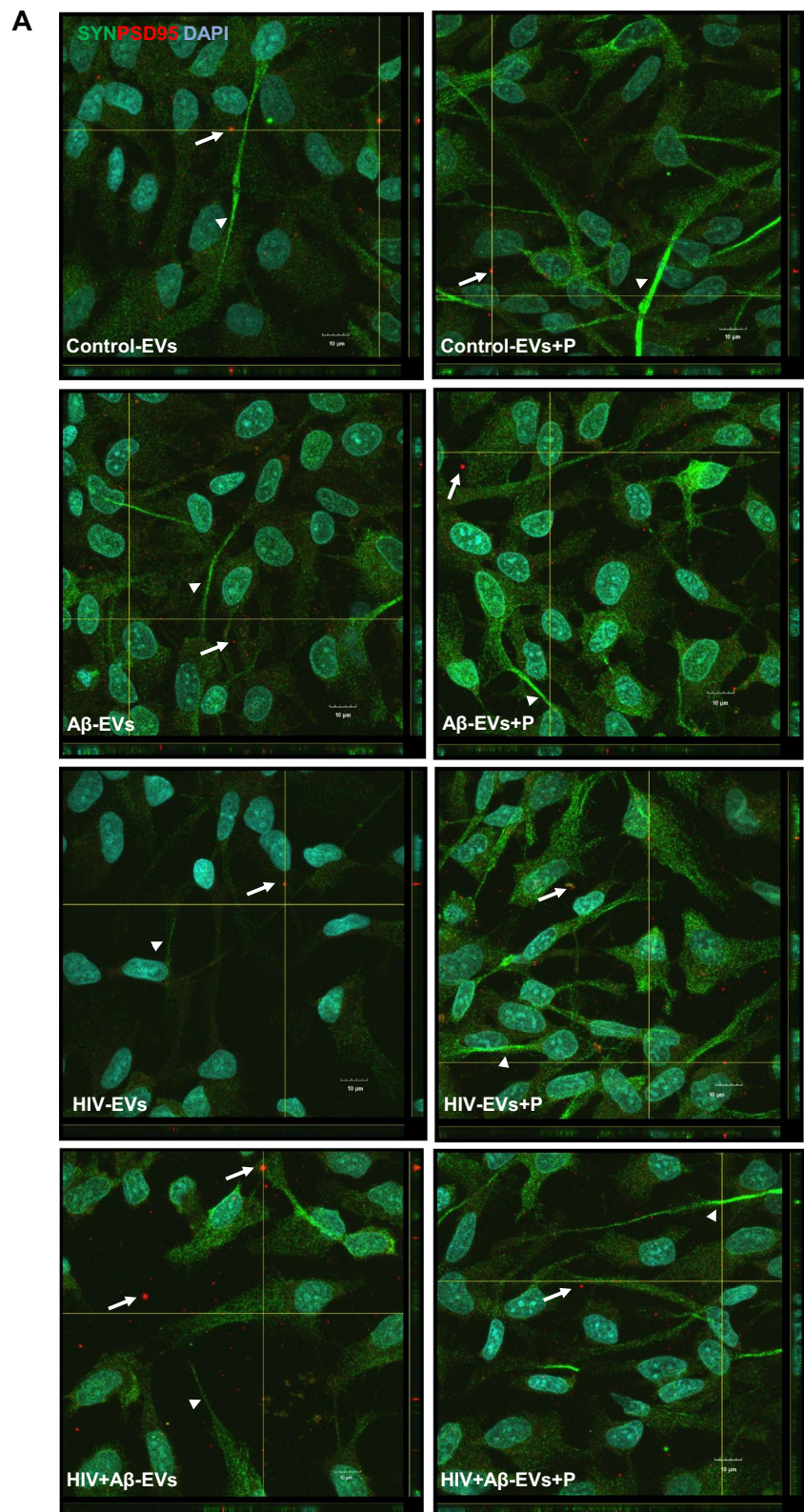
postsynaptic density protein 95 (PSD95) and synaptophysin [44, 45]. In addition, Serpine-1 was demonstrated to be neuroprotective against NMDA-induced neuronal death [46]. In contrast, Serpine-1 was reported to inhibit tPA-mediated neurite outgrowth in NPCs [47].

In order to get a better understanding of the involvement of Serpine-1 in synaptic protein expression in differentiating NPCs, we evaluated PSD95 and synaptophysin expression pattern in NPCs exposed to HBMEC-derived EVs in the presence or absence of PAI039. As illustrated on Fig. 7A and in Supplementary Fig. 2, synaptophysin exhibited a fine punctate immunoreactivity pattern (green fluorescence) in all NPC groups with strikingly stronger fluorescence in segments of the developing NPC projections (arrow heads). PSD95 showed a similar punctate immunoreactivity pattern (red fluorescence). In addition to this fine punctate pattern, PSD95 fluorescence was frequently observed in vesicular structures both inside and outside of NPCs in all groups (arrows) as revealed by z-stacking confocal microscopy (Fig. 7A). This was rarely the case for synaptophysin. It is possible that PSD95 might be also released via EVs, as it was shown before [48], from the developing NPCs to be delivered to the developing synapses.

Quantification of total synaptophysin and PSD95 immunoreactivity from confocal z-stack images showed that both synaptic protein levels significantly decreased in NPCs exposed to HIV-EVs as compared to the Control-EVs treated group. This effect was reversed by PAI039 only for synaptophysin, indicating that PAI039 differentially affected synaptic protein levels overall (Fig. 7B). In addition, total PSD95 level was significantly decreased in NPCs after A β -EV exposure and PAI039 did not block this effect (Fig. 7B, right graph).

