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**Extracellular vesicles derived from
adipose mesenchymal
stem cells (AMSCs) reduce autophagic
flux via microRNA-25-3p transfer after
ischemic stroke**

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Abbreviations

3-MA	3-Methyl-adenin
ANOVA	Analysis of variance
AMSC	adipose mesenchymal stem cells
BafA1	bafilomycin A1
BMSC	bone marrow-derived MSCs
BNIP3	B-cell lymphoma 2–interacting protein 3
CNS	central nervous system
DAPI	4',6-diamidino-2-phenylindole
DNA	deoxyribonucleic acid
EVs	extracellular vesicles
Hrs	Hepatocyte growth factor–associated tyrosine kinase
LC3	Microtubule-associated protein 1A/1B-light chain 3
MCAO	middle cerebral artery occlusion
MSC	mesenchymal stem cells
MTT	3-(4,5-Dimethylthiazol-2-yl)-2,5-Diphenyltetrazolium Bromide
MVs	microvesicles
NTA	nanoparticle tracking analysis
OGD	oxygen-glucose-deprivation
PEG	polyethylene glycol
qRT-PCR	Real-time polymerase chain reaction
RNA	Ribonucleic Acid
TEM	transmission electron microscopy
TTC	2,3,5-Triphenyltetrazolium Chloride
PBS	phosphate-buffered saline
DMEM	Dulbecco's Modified Eagle Medium
CM	conditioned medium

1 Introduction

1.1 Epidemiological and clinical aspects of ischemic stroke

Stroke is the second leading cause of death and the largest increase in deaths since 2000 according to data obtained from the World Health Organization. The causes of stroke can be divided into two types: hemorrhagic strokes and ischemic strokes. Stroke is a disease of the elderly in which one or more risk factors such as hypertension, hyperlipidemia, carotid stenosis, and smoking are present (Boehme et al. 2017). Ischemic strokes can be further differentiated according to their pathophysiology, in which there is embolism or thrombus formation as well as generalized hypoperfusion. However, local thrombus formation is the most common type of ischemic stroke. About 85 % of strokes are ischemic strokes, which are clinically more significant than hemorrhagic strokes (Musuka et al. 2015). Therefore, the current thesis focuses on ischemic stroke only.

Ischemic stroke is a cerebrovascular disease caused by an acute loss of blood supply to the brain. Since the brain relies upon constant blood supply, cell injury begins with minutes upon stroke onset. Significant advances in the treatment of ischemic stroke have been made in the last few decades thanks to the establishment of systemic thrombolysis within 4.5 hours and endovascular therapy (Harder and Klinkhardt 2000). However, most patients do not qualify for either treatment modality due to a limited time window or significant side effects (Seitz 2016). As a result, many patients have a poor quality of life after stroke and are in urgent need of different treatment paradigms. Although neuroregenerative approaches such as mesenchymal stem cell (MSC) transplantation have been successfully applied in preclinical stroke models, they have not been successfully translated into the clinic until recently (Bang et al. 2005; Gladstone et al. 2002). New strategies for stroke treatment are therefore urgently needed.

1.2 Mesenchymal stem cell (MSC)-based experimental stroke therapy is mediated by extracellular vesicles (EVs)

MSCs are a class of stem cells with self-renewal potential and multidirectional differentiation capacity (Andrzejewska et al. 2019). They display rapid proliferation rates, are easy to culture *in vitro*, and show low immunogenicity (Kim HJ and Park 2017). Previous studies have demonstrated neuroprotection and increased neurological recovery *in vivo* after MSC transplantation in rodent ischemic stroke models (Chen J et al. 2001; Gervois et al. 2016; Maria Ferri et al. 2016). Notably, transplanted MSCs are not integrated into residing neural networks but work indirectly (Baraniak and McDevitt 2010; Liang et al. 2014).

Recently, lipid bilayer-structured particles that are naturally released from cells were defined as extracellular vesicles (EVs). EVs are considered to be an important mediator for cell-cell communication since the first report (Tetta et al. 2013). It was secreted by the vast majority of endocytic types in humans and carried a large amount of proteins, lipids, and nucleic acids (Zaborowski et al. 2015). EVs range in diameter from 30 nm to 1000 nm (most of them are less than 200 nm) and carry a cargo of proteins, lipids, mRNAs, DNAs, and even organelles from the parent cell (Kim DK et al. 2013; Mathivanan et al. 2012; Theyry et al. 2018). Indeed, secreted EVs containing biologically active molecules are thought to transduce specific intracellular signaling in physiological or pathological conditions and mediate the transfer of biological information by fusing and pouring contents such as microRNA directly into the recipient cell (Russell et al. 2019). Previous work of my research group demonstrated that EVs obtained from conditioned medium of cultured MSCs under standard cell culture conditions equally increased long-term neuronal survival and reduced motor-coordination deficits when administered intravenously 24 hours after middle cerebral artery occlusion (MCAO). The therapeutic potential of these EVs derived from MSCs does not appear to be inferior compared to that of the host cells (Doepfner et al. 2015). This raises the questions of how EVs might actually work

and what the neuroprotective cargo inside these MSC-EVs might be that transfers the neuroprotective effect under stroke conditions.

1.3 Key substances in EVs

EVs are small membrane vesicles secreted by almost all kinds of cells found in approximately every fluid compartment of the body like saliva, cerebrospinal fluid, urine, and blood (Doyle and Wang 2019). EVs affect signal cascades and biological behavior of target cells by transferring a large number of bioactive molecules including proteins, RNAs, small non-coding RNAs, or DNAs to the recipient cells (Yanez-Mo et al. 2015). As a crucial means of intercellular communication, plenty of reports came up with the idea that the biological behavior of cardiovascular, cerebrovascular, and metabolic diseases are highly influenced by surrounding EVs, in particular, by microRNAs located inside EVs (O'Brien J et al. 2018; Pfeifer et al. 2015; Yoshikawa et al. 2019). Increasing evidence points towards microRNA-containing EVs that may lead to altered phenotypes and protein expression patterns of recipient cells under pathological conditions (O'Neill et al. 2019). The notion that EVs might work as a carrier being able to regulate gene expression through genetic information transfer to recipient cells opened up a new intercellular communication mechanism. There is growing evidence that the effects of EVs on recipient cells predominantly rely on the intravesicular microRNA transfer (Fernandez-Messina et al. 2015; Schwarzenbach and Gahan 2019). By transferring microRNAs to recipient cells, EVs have been established as powerful mediators of intercellular gene regulation (Maas et al. 2017).

MicroRNAs are a class of small (containing about 22 nucleotides) single-stranded non-coding molecules found in animals, plants, and some viruses (Felekakis et al. 2010). They control physiological and pathological processes by base-pairing with mRNA complementary sequences located in the untranslated or coding regions of mRNAs (Xu et al. 2018). As a result, these mRNA molecules are silenced by destabilizing of the mRNA through shortening of its poly(A)

tail, cleavage of the mRNA strand into two pieces, or less efficient translation of the mRNA into proteins by ribosomes (Lim et al. 2005; Vemuganti 2010).

