



Cathepsin K Knockout Exacerbates Haemorrhagic Transformation Induced by Recombinant Tissue Plasminogen Activator After Focal Cerebral Ischaemia in Mice

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Abstract

Severe haemorrhagic transformation (HT), a common complication of recombinant tissue plasminogen activator (rtPA) treatment, predicts poor clinical outcomes in acute ischaemic stroke. The search for agents to mitigate this effect includes investigating biomolecules involved in neovascularization. This study examines the role of Cathepsin K (Ctsk) in rtPA-induced HT after focal cerebral ischaemia in mice. After knockout of *Ctsk*, the gene encoding Ctsk, the outcomes of *Ctsk*^{+/+} and *Ctsk*^{-/-} mice were compared 24 h after rtPA-treated cerebral ischaemia with respect to HT severity, neurological deficits, brain oedema, infarct volume, number of apoptotic neurons and activated microglia/macrophage, blood–brain barrier integrity, vascular endothelial growth factor (VEGF) expression and Akt-mTOR pathway activation. We observed that haemoglobin levels, brain oedema and infarct volume were significantly greater and resulted in more severe neurological deficits in *Ctsk*^{-/-} than in *Ctsk*^{+/+} mice. Consistent with our hypothesis, the number of NeuN-positive neurons was lower and the number of TUNEL-positive apoptotic neurons and activated microglia/macrophage was higher in *Ctsk*^{-/-} than in *Ctsk*^{+/+} mice. *Ctsk* knockout mice exhibited more severe blood–brain barrier (BBB) disruption, with microvascular endothelial cells exhibiting greater VEGF expression and lower ratios of phospho-Akt/Akt and phospho-mTOR/mTOR than in *Ctsk*^{+/+} mice. This study is the first to provide molecular insights into *Ctsk*-regulated HT after cerebral ischaemia, suggesting that *Ctsk* deficiency may disrupt the BBB via Akt/mTOR/VEGF signalling, resulting in neurological deficits and neuron apoptosis. *Ctsk* administration has the potential as a novel modality for improving the safety of rtPA treatment following stroke.

Keywords Cathepsin K · Haemorrhagic transformation · Ischaemic stroke · Akt · mTOR · VEGF

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Introduction

Stroke is a leading cause of morbidity and mortality in developed countries, with an immense number of medical care resources devoted to combatting its debilitating consequences. Reperfusion with the thrombolytic agent recombinant tissue plasminogen activator (rtPA) remains the standard treatment for acute ischaemic stroke (AIS) (Morris 2008; Wardlaw et al. 2013). However, the use of rtPA is severely limited by its association with haemorrhagic transformation (HT), the most significant cause of morbidity and mortality in AIS patients (Lansberg et al. 2007; Pena et al. 2017). Indeed, less than 5% of all AIS patients are eligible for rtPA treatment (Khurana et al. 2017). The development of a new strategy for attenuating the severity of HT would significantly increase the overall efficacy of rtPA thrombolytic therapy, ultimately improving the outcomes of AIS patients.

HT is caused by the breakdown of the blood–brain barrier (BBB) after vessel reperfusion following stroke (Sumii and Lo 2002). Further rupture of the BBB damages the entire neurovascular unit, including the extracellular matrix, endothelial cells, astrocytes, neurons and pericytes (Wang and Lo 2003). While BBB breakdown can occur in the absence of intervention (Hong et al. 2014; Jickling et al. 2014; Nour et al. 2013), treatment with rtPA can aggravate BBB disruption and increase the risk of HT (Wang et al. 2015). While the precise mechanism whereby rtPA induces HT is still unclear, studies implicate multiple factors, including tissue-remodelling proteases (Khurana et al. 2017). Among these proteases are cathepsins, a family of lysosomal enzymes involved in vascular remodelling, angiogenesis and atherosclerosis (Bohley and Seglen 1992). In response to disturbed blood flow, cathepsins are activated to remodel the extracellular matrix and modify arterial walls (Platt and Shockey 2016). Hypoxia also has been observed to induce the activation of cathepsins that degrade the vascular matrix (Fukuda et al. 2004). Thus, the involvement of cathepsins in reperfusion injury and HT after stroke is of interest.

Cathepsin K (Ctsk), an essential enzyme in extracellular matrix metabolism, possesses both type I collagenase and elastase activity (Novinec et al. 2007). Highly expressed in osteoclasts, Ctsk is implicated in the pathogenesis of osteoporosis and other bone diseases. In addition, the expression of Ctsk, the gene encoding Ctsk, is upregulated in several vascular diseases, including arteriosclerosis and aortic abdominal aneurysms. An imbalance between Ctsk and its inhibitor is suspected of promoting cerebral aneurysm rupture via excessive breakdown of the extracellular matrix in arterial walls (Aoki et al. 2008). Ctsk is expressed in the human brain (Bernstein et al. 2007), and

Ctsk activity is important in the development and maintenance of the CNS in mice (Dauth et al. 2011, 2012). In response to hypoxia, Ctsk was observed to activate Notch1, a protein critical to angiogenesis, resulting in neovascularisation after femoral artery ligation-induced ischaemia in mice (Jiang et al. 2014). Together, these observations warrant investigation of the involvement of Ctsk in vessel remodelling after stroke.