Next, we quantified synaptic protein fluorescence intensity in the NPC projections (Figs. 7C–E). Because these were developing NPCs and not mature neurons, their projections were short (Fig. 7C) and their synaptic protein immunofluorescence varied greatly from no signal to very high intensity signal even within the same neurite (Figs. 7A and D). Synaptophysin (green fluorescence) and PSD95 (red fluorescence) showed a similar punctate immunoreactivity pattern in the NPC projections, visible at high magnification (Fig. 7D). Interestingly, there appeared to be little overlap between synaptophysin and PSD95 signals in neurites on these high magnification images. Using an unbiased approach, we highlighted identical rectangular areas randomly on NPC projections visible on brightfield images as depicted on Fig. 7C. Then, we measured synaptophysin (green) and PSD95 (red) fluorescence intensity from the same areas. Quantification of these results are shown on Fig. 7E. Synaptophysin levels significantly increased in the Control-EV + PAI039 group as compared to the Control-EV group and in the HIV-EV + PAI039 group when compared

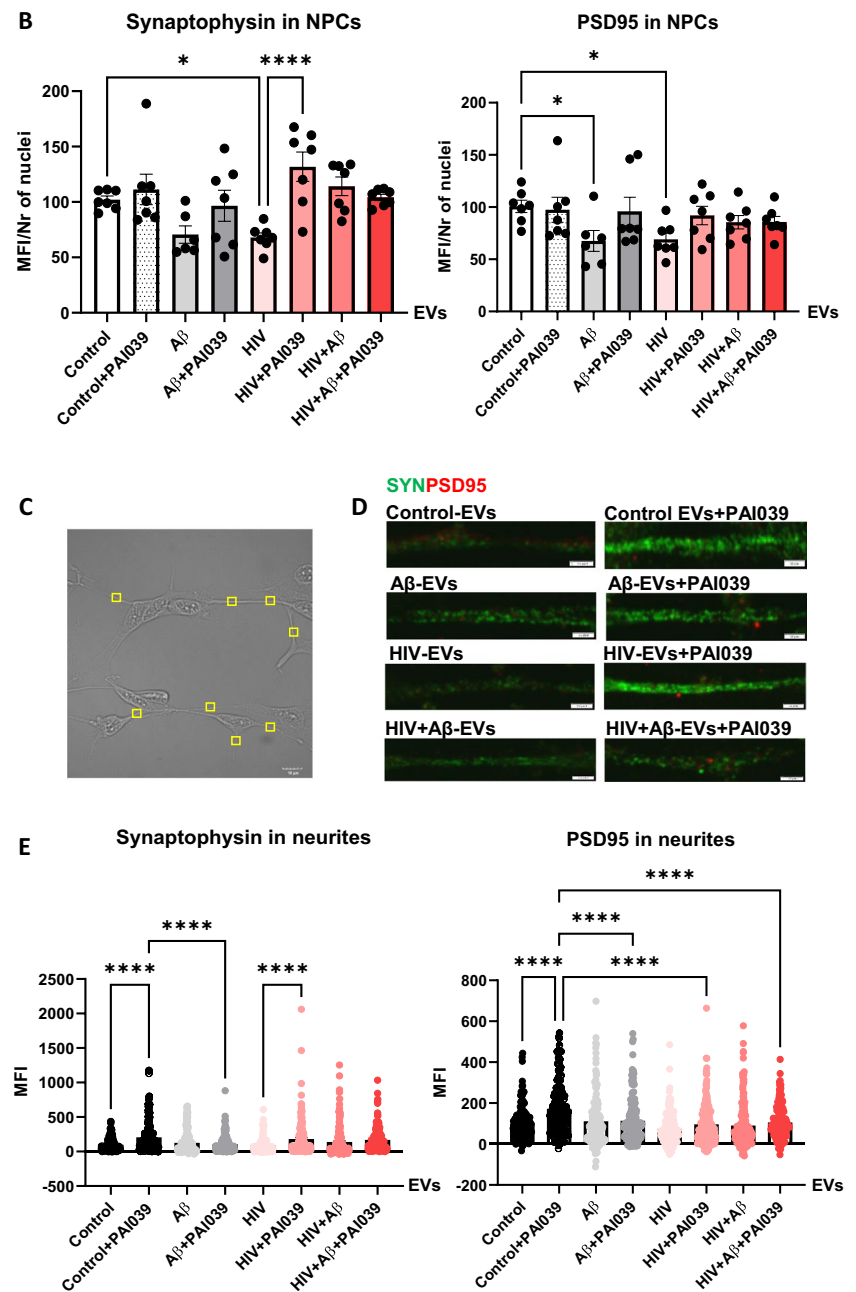
Fig. 7 Impact of HBMEC-derived EVs on synaptic protein expression in NPCs. Non-transfected HBMEC were treated with HIV and/or A β and EVs were isolated as in Fig. 3. Then, human NPCs were exposed to HBMEC-derived EVs for 24 h, with selected cultures additionally treated with 2 μ M PAI039 (P) as in Fig. 6. **A** Synaptophysin (green; arrow heads) and PSD95 (red; arrows) immunoreactivity as imaged by confocal microscopy. DAPI staining (blue) visualizes the NPC nuclei. The combined z-stack images with maximum intensity projection are representative from three experiments. Scale bar: 10 μ m. Individual panels are shown in Supplementary Fig. 2. **B** Total intensity of synaptophysin (left graph) and PSD95 (right graph) immunoreactivity as quantified from the confocal images. Values are mean \pm SEM, $n = 7$. One-, two- and three-way ANOVA with Šídák's multiple comparisons tests. **C** Example of a brightfield confocal image with random identical rectangular areas on NPC projections. **D** Synaptophysin (green) and PSD95 (red) immunoreactivity on representative confocal images of NPC projections. Scale bar: 2 μ m. **E** Intensity of synaptophysin and PSD95 immunoreactivity as quantified from random projection areas. Values are mean \pm SEM, $n = 160$ –180. One-, two- and three-way ANOVA with Šídák's and Tukey's multiple comparisons tests. *Statistically significant at $p < 0.05$, **** $p < 0.0001$