MicroRNAs are evolutionarily conserved and appear to target about 70 % of the genes, which implies their fundamental biological functions. Interestingly, autophagy can be affected by microRNAs (Akkoc and Gozuacik 2020). Autophagy is a self-degrading process that is important for balancing energy sources during development and in response to nutritional stress (Glick et al. 2010). ATG1/ULK1, which belongs to the serine/threonine class of protein kinases, is the key protein kinase that initiates autophagy (Hosokawa et al. 2009). The initial stages of autophagy are the formation of autophagic membranes. The latter further expand and wrap the substrate, eventually forming autophagosomes (He and Klionsky 2009). This process requires the involvement of multiple autophagy-related proteins, and molecules affecting the transcription of autophagy-related genes lead to reduced or activated autophagy (Reggiori and Klionsky 2002). Recent work has highlighted the role of autophagy as a critical regulator of ischemic stroke (Wang P et al. 2018), revealing a new set of potential therapeutic targets for neuroprotection. Therefore, whether microRNAs in AMSC-derived EVs could play a neuroprotective role in stroke by regulating autophagy needs further exploration.

1.4 Ischemic stroke and autophagy

Autophagy is a natural and regulated process that removes dysfunctional or unnecessary intracellular components for recycling, especially under harsh survival conditions such as hypoxia, infection, and nutrient deficiencies (Yoshimori 2007). Ischemic stroke is a starvation and nutrient deficiencies situation caused by acute cerebral blood vessel occlusion. This condition will cause type II programmed cell death—also named autophagic cell death (Adhami et al. 2006).

In the rodent stroke model, the role of autophagy is controversial. Some studies suggest that suppression of excessive autophagy protects neurons against stroke-induced cell injury, while others indicate autophagy itself has a protective effect against ischemia (Carloni et al. 2010; Dai

et al. 2017; Wang P et al. 2012). It is generally believed that autophagy is a double-edged sword under conditions of ischemic stroke (Chen W et al. 2014). Moderate autophagy activation may prevent neuronal damage to a certain degree, but it may also cause neuronal death after ischemia when excessive autophagic activity occurs (Shi R et al. 2012). Due to the dual action of autophagy, the consequence of autophagy activation may vary according to the intensity and duration of ischemia in infarct brains. Therefore, controlling the post-ischemic autophagy level within a reasonable range may be an effective way to rescuing neurons from autophagic cell death.

1.5 MicroRNA and autophagy

As mentioned before, the severity and longevity of autophagic activation may determine the fate of neurons after ischemia and hypoxia. However, the level of cellular autophagy can be manipulated to some extent, but the process must be strictly controlled to prevent a lethal outcome (Mizushima et al. 2010). Substantial progress recently made contributions to our understanding of the molecular mechanisms of autophagy, discovering that microRNAs made collection to nearly all known fundamental biological pathways (Bhaskaran and Mohan 2014). Regulation of the genetic suppressors can contribute to the outcome of various diseases, including ischemic stroke. Recent research revealed that microRNA-30d-5p prevents neuronal death by inhibiting autophagy-mediated microglial activation via exosome transfer (Jiang et al. 2018). This provides a theoretical basis for my hypothesis that naturally secreted MSC-EVs regulate post-ischemic autophagy via microRNA transfer.

2 Aim of the study

The specific mechanisms underlying the neuroprotective effects of MSC-secreted EVs in stroke conditions have not been clearly elucidated to date, which is a prerequisite before clinical trials appear to be reasonable. In the present study, I therefore aimed to verify whether or not the application of AMSC-EVs under conditions of both *in vitro* hypoxia and *in vivo* cerebral ischemia affects autophagic signaling pathways. My studies focused on analyzing post-stroke autophagy related mechanisms that may contribute to neuronal survival after application of AMSC-EVs in preclinical stroke models. .

3 Materials and Methods

3.1 Adipose mesenchymal stem cell (AMSC) isolation

Adipose tissue was collected from mature wild-type C57BL/6 mice, and the tissue was mechanically minced prior to digestion with collagenase I (Gibco, Darmstadt, Germany) for 30 minutes. About 10 times the amount of collagenase I compared to the tissue was added to the tissue and incubated in a water bath at 37°C. The process was stopped by adding an equal amount of Dulbecco's modified Eagle's medium (DMEM; Merck Group, Darmstadt, Germany) containing 10 % fetal bovine serum (FBS; Merck Group, Darmstadt, Germany). After centrifugation to harvest the immature adipocytes, the cell precipitation was resuspended and cultured under standard cell culture conditions (37°C with 5 % CO₂) for two days. The cells were trypsinized for passage or cell characterization. The details about the isolation method and cell characterization can be found in Publication 1.

3.2 Preparation of primary cortical neurons

Extraction of primary cortical neurons begins with killing pregnant C57BL/6 mice by inhalation of CO₂ on embryonic day 17. The cerebral cortex of fetal mice was then dissected and subjected to 14-minute trypsin (Merck Group, Darmstadt, Germany) treatment. After digestion, the harvested cells were seeded on Poly-L-Ornithin pre-coated plates for different experimental purposes. After 4 days of incubation, the cells were used for subsequent experiments.

3.3 AMSC-EV isolation and characterization

When AMSCs reached approximately 90 % confluence after two days of culture, the high glucose medium (DMEM containing 10 % fetal bovine serum) was replaced with serum-free DMEM medium for 24 hours. The conditioned AMSC medium was collected, and EVs were enriched from the FBS-free DMEM conditioned medium (CM) using 1 hour of

ultracentrifugation (UC) (Optima XPN-80 Ultracentrifuge, BECKMAN COULTER, California, United States) at 110,000 g. EV pellets were resuspended in 200 μ l PBS and stored at -80°C . For the polyethylene glycol (PEG) precipitation method (Zheng et al. 2021), CM from AMSC was incubated with 10 % PEG 6,000 (50 % wt/vol; Merck Group, Darmstadt, Germany) for at least 12 hours after cell debris removal. The pre-enriched EVs were concentrated by centrifugation for 30 minutes at 4,500 g followed by UC for 1 hour at 110,000 g. EVs were also diluted in 200 μ l of PBS. The details of EV enrichment protocols can be found in Publication 1. For EV characterization, exosomal enriched proteins including CD9, CD63, TSG101, Alix, and negative control TOM20, albumin, and Histones were analyzed by Western blot. The details about EV characterization and Western blots can be found in Publication 1.

3.4 AMSC-EV uptake and inhibition

To verify that the isolated AMSC-EVs were taken up by cultured primary neurons and the ischemic mouse brain, the EVs were incubated with a tracker named DiI (Invitrogen, Carlsbad, USA) for 1 hour to tag the EVs and then washed with PBS. Primary neurons were incubated with the labeled EVs under hypoxic conditions for 2 hours, and the EV uptake was observed under a fluorescence microscope (Axioplan 2, Zeiss, Göttingen, Germany). To monitor the EV biodistribution *in vivo*, the labeled EVs were injected into the femoral vein of mice right after reperfusion, and the mice were sacrificed 2 hours after injection. The EV biodistribution was analyzed by immunofluorescence counterstained with 4',6-diamidino-2-phenylindole (DAPI) and NeuN⁺ in cryosections (14 μ m thick) of brains under a fluorescence microscope.