This study investigates whether Ctsk plays a role in rtPA-induced HT. We compare Ctsk knockout and wild-type mice after rtPA treatment of experimentally induced cerebral ischaemia with respect to the severity of HT, neurological deficits, brain oedema and infarct volume, neuron and IBA-1-positive cell expression, BBB integrity and permeability and Akt/mTOR/VEGF signalling pathway activity.

Materials and Methods

Animals

Ctsk knockout (Ctsk^{-/-}) mice were generated and bred as previously described (Saftig et al. 1998). Ctsk knockout mice survive and are fertile; they manifest no overt phenotypic abnormalities until the age of 10 months (Saftig et al. 1998). The suppressed growth seen in pycnodysostosis patients was not observed in cathepsin K-deficient mice (Saftig et al. 1998). A total of 32 Ctsk^{-/-} and 32 wild-type (Ctsk^{+/+}) adult male mice (8–10 weeks old) were used in the cerebral ischaemia and rtPA-induced HT mouse model. In addition, 32 wild-type adult male mice (8–10 weeks old) were used as the sham-surgery group. The experiments were done in a blinded manner. Mice were selected by one experimenter and no other experimenters were informed about the species and grouping of mice; another experimenter made the animal model and another two experimenters performed post-operative evaluation and sample collection and analysis. All mice were maintained at the Animal Center of Shanghai Ninth People's Hospital according to the Guidelines for the Care and Use of Laboratory Animals (National Institutes of Health). All experiments related to the animal studies were approved by the Institutional Animal Care and Use Committee (IACUC) at Shanghai Ninth People's Hospital Animal Center (No. HKDL2017411).

Cerebral Ischaemia and rtPA-Induced HT Mouse Model

Cerebral ischaemia was induced as described previously (Shahjouei et al. 2016). In brief, transient right middle cerebral artery occlusion (MCAO) was induced by placing a 7.0-siliconised filament through the external carotid artery. After 60 min, the filament was removed to allow reperfusion.

Mice were anaesthetised with 1.5% isoflurane in 30% oxygen. Body temperature was maintained at 37 ± 0.5 °C using a heating pad. Laser Doppler Flowmetry was used in all mice to confirm the induction of ischaemia and reperfusion. To increase the extent of cerebral haemorrhage, D-glucose (6 mL/kg at 50% wt/vol) was injected intraperitoneally 15 min before MCAO. Glucose was used because animal and human studies have shown that hyperglycaemia increases the risk of rtPA-induced cerebral haemorrhage after ischaemic stroke (Wang et al. 2013). After 50 min of MCAO, rtPA (10 mg/kg) was administered as an intravenous bolus injection via femoral vein (1 mg/kg) followed by a 9-mg/kg infusion administered with a syringe infusion pump (World precision Instruments) for 30 min.

To examine the influence of Ctsk on rtPA-induced HT after cerebral ischaemia, MCAO and HT were induced in wild-type (Ctsk^{+/+}/MCAO/rtPA) and Ctsk^{-/-} (Ctsk^{-/-}/MCAO/rtPA) mice. The sham-surgery group was subjected to the same surgical procedure but without the filament thread insertion.

Quantification of Cerebral Haemorrhage

At 24 h after MCAO, the mice in all three groups ($n = 8$ each) were killed by overdose of chloral hydrate and perfused transcardially with ice-cold PBS. The haemoglobin content was determined to assess the severity of haemorrhage. Drabkin reagent (500 µL) was added to the ischaemic brain tissue of each mouse, followed by homogenisation and centrifugation at 13,000 rpm for 30 min. The supernatant optical density was measured by spectrophotometry (Thermo BioMate 3S; Thermo Scientific) at 540 nm (Wang et al. 2013).

Neurologic Deficit Score

At 24 h after MCAO, the three groups of mice ($n = 12$ each) were examined and scored blindly for neurologic deficits as follows: 0—no neurological deficit; 1—failure to fully extend left forepaw; 2—reduced resistance to lateral push; 3—spontaneous circling to left; 4—absence of spontaneous movement or unconsciousness (Wang et al. 2013).