to the HIV-EV only group. The Control-EV + PAI039 group was also significantly higher than the A β -EV + PAI039 group, suggesting that A β -EV treatment may have blocked

the PAI039 effect (Fig. 7E, left graph). Changes in PSD95 fluorescence were partly similar, with PSD95 levels being significantly increased in the Control-EV + PAI039 group

Fig. 7 (continued)



as compared to the Control-EV group. Interestingly, the Control-EV + PAI039 group was also significantly higher than the A β -EV + PAI039, HIV-EV + PAI039 and HIV + A β -EV + PAI039 groups, suggesting that A β and HIV may have blocked the PAI039 effect (Fig. 7E, right graph).

Discussion

EVs are recognized as important contributors to A β pathology [49–55], including elevated A β deposition in HIV-1 infection [20, 22]. We have shown previously that EV-A β

can be transferred to cells of the neurovascular unit, including neural progenitor cells (NPCs) [20]; however, the mechanisms of EV-mediated A β pathology remain elusive. In an effort to gain more insight into this process, we applied proteomics analysis to better characterize the EV protein cargo in the context of HIV-1, and Serpine-1 was identified as a main connecting “hub” on several EV protein–protein interaction maps [23]. This finding is important because Serpine-1 was previously described as a key player in A β pathology [30, 31] and was linked to HIV-1 infection as well [29, 56]. However, the role of EV-associated Serpine-1 in the A β pathology is not well understood. Since Serpine-1

was linked to A β pathology, HIV-1 comorbidities, and was found in EVs, we hypothesized that brain endothelial EV-Serpine-1 can be involved in HIV-1 and/or A β -induced NPC alterations. To the best of our knowledge, there are no reports in the literature on this subject.

The main physiological role of Serpine-1 is inhibition of tPA leading to inhibition of plasmin, which places Serpine-1 as the critical regulator of fibrinolysis pathways [24]. However, this process also affects A β levels because plasmin can degrade both APP and A β . Indeed, elevated levels of Serpine-1 favor a more procoagulant state by decreasing tPA activity, which, in turn, hinders plasminogen conversion to active plasmin leading to diminished A β degradation [28]. In support of this claim, Serpine-1 involvement in A β deposition in AD has been documented [30, 31]. For example, cerebral blood vessels were shown to be Serpine-1 positive in AD transgenic mice overexpressing Tau [57]. Knock-out of Serpine-1 gene or inhibition of Serpine-1 significantly reduced brain A β load in the APP/PS1 AD mouse model. Oral administration of TM5275, a small molecule inhibitor of Serpine-1, increased the activities of tPA, uPA and plasmin, leading to decreased A β levels in the hippocampus and cortex with improved learning and memory functions [31]. Moreover, Serpine-1 inhibition with PAI039 restored A β -induced decreased tPA activity and altered neurovascular coupling. These effects were associated with reduced perivascular amyloid deposition and improved cognition [58]. In related studies, tPA was proposed to protect against elevated A β levels by accelerating A β degradation and inhibition of A β -mediated neurodegeneration [28]. Overall, the literature data provide evidence that inhibition of Serpine-1 and restoration of tPA activity could be of substantial therapeutic value in AD.

Interestingly, platelets are the main source of A β in the blood [59] establishing a strong relationship between coagulation and amyloid pathology. However, A β in peripheral blood can also originate from the brain being transported across the BBB by transporters, such as the lipoprotein receptor related protein-1 (LRP1). LRP1 may also be responsible for the removal of the vast majority of A β from the circulation via the liver route [16]. In addition, neuronal EVs were demonstrated to clear A β in the brain as intracerebral infusion of these EVs resulted in a decrease in A β levels and amyloid deposition in the brains of APP transgenic mice [54]. It is possible that these EVs may also carry A β from the brain across the BBB into the circulation.

Regarding a link to HIV infection, antiretroviral therapy (ART) was shown to affect plasma Serpine-1 levels, which appeared to be a marker for HIV-1 related comorbidities. For instance, HIV-1 infected patients on protease inhibitors had higher plasma Serpine-1 levels [56]. Moreover, high plasma levels of Serpine-1 were associated with high risk of myocardial infarction in HIV-1 infected people [29].

Elevated expression of Serpine-1 was proposed to be one of the mechanisms of HIV-1 protein Tat-induced inflammation in vascular cells [60], and synthetic Tat-derived peptides were demonstrated to increase production of Serpine-1 in human umbilical vascular endothelial cells [61]. In addition, monocytes from asymptomatic HIV positive viremic donors were characterized by significantly increased Serpine-1 protein levels as compared to HIV negative donors [62]. It is of note that HIV-1 did not appear to increase serum total A β levels in the circulation in patients less than 60 years of age when compared to the uninfected controls. In addition, no correlation was found between A β levels in the serum and CSF [63].