For EV inhibition studies, two approaches were selected, i.e., implied the pharmacological inhibition with GW4869 and the genetic inhibition by Hrs gene knockdown for exosomal inhibition. For GW4869 inhibition, AMSCs were cultured to 90 % confluence, and the cell culture medium was replaced with DMEM containing 10 μ M GW4869 (Sigma-Aldrich, St. Louis, MO, USA) in order to deplete EV isolation. For the genetic exosome inhibition, Hrs

siRNA and their control siRNA (RiboBio, Guangzhou, China) were transfected into AMSCs after 24 hours of seeding using Turbofect Transfection reagents (Thermo Fisher Scientific, Waltham, Massachusetts, USA). After one day of transfection, the medium was replaced with DMEM for further EV enrichment.

3.5 Oxygen-glucose-deprivation (OGD) model and cell survival assay

For the OGD model establishment, primary neurons in 6 or 24-well plates were washed with PBS once and incubated in a hypoxic chamber (less than 1 % O₂, 5 % CO₂) at 37°C in a glucose-free cell culture medium. The duration of the OGD varied according to the needs of the experiment. After the OGD, primary neurons were pre-incubated under standard cell culture conditions with a normal cell culture medium (DMEM containing 10 % FBS) for 24 hours (reoxygenation). Cell viability was determined by a colorimetric-based MTT assay (Thiazolyl Blue Tetrazolium, Sigma-Aldrich, St. Louis, MO, USA) according to the manufacturer's instructions. Cell mortality was also determined by fluorescence microscopy using the LIVE/DEAD kit (Lonza, Basel, Switzerland) according to the manufacturer's instructions.

3.6 Autophagic flux

Two methods were used for autophagic flux assessments. For autophagic flux assessment in primary neurons, cells were cultured on coverslips for 3 days and transfected with an RFP-GFP-LC3B containing plasmid using an RFP-GFP-LC3B kit (Thermo Fisher Scientific, Massachusetts, USA). One day after transfection, the cells were cultured under different experimental conditions. After all the experimental procedures, like OGD and EV treatment, primary neurons were fixed with 4 % paraformaldehyde (PFA). In three independent experiments, the number of autophagosomes and autolysosomes in each cell was quantified by fluorescence microscopy. For autophagic flux assessments by LC-3 II, primary neurons were treated with Bafilomycin A1 (Sigma-Aldrich, St. Louis, MO, USA) 3 hours before harvesting the protein. For autophagic flux assessments in mice, Bafilomycin A1 was injected

intraperitoneally 3 hours before the mice were sacrificed. Western blots were performed after the treatment.

3.7 Real-time qRT-PCR

Total RNA was isolated from primary neurons using Trizol reagent (Invitrogen, St. Louis, MO, USA), stranded cDNA synthesis (Sigma-Aldrich, St. Louis, MO, USA) followed by qRT-PCR that was performed using the SYBR[®] LC480 Kit according to manufacturer's instructions. Relative target microRNA (microRNA-98, microRNA-25-3p, microRNA-30, microRNA-125a-5p, microRNA-125b-5p, and microRNA-214-3p) expression was normalized by the expression of the internal control genes let-7a and U6. The calculation was done by using the standard $2^{-\Delta\Delta C_t}$ method.

3.8 *In vivo* experimental paradigm

Rodent experiments were performed after approval from local authorities, and EU guidelines for the care and protection of laboratory animals were strictly followed. All mice had free access to food and water. For the establishment of middle cerebral artery occlusion (MCAO), male C57BL/6 mice aged 10-12 weeks (Janvier Labs, Le Genest-Saint-Isle, France) were anesthetized, and a silicon-coated monofilament (Doccol Corp., Sharon, MA, USA) was inserted into the right common carotid artery (CCA), which was gradually moved forward to the right middle cerebral artery (MCA) according to a previously established protocol (Doepfner et al. 2015). The reperfusion was initiated one hour after monofilament insertion by monofilament removal. Herein, I first investigated the effect of AMSC-EVs on infarct size. This set of mice underwent MCAO followed by femoral vein injection of an equal amount of PBS (control) or AMSC-EVs (PBS diluted EVs from 2×10^6 AMSCs, 200 μ l) either at the beginning of the reperfusion or 12 hours after reperfusion. The mice were sacrificed 24 hours after stroke induction, and the brains were removed and used for infarct volume analysis by TTC staining (Sigma-Aldrich, St. Louis, MO, USA).

To analyze autophagy after MCAO, mice underwent 1 hour of MCAO followed by different survival periods (6, 24, 48, or 72 hours of reperfusion). The third set of mice was subjected to sham or MCAO surgery followed by PBS (control) or AMSC-EV injection immediately at the end of the operation or 12 hours after reperfusion. The ischemic hemisphere of the brains was also used for Western blot analysis.

To illustrate the neuronal damage caused by excessive autophagy after ischemic stroke, mice were subjected to MCAO followed by intraperitoneal injections of PBS (control) or by the autophagy inhibitor 3-MA immediately at the end of the surgery or 12 hours after reperfusion. The mice were sacrificed 24 hours after MCAO and used for Western blot analysis. The mice, which were sacrificed after 14 days, were used for histochemical studies of brain injury and behavioral analyses. The specific methods of behavioral analyses is based on a previous publication from our group (Doepfner et al. 2015).

To find out the key molecules that are responsible for AMSC-EV therapy, mice were administered with PBS (control), AMSC-EVs, AMSC-EV^{anti-microRNA-25} (microRNA-25 siRNA was transfected into AMSCs before EV isolation), and AMSC-EV^{NC} (control siRNA was transfected into AMSCs before EV isolation) 12 hours after reperfusion in the anesthetized mice. The brains of the mice were used for Western blot and immunohistochemistry studies. The details of the *in vivo* experiment paradigm in table can be found in Publication 1.

Table. Experimental groups of mice.

Groups	Behavior tests and immunofluorescence		Western blot analysis		TTC staining	
	Number	Duration	Number	Duration	Number	Duration
Sham			12	1 day		
PBS	10	14 days	12	1 day	8	1 day
3-MA 0 h	10	14 days	12	1 day		
3-MA 12 h	10	14 days	12	1 day		
AMSC-EVs 0 h	10	14 days	12	1 day	8	1 day
AMSC-EVs 12 h	10	14 days	12	1 day	8	1 day
AMSC-EVs ^{NC}	10	14 days	12	1 day		
AMSC-EVs ^{anti-25}	10	14 days	12	1 day		
Reperfusion 6 h			6	6 h		
Reperfusion 24 h			6	24 h		
Reperfusion 48 h			6	48 h		
Reperfusion 72 h			6	72 h		

3.9 Statistical analysis

GraphPad Prism 8.0 software (GraphPad Software Inc., San Diego, CA, USA) was used for statistic analysis. The two-tailed independent Student's t-test evaluated data for comparison of 2 groups. For the comparison between 3 or more groups, a one-way analysis of variance (ANOVA) followed by the Tukey's post-hoc-test was used. The results are shown as mean values with SD unless otherwise stated. A p -value of <0.05 was considered statistically significant.

4 Results and discussion

4.1 AMSC-EVs alleviate primary neuronal injury after OGD

Previous work from our group has shown that EVs isolated from bone marrow-derived MSCs (BMSCs) improved neurological recovery in rodent ischemic stroke models (Doeppner et al. 2015). Since adipose-derived MSCs (AMSCs) have advantages in terms of isolation and expansion, AMSCs have been introduced as an alternative to BMSCs in stem cell-based therapy. In this study, I aimed to analyze the underlying mechanisms of AMSC-EV-induced neuroprotection using both primary neurons exposed to OGD and an ischemic stroke mouse model. As such, I first established an OGD model in primary neurons, demonstrating that the extent of cell injury significantly correlated in a time-dependent fashion with the duration of the OGD itself (Figure 2a in Publication 1).

