Coronin 1A Expression in Human Astroglia, its Function in Physiology and Astrogliosis in HIV-1 Neuropathogenesis

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Research

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Abstract

In most neurodegenerative disorders, including neuroAIDS, reactive astroglia are detrimental to the neuronal population. Calcium and its downstream regulators play a central role in mediating glial activation. Coronin 1A, an acting binding protein, majorly reported in cells of hematopoietic origin, regulates cell activity in a calcium-dependent manner, but its role in astroglial physiology and astrogliosis is largely unknown. Using a well-characterized primary culture of human astroglia and neurons, we explored the roles of Coronin 1A in astroglia physiology and the mechanisms by which it facilitates reactive astrogliosis. In this study, we report for the first time, that human primary astroglia express Coronin 1A, and it plays activity-dependent roles in events such as PLCγ1 phosphorylation followed by Calcium mobilization from the intracellular stores. HIV-1 Tat, a potent neurotoxicant that induces astrogliosis, enhances the expression of Coronin 1A, apart from affecting GFAP and pro-inflammatory molecules. Downregulation of Coronin 1A ameliorated the HIV-1 Tat-induced deleterious effects of reactive astroglia, measured as enhanced GFAP expression and release of IL-6, and Glutamate and thus reduced glia-mediated neurodegeneration. Our findings also suggest that out of a pool of dysregulated miRNAs studied by us, hsa-miR-92b-5p regulates Coronin 1A expression which further facilitates reactive astrogliosis under the effect of HIV-1 Tat. These findings highlight the novel roles of Coronin 1A in regulating the astroglial physiology and astrogliosis observed in HIV-1 neuropathogenesis.

Introduction

Extensive studies on glial cells conducted over the past 30 years, strongly advocate that astroglia are more than mere supporting cells as they perform many essential functions ranging from neurodevelopment to synaptic plasticity (reviewed in [1,2]). Also, growing evidence point towards the fact that astroglia undergo morphological (including hypertrophy) and molecular changes in various inflammatory or pathological conditions. These changes can further affect the neuronal population. Astrogliosis, encompasses all such changes including the increase in protein expression of intermediate filaments such as glial fibrillary acidic protein (GFAP) and vimentin followed by the enhanced release of numerous factors, can be beneficial or detrimental for the neurons, based on the context or the neuropathology, is also regulated by a number of factors [3,4]. Many of the released factors can also serve as the mediators of the innate immune system [5].

Astroglia play a significant role in human immunodeficiency virus (HIV)-1 neuropathogenesis [6-8]. In fact, in the post-HAART era, the glia-mediated indirect neuronal death is prominent over the direct death of the neurons caused by the virus or the viral proteins. Astroglia contribute significantly to the indirect neuronal death and thus amplify the toxic effects brought about by the activated and/or infected microglia. Up to 19% of astroglia are found to be positive for HIV-1 DNA in subjects with HIV-associated dementia [9]. The extent of HIV-1 infection in astrocytes is correlated with the severity of the neurological problems associated with AIDS patients. HIV-1 infection in astrocytes is non-productive due to the reduced translation of Gag, Env, and Nef mRNAs, whereas the translation of, transactivator of transcription (Tat) mRNA remains unaltered. Studies in mice model show that HIV-1 Tat is sufficient to
induce progressive cognitive abnormalities such as impaired learning and memory deficits [10-12]. Astroglial activation, induced either by HIV-1 infection, the viral proteins, or factors released by other infected cell types, is a key factor in mediating neuronal loss by various mechanisms. Hence, there is an urgent need to better understand the mechanisms which make this supporting cell type reactive and detrimental in the context of neuroAIDS.

Calcium is an important regulator of astroglial activity. Calcium signaling in astroglia once doubted to be significant to bring forth changes in neuronal functions, is now well known to be significantly affecting neuronal physiology [13]. Astroglial [Ca^{2+}]_i flux triggered by neurotransmitters leads to the release of gliotransmitters such as glutamate [14], D-serine [15], and adenosine triphosphate (ATP) [16], which after binding to their receptors can cause neuronal excitation [17]. HIV-1 Tat induces Ca^{2+} mobilization in astroglia from the extracellular milieu or the intracellular stores. These events dysregulate the [Ca^{2+}]_i and facilitate altered cell physiology and neuronal death [18].

We investigated the role of coronin 1A in modulating calcium signaling and astroglial activation in the context of neuroAIDS. Coronin 1A belongs to the coronin family of proteins. The presence of WD repeats in the N-terminal is the hallmark of this family. Coronin 1A is encoded on chromosome 7 in mouse and 16 in humans. This 57 kDa protein is expressed majorly by cells of hematopoietic origin and also in nervous tissue [19-22]. By mediating outside to inside signal transduction, it regulates a number of processes such as phagocytosis in various cell types, and activation of T cells [23-25]. The expression and functions of coronin 1A in different brain cell types are still underexplored and await detailed molecular investigations.

In this study, we delineated the activity-dependent roles of coronin 1A and its involvement in mediating astroglial activation in the presence of neuroinflammatory agent such as HIV-1 Tat. Our experiments were focused to investigate the mechanisms of coronin 1A regulations and how it contributes to the enhanced activity of astroglia which leads to the indirect neuronal toxicity observed in HIV-1 neuropathogenesis. This study explores the activity-dependent roles of coronin 1A in cells of non-hematopoietic origin and highlights its role in astrogliosis observed in neuroAIDS.

**Materials And Methods**

**Human Fetal Brain-Derived Neuronal Precursor Cell Culture**

Human brain tissues collected from elective abortions with the informed consent of mothers, were processed as per the protocols laid down by the Institutional Human Ethics and Stem Cell Research Committee under strict compliance with the guidelines of ICMR, India. Neural Precursor Cells (NPCs) were derived from the telencephalon region of the aborted fetuses of age 10-14 weeks. NPCs were cultured on poly-D-lysine (Sigma-Aldrich, Missouri, USA) coated flasks in Neurobasal Media (Invitrogen, California, USA) supplemented with N2 supplement (Invitrogen, USA), Neural Survival Factor-1 (Lonza, Charles City, IA), 20 ng/mL EGF (Sigma-Aldrich, USA) and 25 ng/mL bFGF (Sigma-Aldrich, USA). NPCs were further
assessed for the expression of markers such as Nestin and SOX2 and 99% of the cells were found to be positive. They were also assessed for their ability to form neurospheres and differentiate into Astrocytes and Neurons.