Earlier reports described the presence of Serpine-1 in microparticles in the blood [64], which might have been EVs as several recent publications reported its presence in EVs. For instance, proteomic analysis from trophoblast-derived EVs identified the presence of Serpine-1 [65]. EVs from nasal olfactory mucosa mesenchymal stem cells and EVs derived from ascites cell cultures also contained Serpine-1 [66, 67]. In line with these reports, we have confirmed that primary human brain endothelial cell-derived EVs (HBMEC-EVs) contained Serpine-1 [23] implicating the brain endothelium and the BBB as important contributors to the Serpine-1 pool. In experiments in which HBMEC were double-transfected with Serpine-1 GFP and CD63 RFP, we found that these cells concentrated and released Serpine-1 via EVs (Figs. 1 and 2). Interestingly, the isolated Serpine-1-positive EVs were rarely positive for CD63 RFP (Fig. 2B), suggesting that different sets of EVs with different cargoes were released by similar pathways from the parent cells, and Serpine-1 may be released in a specific set of EVs. Nevertheless, there is also a possibility that Serpine-1 GFP transfection efficiency was higher than that of CD63 RFP, although the amount of CD63 RFP plasmid DNA used was double of that of Serpine-1 GFP plasmid. However, this was not the case, as the transfection efficiency of the parent cells after 72 h was 90–95% for both constructs. We also observed that Serpine-1 GFP was associated with EVs of different sizes (Fig. 2A), underscoring the importance of evaluating total EVs in the HIV-related A β pathology as opposed to just a particular size-range EVs. Another interesting observation was that almost all Serpine-1 positive EVs were DAPI positive, which demonstrates their nucleic acid cargo (Fig. 2A). Strikingly, the number of CD63 RFP positive EVs was significantly lower in the HIV and HIV + A β groups as compared to the A β only group (Fig. 2B). A similar pattern was observed for the Serpine-1 GFP/CD63 RFP double positive EVs (Fig. 2B), suggesting that HIV-1 exposure of the parent cells alters the protein cargo of the released EVs.

In general, only low levels of Serpine-1 protein were described in the brain and our results are consistent with these observations as only low levels of Serpine-1 were

detected in HBMEC. However, our important results indicate that Serpine-1 was concentrated in EVs, achieving $\sim 30\times$ higher levels when compared to the parent cells (Figs. 3A–B). When comparing the treatment groups, EV-Serpine-1 levels were significantly higher in the HIV-1 and the HIV + A β groups than in controls when normalizing the data to cell culture volume (Fig. 3A). These results were consistent with the activity data (Fig. 3C). On the other hand, there were no differences between the groups when EV-Serpine-1 levels were normalized to EV protein content (Fig. 3A) because exposure to HIV-1 increases the overall EV number as published before [20]. Overall, these observations suggest that one of the main Serpine-1 pools in the brain may be found in endothelial-EVs, pointing to a critical role of the BBB in Serpine-1 related brain pathologies such as AD or stroke or even systemic pathologies as BBB-EVs may also be released into the peripheral circulation. Moreover, higher Serpine-1 levels in the HIV groups implicate EVs to creating a pro-coagulant environment in the vicinity of the BBB of HIV-1-infected brains. These observations are in line with the report that HIV-1 infection was associated with a more pro-coagulant state associated with high Serpine-1 levels in the plasma possibly increasing the risk of myocardial infarction [29]. These findings are also consistent with tPA activity being significantly reduced in the EV groups originating from HIV-1 and/or A β -treated HBMEC as compared to control EVs (Fig. 3D). Indeed, a decrease in tPA activity was observed to be consistently associated with elevated amyloid deposition in the brain [68].

One of the main biological functions of EVs is intercellular communication, which is executed by transferring cargo between different cells and cell types. We have shown before that endothelial-derived EVs can transfer A β to NPCs affecting their neurogenesis [22]. Knowing that Serpine-1 levels are concentrated in EVs, we next evaluated if EVs can serve as carriers to deliver Serpine-1 from endothelial cells to NPCs and what are the outcomes of this process. The rationale for these experiments is the fact that a large pool of NPCs is in the neurogenic niches of the perivascular space in direct proximity to the brain endothelium [42]. We confirmed that Serpine-1 could be transferred to NPCs via EVs (Fig. 4). Moreover, our results indicated that Serpine-1 is transferred together with A β cargo, potentially impacting the A β fate in the acceptor NPCs. Consistent with previously reported results, this process appeared to be enhanced in EVs derived from HIV-1-exposed HBMEC [20]. Nevertheless, Serpine-1 levels in acceptor NPCs were very low, suggesting that secreted EV-Serpine-1 acts on NPCs mostly extracellularly. This mode of action is consistent with the biological impact of Serpine-1 and tPA as both proteins are secreted and act in the extracellular environment.

To evaluate the interactions between Serpine-1 and A β , we examined whether Serpine-1 inhibition can affect EV-A β

transfer to NPCs. Consistent with literature data [69], NPCs appeared to be sensitive to PAI039 toxicity, as 10 μ M caused substantial toxicity (Fig. 5B) and 20 μ M caused massive cell death; therefore, PAI039 was used at 2 μ M. Inhibition of Serpine-1 had an unexpected effect on A β transfer to NPCs, enhancing this process in the HIV + A β -EV + PAI039 group as compared to the HIV + A β -EV group (Fig. 5C). The inhibitor did not affect A β transfer in other groups, suggesting a specific impact of Serpine-1 on the transfer of EV-derived A β cargo in the context of HIV-1. Although this was a unique complex system, where EVs interacted with NPCs, separate observations that PAI039 exposure with EVs alone increased Serpine-1 activity in the HIV groups (Supplementary Fig. 1A) suggest that this phenomenon may have also affected EV-A β transfer.