To support the cells under stress, autophagy is regarded as an adaptive response for cells to starvation, stress, and inflammation. However, autophagy appears to also boost cell injury and death in various pathological conditions (Bialik et al. 2018). Recent studies highlight a role for autophagy to be involved in post-stroke neuronal injury, although published data are contradictory. Whether or not autophagy is harmful or beneficial under conditions of ischemic stroke is a matter of ongoing scientific discourse (Chen W et al. 2014; Mo et al. 2020; Wei et al. 2012). As a proof of concept, I first analyzed autophagy levels in the aforementioned OGD model, for which the measurement of microtubule-associated protein light chain 3 (LC3) is regarded as one marker used to monitor autophagy. In this context, the conversion of LC3 (LC3-I to LC3-II) correlates with the number of autophagosomes (Mizushima and Yoshimori 2007). Indeed, LC3-II abundance was significantly increased and correlated with the expanded duration of OGD (Figure 2b-c in Publication 1), indicating increased levels of autophagy when neurons were exposed to OGD. The latter included an experimental paradigm with 10 hours of OGD followed by 24 hours of reoxygenation, yielding sufficiently high cell injury rates as well

as adequate levels of autophagy. These settings were therefore used for the remainder of the study.

Before using AMSC-EVs in the neuronal OGD model, AMSC-EVs were enriched from cell culture conditioned medium of AMSCs followed by further characterization. Western blot analysis was used for identifying the EV surface marker expression patterns. The protein abundance of Tsg101, CD9, Alix, and CD63 were high in these AMSC-EVs (Figure 1a in Publication 1). The nanosight tracking analysis (NTA) demonstrated that most AMSC-EVs are about 100 nm in size (Figure 1b in Publication 1). Transmission electron microscopy (TEM) revealed two EV sub-populations based upon their size; I found both exosomes and microvesicles (MVs) in the respective samples (Figure 1c in Publication 1). Details of NTA and TEM can be found in publication 1. Thereafter, the so validated EVs were used in the primary neuron OGD model. Therein, treatment of OGD-exposed primary neurons with AMSC-EVs at the onset of hypoxia and during reoxygenation resulted in significantly reduced cellular damage compared to PBS-treated controls (Figure 2d in Publication 1). Interestingly, AMSC-EV treatment was not inferior to the cell therapy (AMSC co-culturing) with regard to neuronal survival after OGD (Figure 2d in Publication 1).

4.2 AMSC-EVs protect primary cortical neurons from OGD-induced damage via autophagy modulation

To verify the hypothesis that AMSC-EVs decrease OGD-induced neuronal injury by regulating autophagic activities, OGD exposed primary cortical neurons were co-cultured with either AMSCs or AMSC-derived EVs at both the onset of hypoxia and during reoxygenation. As shown by Western blot analysis, the protein levels of LC3-II were significantly increased after OGD. However, the protein abundance of LC3-II was reversed when neurons were treated with either AMSCs or AMSC-EVs (Figure 2e in Publication 1). Since autophagy is a highly complex and dynamic process, it needs to be assessed accurately.

Analysis of autophagy is a dynamic process, and autophagosome accumulation may indicate either enhanced autophagy or a blockage of downstream steps such as decreased levels of autolysosomes and others (Zhang XJ et al. 2013). The mere detection of LC3-II levels only therefore inadequately represents an overall assessment of the whole autophagic system. Hence, the so-called “autophagic flux” is used to demonstrate autophagy dynamics, by using an autophagy inhibitor like bafilomycin-A1 (BafA1) or chloroquine (du Toit et al. 2018). Inhibiting the fusion of autophagosomes and autolysosomes as well as the autophagic flux by means of BafA1 . thus ensures that accumulation of LC3-II indicates the amount of autophagosome production at a given time point. As shown in figure 3, publication 1, application of BafA1 increased the levels of LC3-II in all experimental groups. The protein levels of BafA1-induced LC3-II was significantly higher in primary cortical neurons cultured with PBS as compared to AMSCs or AMSC-EVs treated cells under OGD conditions. Furthermore, autophagic flux was monitored by a fluorescence RFP-GFP-LC3B (alternative name for LC3-II) reporter in OGD-exposed primary neurons. Co-culturing primary cortical neurons with AMSC-EVs under OGD conditions significantly decreased the number of both autophagosomes and autolysosomes (Figure 3c-d in Publication 1).

To discover the intrinsic relationship between autophagy and AMSC-EVs, the autophagy activator rapamycin as well as the autophagy inhibitor 3-MA were used. As shown in Figure 3e, Publication 1, treating the cells with 3-MA significantly decreased neuronal injury in a concentration-dependent manner. However, the neuroprotective effects of the autophagy inhibitor 3-MA were lost when it was utilized in a relatively high dosage (5 mM), indicating that complete elimination of autophagy in a starvation condition may be detrimental. On the contrary, treatment of primary neurons with different dosages of rapamycin combined with the same dose of AMSC-EVs resulted in a reversal of AMSC-EV-induced neuroprotection, suggesting that AMSC-EVs act in the opposite way to autophagy activators (Figure 3 in

Publication 1). These results indicate that a moderately decreased autophagy activity confers significant protection in primary neurons against OGD-induced cell death.

Autophagy was strongly activated upon induction of OGD. A significant increase of autophagy flux under OGD conditions represented a dramatic activation of autophagy. Notably, a moderate down-regulation of autophagy significantly improved neuron viability in this *in vitro* stroke model. AMSC-EV treatment could mimic the protective effect of the autophagy inhibitor 3-MA. However, this protective effect was eliminated by treating the cells together with the autophagy activator rapamycin. This suggests that the protective effect of AMSC-EVs on neurons was induced by modulating the level of autophagy. According to previous studies, basal levels of autophagy are essential for maintaining neuronal function, and it is reasonable to explain that at the beginning of hypoxia and nutrient deprivation, a moderate increase in autophagic activity may be correlated with neuroprotection (Kim KA et al. 2018). In line with the present research, more severe noxious stimuli, such as prolonged hypoxia and starvation, are followed by a significant increase in autophagic flux that may eventually lead to autophagic cell death (Ginet et al. 2014; Liu Y and Levine 2015; Zhang A et al. 2021). Therefore, a well-balanced inhibition of autophagy seems to be particularly important in the context of stroke (Shi R et al. 2012).

4.3 P53 and B-cell lymphoma 2–interacting protein 3 (BNIP3) signaling is involved in AMSC-EVs regulating autophagy

To investigate the mechanisms by which AMSC-EVs regulate OGD-induced autophagy, autophagy-associated signaling cascades were further analyzed. At least four signaling pathways are involved in autophagy regulation, among which are p53 and BNIP3 (Wang EY et al. 2013). Interestingly, the magnitude of p53 and BNIP3 was significantly enhanced in OGD-exposed primary neurons compared to standard cultured controls (Figure 3f in Publication 1). The OGD-induced upregulation of p53 and BNIP3 was reversed when co-culturing primary

neurons with either AMSCs or AMSC-EVs (Figure 3f in Publication 1), indicating that AMSC-EVs regulate autophagy through the p53-BNIP3 axis.