Brain Infarct Volume and Oedema Volume

The infarct volume was determined 24 h after MCAO. The brains of the mice in all three groups ($n = 6$ each) were quickly removed, sectioned coronally at 2-mm intervals and stained by immersion in the vital dye 2% 2,3,5-triphenyltetrazolium hydrochloride at 37 °C for 30 min. The extents of the normal and infarcted areas were analysed using ImageJ (National Institutes of Health, Bethesda, MD, USA) and determined indirectly as follows: Infarct

volume = (contralateral hemisphere volume – non-ischaemic ipsilateral hemisphere volume)/contralateral hemisphere volume $\times 100$; Oedema volume = (ipsilateral hemisphere volume – contralateral hemisphere volume)/contralateral hemisphere volume $\times 100$.

Evaluation of BBB Permeability

At 24 h after MCAO, the mice in all three groups ($n = 6$ each) were injected intravenously with 2% Evans blue dye (2 mL/kg body weight) and allowed to circulate for 60 min. Animals were perfused transcardially with PBS, and their brains were removed and placed in formamide for 72 h. The amount of extravasated dye was evaluated by spectrophotometry (Thermo BioMate 3S; Thermo Scientific, Waltham, MA) at 620 nm.

Western Blot Analysis

Ischaemic hemispheric brain tissues and matching tissues from sham-surgery mice ($n = 6$) were dissected. The brain tissues were homogenised in RIPA lysis buffer including a protease inhibitor cocktail (Roche Diagnostics, Mannheim, Germany). After centrifugation, the protein concentration was determined by BCA protein assay. Equal amounts of protein were loaded onto a 10% Tris–glycine gel, electrophoresed and transferred onto polyvinylidene difluoride membranes. Membranes were blocked with 5% non-fat dry milk in Tris-buffered saline with 0.1% Tween-20 and probed with primary antibodies, followed by incubation at 4 °C overnight with primary antibodies at the following dilution: NeuN (rabbit monoclonal antibody, MAB377, Millipore, Massachusetts, USA), 1:1000; Occludin (rabbit monoclonal antibody, ab167161, Abcam, Cambridge, UK), 1:1000; Claudin-5 (rabbit monoclonal antibody, 35-2500, Thermo Fisher Scientific), 1:1000; VEGF (rabbit monoclonal antibody, ab1316, Abcam), 1:1000; Akt (rabbit polyclonal antibody, ab8805, Abcam), 1:1000; p-Akt (rabbit polyclonal antibody, ab66138, Abcam), 1:1000; mTOR (rabbit polyclonal antibody, ab2732, Abcam), 1:1000; p-mTOR (rabbit monoclonal antibody, ab109268, Abcam), 1:1000 and beta-actin (rabbit monoclonal antibody, #4970, Cell Signaling Technology, Beverly, MA), 1:1000. For each primary antibody, the appropriate horseradish peroxidase-conjugated secondary antibody was used. The target bands were visualised using enhanced chemiluminescence reagent and quantified by scanning densitometry using Bio-Image Analysis System. The intensity of target protein bands was standardised to the intensity of β -actin bands.

Immunofluorescence Staining

Twenty-four hours after MCAO, the three groups of mice ($n=6$ each) were killed by overdose of chloral hydrate and perfused transcardially with ice-cold PBS and 4% paraformaldehyde. The brains were removed and cryoprotected in 30% sucrose in PBS. Brain cryosections (20 μm) were blocked with 1% bovine serum albumin and 5% normal donkey serum in PBS and incubated with anti-NeuN (rabbit monoclonal antibody, MAB377, Millipore, Massachusetts, USA) or anti-IBA-1 (rabbit monoclonal antibody, ab178847, Abcam) or a combination of anti-VEGF (rabbit monoclonal antibody, ab1316, Abcam) and anti-CD31 antibodies (rat monoclonal antibody, PECAM-1, BD Pharmingen) overnight at 4 °C. The sections were then rinsed and incubated with Alexa Fluor 594 goat anti-rabbit IgG (#8889, Cell Signaling Technology) or Alexa Fluor 488 goat anti-rabbit IgG (#4412, Cell Signaling Technology), or a mixture of Alexa Fluor 594 goat anti-rabbit IgG and Alexa Fluor 488 donkey anti-rat IgG (A21208, Invitrogen) for 1 h at room temperature. Images were obtained using a Leica microscope. For the quantification of NeuN-positive cells and IBA-1-positive cells and the area of double-positive CD31 and VEGF cells, 5 non-overlapping images in the peri-infarct area were digitised using a $\times 40$ objective, two of them were located in the cortex and three in the basal ganglia. The images were selected by a researcher who did not know the grouping information. The images were processed by NIH ImageJ and contrast enhanced to clearly differentiate positivity from background. For each brain, three sections from the ischaemic hemisphere were analysed.

TUNEL Assay

Apoptotic cells were identified by TUNEL assay (11684817910, Roche Diagnostics, Mannheim, Germany), and the total cell number was counted after DAPI (4',6-diamidino-2-phenylindole) staining. Coronal cryosections from at least five mice in each group were obtained. Five random fields in the peri-infarct area of each section were digitised using a $\times 40$ objective. The results are expressed as the percentage of TUNEL⁺/DAPI⁺ cells in the section.