The role of Serpine-1 in neuronal dysfunction is controversial with conflicting reports in the literature. It was observed that Serpine-1 might be neuroprotective against A β -induced neurotoxicity, preserving neuronal networks, and promoting synaptogenesis by increasing PSD95 and synaptophysin [44, 45]. Similarly, Serpine-1 was demonstrated to be neuroprotective against NMDA-induced neuronal death [46]. On the other hand, Serpine-1 can inhibit neuroprotective impact of tPA, which was shown to control neurite outgrowth in cortical neurons after stroke or in NPCs [47, 70]. This is an important observation, because both A β and HIV pathologies are linked to increased incidents of strokes and administration of tPA is an approved intervention to restore blood flow to brain regions affected by a stroke. By blocking these beneficial effects, Serpine-1 can exert undesirable neurotoxic impact. To address these problems, we evaluated the effects of EVs carrying Serpine-1 and/or A β on the NPC mitochondrial networks, bioenergetics, and synaptic integrity. The results clearly demonstrated altered mitochondrial morphology in NPCs exposed to HBMEC-derived EVs in the HIV + A β -EV treated group, with PAI039 reversing the majority of these alterations (Fig. 6B). The characteristics of these changes related to long mitochondria branches and an increase in total branch length may point to mitochondrial hyperfusion. In support of a role of EVs and HIV-1 in these mitochondrial alterations, EVs from latent HIV-infected T cells were shown to enhance mitochondrial superoxide production, reduce mitochondrial membrane potential, and induce mitochondrial hyperfusion in primary human brain microvascular endothelial cells [71]. In addition, the obtained results were consistent with the observations that A β may cause NPC damage with mitochondrial alterations, which, in turn, may affect their functions [43]. Regarding NPC mitochondrial bioenergetics, changes caused by inhibition of Serpine-1 were minimal and appeared to impair the cells' ability to respond to stress or metabolic demand after A β -EV exposure as demonstrated by a decrease in spare respiratory capacity in this treatment group (Supplementary Fig. 1I).

In order to assess the impact of A β and/or HIV-1 EVs on synaptic integrity, we evaluated the levels of synaptophysin and PSD95 in differentiating NPCs. Synaptophysin expression served as a marker of presynaptic plasticity and synaptogenesis [72]. Loss of synaptophysin was found in AD [73] and learning and memory deficits have been demonstrated in synaptophysin knockout mice [74]. PSD95 is a postsynaptic marker of synaptic integrity and its decrease can also lead to learning and memory impairment [75, 76]. In contrast, up-regulation of PSD-95 was shown to improve memory [77], underscoring the importance of PSD95 in these key brain functions. We detected that even in early developmental stage (namely, three days of differentiation), NPCs expressed a fine punctate immunoreactivity for both synaptophysin and PSD95 in the developing neurite segments (Fig. 7D). In addition, PSD95-positive, but not synaptophysin-positive, fluorescence was often demonstrated in vesicular structures both intra- and extracellularly on z-stacking confocal microscopy images (Fig. 7A), suggesting that it could be secreted via EVs from the developing NPCs. Exposure to EVs derived from HIV-exposed HBMEC decreased both synaptophysin and PSD95 in NPCs, which is consistent with neurotoxicity and aberrant neurogenesis in HIV infected brains [78, 79]. In addition, loss of synaptophysin was observed in HIV-1 infection in humanized mice [80] and exposure to HIV-1 proteins, like Tat and gp120, markedly decreased PSD95 in hippocampal neurons [81–83]. The levels of synaptophysin and PSD95 were also decreased in mice following exposure to HIV-1 Tat and methamphetamine [84]. Therefore, it was important that inhibition of Serpine-1 protected against HIV-EV-induced alterations in synaptophysin (Fig. 7B) and increased both synaptic proteins in the projections (Fig. 7E) of differentiating NPCs. These observations are consistent with the reports advocating for a beneficial effect of Serpine-1 inhibition on synaptic protein expression and neurite development, although EV-Serpine-1 activity increase evoked by PAI039 (Supplementary Fig. 1A) may have also contributed to these effects. Overall, these data may be relevant in the context of HIV-1 associated neurocognitive impairments.

Conclusions

The results of the present study indicate that brain endothelial EVs contain active Serpine-1 cargo, which can be delivered to the recipient NPCs along with A β . These findings represent a novel concept that endothelial-derived EVs constitute a major Serpine-1 pool in the brain, which can create a pro-coagulant environment at the BBB and lead to mitochondrial and synaptic alterations in NPCs. These processes may further contribute to HIV-1 associated neurocognitive disorders (HAND), especially in older brains, which are characterized by elevated A β depositions. Overall, EV-associated Serpine-1 may be an

important player in vascular A β pathology in HIV-infection. This may guide potential therapeutic avenues in an effort to mitigate neuropathology in aging patients with HIV-1.

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Data Availability All source data supporting the findings of this manuscript are available from the corresponding authors upon request.

Declarations

Ethics Approval and Consent to Participate Not applicable.

Consent for Publication Not applicable.