Using the exosome secretion inhibitor GW4869, I observed that co-culturing primary neurons with AMSC-EVs decreased the activation of the p53-BNIP3 axis after OGD. In contrast, co-culturing primary neurons with either AMSCs pretreated with GW4869 or with EVs obtained from GW4869 pre-treated AMSCs failed to inhibit the p53-BNIP3 axis (Figure 4e-f in Publication 1). Thus, the critical subgroup of EV-exosome appears to be crucial in the EV-mediated anti-autophagic effects of AMSCs in primary neurons exposed to OGD.

Inhibition of autophagy alone does not fully account for the therapeutic effect of AMSC-EVs. A successful therapeutic intervention also depends on the precise gene or signaling target that should be suppressed. Several studies have shown that upregulation of p53 after hypoxia or cerebral ischemia can be mediated by activation of HIF-1 α (Althaus et al. 2006; Li et al. 2013). The HIF-1 α enhanced p53 further promotes the expression of BNIP3 (Wang EY et al. 2013; Xin XY et al. 2011). Increased BNIP3, which contains a single Bcl-2 homology 3 (BH3) domain, competes with beclin-1 to bind to Bcl-2. Released beclin-1, in turn, then activates autophagy and cell death (Glick et al. 2010; He and Klionsky 2009). What is more, BNIP3 is a regulator of a specific type of autophagy — mitophagy (mitochondrial autophagy). It is well known that mitochondria are energy factories, and their proper functioning is essential in energy-intensive organs such as the heart and brain. The quantity and quality of cellular mitochondria are necessary for the functioning of these organs. However, under pathological conditions such as ischemic stroke, excessive autophagy, particularly mitophagy, can kill cells that are already in an energy-depleted state (Yuan et al. 2017). Worse still, ATP production through the respiratory chain is accompanied by the production of reactive oxygen species (ROS), excessive accumulation of DNA (mtDNA), lipids, and protein, which in turn aggravates mitochondrial dysfunction in a feedforward manner (Cenini et al. 2020).

Hereto, the protein levels of p53 and BNIP3 were enhanced upon hypoxia, and treatment with AMSC-EVs reversed this over-activation trend. Although it cannot be excluded entirely that other related signaling of autophagy is being regulated by AMSC-EVs, this study shows that autophagy inhibition through p53-BNIP3 signaling plays a vital role in AMSC-EV-based ischemic stroke treatment.

4.4 Post-stroke autophagic regulation and neuroprotection by AMSC-EVs depends on microRNA-25-3p

To identify the EV compounds responsible for AMSC-EV-induced regulation of autophagy, the p53-BNIP3 signaling involved microRNAs were first to be noticed. According to published data, there are more than ten microRNAs known to directly target p53 (Liu J et al. 2017), six candidate microRNAs (microRNA-98, microRNA-25-3p, microRNA-30, microRNA-125a-5p, microRNA-125b-5p, and microRNA-214-3p) which have a closer relationship with stroke, ischemia, neurons, autophagy or cell death were selected for further exploration. The results from qRT-PCR analyses revealed that microRNA-25-3p was one of the most abundant p53 target microRNAs in AMSC secreted EVs, microRNA-214 ranked second in terms of content (Figure 5a in Publication 1). Therefore, the focus was set on these two microRNAs. I checked the intracellular concentration of microRNA-25-3p in OGD exposed primary neurons, and consistent with the hypothesis, the level of microRNA-25-3p decreased dramatically when these cells were exposed to OGD. On the contrary, AMSC-EV treatment reversed this trend (Figure 5c in Publication 1).

To further confirm a functional role of microRNA-25-3p in AMSC-EV-induced autophagy regulation and neuroprotection in OGD-exposed primary cortical neurons, knockdown experiments were initiated in AMSCs. EVs were obtained from AMSCs pretreated with microRNA-25 siRNA or a scrambled construct (NC). The cell injury was significantly lower in primary neurons cultured with EVs^{NC} when compared to PBS-treated controls. However, these

protective effects of AMSC-EVs^{NC} were reversed when the cells were treated with EVs^{anti-microRNA-25} (Figure 5d in Publication 1). The level of autophagic flux was also significantly lower in primary neurons cultured with EVs^{NC} when compared to PBS-treated cells under OGD conditions, whereas the autophagic flux was again reversed when the neurons were incubated with EVs^{anti-microRNA-25} (Figure 5e-f in Publication 1). Likewise, by using a luciferase system, a significant decrease in the number of autophagosomes and autolysosomes was detected in neurons incubated with EVs^{NC} compared to EVs^{anti-microRNA-25} (Figure 5g-h in Publication 1).

To better visualize the role of microRNA-25 in primary neurons after OGD, the neuroprotection and autophagy regulation effects of microRNA-25 were further analyzed by direct overexpression or down-regulation of microRNA-25. Forced expression of microRNA-25-3p in hypoxic neurons duplicated the inhibitory effects on autophagy regulation (Figure S3a-b in Publication 1) and autophagic flux (Figure S3c-d in Publication 1) compared with the nonsense transfected one. However, a loss of function study by microRNA-25-3p siRNA transfection yielded opposite results (Figure S3g-h in Publication 1). Being consistent with the autophagic regulation, the protein levels of p53 and BNIP3 were diminished dramatically in response to the forced expression of microRNA-25-3p (Figure S3i-j in Publication 1). In contrast, the amount of these proteins was enhanced when the primary neurons were treated with the microRNA-25 siRNA (Figure S3k-l in Publication 1). Finally, microRNA-25 overexpression in primary cortical neurons resulted in increased cell viability (Figure S3m in Publication 1), whereas microRNA-25 knockdown acts in the opposite way, decreasing neuronal viability after OGD (Figure S3n in Publication 1).

The role of EVs and their therapeutic potential for ischemic stroke treatment has only recently become a notable topic of research (Cunningham et al. 2018; Webb et al. 2018). Previous work from different groups has shown that MSC-EVs or NPC-EVs have both anti-inflammatory and pro-angiogenic properties (Dabrowska et al. 2019; Doepfner et al. 2015; Keshtkar et al. 2018; Xin H et al. 2013). These EVs carry biological information for cell communication and contain

DNA, proteins, non-coding RNAs, and microRNAs for cell regulation (Bertoli et al. 2015). The latter guides themselves to regions of partial complementary sequences within mRNA molecules via base-pairing. Most of the sequences were mainly within 3'-untranslated regions (UTRs) of target mRNA, and the pairing results in translational inhibition and RNA silencing (Bartel 2009). Different groups have demonstrated that cell-secreted EVs can influence the performance of recipient cells through microRNAs transfer *in vitro* (O'Brien K et al. 2020; Thomou et al. 2017; Wang L et al. 2019). Since the autophagic pathway is also known to be regulated by microRNAs (Abels and Breakefield 2016), it is fair enough to assume that AMSC-EVs yield autophagic regulation and neuroprotection using this mode of action. I herein showed that AMSC-EVs improve neurological survival by regulating autophagy. Moreover, the critical cytoprotective and anti-autophagic molecule of AMSC-EV cargo may be the p53 targeted microRNA-25. Previous studies have primarily focused on the biological function of microRNA-25 under conditions of cancer (Zeng et al. 2018; Zhang J et al. 2019). The present results demonstrate for the first time that microRNA-25-3p may contribute to the pathogenesis of ischemic stroke *in vitro*.