**Human Primary Astroglia and Neuronal Culture**

NPCs were differentiated into astroglia by replacing the media with Eagle's Minimal Essential Media (MEM) (Sigma-Aldrich, USA) containing 10% Fetal Bovine Serum (Gibco, California, USA). These cells were cultured for 3 weeks, with half media changes on alternate days, after which they were assessed for the expression of GFAP and Vimentin, for which more than 95% of cells were found immunopositive (Supporting Information Fig. S1). To differentiate NPCs into neurons, EGF and FGF from NPC media were replaced with 10 ng/mL BDNF (Peptrotech, New Jersey, USA) and 10 ng/mL PDGF-AB (Peptrotech, New Jersey, USA). After maintaining them for 3 weeks, with half media changes on alternate days, they were assessed for expression of Tuj-1 and MAP2, for which more than 95% of cells were immunopositive.

**[^Ca^{2+}]_i measurement / Live cell imaging**

0.2 million cells were seeded in 35-mm dishes and incubated with 2.5 mM Fluo-4-AM (Invitrogen, USA) in HEPES buffer at 37 °C for 30 min. After 3 washes, the dish was mounted on the stage for live-cell imaging using a spinning disc confocal microscope (Zeiss, Oberkochen, Germany) 10 x objective (Zeiss, Germany) was used for the measurement of fluorescent intensity as a measure of the change in[^Ca^{2+}]_i. Baseline fluorescence intensity was recorded for the initial 1 minute, thereafter, 10 mM ATP (Sigma-Aldrich, USA) was added to the static bath followed by 5 min of recording. Change in fluorescence intensity was calculated by subtracting fluorescence at t=0 from all the fluorescence intensity values. ZEN software (Zeiss, Germany) was used for data acquisition and analysis. Areas having a significant number of cells (≥50) were randomly selected for the recordings after which individual cells were marked as the region of interests and fluorescence intensity was measured across time.

**Immunocytochemistry**

Astrocytes were seeded in a 4-well chamber slide (Nunc, Kamstrupvej, Denmark) at a density of 30,000 cells per well. After 24 h, the cells were fixed using 4% paraformaldehyde, washed thrice with 1X PBS, blocked and permeabilized using 4% BSA containing 0.4% Triton-X 100. The cells were then incubated with antibodies, mouse anti-Coronin 1A (Santa Cruz Biotechnology, Texas, USA, 1:100), moues anti- Ki 67 (Millipore, Billerica, USA, 1:1000), mouse anti-Vimentin (Santa Cruz Biotechnology, USA, 1:1000), overnight at 4 °C or Anti-GFAP (Dako, California, USA, 1:1000) for 1 h at 25°C. The cells were then washed thrice with 1X PBS and incubated with appropriate fluorophore tagged secondary antibodies (Invitrogen, USA, 1:1000). The wells were washed thrice with 1X PBS and mounted using Hardset mounting media with DAPI (Vector Labs, Burlingam, USA). For each group, a minimum of five images were captured from random fields using the AxioImager Z1 microscope (Zeiss, Germany).

**ATP Stimulations**
Astroglia knocked down for coronin 1A for 48h were serum-starved for 2-3 h and then stimulated with 10 mM ATP (Sigma-Aldrich, USA). ATP stimulations were given for either 0, 45, 120, and 300 sec or 5, 10, 30, 60, 120 min, after which the cells were harvested using lysis buffer. The lysates were further processed for protein analysis.

**HIV-1 Tat Transfection**

Astroglia cultures with 80% confluence were used for transfections with expression vector pcDNA3.1 expressing full-length HIV-1 Tat B, which was a kind gift from Prof. Udaykumar Ranga, Jawaharlal Nehru Centre for Advanced Scientific Research (JNCASR), India. Transfections were done using Lipofectamine 3000 (Invitrogen, USA) as per the manufacturer’s protocol. After 24 h of transfection, cells were further harvested as per the experimental requisite.

**Small Interfering RNA (siRNA) mediated knockdown of coronin 1A**

Human primary astrocytes were knocked down using siRNA against coronin 1A. 10 nM coronin 1A siRNA (Dharmacon, Colorado, USA) or control scrambled siRNA (Dharmacon, USA) was transfected using lipofectamine RNAiMax (Invitrogen, USA) following manufacturer's protocol. The cells were either harvested after 48h of transfection to check the knockdown efficiency or carried over for further treatments or transfections.

**Western Blotting**

The cells were harvested using lysis buffer, which consisted of 50 mM Tris-HCl (pH 7.5), 150 mM NaCl, 50 mM Sodium fluoride, 1 mM sodium orthovanadate, 1 mM EDTA (pH 8.0), 2% SDS, and protease inhibitor cocktail (Roche, Basel, Switzerland). Estimation of protein concentration was done using 4% copper sulfate and bicinchoninic acid (Sigma-Aldrich, USA). Proteins were resolved on 10-13% SDS-PAGE and transferred onto a nitrocellulose membrane (MDI, Ambala, India). The blots were blocked using 5% skimmed milk in PBS-Tween 20 (PBST) followed by incubation with primary antibodies, mouse anti-Coronin 1A (Santacruz biotech, USA, 1:500), mouse anti-GAPDH (Santacruz biotech, USA, 1:1000), rabbit anti-Phospho-p44/42 MAPK (Cell Signaling Technology, USA, 1:2000), rabbit anti-p44/42 MAPK (Cell Signaling Technology, USA, 1:2000), rabbit anti-Phospho-PLCγ1 (Cell signaling, USA, 1:1000), rabbit anti-PLCγ1 (Cell signaling, USA, 1:1000), rabbit anti-GFAP (Dako, USA, 1:2000). Blots were washed thrice with TBS-Tween 20 (TBST) and then incubated with appropriate HRP labeled secondary antibodies (Vector Labs, USA, 1:4000) for 2 h at room temperature. The blots were washed thrice with TBST and then developed using Chemiluminescent Reagent (Millipore, USA) and were imaged using Gel Documentation System (Uvitech, Cambridge, UK). Densitometry of the protein bands was done using ImageJ software (NIH, Maryland, USA).