Competing Interests The authors declare that they have no competing interests.

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References

- Xu J, Ikezu T (2009) The comorbidity of HIV-associated neurocognitive disorders and Alzheimer's disease: a foreseeable medical challenge in post-HAART era. *J Neuroimmune Pharmacol* 4(2):200–212
- Soontornniyomkij V et al (2012) Cerebral beta-amyloid deposition predicts HIV-associated neurocognitive disorders in APOE epsilon4 carriers. *AIDS* 26(18):2327–2335
- Green DA et al (2005) Brain deposition of beta-amyloid is a common pathologic feature in HIV positive patients. *AIDS* 19(4):407–411

4. Steinbrink F et al (2013) Cognitive impairment in HIV infection is associated with MRI and CSF pattern of neurodegeneration. *Eur J Neurol* 20(3):420–428
5. Esiri MM, Biddolph SC, Morris CS (1998) Prevalence of Alzheimer plaques in AIDS. *J Neurol Neurosurg Psychiatry* 65(1):29–33
6. Rempel HC, Pulliam L (2005) HIV-1 Tat inhibits neprilysin and elevates amyloid beta. *AIDS* 19(2):127–135
7. Brew BJ et al (2009) Neurodegeneration and ageing in the HAART era. *J Neuroimmune Pharmacol* 4(2):163–174
8. Achim CL et al (2009) Increased accumulation of intraneuronal amyloid beta in HIV-infected patients. *J Neuroimmune Pharmacol* 4(2):190–199
9. Pulliam L (2009) HIV regulation of amyloid beta production. *J Neuroimmune Pharmacol* 4(2):213–217
10. Morgello S et al (2021) HIV disease duration, but not active brain infection, predicts cortical amyloid beta deposition. *AIDS* 35(9):1403–1412
11. Gross AM et al (2016) Methylome-wide Analysis of Chronic HIV Infection Reveals Five-Year Increase in Biological Age and Epigenetic Targeting of HLA. *Mol Cell* 62(2):157–168
12. Lew BJ et al (2021) Reductions in Gray Matter Linked to Epigenetic HIV-Associated Accelerated Aging. *Cereb Cortex* 31(8):3752–3763
13. Cohen J, Torres C (2017) HIV-associated cellular senescence: A contributor to accelerated aging. *Ageing Res Rev* 36:117–124
14. Montano M et al (2022) Biological ageing with HIV infection: evaluating the geroscience hypothesis. *Lancet Healthy Longev* 3(3):e194–e205
15. Schank M et al (2021) The Impact of HIV- and ART-Induced Mitochondrial Dysfunction in Cellular Senescence and Aging. *Cells* 10(1):174
16. Deane R, Zlokovic BV (2007) Role of the blood-brain barrier in the pathogenesis of Alzheimer's disease. *Curr Alzheimer Res* 4(2):191–197
17. Ramanathan A et al (2015) Impaired vascular-mediated clearance of brain amyloid beta in Alzheimer's disease: the role, regulation and restoration of LRP1. *Front Aging Neurosci* 7:136
18. Deane R et al (2003) RAGE mediates amyloid-beta peptide transport across the blood-brain barrier and accumulation in brain. *Nat Med* 9(7):907–913
19. Andras IE et al (2010) HIV-1-induced amyloid beta accumulation in brain endothelial cells is attenuated by simvastatin. *Mol Cell Neurosci* 43(2):232–243
20. Andras IE et al (2017) Extracellular vesicles of the blood-brain barrier: Role in the HIV-1 associated amyloid beta pathology. *Mol Cell Neurosci* 79:12–22
21. Dardet JP et al (2022) Overcoming Blood Brain Barrier Resistance: Implications for Extracellular Vesicle-Mediated Drug Brain Delivery. *Front Drug Deliv* 2:855017. <https://doi.org/10.3389/fddev.2022.855017>
22. Andras IE et al (2020) Extracellular vesicle-mediated amyloid transfer to neural progenitor cells: implications for RAGE and HIV infection. *Mol Brain* 13(1):21
23. Andras IE, Sewell BB, Toborek M (2020) HIV-1 and Amyloid Beta Remodel Proteome of Brain Endothelial Extracellular Vesicles. *Int J Mol Sci* 21(8):2741
24. Urano T et al (2019) Recognition of Plasminogen Activator Inhibitor Type 1 as the Primary Regulator of Fibrinolysis. *Curr Drug Targets* 20(16):1695–1701
25. Benjamin LA et al (2012) HIV infection and stroke: current perspectives and future directions. *Lancet Neurol* 11(10):878–890
26. Baker JV et al (2013) HIV replication alters the composition of extrinsic pathway coagulation factors and increases thrombin generation. *J Am Heart Assoc* 2(4):e000264
27. Ismael S et al (2020) HIV Associated Risk Factors for Ischemic Stroke and Future Perspectives. *Int J Mol Sci* 21(15):5306
28. Melchor JP, Pawlak R, Strickland S (2003) The tissue plasminogen activator-plasminogen proteolytic cascade accelerates amyloid-beta (A β) degradation and inhibits A β -induced neurodegeneration. *J Neurosci* 23(26):8867–8871
29. Knudsen A et al (2014) Plasma plasminogen activator inhibitor-1 predicts myocardial infarction in HIV-1-infected individuals. *AIDS* 28(8):1171–1179
30. Liu RM et al (2011) Knockout of plasminogen activator inhibitor 1 gene reduces amyloid beta peptide burden in a mouse model of Alzheimer's disease. *Neurobiol Aging* 32(6):1079–1089
31. Akhter H et al (2018) A Small Molecule Inhibitor of Plasminogen Activator Inhibitor-1 Reduces Brain Amyloid-beta Load and Improves Memory in an Animal Model of Alzheimer's Disease. *J Alzheimers Dis* 64(2):447–457
32. Weksler BB et al (2005) Blood-brain barrier-specific properties of a human adult brain endothelial cell line. *FASEB J* 19(13):1872–1874
33. Paris D et al (2002) Pro-inflammatory effect of freshly solubilized beta-amyloid peptides in the brain. *Prostaglandins Other Lipid Mediat* 70(1–2):1–12
34. Elokda H et al (2004) Tiplaxtinin, a novel, orally efficacious inhibitor of plasminogen activator inhibitor-1: design, synthesis, and preclinical characterization. *J Med Chem* 47(14):3491–3494
35. Gerenu G et al (2017) Modulation of BDNF cleavage by plasminogen-activator inhibitor-1 contributes to Alzheimer's neuropathology and cognitive deficits. *Biochim Biophys Acta Mol Basis Dis* 1863(4):991–1001
36. Gorlatova NV et al (2007) Mechanism of inactivation of plasminogen activator inhibitor-1 by a small molecule inhibitor. *J Biol Chem* 282(12):9288–9296
37. Valente AJ et al (2017) A simple ImageJ macro tool for analyzing mitochondrial network morphology in mammalian cell culture. *Acta Histochem* 119(3):315–326
38. Osborne OM, Kowalczyk JM, Pierre Louis KD, Daftari MT, Colbert BM, Naranjo O, Torices S, András IE, Dykxhoorn DM, Toborek M (2022) Brain Endothelium Derived Extracellular Vesicles Containing Amyloid-Beta Induce Mitochondrial Alterations in Neural Progenitor Cells. *Extracell Vesicles Circ Nucl Acids* 3(4):340–362
39. Kuszczak MA et al (2014) 1MeTIQ provides protection against A β -induced reduction of surface expression of synaptic proteins and inhibits H $_2$ O $_2$ -induced oxidative stress in primary hippocampal neurons. *Neurotox Res* 25(4):348–357
40. Abbott NJ, Friedman A (2012) Overview and introduction: the blood-brain barrier in health and disease. *Epilepsia* 53:1–6
41. Shen Q et al (2004) Endothelial cells stimulate self-renewal and expand neurogenesis of neural stem cells. *Science* 304(5675):1338–1340
42. Shen Q et al (2008) Adult SVZ stem cells lie in a vascular niche: a quantitative analysis of niche cell-cell interactions. *Cell Stem Cell* 3(3):289–300
43. Meshrkey F et al (2021) Quantitative analysis of mitochondrial morphologies in human induced pluripotent stem cells for Leigh syndrome. *Stem Cell Res* 57:102572
44. Cho H et al (2013) Plasminogen activator inhibitor-1 promotes synaptogenesis and protects against A β (1–42)-induced neurotoxicity in primary cultured hippocampal neurons. *Int J Neurosci* 123(1):42–49
45. Soeda S et al (2008) Anti-apoptotic roles of plasminogen activator inhibitor-1 as a neurotrophic factor in the central nervous system. *Thromb Haemost* 100(6):1014–1020
46. Docagne F et al (2002) Smad3-dependent induction of plasminogen activator inhibitor-1 in astrocytes mediates neuroprotective activity of transforming growth factor-beta 1 against NMDA-induced necrosis. *Mol Cell Neurosci* 21(4):634–644

47. Lee SH et al (2014) tPA regulates neurite outgrowth by phosphorylation of LRP5/6 in neural progenitor cells. *Mol Neurobiol* 49(1):199–215
48. Hughes AN, Appel B (2019) Oligodendrocytes express synaptic proteins that modulate myelin sheath formation. *Nat Commun* 10(1):4125
49. Vella LJ et al (2008) The role of exosomes in the processing of proteins associated with neurodegenerative diseases. *Eur Biophys J* 37(3):323–332
50. Kalani A, Tyagi A, Tyagi N (2014) Exosomes: mediators of neurodegeneration, neuroprotection and therapeutics. *Mol Neurobiol* 49(1):590–600
51. Gupta A, Pulliam L (2014) Exosomes as mediators of neuroinflammation. *J Neuroinflammation* 11:68
52. An K et al (2013) Exosomes neutralize synaptic-plasticity-disrupting activity of A β assemblies in vivo. *Mol Brain* 6:47
53. Yuyama K et al (2014) Decreased amyloid-beta pathologies by intracerebral loading of glycosphingolipid-enriched exosomes in Alzheimer model mice. *J Biol Chem* 289(35):24488–24498
54. Yuyama K et al (2015) A potential function for neuronal exosomes: sequestering intracerebral amyloid-beta peptide. *FEBS Lett* 589(1):84–88
55. Dinkins MB et al (2015) The 5XFAD Mouse Model of Alzheimer's Disease Exhibits an Age-Dependent Increase in Anti-Ceramide IgG and Exogenous Administration of Ceramide Further Increases Anti-Ceramide Titers and Amyloid Plaque Burden. *J Alzheimers Dis* 46(1):55–61
56. Koppel K et al (2002) Hypofibrinolytic state in HIV-1-infected patients treated with protease inhibitor-containing highly active antiretroviral therapy. *J Acquir Immune Defic Syndr* 29(5):441–449
57. Bennett RE et al (2018) Tau induces blood vessel abnormalities and angiogenesis-related gene expression in P301L transgenic mice and human Alzheimer's disease. *Proc Natl Acad Sci U S A* 115(6):E1289–E1298
58. Park L et al (2020) tPA Deficiency Underlies Neurovascular Coupling Dysfunction by Amyloid-beta. *J Neurosci* 40(42):8160–8173
59. Chen M et al (1995) Platelets Are the Primary Source of Amyloid β -Peptide in Human Blood. *Biochem Biophys Res Comm* 213(1):96–103
60. Albin A et al (1996) The angiogenesis induced by HIV-1 tat protein is mediated by the Flk-1/KDR receptor on vascular endothelial cells. *Nat Med* 2(12):1371–1375
61. Ismail M et al (2006) Identification of HIV-1 Tat peptides for future therapeutic angiogenesis. *Eur J Haematol* 77(2):157–165
62. Gekonge B et al (2012) Retinoblastoma protein induction by HIV viremia or CCR5 in monocytes exposed to HIV-1 mediates protection from activation-induced apoptosis: ex vivo and in vitro study. *J Leukoc Biol* 92(2):397–405
63. de Almeida SM et al (2020) Blood amyloid- β protein isoforms are affected by HIV-1 in a subtype-dependent pattern. *J Neurovirol* 26(1):3–13
64. Huisse MG et al (2009) Microparticle-linked tissue factor activity and increased thrombin activity play a potential role in fibrinolysis failure in ST-segment elevation myocardial infarction. *Thromb Haemost* 101(4):734–740
65. Familiari M et al (2017) Placenta-derived extracellular vesicles: their cargo and possible functions. *Reprod Fertil Dev* 29(3):433–447
66. Xun C et al (2020) Insight into the proteomic profiling of exosomes secreted by human OM-MSCs reveals a new potential therapy. *Biomed Pharmacother* 131:110584
67. Bortot B et al (2021) Small extracellular vesicles from malignant ascites of patients with advanced ovarian cancer provide insights into the dynamics of the extracellular matrix. *Mol Oncol* 15(12):3596–3614
68. Bi Oh, S., et al (2015) Impacts of aging and amyloid-beta deposition on plasminogen activators and plasminogen activator inhibitor-1 in the Tg2576 mouse model of Alzheimer's disease. *Brain Res* 1597:159–167
69. Daniel AE et al (2015) Plasminogen Activator Inhibitor-1 Controls Vascular Integrity by Regulating VE-Cadherin Trafficking. *PLoS ONE* 10(12):e0145684
70. Xin H et al (2010) Increasing tPA activity in astrocytes induced by multipotent mesenchymal stromal cells facilitate neurite outgrowth after stroke in the mouse. *PLoS ONE* 5(2):e9027
71. Chandra PK et al (2021) Latent HIV-Exosomes Induce Mitochondrial Hyperfusion Due to Loss of Phosphorylated Dynamin-Related Protein 1 in Brain Endothelium. *Mol Neurobiol* 58(6):2974–2989
72. Ujike H et al (2002) Gene expression related to synaptogenesis, neurogenesis, and MAP kinase in behavioral sensitization to psychostimulants. *Ann N Y Acad Sci* 965:55–67
73. Sze CI et al (1997) Loss of the presynaptic vesicle protein synaptophysin in hippocampus correlates with cognitive decline in Alzheimer disease. *J Neuropathol Exp Neurol* 56(8):933–944
74. Schmitt U et al (2009) Detection of behavioral alterations and learning deficits in mice lacking synaptophysin. *Neuroscience* 162(2):234–243
75. Fitzgerald PJ et al (2015) Durable fear memories require PSD-95. *Mol Psychiatry* 20(7):901–912
76. Migaud M et al (1998) Enhanced long-term potentiation and impaired learning in mice with mutant postsynaptic density-95 protein. *Nature* 396(6710):433–439
77. Li C et al (2004) Estrogen alters hippocampal dendritic spine shape and enhances synaptic protein immunoreactivity and spatial memory in female mice. *Proc Natl Acad Sci U S A* 101(7):2185–2190
78. Hahn YK et al (2012) HIV-1 alters neural and glial progenitor cell dynamics in the central nervous system: coordinated response to opiates during maturation. *Glia* 60(12):1871–1887
79. Lee MH et al (2011) Rescue of adult hippocampal neurogenesis in a mouse model of HIV neurologic disease. *Neurobiol Dis* 41(3):678–687
80. Dash PK et al (2011) Loss of neuronal integrity during progressive HIV-1 infection of humanized mice. *J Neurosci* 31(9):3148–3157
81. Kim HJ, Martemyanov KA, Thayer SA (2008) Human immunodeficiency virus protein Tat induces synapse loss via a reversible process that is distinct from cell death. *J Neurosci* 28(48):12604–12613
82. Kim HJ, Shin AH, Thayer SA (2011) Activation of cannabinoid type 2 receptors inhibits HIV-1 envelope glycoprotein gp120-induced synapse loss. *Mol Pharmacol* 80(3):357–366
83. Zhang X, Thayer SA (2018) Monoacylglycerol lipase inhibitor JZL184 prevents HIV-1 gp120-induced synapse loss by altering endocannabinoid signaling. *Neuropharmacology* 128:269–281
84. Nookala AR et al (2018) Methamphetamine augment HIV-1 Tat mediated memory deficits by altering the expression of synaptic proteins and neurotrophic factors. *Brain Behav Immun* 71:37–51

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