4.5 AMSC-EVs reduce brain injury and induce sustained neurological recovery after ischemic stroke in mice

Based on the above *in vitro* data on primary cortical neurons, I next investigated whether AMSC-EV administration could improve neurological recovery after ischemic stroke by modulating autophagic activity *in vivo*. I first examined the biodistribution pattern of AMSC-EVs under ischemic conditions. Consistent with published data (Zagrean et al. 2018), AMSC-derived EVs reached the ischemic infarct core (Figure 6a in Publication 1). Following our established protocol (Doepfner et al. 2015), EVs isolated from 2×10^6 AMSCs were diluted in PBS and systemically injected immediately at the end of the surgery or 12 hours after reperfusion. Consistent with our previous results, mice that received EV injections immediately at the end of MCAO had significantly smaller infarct volumes than PBS controls (Figure 6b in Publication

1). Besides, the infarct volume was also significantly reduced in those animals treated with EVs 12 hours after reperfusion compared to the control group (Figure 6b in Publication 1). Along with the reduction in acute brain injury and cell death, behavior analysis of mice also showed better performance when mice were treated with AMSC-EVs at either time point compared to the PBS control group (Figure 6d-e in Publication 1). Importantly, this better test score in behavior tests (both the tight rope test and the corner turn test) was stable and long-lasting from the third day of the treatment until the end of the test. Analyzing neuronal density in the striatum 14 days after ischemic stroke revealed I observed an increase in neuronal density in both EV treatment groups in stroke mice (Figure 6f in Publication 1). Conclusively, these data reveal that AMSC-secreted EVs reduce post-stroke brain injury at both functional and histological levels under experimental stroke conditions.

Ischemic strokes have a high morbidity and mortality rate, and most stroke patients are left with severe and permanent neurological deficits. The time window for stroke treatment is concise; there are few recovery options once neurological deficits are fixed. Any treatment that results in even minor restoration of neurological deficits is likely to be beneficial. In this study, we demonstrated that intravenous administration of AMSC-EVs improved functional outcomes and showed a neuroprotection tendency against cerebral ischemia.

Although stem cell-based therapy could not directly replace the neurons from infarcted area, it can still provide trophic support and regulate the cellular status of the infarcted site. Various types of cells ranging from neuronal cell lines to embryonic stem cells are candidates for cell therapy in ischemic stroke, clinical trials of neural progenitor cells (Savitz et al. 2005), and autologous bone marrow-derived MSCs (Bang et al. 2005) have been conducted. Moreover, these stem cells also secrete trophic and growth factors, cytokines, and other biological information like microRNAs, which activate angiogenesis and neurogenesis resulting in improved neurological functions (Hofer and Tuan 2016).

MSC-derived EVs have been repeatedly shown to induce neuroprotection by different experimental groups (Doepfner et al. 2015; Jiang et al. 2018; Safakheil and Safakheil 2020). But some studies showed that administration of MSC-EVs did not change the infarct volume in rodent models (Otero-Ortega et al. 2017; Zhang Y et al. 2015). The present data showed a significant reduction in infarct volume when AMSC-EVs were administered in mice after stroke, suggesting AMSC-EVs are potentially protective and thus therapeutically relevant in stroke treatment.

4.6 P53-BNIP3 signaling participates in AMSC-EV-reduced autophagic flux after ischemic stroke

To further confirm whether autophagy inhibition contributes to functional improvement and neuroprotection in mice after MCAO, I first used the autophagy inhibitor 3-MA to downregulate autophagy levels after MCAO and then assessed neurological recovery as well as neuronal survival in mice. Mice treated with the autophagy inhibitor 12 hours after the induction of stroke were found to have a better behavior performance until the end of the observation period of 14 days, confirming the results above and further supporting the hypothesis that autophagy inhibition contributes to neurological recovery and neuronal survival after stroke (Figure 6d-e in Publication 1). In contrast, immediate delivery of the autophagy inhibitor 3-MA at the onset of reperfusion only partially enhanced neurological recovery (Figure 6d-e in Publication 1) and did not affect cell injury at all (Figure 6f-g in Publication 1). Analysis of neuronal survival 14 days after stroke showed similar results, with an increase in neuronal density in mice treated with 3-MA 12 hours after reperfusion.

To further investigate the correlation between neuroprotection induced by AMSC-EVs and the regulation of autophagy, time-dependent patterns of autophagy levels in mice with non-therapeutic stroke were evaluated. Using LC3-II to analyze autophagy levels, Western blotting results showed that protein abundance peaked 24 hours after stroke compared to sham-operated

mice (Figure S4a-b in Publication 1). Treatment of mice with AMSC-EVs significantly reduced autophagy activation measured one day after stroke when EVs were given immediately at the beginning of reperfusion (EVs 0 hour) or 12 hours after stroke (EVs 12 hours) (Figure S4c-d in Publication 1). Similarly, treatment of mice with 3-MA immediately at the onset of reperfusion did not affect autophagic flux, which is in line with the results mentioned before. In contrast, treatment of stroke mice with 3-MA 12 hours after reperfusion significantly reduced autophagic flux when BafA1 was injected in these animals 4 hours before sacrifice (Figure S4e-f in Publication 1). These results suggest that the administration of 3-MA and AMSC-EVs can reverse stroke-induced upregulation of autophagic flux. It also indicates that appropriate time is a prerequisite for properly modulating over-activated autophagic activity in the rodent MCAO model.

Thereafter, I measured the expression of p53 and BNIP3 *in vivo*. In agreement with the *in vitro* results, the relative expression of p53 and BNIP3 was increased in the infarcted hemisphere of MCAO mice. Administration of AMSC-EVs at the two injection time points described above (0 hour and 12 hours after MCAO) reversed the up-regulation trend of both p53 and BNIP3 (Figure 4g-h in Publication 1).

Although 3-MA and AMSC-EV both inhibit post-stroke autophagy, they show different therapeutic effects in mice. Due to short half-life time and metabolization, pharmacological agents such as 3-MA may only transiently affect some of the vital autophagic molecules within about 2-3 hours (Sweet et al. 1981). This may explain why the use of 3-MA at the beginning of reperfusion did not have a therapeutic effect. Still, biological molecules such as EVs can induce long-term effects in various situations (Lai et al. 2013; Tian et al. 2018). In our study, the application of AMSC-derived EVs resulted in a significant downregulation of ischemia-induced autophagy under both *in vitro* and *in vivo* conditions, contributing to neuroprotection and neurological recovery. This result has also been shown in other models, confirming that the beneficial effect of MSC transplantation after myocardial infarction is due, at least in part, to the

improved autophagic flux through exosome secretion (Xiao et al. 2018). Similarly, there is also a study showing that exosomes derived from AMSCs significantly reduce neuronal injury by inhibiting autophagy and promote microglia polarization after stroke (Jiang et al. 2018). These results together suggest that MSC-derived EVs can prevent brain damage by inhibiting autophagy.