**Quantitative Real-Time PCR**

The cells were harvested for RNA isolation using Trizol reagent (Ambion, Texas, USA) following the manufacturer's protocol. The purified RNA was utilized for synthesizing cDNA using High-capacity cDNA
Reverse Transcriptase Kit (Applied Biosystems, California, USA). qPCR was performed using SYBR Green Master Mix (Applied Biosystems, USA) using the following primers: CORO1A forward 5'-CACCAACATCGTCTACCTCTG-3' and reverse 5'-ACTCCTTTGGAACACTGGAG-3', GAPDH forward 5'-CAAGAGCACAAGAGGAAGAGAG-3' and reverse 5'-CTACATGGCAACTGTGAGGAG-3', GFAP forward 5'-ACCTGCGATTTGGAACACCAG-3' and reverse 5'-TAATGACCTCCATCCGCATC-3', Vimentin forward 5'-AAGTCCGCACATTGAGCAGCA-3' and reverse 5'-CTACCAACTTTACAGCTGGGC-3', VCAM1 forward 5'-GGGAGCCGATCAGCTCA-3' and reverse 5'-TCCTGTCTGCTCTCCAGA-3', ICAM1 forward 5'-CCGCAGTCATAATGGGCACT-3' and reverse 5'-GGTTTCATGGGGGTCCCTTT-3', PTGS2 forward 5'-TGTAATGAGTGGGATTTGACC-3' and reverse 5'-TGTTTGGAGTGGGTTCAG-3', NOS2 forward 5'-CTTTGCCTGTATGCTGATGC-3' and reverse 5'-GCCTCTGATTCCCTGTCTATG-3'.

miRNA along with total RNA was isolated using the miRNEASY mini kit (Qiagen, Hilden, Germany) following the manufacturer's protocol. Subsequently, the cDNA was synthesized using the miScript II RT kit (Qiagen, Germany) as per the manufacturer's protocol. The qPCR assay was done using the miScript SYBR Green PCR kit (Qiagen, Germany) following the manufacturer's protocol. The qPCR was done in ViiA7™ (Applied Biosystems, USA). The specificity of the primers was confirmed via analyzing the melt curve.

**IL-6-release measurement by Flow-cytometry**

Astroglia-conditioned-media (ACM) was collected from astroglia knocked down for coronin 1A for 48 h and transfected with HIV-1 Tat for 24 h. The conditioned-media thus collected was then analyzed for the levels of released pro-inflammatory cytokine interleukin-6 (IL-6) by using flow-cytometry based Cytometric bead array (CBA)-human inflammatory cytokines kit (BD Biosciences, San Diego, USA), and following manufacturer’s protocol.

**Glutamate-release measurement**

ACM was collected from astroglia knocked down for coronin 1A for 48 h and transfected with HIV-1 Tat for 24 h. The conditioned media thus collected was analyzed for the levels of released glutamate using an enzyme kinetics-based glutamate determination kit (Sigma-Aldrich, USA), following the manufacturer's protocol.

**Assessment of Neuronal Survival by TUNEL Assay**

ACM was collected from astroglia knocked down for coronin 1A for 48 h and transfected with HIV-1 Tat for 24 h. Neurons at a density of 20,000 cells/well were seeded in Poly-D-Lysine coated 8 well-chambered slides. The next day the cells were treated with 50% ACM and Neuronal Media for 24 h. Cells were then fixed using 4% paraformaldehyde. Washed thrice with 1x PBS, blocked and permeabilized using 4% BSA containing 0.1% Triton-x 100. Cells undergoing apoptosis were detected using in situ Cell Death Detection Kit, TMR red (Roche, Switzerland), following the manufacturer’s protocol. Cells were washed thrice with 1x...
PBS, and the slide was mounted using Hardset mounting media with DAPI (Vector Labs, USA). Five images were captured for each group using the AxiosImager Z1 microscope (Zeiss, Germany).

**miRNA Target Prediction**

miRWalk (http://mirwalk.umm.uni-heidelberg.de/) online prediction tool was used to filter miRNAs targeting 3’ UTR of CORO1A.

**miRNA Mimic and Inhibitor Transfection**

Transfection was done on astroglia, in 80% confluent T25 flasks using RNAi MAX (Invitrogen, USA) as per the manufacturer's protocol. 62.5 nM of Syn-hsa-miR-92b-5p miScript miRNA mimic (Qiagen, Germany) was used for mimic transfection. All-Stars Negative Control siRNA (Qiagen, Germany) was used as mimic control. miRNA inhibition was done using 62.5 nM of miRCURY LNA miRNA inhibitor against has-miR-92b-5p (Qiagen, Germany) and miRCURY LNA miRNA Inhibitor Control (Qiagen, Germany) was used as the inhibitor control. Transfections were done using Opti-MEM media (Invitrogen, USA) for 5 h and replaced with complete MEM. Cells were processed after 48 hours of transfection for RNA and Protein analysis. Effective doses of mimic and inhibitor miRNA were standardized via qPCR and protein studies.

**Cloning of CORO1A 3’ UTR and Luciferase Assay**

The 3’ UTR sequence of coronin 1A, being very short in length (111 bp), was outsourced in the form of 2 single-stranded nucleotide chains having restriction sites for SpeI and HindIII respectively. Both the chains were annealed, digested, and was cloned into pMIR-Report plasmid between SpeI and HindIII restriction sites. The cloned plasmid was then sequenced to confirm the successful insertion.

CORO1A 3’UTR luciferase reporter plasmid was co-transfected with 12.5 nM of mimic-92b-5p into HeLa cells, using Lipofectamine 3000 as per manufacturer's protocol. After 24 h the samples were harvested for luciferase detection, using Luciferase Detection Kit (Promega, Wisconsin, USA) following the manufacturer’s protocol, in Tecan SPARK multiplate reader (Tecan, USA). The readings were normalized using total protein content.

**Statistical Analysis**

Experiments of all kinds were performed independently three to five times; Student’s t-test was used to assess statistical significance between control and experimental groups. P values <0.05 were considered statistically significant. * represents p<0.05, **p<0.005, and ***p<0.0005.

**Results**

**Human Neural Progenitors, Astroglia, and Neurons express coronin 1A**

Coronin 1A expression has been reported majorly in cells of hematopoietic origin [20,23,26] and neurons [22] but its expression in astroglia was not explored till this study. In this regard, human brain cells were
harvested to assess the protein levels of coronin 1A through western blotting. Jurkat T cells were taken as a positive control for coronin 1A expression. We found coronin 1A expression in all the brain cell types viz NPCs, astroglia, and neurons (Fig. 1a). The levels of the GAPDH protein varied in different lanes, this might be because of its differential expression across cell types assessed. Also, to assess differentiating astroglia for coronin 1A expression, cells were harvested at 1-, 2-, 3-weeks of differentiation, and it was found that cells at all the 3 stages showed decent expression (Fig. 1b). Additionally, immunocytochemistry was performed which confirmed coronin 1A (red) localization in the cells along with a marker for astroglia, GFAP (green) (Fig. 1c).

Coronin 1A is dispensable in resting-state human astroglia

The functional roles of coronin 1A in astroglia were unexplored, therefore, to study the role of coronin 1A in astroglia, we carried out a loss of function experiments on human astroglia. We performed siRNA mediated knockdown and checked the knockdown efficiency after 48 h; the siRNA was found to reduce the mRNA levels by 83%±4% (p<0.0005) and protein levels by 41%±2% (p<0.0005) (Fig. 2a). To check if the coronin 1A knockdown results in astroglial cell death, we performed a live-dead cell assay on coronin 1A siRNA transfected astroglia. We found that there was no significant difference in the live-dead cell proportions in the siRNA transfected group compared to the control scrambled siRNA transfected group (Fig. S2a). Mature astroglia are low-proliferating cells though, but whatever the rate of proliferation be, was found to be unaffected by coronin 1A knockdown, as assessed through immunocytochemistry using Ki-67 protein as a marker for proliferation (Fig. S2b). Further, to assess the role of coronin 1A in resting-state cell physiology, if any, we checked the expression of intermediate filament proteins such as GFAP and Vimentin. It was found that downregulating coronin 1A neither affected GFAP (although mRNA significantly affected) (Fig. 2b) nor vimentin protein expression (Fig. 2c). These results signify that coronin 1A is dispensable in resting-state astroglia and hence prompted us to explore if its functions are activity-dependent.

Coronin 1A facilitates Calcium mobilization and Ca\(^{2+}\) dependent pathways in stimulated human astroglia

Ca\(^{2+}\) mobilization upon stimulation with a ligand is a good measure of the astroglia activity and can be correlated with the effects of astroglial activity on neuronal population [13,27]. ATP is a well-known and widely used gliotransmitter that can induce calcium mobilization [28,29]. To investigate the role of coronin 1A in stimulated astroglia, fluorescence-based live-cell calcium imaging was performed upon stimulation with ATP. An increase in the fluorescence of the calcium-sensitive dye Fluo-4 was taken as a measure of Ca\(^{2+}\) flux. An increase in intracellular calcium upon ATP stimulation was recorded in HEPES buffer (+/- CaCl\(_2\)). Basal fluorescence was recorded for 1 min and then 10 µM ATP was applied to the bath and recordings were performed for an additional 5 min. ATP stimulation induced the Ca\(^{2+}\) flux and fluorescence intensity increased. In the absence of CaCl\(_2\) in the HEPES buffer (i.e. source of Ca\(^{2+}\) is just intracellular), astroglia downregulated for coronin 1A showed a 2.67-fold decrease in fluo-4 fluorescence as compared to the control group (p<0.05), suggestive of its role in calcium mobilization in stimulated
conditions (Fig. 3a). The Ca\(^{2+}\) flux compromise observed when the HEPES buffer with CaCl\(_2\) was used (i.e. source of Ca\(^{2+}\) is both, intra- and extracellular), was not statistically significant (Fig. 3b). This indicated that coronin 1A downregulation affected the calcium mobilization from the intracellular stores.

The reduced Ca\(^{2+}\) flux in coronin 1A knocked down cells could be an outcome of the altered PLC\(\gamma\)1 phosphorylation. To assess this possibility, astroglia transfected with control scrambled siRNA or coronin 1A targeting siRNA were stimulated with 10 \(\mu\)M ATP for varying time points (0, 45, 120, and 300 sec) after which the cells were harvested and the whole cell lysate was analyzed through western blotting for the phosphorylated and the total forms of PLC\(\gamma\)1. It was observed that coronin 1A knockdown resulted in reduced PLC\(\gamma\)1 phosphorylation as compared to the control group (at t=120 sec; p<0.05) (Fig. 3c). We also assessed the ERK1/2 phosphorylation pattern after ATP stimulations for varying time-points (0, 5, 10, 30, 60, and 120 min) and observed a slight decrease in ERK1/2 phosphorylation, although not significant (Fig. 3d). The above results indicate that, although coronin 1A is dispensable in resting-state but plays activity-dependent roles by facilitating calcium mobilization from the intracellular stores upon stimulation.

**Coronin 1A is upregulated in human astroglia activated by neuroinflammatory agent HIV-1 Tat**

Neuroinflammatory agent HIV-1 Tat is a well-known neurotoxicant that induces astrogliosis and hence causes astroglia-mediated neuronal death in HIV-1 neuropathogenesis [30,7]. To further check the possibility of activity-dependent roles of coronin 1A, we checked whether HIV-1 Tat can modulate coronin 1A expression in astroglia. The real-time PCR assay and western blot analysis revealed a 1.46-fold and 1.43-fold increase in the level of coronin 1A mRNA and protein expression, respectively, in astroglia transfected with HIV-1 Tat expression vector compared to cells transfected with vehicle control (p<0.0005) (Fig. 4a, b). HIV-1 Tat, indeed, turned the resting state astroglia reactive as the cells showed enhanced expression of GFAP (1.66-fold; p<0.005) (Fig. 4c) and pro-inflammatory gens such as VCAM1 (1.39-fold; p<0.0005), ICAM1 (1.2-fold; p<0.0005), PTGS2 (2.11-fold, p<0.05) and NOS2 (1.59-fold; p<0.05) (Fig. 4d). These findings, in conjunction with the previous results, further indicate that with the increase in its expression in the inflammatory environment, coronin 1A might be involved in astrogliosis and glia-mediated neurodegeneration.

**Downregulation of coronin 1A attenuates HIV-1 Tat-induced GFAP, IL-6, and Glutamate release and supports Neuronal survival**

Next, we hypothesized if coronin 1A is important for calcium mobilization and the literature suggests that calcium plays a key role in astroglial activity and reactivity [14,31], downregulating coronin 1A might ameliorate the detrimental effects of reactive astroglia observed in various neurodegenerative disorders, including neuroAIDS. So, to investigate this hypothesis, astroglia were knocked down using siRNA against coronin 1A for 48 h followed by transfection with HIV-1 Tat expression vector for 24 h. Quantitative PCR and western blotting analyses revealed that the Tat-induced expression of GFAP ameliorated in the experimental group where coronin 1A was downregulated before-hand. Scrambled +
HIV-1 Tat group showed a 1.51-fold (p<0.0005) increase in GFAP levels, whereas coronin 1A siRNA + Tat group showed GFAP expression similar to the scrambled + vector control group (Fig. 5a). Astrocytes have also been reported to be immunocompetent and possess immunological properties [32]. To check the immune status of the cells, mRNA levels of several pro-inflammatory genes were also checked in the current setting. Downregulating coronin 1 could ameliorate the levels of some of the Tat-induced pro-inflammatory genes (refer Fig. 4). It was observed that the scrambled + HIV-1 Tat group showed up-regulation in PTGS2 (1.67-fold, p<0.05), NOS2 (1.73-fold, p<0.05) whereas coronin 1A siRNA + Tat group had PTGS2 and NOS2 levels similar to the scrambled + vector control group (Fig. 5b).

Reactive astroglia release excess of glutamate, cytokines, and/or chemokines which may cause excitotoxicity or neurotoxicity to the neurons present in the vicinity [32-34,5]. In this regard and to validate the findings of the previous experiments, several functional assays were performed using the conditioned-media derived from astroglia after coronin 1A knockdown for 48 h followed by HIV-1 Tat transfection for 24 h. The freshly collected cell supernatants were analyzed for the levels of released Interleukin-6 (IL-6) and glutamate. The FACS analysis supported the hypothesis even further as it was observed that the enhanced release of the pro-inflammatory cytokine, IL-6 observed under the effect of HIV-1 Tat (1.17-fold, p<0.05), ameliorated when the cells were knocked down for coronin 1A, beforehand (Fig. 6a). Also, the glutamate release assay revealed that the HIV-1 Tat could not induce the enhanced glutamate release when the cells were knocked down for coronin 1A. HIV-1 Tat alone could indeed cause a 1.24-fold (p<0.05) increase in the levels of released glutamate (Fig. 6b).

The deleterious effects of HIV-1 Tat-induced reactive astroglia are also manifested in the form of neuronal death. So, to further investigate whether HIV-1 Tat-induced reactive astroglia-mediated neuronal death could be rescued by downregulating coronin 1A, we transfected astroglia with coronin 1A siRNA for 48 h followed by transfection with the HIV-1 Tat expression vector for 24 h. Astrocyte-conditioned media (ACM) were collected from these experimental sets and added to the neuronal cultures to inspect the indirect effects. ACM from the Tat alone group decreased neuronal survival to 75.5% (p<0.05) whereas ACM collected from the group which was transfected with coronin 1A siRNA before HIV-1 Tat transfection, showed rescue in neuronal death (Fig. 6c). These results signified that downregulating coronin 1A is protective as it slows astroglial reactivity and thus supported our hypothesis that coronin 1A facilitates HIV-1 Tat-induced astrogliosis and subsequent neuronal death.

**miR-92b-5p regulates coronin 1A in HIV-1 Tat-induced reactive astroglia**

HIV-infected brains or brain cells also present a dysregulated cellular miRNA profile which might contribute to the virus-induced neurological complications [35-37]. To check the miRNA regulatory machinery, a small RNA sequencing was outsourced which was performed on astroglia transfected with HIV-1 Tat expression vector or vector control. A heat map was generated for the miRNAs which were differentially regulated (by ± 1.3 to ± 2.5-fold) under the effect of HIV-1 Tat. Heatmap function of display ‘R’ tool (https://www.displayr.com) was used to generate the heatmap by using fold-change derived from the absolute reads of the sequencing data (Fig. S3). Out of all the differentially regulated miRNAs, we
narrowed down to the downregulated ones, and further, bioinformatics tool miRWalk, was employed to filter down to miRNAs which are predicted to target 3’ UTR of CORO1A (Fig. 7a). Individual qPCR assay performed for these miRNAs confirmed the HIV-1 Tat-induced downregulation. It was observed that HIV-1 Tat downregulated miR-92b-5p the most (-1.57-fold, p<0.0005) (Fig. 7b), hence further confirmatory assays were performed only for miR-92b-5p.

To confirm whether miR-92b-5p regulates CORO1A, 111 bp long 3’ UTR of CORO1A was cloned in pMIR-REPORT and was used along with mimic-92b-5p for co-transfection into Hela cells. Luciferase assay was performed following the manufacturer’s protocol and relative luciferase units were measured after 24h of transfection. It was observed that mimic-92b-5p reduced the luminescence by 1.85-fold (p<0.005) confirming that it indeed targets 3’ UTR of CORO1A (Fig. 7c).

To further validate the findings, mimic, and inhibitor against miR-92b-5p were employed. Mimic 92b-5p transfection into human astroglia for 48 h followed by qPCR analysis revealed that miR-92b-5p level increased by 247-fold (p<0.05) compared to the cells transfected with mimic control (Fig. 8a). The transcript and protein levels of coronin 1A showed a significant decrease by -2.36-fold (p<0.0005) and -1.35-fold (p<0.005) respectively, because of the mimic-92b-5p transfection, compared to the control transfected group (Fig. 8b, c). In contrast, transfection of inhibitor against miR-92b-5p resulted in significant inhibition of miR-92b-5p levels (-2.33-fold, p<0.05) compared to the control group (Fig. 8d). The transcript and the protein levels of coronin 1A showed a significant increase by 1.6-fold (p<0.05) and 1.57-fold (p<0.0005) respectively, because of the transfection with an inhibitor against miR-92b-5p, compared to the control group (Fig. 8e, f).

**Discussion**

Despite being heavily explored, the context-dependent behavior of the astroglial cells is yet to be thoroughly understood. Apart from performing many supportive roles, these cells exhibit varied behavior in pathophysiological conditions. Astroglial activation is a common phenomenon in a variety of neurological disorders [1,3]. The reactive astroglia become neuroinflammatory in several contexts and thus become detrimental for the neurons present in the vicinity, and hence cause mild to severe neuronal death. Reactive astroglia-mediated neuronal death is a leading cause of neuronal loss observed in various neurological disorders including neuroAIDS. NeuroAIDS encompasses the neurological complications associated with HIV-1 infected individuals, around 50% of such individuals manifest some degree of HIV-1 associated neurocognitive disorder (HAND) [38-40].

Calcium excitability of the astroglia is a well-established phenomenon in the field of glia biology. In fact, Calcium regulates astroglial secretions including glutamate [31,14,27]. Hence, it plays a central role in inducing astroglial activity both in physiological as well as pathophysiological conditions.

Coronin 1A has been found to regulate T cell differentiation and activation events [20], it also regulates their survival by mediating calcium mobilization [23] and links cytoskeletal dynamics to receptor signaling [26,21]. In the brain, it has been found to influence NGF-TrkA endosomal trafficking as well as
the signaling events. Also, by promoting cAMP/PKA signaling, it regulates synaptic plasticity and hence influences cognition and behavior [41,22]. The role of this protein in different cell types and whether its functions are context-dependent still needs extensive exploration. Astroglia, indeed, play significant roles in regulating neuronal activity and astroglial activation directly influences neuronal health and functioning in pathological contexts. Whether these cells express coronin 1A, and what role it plays is not known at all. To address this, and to identify the roles of this protein in different cell types and contexts, we investigated the role of coronin 1A in astroglia physiology, and astroglial activation observed in the presence of a potent neurotoxicant HIV-1 Tat B. For the current study, we used a well-characterized model of primary astrocytes and neurons that were differentiated from human fetal brain-derived neural precursor cells.

Coronin 1A has mostly been studied in cells of hematopoietic origin, so, at first, we checked for the coronin 1A expression in various brain cell types such as neural precursor cells (NPCs), astroglia, and neurons; Jurkat T cells were taken as a positive control. All the brain cell types were found to express coronin 1A. We also assessed the differentiating astrocytes for coronin 1A protein expression at day 0, 7,14, and 21. Immunocytochemistry performed to confirm the intracellular localization of coronin 1A in astroglia revealed its expression throughout the cytoplasm of the cell (Fig. 1). The astroglial culture was found to be positive for GFAP and Vimentin in more than 95% of the cells (Supporting Information Fig. S1).

To explore the functions of this protein in resting-state astroglia, we deployed a siRNA-mediated loss-of-function approach. A 48 h of knockdown was carried out and the cells were then assessed for knockdown efficiency and the transcript and protein levels of GFAP and Vimentin. This experiment led us to an important conclusion that coronin 1A was dispensable in the resting-state astroglia as the knockdown does not affect the GFAP and vimentin protein levels (Fig. 2). GFAP mRNA, although was significantly affected which might not be enough to bring about substantial change in the protein levels. Also, the above experimental condition did neither alter the cell-proliferation nor the cell-viability (Supporting Information Fig. S2). The literature about the role of this protein in T cell activation [20,23] provided us the momentum to investigate its involvement, if any, in stimulated conditions.

Astroglia possess calcium excitability and gliotransmitters such as ATP, glutamate can stimulate astroglia and cause calcium mobilization followed by downstream calcium signaling [28,29,27]. To explore whether coronin 1A plays activity-dependent roles, we measured ATP-induced calcium mobilization. Interestingly, we found that coronin 1A knockdown compromised the calcium mobilization significantly, especially when the extracellular buffer used was devoid of CaCl$_2$ i.e. when the source of calcium was only the intracellular stores. To check what caused compromised calcium mobilization, we assessed the phosphorylation pattern of PLC$_{\gamma}$1, which is the most common and widely studied enzyme involved in calcium signaling. Astroglial cells knocked down for coronin 1A were stimulated with ATP for several time-points and further harvested to check the phosphorylation pattern of PLC$_{\gamma}$1. PLC$_{\gamma}$1 phosphorylation was found to be significantly compromised at one time-point (at t=120 sec). We also assessed ERK1/2 phosphorylation, which is downstream to Ca$^{2+}$, and showed slightly compromised
phosphorylation but the differences were not significant (Fig. 3). These observations indicated the activity-dependent roles of coronin 1A in astroglial cells and therefore prompted us to look for its involvement in astrogliosis.