Statistical Analysis

All continuous variables are presented in bar graphs (mean \pm SEM). For each figure, the error bars indicated SEM. Differences were assessed using Student's *t* test for 2-group comparisons. Ranked data (neurological deficit scores) are presented as median (interquartile range, IQR) and in box-plot figures, and non-parameter Mann–Whitney U test is used for 2-group comparisons. Two-tailed $p < 0.05$ was considered statistically significant. All graphing and analyses were performed using SAS 9.4 software (SAS Institute Inc., Cary, NC, USA.).

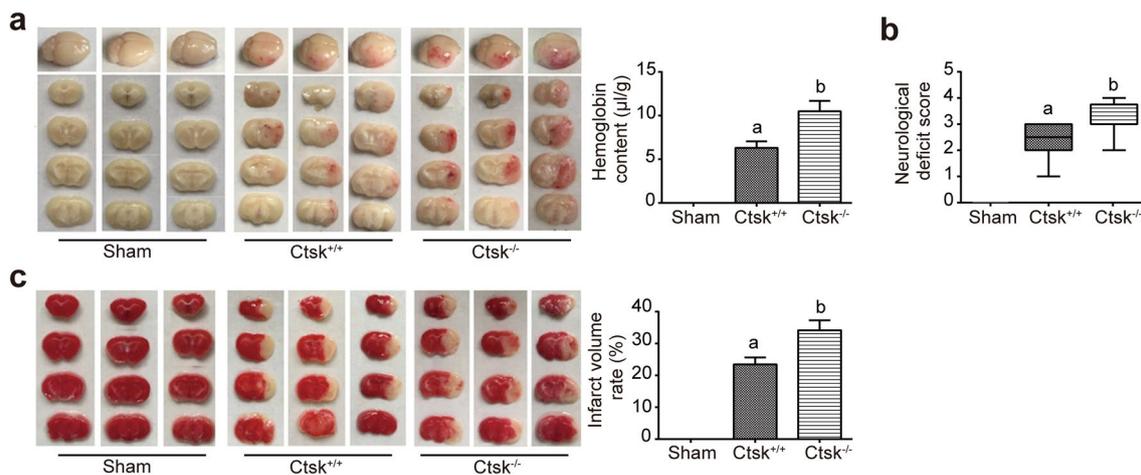


Fig. 1 Greater degree of HT and worse outcomes in Ctsk^{-/-}/MCAO/rtPA mice. **a** *Left panel*, images of whole brain sections of mice from each group. *Right panel*, haemoglobin content was greater in Ctsk^{-/-}/MCAO/rtPA (Ctsk^{-/-}) mice than in Ctsk^{+/+}/MCAO/rtPA (Ctsk^{+/+}) mice. **b** Ctsk^{-/-}/MCAO/rtPA mice showed more severe neurologic

deficits than Ctsk^{+/+}/MCAO/rtPA mice. **c** *Left panel* TTC staining of HT in brain sections from mice of each group. *Right panel* larger infarct volume observed in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice. ^a $p < 0.05$, sham versus Ctsk^{+/+}; ^b $p < 0.05$, Ctsk^{-/-} versus Ctsk^{+/+}

Results

Ctsk Deficiency Increased the Neurological Deficits and Infarct Volume in MCAO/rtPA-Induced HT Mice

To examine the influence of Ctsk expression on rtPA-induced HT after cerebral ischaemia, the degree of HT after cerebral ischaemia (Fig. 1a) was compared between Ctsk^{+/+} and Ctsk^{-/-} mice. Spectrophotometric haemoglobin assay showed that haemoglobin concentrations were significantly higher in Ctsk^{-/-}/MCAO/rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 1a) at 24 h after stroke. Ctsk^{-/-}/MCAO/rtPA mice showed more severe neurological deficits as assessed by neurological scores than did Ctsk^{+/+}/MCAO/rtPA mice (Fig. 1b). Analysis of TTC-stained brain sections showed that the infarct volume was significantly larger in Ctsk^{-/-}/MCAO/rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 1c).

Ctsk Knockout Increased the Number of Apoptotic Neurons and Activated Microglia/Macrophage in MCAO/rtPA-Induced HT Mice

To study the role of Ctsk in neuron expression and microglia/macrophage activation, the expression of NeuN and IBA-1 in the peri-infarct area was determined. Ctsk^{+/+}/MCAO/rtPA mice had significantly fewer NeuN-positive cells and lower NeuN protein expression levels than did sham-surgery mice (Fig. 2a, b). However, Ctsk^{+/+}/MCAO/rtPA mice had more NeuN-positive cells and greater NeuN protein expression than did Ctsk^{-/-}/MCAO/rtPA mice (Fig. 2a, b). The number of TUNEL-positive cells was significantly greater in Ctsk^{+/+}/MCAO/rtPA mice than in sham-surgery mice, with more TUNEL-positive cells in Ctsk^{-/-}/MCAO/rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 2c). Immunostaining revealed that the number of IBA-1-positive cells (microglia cells/macrophages) was significantly higher in Ctsk^{+/+}/MCAO/rtPA mice than in sham-surgery mice, with significantly more IBA-1-positive cells in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 2d).

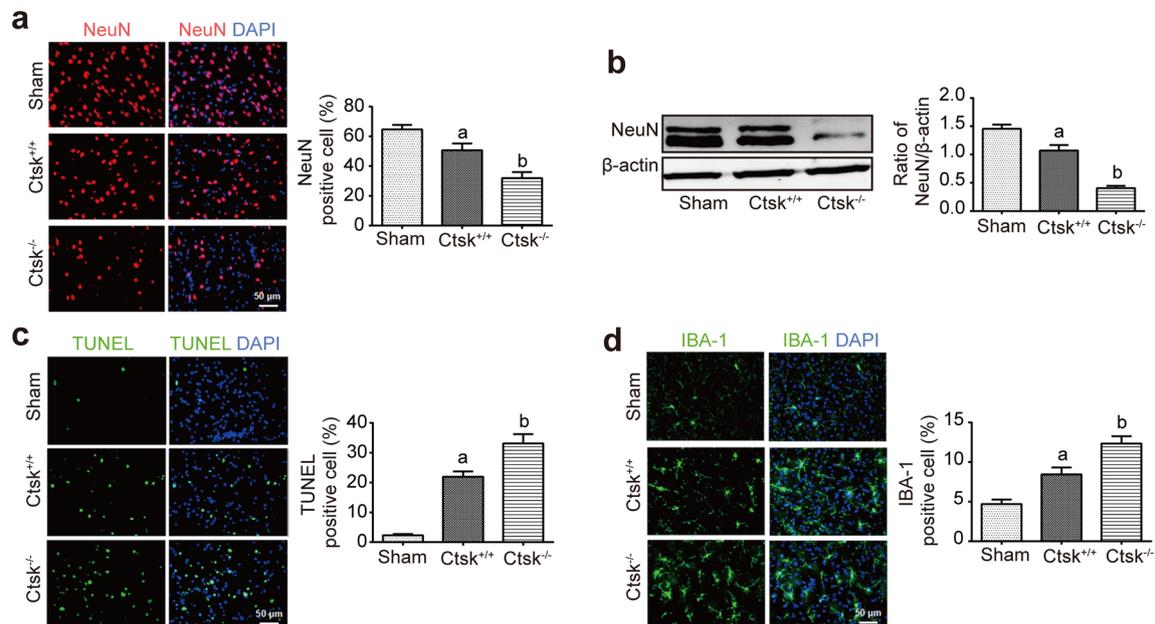


Fig. 2 Neuron damage and IBA-1 positive cell activation elevated in Ctsk^{-/-}/MCAO/rtPA mice. **a** Comparison of brain coronal sections immunostained with anti-NeuN (red) and DAPI (blue). Neurons were identified as NeuN⁺ and DAPI⁺ cells in brain sections. The number of neurons was lower in Ctsk^{+/+}/MCAO/rtPA (Ctsk^{+/+}) mice compared with the sham-surgery group and lower in Ctsk^{-/-}/MCAO/rtPA than Ctsk^{+/+}/MCAO/rtPA mice. **b** Western blot of NeuN and β-actin loading control in ischaemic hemispheres lysates from mice of different groups. Quantification of NeuN from western blots shows lower expression in Ctsk^{+/+}/MCAO/rtPA mice than in sham mice, with lower expression in Ctsk^{-/-}/MCAO/rtPA than Ctsk^{+/+}/MCAO/rtPA mice. **c** The number of apoptotic cells was determined by counting

TUNEL+ (green) and DAPI+ (blue) cells. The number of apoptotic cells was higher in Ctsk^{+/+}/MCAO/rtPA mice than in sham-surgery mice, with more apoptotic cells in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice. **d** Comparison of brain coronal sections immunofluorescently stained with anti-IBA (green) and DAPI (blue) between mice of different groups. Activated microglia or microphage were identified as IBA⁺/DAPI⁺ cells in brain sections. The number of activated microglia or microphage was greater in Ctsk^{+/+}/MCAO/rtPA mice than in sham-surgery mice, with more activated microglia or microphage in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice. ^a*p* < 0.05, sham versus Ctsk^{+/+}; ^b*p* < 0.05, Ctsk^{-/-} versus Ctsk^{+/+}

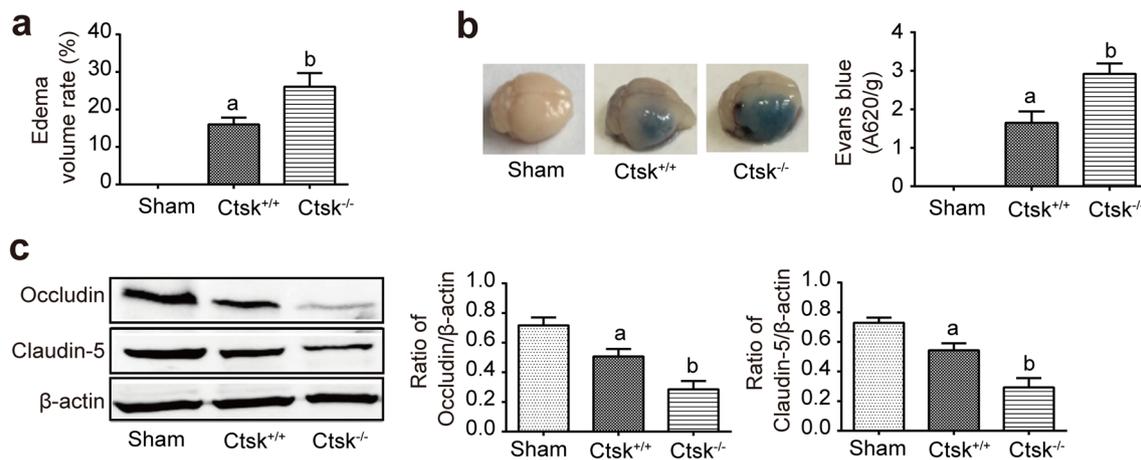


Fig. 3 Greater BBB disruption in Ctsk^{-/-}/MCAO/rtPA mice. **a** The oedema volume was greater in Ctsk^{-/-}/MCAO/rtPA (Ctsk^{-/-}) than in Ctsk^{+/+}/MCAO/rtPA (Ctsk^{+/+}) mice, as evidenced by TTC staining. **b** *Left panel* images of whole brains from mice of different groups after EB treatment. *Right panel* significantly more extravasated EB dye was seen in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice.

c Expression of occludin, claudin-5 and β-actin loading control in the ischaemic hemisphere lysates from mice of different groups were analysed by western blot. Occludin and claudin-5 expression was higher in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice. ^a*p* < 0.05, sham versus Ctsk^{+/+}; ^b*p* < 0.05, Ctsk^{-/-} versus Ctsk^{+/+}

Ctsk Knockout Disrupts BBB Integrity and Permeability in MCAO/rtPA-Induced HT Mice

The oedema volume was greater in Ctsk^{-/-}/MCAO/rtPA than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 3a). Studies of BBB function, as evaluated using the Evans blue assay 24 h after MCAO, showed marked extravasation of EB dye in the ischaemic hemispheres of Ctsk^{+/+}/MCAO/rtPA mice as compared to that of sham-surgery mice, indicating disruption of the BBB. Significantly more extravasated EB dye was seen in Ctsk^{-/-}/MCAO/rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 3b), indicating that Ctsk deficiency aggravates BBB disruption after cerebral ischaemia.

Western blot analysis of ischaemic brain conducted 24 h after MCAO indicates that expression of the tight junction proteins occludin and claudin-5, essential for maintaining BBB integrity, was suppressed in Ctsk^{+/+}/MCAO/rtPA mice as compared to sham-surgery mice. Ctsk^{-/-}/MCAO/rtPA mice expressed much less occludin and claudin-5 than did Ctsk^{+/+}/MCAO/rtPA mice (Fig. 3c).

Ctsk Knockout Induces VEGF Expression in MCAO/rtPA-Induced HT Mice and the Expression of Akt/mTOR Pathway

Because increased VEGF expression is associated with BBB leakage in the acute ischaemic brain, we examined VEGF expression in rtPA-induced HT after MCAO. Western blot analysis revealed that cerebral ischaemia significantly increased VEGF expression over that of sham-surgery mice and that VEGF expression was higher in Ctsk^{-/-}/MCAO/

rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 4a). Double immunostaining revealed that VEGF primarily colocalised with CD31-positive blood vessels at 24 h after stroke. Consistent with the western blot results, VEGF immunoreactivity in CD31-positive vascular endothelium in the peri-infarct area was elevated in Ctsk^{-/-}/MCAO/rtPA mice as compared to Ctsk^{+/+}/MCAO/rtPA mice (Fig. 4b).

To test the expression of Akt-TOR pathway, we compared the expression of Akt, p-Akt, mTOR and p-mTOR between the different mice. Western blot analysis showed that cerebral ischaemia reduced Akt phosphorylation (p-Akt/Akt) and mTOR phosphorylation (p-mTOR/mTOR). This effect was significantly greater in Ctsk^{-/-}/MCAO/rtPA mice than in Ctsk^{+/+}/MCAO/rtPA mice (Fig. 4c, d).

Discussion

rtPA, the only approved drug for the treatment of AIS, has provided excellent outcomes for many patients but imposes an increased risk of HT, a life-threatening complication. Here we observed greater HT, cerebral oedema, BBB disruption, neurological deficits and infarct volume in Ctsk knockout mice than in wild-type mice. These results suggest that Ctsk protects against rtPA-induced HT after cerebral ischaemia.

Although the precise mechanism whereby rtPA induces HT after ischaemia remains elusive, experimental and clinical evidence indicates that neurovascular unit disruption is likely a major contributor (Fagan et al. 2004; Wang and Lo 2003). The BBB is a vital component of the neurovascular

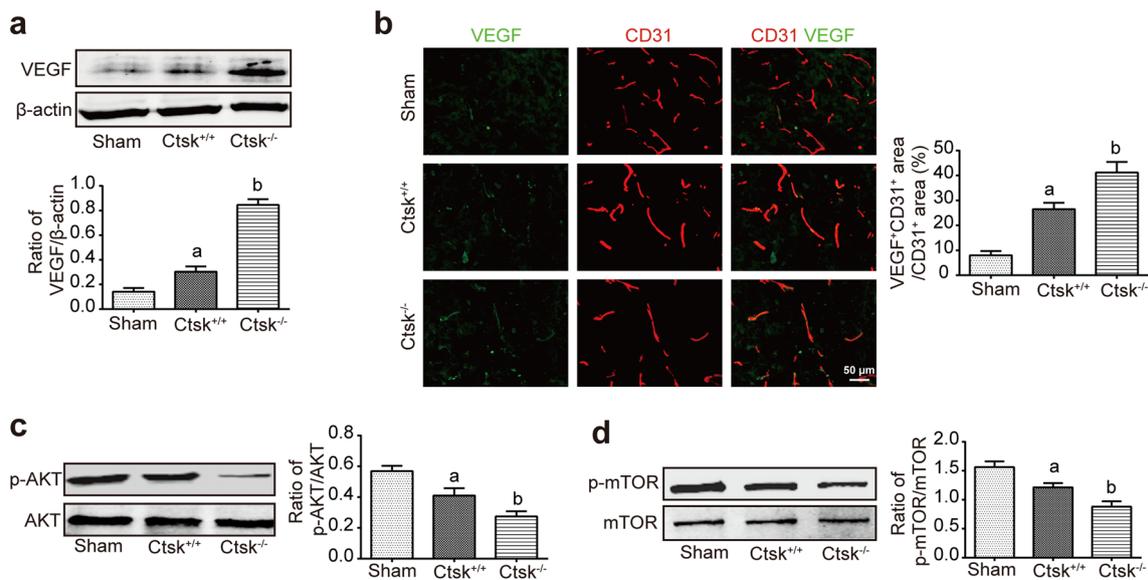


Fig. 4 Elevated VEGF expression in $Ctsk^{-/-}$ /MCAO/rtpA mice via Akt/mTOR pathway activation. **a** Western blot analysis of VEGF expression in ischaemic hemisphere lysates from mice of different groups. VEGF expression was greater in $Ctsk^{+/+}$ /MCAO/rtpA ($Ctsk^{+/+}$) mice than in sham-surgery mice, with the expression level greater in $Ctsk^{-/-}$ /MCAO/rtpA ($Ctsk^{-/-}$) than in $Ctsk^{+/+}$ /MCAO/rtpA mice. **b** Double immunostaining of mouse brain coronal sections with anti-VEGF (green) and anti-CD31 (red). The number of VEGF immunoreactive cells in vascular endothelium was determined

by counting VEGF $^{+}$ /CD31 $^{+}$ cells in brain sections. The number of VEGF $^{+}$ /CD31 $^{+}$ cells was higher in $Ctsk^{+/+}$ /MCAO/rtpA than in sham-surgery mice, with more VEGF $^{+}$ /CD31 $^{+}$ cells in $Ctsk^{-/-}$ /MCAO/rtpA than in $Ctsk^{+/+}$ /MCAO/rtpA mice. **c**, **d** Western blot of Akt, p-Akt, mTOR and p-mTOR in ischaemic hemisphere lysates from mice of different groups. p-Akt/Akt and p-mTOR/mTOR ratios were higher in $Ctsk^{+/+}$ /MCAO/rtpA than in sham-surgery mice, and lower in $Ctsk^{-/-}$ /MCAO/rtpA than in $Ctsk^{+/+}$ /MCAO/rtpA mice. $^a p < 0.05$, sham versus $Ctsk^{+/+}$; $^b p < 0.05$, $Ctsk^{-/-}$ versus $Ctsk^{+/+}$

unit, containing tight junctional (TJ) proteins and transporters of ions and nutrients that maintain normal brain physiology. Ischaemic stroke-induced BBB breakdown promotes complications such as cerebral oedema and HT, especially in association with therapeutic recanalisation of occluded vessels (Horsch et al. 2015; Kono et al. 2014). Further rupture of the BBB damages the entire neurovascular unit, including the extracellular matrix, endothelial cells, astrocytes, neurons and pericytes. The involvement of *Ctsk* in protecting against this process is supported by our observation that *Ctsk* knockout mice exhibited more brain oedema, greater BBB disruption and greater numbers of apoptotic neurons and IBA-1 positive cell than did $Ctsk^{+/+}$ /MCAO/rtpA mice. Thus, *Ctsk* deficiency might cause greater HT by altering the neurovascular unit and disrupting the BBB. Our findings are consistent with those of Jiang et al., who report that functional recovery after femoral artery ligation-induced ischaemia is impaired in *Ctsk*-deficient mice (Jiang et al. 2014). These authors observed that in response to hypoxia, *Ctsk* activates Notch-1, leading to downstream activation of the Akt/mTOR/VEGF pathway. Thus, the absence of such signalling is implicated in the greater degree of HT observed in *Ctsk*-deficient mice.

As a potent mediator of endothelial proliferation and migration, VEGF is associated with angiogenesis,

neurogenesis, axon plasticity, neuron survival and vascular permeability. VEGF also plays an important role in the development of brain oedema during ischaemic stroke (Greenberg and Jin 2013; Zhang et al. 2017b). VEGF expression is upregulated in peri-infarct regions within 3 h of the onset of ischaemia (Hayashi et al. 1997). Early administration of VEGF to ischaemic rat brains is reported to increase BBB leakage, induce HT, increase the infarct size in the brain (Zhang et al. 2000), as well as induce the apoptotic death of neurons. In rats, inhibition of the VEGF signalling pathway was observed to attenuate HT after rtpA treatment (Kanazawa et al. 2011). In the absence of *Ctsk*, we observed elevated levels of VEGF after rtpA treatment of MCAO mice. Thus, we propose that *Ctsk* may protect against HT by inhibiting VEGF expression.

VEGF expression is regulated by the Akt/mTOR signalling pathway (Park et al. 2016; Sarnelli et al. 2016; Zeng et al. 2017; Zhang et al. 2017a). Studies suggest that the degree and duration of mTOR signalling may be an essential factor in neuroprotection (Maiese et al. 2013). mTOR activation can foster long-term potentiation and synaptic plasticity in models of Alzheimer's disease (Ma et al. 2010) and block inflammatory cell death during toxic β -amyloid exposure (Shang et al. 2012). However, sustained activation of mTOR can result in impaired neuronal

stem cell maturation (Uzdensky et al. 2013) and clinical disability with dyskinesia in Parkinson's disease patients (Chong et al. 2011). Prolonged inhibition of mTOR can lead to neuron death (Jessen et al. 2010). Consistent with these findings, we observed that Ctsk knockout results in significantly higher VEGF expression and significantly lower levels of phospho-Akt and phospho-mTOR compared to wild-type mice. These findings suggest that suppression of Akt/mTOR signalling due to the absence of Ctsk induces apoptotic cell death in the peri-infarct area of Ctsk^{-/-}/MCAO/rtPA mice, disrupting BBB integrity and altering its permeability. In response, VEGF expression increases, leading to apoptotic cell death via VEGF/VEGFR2 signalling.

This study has several limitations. The extent and mechanism underlying the protective effect of Ctsk is unclear and requires further study. Determining the mechanism whereby Ctsk affects Akt-mTOR-VEGF signalling will require studies using pathway inhibitors. The relevance of these findings to clinical applications must be determined by examining the expression of Ctsk in AIS patients and investigating the relationship between Ctsk expression level and the occurrence of rtPA-induced HT.

In conclusion, Ctsk knockout resulted in damage to the neurovascular unit and induced BBB breakdown, likely by affecting the Akt-mTOR-VEGF signalling cascade. These results suggest that Ctsk administration has the potential as a novel modality for improving the safety of rtPA following stroke.

Author Contributions RZ, XWH, YHS and YSL performed the major experiments and wrote the draft manuscript. FDL, YH, MTZ, XYF, LZ, BQZ and HQL performed the experiments and acquired the data, participated in drafting of the manuscript, created the figures and revised the manuscript. GPS and JRL designed the study and revised the manuscript for important intellectual content. All authors read and approved the final manuscript.

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Compliance with Ethical Standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical Approval All applicable international, national and/or institutional guidelines for the care and use of animals were followed.

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