Accumulating evidence suggests that mitochondria play a central role in the development of ischemia-induced cell death (Kuznetsov et al. 2019; Ong and Gustafsson 2012), and maintaining mitochondrial function is thus critical for neurological recovery and neuronal survival. BNIP3 is a member of unique mitochondrial proteins, which induce cell death by promoting early mitochondrial damage (Gao et al. 2020; Zhang J and Ney 2009). In my study, BNIP3 is markedly upregulated after cerebral ischemia, which is reversed when treating the animals with 3-MA or AMSC-EVs. Several studies have demonstrated a link between BNIP3, mitophagy and cell death, showing that inhibiting BNIP3 completely abrogated cell injury via regulating mitochondrial signaling (Dhingra et al. 2014; Shi RY et al. 2014). Moreover, the role of BNIP3 is not limited to its involvement in mitophagy, but it also mediates other types of cell death, like apoptosis and necrosis (Dhingra et al. 2014; Ma et al. 2017; Vande Velde et al. 2000). Therefore, it cannot be excluded that AMSC-EVs are also involved in regulating other signaling pathways by regulating the expression of BNIP3, thus yielding a neuroprotective effect.

4.7 Autophagic regulation and post-stroke neuroprotection by AMSC-EVs depend on microRNA-25

To confirm that microRNA-25-containing EVs mediate the reduction in autophagic flux associated with AMSC-EV administration, mice were injected with PBS (negative control), AMSC-EVs (positive control), AMSC-EVs derived from AMSC pretreated with control oligonucleotides (EVs^{NC}), and AMSC-EVs derived from AMSC pretreated with anti-microRNA25 (EVs^{anti-microRNA25}) 12 hours after the induction of MCAO. One day after

reperfusion, the autophagic flux in the ischemic striatum was significantly lower in both the AMSC-EV and AMSC-EV^{NC} groups compared to the PBS controls (Figure 7a-b in Publication 1). However, the AMSC-EV^{anti-microRNA25} treatment failed to reach similar effects on autophagy regulation (Figure 7a-b in Publication 1). Consistently, the behavior test analysis indicated a better test performance in the tight rope and in the corner turn test when mice were treated with EVs^{NC} in comparison to both controls (PBS group) and mice treated with EVs^{anti-microRNA25} (Figure 7c-d in Publication 1). Analysis of the neuronal density showed that mice treated with EVs^{NC} displayed a higher neuronal density than animals treated with EVs^{anti-microRNA25} 14 days after MCAO (Figure 7e-f in Publication 1). Conclusively, AMSC-derived EVs alleviate brain injury on both the functional and the histological level. The autophagy regulation capability associated with ADMSC-EV administration is at least partially mediated by the EV transfer of microRNA-25 after stroke.

Recent findings highlight that EVs selectively carry large amounts of miRNAs and mediate the crosstalk between different cell types by directly transferring miRNAs to recipient cells (Mittelbrunn et al. 2011; Xiao et al. 2018). Experiments in which human CD34+ stem cell-derived EVs carry microRNAs associated with proangiogenesis is a good example in this context (Sahoo et al. 2011). Consistent with my findings, intercellular communication processes of naturally secreted EVs can thus be utilised to deliver exogenous genetic material to further improve the success and effectiveness of gene therapy (Xiao et al. 2018; Zomer et al. 2015).

4.8 Summary

Causal therapy of ischemic stroke remains available to only a minority of patients despite the success of systemic thrombolysis and thrombectomy. Consequently, new adjuvant therapeutic options are urgently needed. MSC transplantation leads to improved neurological recovery in preclinical stroke models. However, MSCs do not directly act on ischemic brain tissue, but rather mediate their effects via the secretion of EVs. The latter are a heterogeneous group of corpuscular structures with a diameter of 30-1000 nm, which contain non-coding RNA, DNA,

microRNA, and proteins. The application of EVs seems to be a more effective and low-risk treatment than stem cell transplantation in experimental stroke models. Hence, MSC-EVs provide an adjunctive approach to the treatment of ischemic stroke. Although the mechanisms by which MSC-EVs act remains elusive, recent evidence suggests that MSC-EVs may be responsible for neuroprotection in pathological stroke conditions. Herein, it was first confirmed that the application of AMSC-EVs is not inferior to the use of AMSCs in an *in vitro* hypoxia model. Incubation of neurons with AMSCs or AMSC-EVs increases the resistance to hypoxic cell injury. Interestingly, the inhibition of EV secretion from AMSCs by GW4869 leads to a loss of the therapeutic effect of AMSCs. This observation again underscores the importance of EVs as a biologically active agent in stem cell treatment of stroke. For the neuroprotective effect of AMSC-EVs under experimental stroke conditions, I for the first time illustrated the interaction of intravesicular microRNA-25-3p with the p53-BNIP3 signaling pathway. The regulation of this pathway by EVs is accompanied by inhibition of the autophagic flux, which in turn leads to reduced neuronal death after OGD. Indeed, the observations described above can be reversed by the use of anti-oligonucleotides and thus reduce the therapeutic effect of EVs. The functional significance of the p53-BNIP3 signaling pathway and the modulation of the autophagic flux via intravesicular transfer of microRNA-25-3p were further investigated in a mouse stroke model. It was shown for the first time *in vivo* that the application of AMSC-EVs via this mechanism leads to a reduction of cerebral tissue damage and better functional recovery of the animals after stroke. The present work makes an important contribution to the analysis of the mechanisms of EV-based therapy of ischemic stroke. Further preclinical work in this field is needed before EVs can be used as adjuvant therapy in stroke patients.

In conclusion, the research results presented herein provide a novel insight into the mechanisms by which AMSC-EVs induce therapeutic actions under experimental stroke settings. This study also provides evidence that native AMSC-EVs yield enhanced neurological recovery and neuroprotection by inhibiting ischemia-induced autophagy. The autophagy inhibition, in turn,

is mainly responsible for EVs transferring microRNA-25-3p from AMSCs to their recipient cells. The latter results in interference with the autophagy regulation signaling pathway, p53-BNIP3. These new observations on EVs derived from AMSC may lead to the development of new therapeutic targets and strategies to treat ischemic stroke.

5 References

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RESEARCH ARTICLE

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Adipose-derived mesenchymal stem cells reduce autophagy in stroke mice by extracellular vesicle transfer of miR-25

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TÜBITAK

Abstract

Grafted mesenchymal stem cells (MSCs) yield neuroprotection in preclinical stroke models by secreting extracellular vesicles (EVs). The neuroprotective cargo of EVs, however, has not yet been identified. To investigate such cargo and its underlying mechanism, primary neurons were exposed to oxygen-glucose-deprivation (OGD) and cocultured with adipose-derived MSCs (ADMSCs) or ADMSC-secreted EVs. Under such conditions, both ADMSCs and ADMSC-secreted EVs significantly reduced neuronal death. Screening for signalling cascades being involved in the interaction between ADMSCs and neurons revealed a decreased autophagic flux as well as a declined p53-BNIP3 activity in neurons receiving either treatment paradigm. However, the aforementioned effects were reversed when ADMSCs were pretreated with the inhibitor of exosomal secretion GW4869 or when Hrs was knocked down. In light of miR-25-3p being the most highly expressed miRNA in ADMSC-EVs interacting with the p53 pathway, further in vitro work focused on this pathway. Indeed, a miR-25-3p oligonucleotide mimic reduced cell death, whereas the anti-oligonucleotide increased autophagic flux and cell death by modulating p53-BNIP3 signalling in primary neurons exposed to OGD. Likewise, native ADMSC-EVs but not EVs obtained from ADMSCs pretreated with the anti-miR-25-3p oligonucleotide (ADMSC-EVs^{anti-miR-25-3p}) confirmed the aforementioned in vitro observations in C57BL/6 mice exposed to cerebral ischemia. The infarct size was reduced, and neurological recovery was increased in mice treated with native ADMSC-EVs when compared to ADMSC-EVs^{anti-miR-25-3p}. ADMSCs induce neuroprotection by improved autophagic flux through secreted EVs containing miR-25-3p. Hence, our work uncovers a novel key factor in naturally secreted ADMSC-EVs for the regulation of autophagy and induction of neuroprotection in a preclinical stroke model.

KEYWORDS

cerebral ischemia, extracellular vesicles, adipose-derived MSCs, neurological recovery, autophagy

1 | INTRODUCTION

Mesenchymal stem cells (MSCs) have been thoroughly analysed in preclinical stroke models, revealing both neuroprotection and increased neurological recovery after transplantation in vivo (Chen et al., 2001; Gervois et al., 2016; Lucia Maria Ferri et al., 2016).

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These grafted stem cells are neither integrated into residing neural networks nor do they take over functional characteristics of lost neurons (Fernández-Susavila, Bugallo-Casal, Castillo, & Campos, 2019). Instead, transplanted cells work through indirect ways (Baraniak & Mcdevitt, 2010; Liang, Ding, Zhang, Tse, & Lian, 2014). Recently, bilayer-structured vesicles ranging in size from 30 nm to 1000 nm and containing a defined set of cargo such as non-coding RNAs, DNAs, and proteins were defined as extracellular vesicles (EVs) (Kim et al., 2013; Mathivanan, Fahner, Reid, & Simpson, 2012; Théry et al., 2018). These EVs derived from MSCs or other stem cell sources appear to be non-inferior with regard to their therapeutic potential against cerebral ischemia when compared to their host cells (Cunningham, Redondo-Castro, & Allan, 2018; Doepfner et al., 2015; Xin et al., 2013). The neuroprotective cargo of MSCs-derived EVs, however, has not been sufficiently identified.

Recent studies have shown that EVs regulate protein expression of recipient cells and modify cell characteristics through microRNA transfer (Phinney et al., 2015; Tkach & Théry, 2016). MicroRNAs in turn bind to untranslated regions and in some cases also to coding regions of mRNAs, thus post-transcriptionally regulating gene expression (Lim et al., 2005; Vemuganti, 2010). Interestingly, autophagy – a self-degradative process – is one target of miRNA regulation (Frankel & Lund, 2012). Recent work emphasized the role of autophagy to be a key regulator of ischemic stroke (Wang et al., 2018), unveiling a new range of potential therapeutic targets for neuroprotection.

Autophagy includes a process of forming an isolation membrane that spherically expands in order to engulf materials designated for degradation, which gives rise to a double-membrane vesicle called the autophagosome. The latter will deliver these engulfed materials to the lysosome for later degradation (Yoshimori, 2007). Under conditions of cerebral ischemia, autophagy acts as a double-edged sword for neuronal survival (Chen, Sun, Liu, & Sun, 2014). A modest induction of autophagy upon induction of hypoxia or cerebral ischemia has been associated with enhanced cell viability and decreased infarct size (Carloni et al., 2010; Dai et al., 2017; Wang et al., 2012). Nevertheless, prolonged oxygen and glucose deprivation or stroke event propels excessive autophagy, turning temporarily activated and protective autophagy to chronic activation, which in turn contributes to significant cell death (He et al., 2012; Mo, Fang, He, & Zhang, 2012; Shi et al., 2012; Xin et al., 2011). The role of autophagy in stem cell-induced and even more so in EV-induced neuroprotection against stroke as well as the underlying mechanisms still remain elusive. Hence, the present work analyzes whether or not the application of ADMSC-EVs affects the autophagic response under conditions of both *in vitro* hypoxia and *in vivo* cerebral ischemia with regard to cell survival and underlying mechanisms.

2 | MATERIALS AND METHODS

2.1 | Experimental *in vivo* paradigm and induction of cerebral ischemia

All studies were performed with governmental approval according to the NIH guidelines for the care and use of laboratory animals, following both the ARRIVE and the STAIR guidelines. Male C57BL/6 mice aged 10–12 weeks (Janvier labs, Le Genest-Saint-Isle, France) were kept under circadian rhythm and had free access to food and water. At all stages of the study, the researchers were blinded from the experimental conditions. The mice were randomly assigned to the various treatment groups. The result of the power calculation was 0.8732706 for Western blot analysis, assuming an effect size of 0.75. The result of the power calculation was 0.9105700 for both behavioural test analysis and immunofluorescence analysis, assuming an effect size of 0.70. Precise numbers of animals used are given for each condition in the figure legends and in the Supplementary Table S1 including survival rates of mice.

The induction of cerebral ischemia was performed via middle cerebral artery occlusion (MCAO), as described previously (Doepfner et al., 2015). In brief, silicon-coated monofilament (Doccol Corp., Sharon, MA, USA) was inserted into the right CCA and then gently moved forward towards the offspring of the right middle cerebral artery (MCA). During the experiment, the laser Doppler flow (LDF) was recorded with a flexible probe (Perimed AB, Järfälla, Sweden) covering the core of the right MCA territory. One hour after monofilament insertion, reperfusion was initiated by monofilament removal, and the LDF recordings were continued for an additional 15 min before the wounds were carefully sutured.

The first set of mice underwent MCAO followed by injection of PBS (control) or ADMSC-EVs. EVs were released by 2×10^6 ADMSCs diluted in 200 μ l of PBS corresponding to 10 μ g of EVs. Injection was done by cannulating the right femoral vein, and the injection rate was 200 μ l per 10 min. Mice received such a treatment either at the beginning of the reperfusion or 12 h after reperfusion, followed by a survival of 24 h (eight mice per group). The brains were removed and used for infarct volume analysis as described later.

The second set of mice underwent sham surgery or MCAO followed by PBS (control) or ADMSC-EV treatment immediately at the beginning of the reperfusion or 12 h after reperfusion. Animals in the sham group underwent all surgical procedures for MCAO induction except for occlusion of the MCA. All these groups were allowed to survive for 24 h post-stroke (six mice per group). These brains were used for Western blot analysis.

The third set of mice (six mice per group) was subjected to 1 h of ischemia (except for the sham group) followed by 6, 24, 48 or 72 h of reperfusion. All these groups were also used for Western blot analysis.

The fourth set of mice was exposed to MCAO followed by intraperitoneal injections of PBS (control) or 3-Methyladenine (3-MA, 15 mg/kg; Merck Group, Darmstadt, Germany) immediately at the beginning of the reperfusion or 12 h after reperfusion. The mice were allowed to survive for one day (six mice per group) or fourteen days (ten mice per group). These mice were used for Western blot analysis, behavioural analyses and histochemical studies of brain injury.

The fifth set of mice was exposed to MCAO followed by administration of PBS (control), ADMSC-EVs, ADMSC-EV^{NC} and ADMSC-EV^{anti-miR-25} (anti-miR-25-3p oligonucleotides or negative control oligonucleotides were transfected into ADMSCs before harvesting EVs) via cannulation of the right femoral vein in the anesthetized mice 12 h after reperfusion. The mice were allowed to survive for one day (six mice per group) or for fourteen days (ten mice per group). These mice were used for Western blot analysis, behavioural analyses and histochemical studies of brain injury.

2.2 | