

## UPDATE Blood-brain barrier pericytes as a target for HIV-1 infection

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Pericytes are multifunctional cells wrapped around endothelial cells via cytoplasmic processes that extend along the abluminal surface of the endothelium. The interactions between endothelial cells and pericytes of the blood-brain barrier are necessary for proper formation, development, stabilization, and maintenance of the blood-brain barrier. Blood-brain barrier pericytes regulate paracellular flow between cells, transendothelial fluid transport, maintain optimal chemical composition of the surrounding micro-environment, and protect endothelial cells from potential harmful substances. Thus, dysfunction or loss of blood-brain barrier pericytes is an important factor in the pathogenesis of several diseases that are associated with microvascular instability. Importantly, recent research indicates that blood-brain barrier pericytes are prone to establish a latent infection, which can be reactivated by a mixture of histone deacetylase inhibitors in combination with TNF. HIV-1 infection of blood-brain barrier pericytes has been confirmed in a mouse model of HIV-1 infection and in human post-mortem samples of HIV-1 target and reservoir in the brain.

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### Introduction

Pericytes are multifunctional cells wrapped around endothelial cells via cytoplasmic processes that extend along the abluminal surface of the endothelium. Most of the pericyte-endothelial interface is separated by the basement membrane (Armulik *et al.*, 2011), which is actively remodelled by both pericytes and endothelial cells during angiogenesis (Kloc *et al.*, 2015), development (Armulik *et al.*, 2011), and tumorigenesis (Baluk *et al.*, 2003). However, selected pericyte populations are also present with incomplete or absent basement membranes and in direct contact with endothelial cells (Armulik *et al.*, 2011). Pericytes can transiently rearrange their contacts with endothelial cells via the peg-socket type of arrangements, and/or modulation of adherent junctions or gap junctions (Winkler *et al.*, 2011) (Fig. 1).

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barrier pericytes (yellow) and endothelial cells (red) share the basement membrane (sky blue) or are in direct contacts. In addition, blood-brain barrier pericytes are surrounded by glial cells (astrocytes, green; microglia, dark blue) and neurons (purple). (**B**) Blood-brain barrier pericytes and endothelial cells communicate with each other by direct contact (gap and adherens junctions) or through signalling pathways, such as platelet-derived growth factor BB (PDGF-BB)/PDGFR $\beta$  and transforming growth factor- $\beta$  (TGF- $\beta$ )/type 2 TGF- $\beta$  receptor (TGF $\beta$ R2).

Pericytes display contractile, cytoskeletal, and surface proteins; however, no single or specific pericyte marker is currently available. Identification of pericytes is frequently achieved by determination of more than one marker (Fig. 2). Platelet-derived growth factor receptor- $\beta$ (PDGFR $\beta$ ) (Armulik *et al.*, 2010), neuron-glial antigen 2 (NG2) (Trost *et al.*, 2013), CD13 (Armulik *et al.*, 2010), alpha-smooth muscle actin ( $\alpha$ SMA) (Trost *et al.*, 2016), regulator of G protein signaling 5 (RGS5) (Mitchell *et al.*, 2008), and desmin (Bergers and Song, 2005) have been suggested to be useful for pericyte characterization based on their dynamics of expression related to developmental status and pathological conditions. For example, a contractile protein  $\alpha$ SMA is differentially expressed in pericytes depending on the type of blood vessel.

Given that  $\alpha$ SMA, NG2, PDGFR $\beta$ , and desmin are also expressed by vascular smooth muscle cells (SMCs), a collective approach of well preserved tissue morphology, counter-labelling of endothelial cells, and expression of two or more pericyte markers has been advocated for pericyte identification *in vivo* (Trost *et al.*, 2013; Yamamoto *et al.*, 2017). Nevertheless, the precise structural and molecular identity of pericytes is still controversial. It was recently proposed that capillary pericytes are strictly defined and separated from other cell types by the lack of  $\alpha$ SMA expression (Hill *et al.*, 2015). However, the majority of literature reports acknowledge that pericytes exhibit heterogeneity with distinct morphologies, expression level of  $\alpha$ SMA, and functions depending on their positioning within the vasculature (Hall *et al.*, 2014; Attwell *et al.*, 2016).

Although the origins of pericytes across all tissues are still not fully understood, peripheral pericytes, including those of the gut, lung, and liver, mostly develop from the mesoderm during embryogenesis via epithelial-to-mesenchymal transition, delamination, and migration (Wilm et al., 2005; Que et al., 2008; Asahina et al., 2011). Similarly, the epicardial mesothelium is thought to give rise to cardiac pericytes (Cai et al., 2008). While pericytes in the spinal cord and brainstem have developed from the mesoderm, the majority of blood-brain barrier pericytes in the CNS are more likely to be derived from the neural crest cells of the neuroectoderm (Korn et al., 2002; Simon et al., 2012). Moreover, a subset of cerebrovascular pericytes was shown to originate from mature macrophages in the early phase of CNS vascular development (Yamamoto et al., 2017).

While pericytes cover ~30% of the abluminal surface of peripheral endothelial cells (Mathiisen *et al.*, 2010), this coverage is closer to 90% in the CNS (Sá-Pereira *et al.*, 2012) and appears to be inversely correlated with endothelial cell proliferation (Díaz-Flores *et al.*, 2009). By covering endothelial cells, pericytes play a critical role in regulation of capillary blood flow (Hamilton *et al.*, 2010). Indeed,



Figure 2 Blood-brain barrier pericyte markers. Standardized characterization of non-infected (**A**) and HIV-1-infected (**B**) pericytes. Infections were performed by incubating pericytes with HIV-1 (NL4-3 strain) in the amount of 60 ng p24/ml for 48 h. *Top*: Phase contrast imaging; *middle* three panels represent immunofluorescence staining for the three distinct pericyte markers; namely, PDGFR $\beta$ , NG2, and  $\alpha$ SMA (all in green). *Bottom*: Merged images of triple stained pericytes for these markers (PDGFR $\beta$ , green; NG2, blue; and  $\alpha$ SMA, red). Nuclei were stained with Hoechst (blue). Scale bars = 200 µm (brightfield) and 50 µm (PDGFR $\beta$ , NG2, and  $\alpha$ SMA).

pericyte-mediated capillary contraction and dilation has been observed both in vivo and ex vivo (Attwell et al., 2010; Itoh and Suzuki, 2012). Recent evidence also suggests that interactions between endothelial cells and blood-brain barrier pericytes are necessary for the proper formation, development, and maintenance of the blood-brain barrier. Blood-brain barrier pericytes were shown to regulate paracellular flow between cells and transendothelial fluid transport, maintain optimal chemical composition of the surrounding microenvironment, and protect endothelial cells from potential harmful substances (Winkler et al., 2011). The intricate cross-talk and functional coupling between pericytes and endothelial cells have been implicated in vessel stabilization and development via several signal transduction cascades (Sweeney et al., 2016), including platelet-derived growth factor-BB (PDGFBB) (Winkler et al., 2010), transforming growth factor- $\beta$ , (TGF- $\beta$ ) (Walshe *et al.*, 2009), Notch (Hofmann and Iruela-Arispe, 2007), sphingosine-1 phosphate (Paik et al., 2004) and angiopoietin (Gaengel et al., 2009) signalling. Multiple ligand-receptor complexes mediate pericyte recruitment to the endothelium, such as PDGF-BB/PDGFB (Tallquist et al., 2003), heparin-binding epidermal growth factor (HB-EGF)/EGF receptors (Stratman et al., 2010), and stromal-derived factor-1 $\alpha$ , (SDF-1 $\alpha$ )/ CXCR4 (Song et al., 2009). Moreover, blood-brain barrier pericytes control the integrity of the blood-brain barrier by regulating endothelial tight and adherent junctions (Bell et al., 2010; Daneman et al., 2010).

Pericytes were demonstrated to actively participate in immune responses. Insult by inflammatory stimuli causes pericytes to release inflammatory mediators, including IL-1  $\beta$ , IL-6, CCL2, and matric metalloproteinases (MMP2/9) (Nakagawa *et al.*, 2012; Guijarro-Muñoz *et al.*, 2014). Blood-brain barrier pericytes also control leucocyte adhesion and transmigration in the CNS (Olson and Soriano, 2011) and can perform macrophagic-like activities to clear tissue debris and cellular by-products in the CNS (Bergers and Song, 2005; Bell *et al.*, 2010). Furthermore, pericytes were proposed to exhibit multipotent stem cell potential, capable of forming neurons, glia and vascular lineage cells (Nakagomi *et al.*, 2015).

loss Dysfunction or of blood-brain barrier pericytes leads to blood-brain barrier breakdown (Bell et al., 2010; Winkler et al., 2012), contributing to the pathogenesis of several diseases that are associated with microvascular instability and blood vessel ruptures (Armulik et al., 2011). Dysfunctions of signalling pathways, such as PDGF-BB/PDGFRB, TGF-B/TGFBR2, Notch, and Ang-1/Tie-2, within pericytes or between pericytes and neighbouring cells were shown to be involved in CNS diseases, such as stroke (Li et al., 2011; Arnold et al., 2014), Alzheimer's disease (Sagare et al., 2013), amyotrophic lateral sclerosis (Winkler et al., 2013), and brain tumours (Bergers et al., 2003; Zhang et al., 2003).

# Infection of blood-brain barrier pericytes by HIV-1

Compelling evidence demonstrates that HIV-1 infection of cells of the CNS, resulting in production of toxic viral proteins, is a critical factor in the development of HIV-associated neurocognitive disorder (HAND). The majority of HIV-1 replication in the brain occurs in microglial cells and perivascular macrophages (Kim *et al.*, 2005; Joseph *et al.*, 2015; Chen *et al.*, 2017). It has also been established that astrocytes can be productively infected and play a role as a HIV-1 reservoir, given their longevity and ability to support HIV-1 reactivation (Li *et al.*, 2016). In addition, our group demonstrated for the first time that blood-brain barrier pericytes can be productively infected by HIV-1 (Nakagawa *et al.*, 2012). These findings were confirmed in subsequent studies by us and others (Persidsky *et al.*, 2016; Cho *et al.*, 2017).

# HIV-1 entry in blood-brain barrier pericytes

Canonical HIV-1 virus entry uses both the main HIV-1 receptor, CD4, and the co-receptors, primarily CXCR4 and CCR5. Pericytes express high levels of HIV-1 co-receptors, CCR5 and CXCR4, and also CD4, albeit at a low level (Nakagawa et al., 2012). Consistent with high expression of CCR5 and CXCR4, blood-brain barrier pericytes can be infected by both X4 and R5 tropic HIV-1 strains. The importance of these co-receptors in the viral entry to blood-brain barrier pericytes was demonstrated by the successful inhibition of infection with maraviroc, a clinically approved inhibitor of CCR5, and AMD3100, a specific inhibitor of CXCR4 (Nakagawa et al., 2012). The exact mechanism of entry for HIV-1 to blood-brain barrier pericytes remains to be elucidated; however, data from myeloid cells show that even with low CD4 levels, receptor mediated fusion can occur (Joseph et al., 2014). In addition, astrocytes, which do not express CD4, can still be infected by HIV-1 (Liu et al., 2004; Chauhan et al., 2014; Li et al., 2015). Given that in vitro infection of blood-brain barrier pericytes with HIV-1 results in a relatively low number of infected cells, it is highly probable that infection could be enhanced by cell-to-cell transmission, as it was demonstrated for astrocytes (Li et al., 2015). This is especially relevant given the high degree of cell-cell contact present within the neurovascular unit. In addition, the extensive coverage of the brain perivascular environment by blood-brain barrier pericytes increases the possibility of direct viral contact or of cell-to-cell transmission with transmigrating peripheral infected cells, such as T cells and/or macrophages (Armulik et al., 2010; Daneman et al., 2010).

A controversial issue is transmission of HIV-1 to bloodbrain barrier pericytes from brain endothelial cells. While endothelial cells can be infected with HIV-1 *in vitro* (Moses *et al.*, 1993), no productive infection of these cells was demonstrated *in vivo*. However, blood-brain barrier pericytes, when co-cultured with brain endothelial cells, enhance the lipopolysaccharide-mediated transendothelial transfer of HIV-1 free virus. This effect occurs without altering transendothelial electrical resistance, suggesting that pericytes affect the transcytotic component of HIV-1 permeation (Dohgu and Banks, 2013).

# Productive HIV-1 infection in blood-brain barrier pericytes

While virus entry is an essential step in infection, it does not necessarily signify that cells can be productively infected. Indeed, several additional factors come into play to enable successful HIV-1 integration and viral replication. By measuring extracellular p24 levels produced by infected cells and reverse transcriptase (RT) activity, we demonstrated that blood-brain barrier pericytes support HIV-1 replication, leading to a peak of extracellular virus production 2-3 days post-infection, followed by a gradual decline to the baseline 7-10 days post-infection (Castro et al., 2016; Cho et al., 2017). Further demonstration of competent replication was obtained using UV-inactivated virus (Cho et al., 2017). Blood-brain barrier pericytes are resistant to HIV-1-induced cytopathic effects, continue to proliferate, and do not display any apparent changes in morphology after infection. A typical experiment demonstrating HIV-1 infection of cultured human pericytes is shown in Fig. 3A and B.

The initial peak of HIV-1 virus production followed by a gradual decline in p24 production, and an increase in integrated HIV-1 genome (Fig. 3C) suggest a potential for the establishment of a latent infection. To confirm these findings, we performed HIV-1 reactivation studies using histone modifiers. Specifically, HIV-1-infected blood-brain barrier pericytes in the latent stage were exposed to mixtures of histone deacetylase (HDAC) inhibitors vorinostat (suberoylanilide hydroxamic acid, SAHA) and apicidin, as well as tumour necrosis factor (TNF) for 3 days. Treatments using HDAC inhibitors in combination with TNF resulted in a significant increase in p24 production and HIV-1 RNA (Fig. 3D and E, respectively). Overall, the results from *in vitro* studies indicate that pericytes can be a target for a productive HIV-1 infection, which can thereafter enter a latent phase and be reactivated, acting as a potential reservoir.

Initial *in vivo* demonstration of HIV-1 infection in blood-brain barrier pericytes was obtained in mice infected with a chimeric HIV-1 strain called EcoHIV-1, which was derived by replacing gp120 with gp80 of murine leukaemia virus (Potash *et al.*, 2005). This change altered the species specificity; thus, EcoHIV-1 infects mice, but not humans. Brain infections with EcoHIV-1 exhibit neuroinvasion, microglial activation, astrocytosis, and increased expression of genes coding for inflammatory cytokines and chemokines typical of human HIV-1 brain disease. We standardized the EcoHIV-1 brain inoculation technique by infusion



**Figure 3 HIV-1** replication and reactivation in human primary blood–brain barrier pericytes. (A) Quantification of p24 release into culture media at the indicated days post-infection with different HIV-1 strains, namely NL4-3, YU-2, or 49.5, in the amount of 60 ng p24/ml. The tropism (X4 or R5) of these strains is indicated on the graph. Day 0 indicates the start of infection. Data represent mean  $\pm$  standard error of the mean (SEM) from three independent experiments, (3 × 10<sup>5</sup> pericytes per replicate, total *n* = 9 per group). No p24 levels were detected in the non-infected (NI) group. (**B**) Representative images of p24 immunoreactivity at Day 2 post-infection with HIV-1 NL4-3 (60 ng p24/ml; orthogonal view in the merged image of HIV-1 NL4-3 group). No p24 levels were detected in NI group. Nuclei (blue, Hoechst staining), p24 (green, HIV-1 marker) and membranes (red, Dil staining). Scale bar = 10 µm. (**C**) Blood–brain barrier pericytes were infected with HIV-1 NL4-3 as in Fig. 3A (3 × 10<sup>5</sup>, 60 ng p24/ml), washed and incubated for 7 days. HIV-1 p24 release from HIV-1-infected pericytes and HIV-1 DNA integration into their genome as quantified by droplet digital PCR (ddPCR). Note that a decrease in active production of p24 is associated with elevated integration of the HIV-1 genome into the host genome. \**P* < 0.05 versus Day 3 post-infection. (**D** and **E**) On Day 8 post-infection, 3 × 10<sup>5</sup> pericytes were exposed to the indicated factors for 3 days and then assayed for either (**D**) HIV-1 p24 by ELISA or (**E**) HIV-1 RNA using RT-qPCR. Results are reflected by minimum and maximum box and whisker plots. The HIV-1 reactivation factors were used at the following concentrations: TNF, 100 U/ml; SAHA, 10 µM; apicidin, 1 µg/ml. \**P* < 0.05 versus HIV-1; \*\**P* < 0.01 versus HIV-1; \*\**P* < 0.001 versus HIV-1. **A**–**C** were adapted from Cho et *al.* (2017).

of 1 µg EcoHIV-1 p24 into the internal carotid artery (Bertrand *et al.*, 2016*a*). This approach mimics bloodborne HIV-1 trafficking into the brain in peak viraemia (Stephens *et al.*, 2003; Wright *et al.*, 2015). Mice were sacrificed 1 week post-infection, and brain microvessels were isolated and stained for NG2 (marker of pericytes) and p24 (marker of active HIV-1 replication). Pericytes positive for HIV-1 were detected on isolated brain microvessels derived from infected animals (Cho *et al.*, 2017), as evidenced by the co-localization of the pericyte markers PDGFR $\beta$  and HIV-1 protein p24, but not endothelial marker CD31 (Fig. 4A and B). Importantly, pericytes



**Figure 4** Infection of blood–brain barrier pericytes by HIV *in vivo*. (**A**) Male C57BL/6 mice were infected with EcoHIV-1/NDK (1  $\mu$ g of p24) via infusion through the internal carotid artery, while control animals (non-infected, NI) received saline. Mice were sacrificed at Day 7 post-infection. Brain microvessels were isolated and immunostained for PDGFR $\beta$  (green, arrow, marker of pericytes) and HIV-1 protein p24

positive for both HIV-1 gag and spliced Rev mRNA were observed using *in situ* PCR (Fig. 4C), indicating active transcription *in vivo*.

To confirm these results in human samples, we examined brain samples from patients with HAND. Human brain microvessels were isolated using the same approach as for mouse brains and the microvessels were stained for the pericyte markers PDGFR $\beta$  and HIV-1 p24. Figure 4D shows the first evidence that blood-brain barrier pericytes are infected in the brains of HIV-1-positive patients. Specifically, immunofluorescence analysis of the microvessels from HIV-1 infected brains demonstrates the presence of the protein p24 associated with blood-brain barrier pericytes. Details of these analyses are provided in the legend to Fig. 4D. At least five infected pericytes were clearly detected per slide that contained on average 100–150 microvessels.

# Blood-brain barrier pericytes as potential HIV-1 reservoir cells

Mounting evidence indicates that secluded HIV-1 populations and reservoirs in the CNS can perpetuate and become distinct from the periphery (Blankson et al., 2002; Strain et al., 2005). Restrictions of treatment efficacy due to the blood-brain barrier, which limits brain entry of several antiretroviral drugs, can result in their suboptimal concentrations, leading to the selection of antiretroviral-resistant mutants (Lambotte et al., 2003; Letendre et al., 2008; Bertrand et al., 2016b). Importantly, these resistant viruses can re-enter the periphery, leading to treatment failure (Hellmuth et al., 2015; Joseph et al., 2016). We propose that blood-brain barrier pericytes may play a pivotal role as CNS reservoirs. Indeed, these cells have a ubiquitous presence on the CNS vasculature (Armulik et al., 2010; Daneman et al., 2010) and they can self-renew. While the pericyte turnover rate in the neurovascular unit is not clearly defined, they may have a potentially long half-life, given the low angiogenesis plasticity in the CNS. The location of pericytes in the neurovascular unit could also facilitate the role of shuttling HIV-1 between the CNS and the periphery. As demonstrated in Fig. 3C-E, pericytes can harbour latent HIV-1, evidenced by the increase in DNA genome copies, decrease in HIV-1 p24 production and

subsequent ability to reactivate. The switch from productive to latent steps and back of the viral cycle observed in pericytes suggests their potential as a new and previously unrecognized HIV-1 CNS reservoir.

### Impact of HIV-1 infection on functions of blood-brain barrier pericytes

As described earlier, blood-brain barrier pericytes play a primordial role in the maintenance of the blood-brain barrier (Armulik et al., 2010). Through their contractile ability they can regulate cerebral blood flow and by paracrine signalling they can influence the tightness of the blood-brain barrier. This delicate balance can be altered by pathological conditions and its disruption can affect progression of several CNS disorders (Persidsky et al., 2016; Ferland-McCollough et al., 2017; Iadecola, 2017). Alterations of pericyte morphology and functions in HIV-1 infected patients have been observed in several recent reports. For example, it was demonstrated that patients infected with HIV-1, with or without antiretroviral treatment, have a significantly lower pericyte coverage of blood-brain barrier capillaries as shown by co-staining for CD13 with endothelial marker CD31 (Persidsky et al., 2016). The results presented in the present study (Fig. 2) indicate that infection with HIV-1 does not result in any apparent decrease in the expression of pericyte markers, suggesting that true loss of blood-brain barrier pericytes, and not merely a loss of marker expression, occurs in HIV-1 infection.

The mechanisms of HIV-1-induced loss of pericytes may be related to the exposure to viral proteins, such as Tat protein (Niu *et al.*, 2014, 2015). Specifically, HIV-1 Tat was shown to induce PDGF-BB expression in pericytes via stimulation of the MAPK/NF- $\kappa$ B pathway. Secreted PDGF-BB resulted in autocrine activation of the PDGF-BB/PDGFR $\beta$  signalling, culminating into increased pericyte migration (Niu *et al.*, 2014, 2015). PDGF-BB that is essential for pericyte generation, in high concentrations can lead to pericyte loss, the effect that was also implicated in HIV-1 Tat-mediated blood-brain barrier pericyte loss (Niu *et al.*, 2014, 2015). In addition, PDGF-BB has been shown to induce changes to pericyte secretome, increasing production of growth factors and proinflammatory cytokine IL-6 (Gaceb *et al.*, 2018).

#### Figure 4 Continued

(red, arrow, marker of active HIV-1 infection). Scale bars = 20  $\mu$ m. (**B**) Triple staining of infected microvessels from (**A**) for PDGFR $\beta$  (green, arrow), p24 (red, arrow), and CD31 (blue, marker for microvessel endothelial cells). Arrows denote area of co-localization solely between p24 and PDGFR $\beta$  but not with CD31, indicating that blood–brain barrier pericytes but not brain endothelial cells, are infected. (**C**) *In situ* RT-PCR assay using fluorescently-labelled primers against spliced HIV-1 *Rev* mRNA (blue, arrow), HIV-1 *gag* (NDK, red, arrow), and *NG2* spliced mRNA (green, pericyte marker). Focal signal indicates area of cDNA and DNA amplification. No signal for spliced Rev and NDK were observed in non-infected mice. (**D**) Brain samples (frontal cortex; 0.5 cm<sup>3</sup> each) from three healthy (non-infected, NI) and three HIV-1-infected patients with HIV encephalopathy were processed to isolate microvessels. Microvessels from each brain sample were spread on around 30 slides, each slide containing ~100–150 microvessels of varying sizes and number of associated pericytes. Samples were then immunostained for PDGFR $\beta$  (green, marker of pericytes) and HIV-1 protein p24 (red, marker of active HIV-1 infection). Approximately five infected pericytes were clearly detected on each slide containing microvessels from infected brains. Arrows indicate area of co-localization of p24 and PDGFR $\beta$ . Scale bars = 10  $\mu$ m. Arrow indicates area of co-localization.

Loss of pericyte coverage is exemplified by a decrease in blood-brain barrier integrity (Persidsky et al., 2016). Indeed, our studies indicate that exposure of brain endothelial cells to conditioned media obtained from HIV-1-infected pericytes results in disturbances in barrier integrity (Nakagawa et al., 2012). Part of this effect may be mediated by increased production of IL-6 (Nakagawa et al., 2012), which can induce proinflammatory responses and decrease integrity of the neurovascular unit (Blecharz-Lang et al., 2018). An increase in pericyte-produced proinflammatory cytokines, along with dysregulation of autophagy, was also observed in response to cocaine exposure (Sil et al., 2018). Loss of pericyte coverage on the brain endothelium can also lead to a reduction of connexin expression, leading to compromised cell-cell communication. Indeed, we demonstrated that HIV-1 infection of pericytes results in an increase in connexin 43 (Cho et al., 2017). While the role of these processes in HIV-1 infection remains to be elucidated, gap junction inhibitors, such as carbenoxolone, resulted in a drastic reduction in virus production by infected pericytes (Cho et al., 2017).

In summary, recent results from *in vitro*, *in vivo*, and human samples clearly indicate that blood–brain barrier pericytes can be productively infected with HIV-1 and may constitute a previously unrecognized HIV-1 reservoir in the CNS. The role of these processes in the pathogenesis of HAND and/or HIV-1 escape from the CNS remain to be elucidated.

#### **Future perspectives**

The present report indicates the need for more research on the role of pericytes not only in the brain infection by HIV-1 but also in brain physiology, the blood-brain barrier development, and the pathology of other neurodegenerative diseases and neuroinfections. The current controversy on defining blood-brain barrier pericytes highlights the critical gaps of knowledge related to these cells. The crucial role of pericytes in the maintenance of a healthy blood-brain barrier determines the paramount necessity to better understand their involvement in disease processes. Such knowledge is important because a high degree of plasticity of pericytes widens their range of responses to disruption of the neurovascular unit. While the data obtained from patients harbouring HIV-1 might potentially be extrapolated to other conditions affecting the blood-brain barrier, little is known about the involvement of pericytes in other neuroinfections. Emerging experimental and clinical evidence indicate a crucial role of pericytes in accelerated ageing, Alzheimer's disease, cerebrovascular diseases, and neurocognitive disorders (Meir-Shafrir and Pollack, 2012; Gross et al., 2016; Rajasuriar et al., 2017).

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### **Competing interests**

The authors report no competing interests.

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