HIV-1 Tat is a potent neurotoxicant and is known to induce astrogliosis, which is characterized by an increase in GFAP expression [7], it also elevates the expression of cell adhesion molecules such as VCAM-1 and ICAM-1 in astroglia [42,43] and thus can facilitate the entry of inflammatory cells into the CNS. HIV-1 Tat also induces the expression of inducible nitric oxide synthase (iNOS) and Cyclooxygenase (COX)-2, an enzyme involved in prostanoid synthesis, which further modulates the expression of several inflammatory cytokines, and chemokines [44,45]. In agreement with the above-mentioned reports, we found enhanced expression of GFAP in human astroglia transfected with HIV-1 Tat expression vector, which confirmed the reactive status of these cells. Also, HIV-1 Tat was found to enhance the mRNA levels of several pro-inflammatory genes such as VCAM-1, ICAM-1, PTGS2, and NOS2. Interestingly, these reactive astroglia also exhibited the enhanced expression of coronin 1A, which further pointed out its activity-dependent roles (Fig. 4).

If coronin 1A has activity-dependent roles, then it might also be involved in the astroglial activation observed in the presence of HIV-1 Tat. Hence, to explore whether it really facilitates astroglial activation, HIV-1 Tat expression vector was transfected in cells downregulated for coronin, sequentially. It was indeed observed that HIV-1 Tat-induced glial activation, assessed by measuring GFAP levels, diminished when the cells were downregulated for coronin 1A. This propelled us to also assess the cell functionality by measuring the levels of release factors. Functional assays performed on the cells and the conditioned-media revealed that downregulating coronin 1A weakened the deleterious outcomes of the reactive astroglia such as enhanced expression of pro-inflammatory genes, PTGS2, and NOS2, the release of IL-6, and glutamate. Cumulatively, the reduced inflammatory response further supported primary neurons, as the neuronal survival was found to be improved in the group treated with conditioned media collected from reactive astroglia downregulated for coronin 1A (Fig. 5 and 6). This confirmed our hypothesis of the involvement of coronin 1A in facilitating HIV-1 Tat-induced astrogliosis.

A closer look at the results obtained from the sequential transfection of coronin 1A siRNA followed by transfection of HIV-1 Tat expression vector, point out that downregulating coronin 1A only (compare blue bars) has some calming effects on astroglia behavior. Downregulating coronin 1A suppressed the levels of PTGS2 transcripts and released IL-6 (Fig. 5 and 6, respectively). This effect also reflected in the neuronal survival assessment where the neurons treated with conditioned media collected from astroglia downregulated for coronin 1A, survived even better than the control group (although not significant). The mechanisms causing these effects, require in-depth exploration in future studies.

Micro-RNAs dysregulated under the effect of HIV-1 often contribute to HIV-1 neuropathogenesis [36,37,35]. Small RNA sequencing data gave us the pool of HIV-1 Tat-induced dysregulated miRNAs (Fig. S3). We further narrowed down to a list of miRNAs that are downregulated under the effect of HIV-1 Tat and target the CORO1A 3’ UTR. The expression of short-listed miRNAs was assessed in the cells
transfected with HIV-1 Tat expression vector, using real-time PCR assay. Out of all the selected miRNAs, miR-92b-5p was found to be affected the most and hence it could act as the potential regulator of coronin 1A expression in reactive astroglia. Utilizing mimic and inhibitor against miR-92b-5p to assess its binding with CORO1A 3’ UTR followed by measuring protein expression further confirmed the involvement of miR-92b-5p in augmenting coronin 1A expression in the HIV-1 Tat-induced reactive astroglia. Mimic-92b-5p significantly reduced the luciferase activity obtained when CORO1A 3’ UTR was cloned in pMIR vector having luciferase gene. The mimic also decreased the transcript and protein levels of coronin 1A. In contrast, inhibitor-92b-5p increased the transcript and the protein levels of coronin 1A (Fig. 7 and 8). This confirmed that miR-92b-5p mediated coronin 1A regulation under the effect of HIV-1 Tat.

This study concludes that coronin 1A, although dispensable in resting-state astroglia, facilitates calcium mobilization upon ATP stimulation in astroglia. Coronin 1A expression is elevated in HIV-1 Tat-induced reactive astroglia. Downregulating coronin 1A ameliorates the deleterious effects of reactive astroglia and hence supports neuronal survival. miR-92b-5p regulates coronin 1A expression by targeting its 3’ UTR. By reducing the levels of miR-92b-5p, HIV-1 Tat might augment coronin 1A protein levels which further facilitates astrogliosis and astroglia-mediated neuronal toxicity. Hence, this study reports the non-canonical, activity-dependent roles of coronin 1A, in stimulated astroglia, and in facilitating HIV-1 Tat-induced astrogliosis.

**Declarations**

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**Competing interests**

The authors declare that they have no competing interests.
Ethics approval

Human brain tissues collected from elective abortions with the informed consent of mothers, were processed as per the protocols laid down by the Institutional Human Ethics and Stem Cell Research Committee under strict compliance with the guidelines of ICMR, India.

Consent to participate

Not applicable.

Consent for publication

Not applicable.

Availability of data and materials

All data generated or analyzed during this study are included in this published article [and its supplementary information files].

Code availability

Not applicable.

Author's contributions

HSP designed, performed experiments, analyzed, and interpreted the data, and formulated the manuscript. RK performed experiments and helped in analyzing data. Bindu helped in performing experiments. PS helped in designing experiments, analyzing data, and finalizing manuscript. All authors read and approved the final manuscript.

References


Figures
Figure 1

Human Neural Progenitors, astroglia, and neurons express Coronin 1A. Coronin 1A protein expression in (a) neural progenitors, astrocytes, neurons, and Jurkat T cells and (b) differentiating astroglia at day-7, -14, and -21, determined by western blotting. (c) Immunocytochemistry was performed on astroglia immunolabeled for Coronin 1A (red), and GFAP (green). Cell nuclei were DAPI stained (blue). Images were acquired using a 20X objective lens and the scale bar represents 50 µm. NPC=Neural precursor cells;
PDA=Progenitor-derived astroglia; PDN=Progenitor-derived neurons; P2A=Progenitor to astroglia differentiation.

**Figure 2**

Coronin 1A knockdown does not affect the resting-state human astroglia. Astroglia were transfected with CORO1A siRNA for 48 h, harvested, and transcript and protein levels were analyzed using real-time qPCR western blotting, respectively. (a) The bar graph represents the fold change in coronin 1A expression at mRNA (upper panel) and protein (lower panel) compared to the control group. (b) Cells downregulated for coronin 1A were assessed for Glial Fibrillary Acidic Protein (GFAP) mRNA (upper panel) and protein (lower panel) expression. (c) Vimentin mRNA (upper panel) and protein (lower panel) expression were also assessed in the same experimental group. Blots are representative and the bar graph shows the densitometric analysis of the blots. GAPDH was used as a normalization control for all the analyses. Data represent mean ± S.D of four independent experiments. *** represents p<0.0005. SCR=scrambled; CORO1A=coronin 1A.
Figure 3

ATP-induced Calcium mobilization and signaling pathways in human astroglia downregulated for coronin 1A. Calcium mobilization in Fluo-4 AM labeled cells, knocked down for coronin 1A, upon stimulation with 10 µM ATP in (a) HEPES buffer without CaCl2, and (b) HEPES buffer having CaCl2. Arrow indicates the time of the ATP application in the buffer. The bar graph (right panel) shows the area under the curve (AUC). Astroglia downregulated for coronin 1A were assessed for (c) PLCγ1 and (d)
ERK1/2 phosphorylation upon stimulation with 10 µM ATP for the indicated time-points. The upper panel shows the representative blots, and the bar graph shows the densitometric analysis of the same. Data represent mean ± S.D of three independent experiments. * represents p<0.05. SCR=scrambled; CORO1A=coronin 1A, ΔF=change in fluorescence intensity.

Figure 4
HIV-1 Tat augments coronin 1A expression and activates human astroglia. Astroglia transfected with HIV-1 Tat expression vector for 24 h were harvested and analyzed for (a) Coronin 1A mRNA levels, using real-time PCR assay and (b) protein level, using western blotting, upper panel shows the representative blots and the lower panel shows the densitometric analysis of the same, compared to cells transfected with vector control. The activated status of the HIV-1 Tat transfected cells was checked by assessing the expression of (c) GFAP protein and (d) pro-inflammatory genes viz VCAM1, ICAM1, PTGS2, and NOS2 compared to the group transfected with vector control. GAPDH was used as a normalization control for all the analyses. Data represents mean ± S.D of four independent experiments. * represents p<0.05, **p<0.005, and ***p<0.0005. VC=vector control.

Figure 5

Coronin 1A downregulation in human astroglia ameliorates expression of HIV-1 Tat-induced GFAP and pro-inflammatory genes in reactive astroglia. Human astroglia downregulated for coronin 1A using siRNA for 48 h followed by transfection with HIV-1 Tat for 24 h, were harvested and analyzed for (a) GFAP
protein expression, using western blotting. The left panel shows representative blot and the right panel shows the densitometric analysis of the same and (b) pro-inflammatory genes viz PTGS2 and NOS2, using real-time PCR assay. GAPDH was used as a normalization control for all the analyses. Data represents mean ± S.D of three independent experiments. *p<0.05, ***p<0.0005. SCR=scrambled; CORO1A=coronin 1A.

Figure 6

Downregulation of Coronin 1A in astroglia ameliorates HIV-1 Tat-induced release of IL-6, glutamate, and the associated indirect-neuronal death. Astroglia-conditioned media (ACM) collected from cells, transfected with CORO1A siRNA for 48 h followed by transfection with HIV-1 Tat expression vector for 24 h, were analyzed for released (a) interleukin-6 (IL-6), using Flow cytometry and (b) glutamate, using
enzyme kinetics based glutamate release assay. Human primary neurons were treated with ACM collected from the above experiment for 24 h and assessed for (c) Neuronal survival by analyzing the TUNEL-positive neuronal cells. Data represent the mean ± S.D of three independent experiments. *p<0.05 with respect to control. SCR=scrambled; CORO1A=coronin 1A.

Figure 7

MiR-92b-5p is downregulated in response to HIV-1 Tat and targets 3' UTR of CORO1A. Astroglia were transfected with HIV-1 Tat expression vector for 24 h and were assessed for the expression of miRNAs. (a) Heatmap prepared using fold-change derived from absolute values of sequencing data, showing miRNAs which are downregulated by -1.3 to -2.5-fold in the presence of HIV-1 Tat, compared to the vector control group, and predicted to target CORO1A. (b) The bar graph represents the levels of selected miRNAs compared to the vector control group, analyzed using real-time PCR assay; U6 was used for normalizing the data. (c) CORO1A 3' UTR was cloned downstream to the luciferase gene in the pMIR-Report vector and was co-transfected along with mimic control or mimic-92b-5p into Hela cells.
Quantitative analysis shows the normalized relative luciferase activity in the indicated groups. Data shown represent four independent experiments (mean ± S.D). *p<0.05, **p<0.005, and ***p<0.0005.

**Figure 8**

MiR-92b-5p mimic downregulates and inhibitor of miR-92b-5p upregulates CORO1A levels. Astroglia were transfected with mimic-92b-5p for 48 h. Cells were then harvested for miRNA, mRNA, and whole-cell protein. The bar graph shows the effect of mimic-92b-5p on levels of (a) miR-92b-5p, (b) CORO1A transcripts, and (c) coronin 1A protein, compared to their respective levels in the control transfected group. Similarly, astroglia transfected with an inhibitor against miR-92b-5p were assessed for the effects on levels of (d) miR-92b-5p, (e) CORO1A transcripts, and (f) coronin 1A, compared to their respective...
levels in the control transfected group. miRNA levels were normalized using U6 whereas mRNA and protein levels were normalized with GAPDH mRNA and protein levels, respectively. Data represents mean ± S.D of at least three to four independent experiments. *p<0.05, **p<0.005, and ***p<0.0005. Inh=inhibitor.

**Supplementary Files**

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- SupplementaryInformation.pdf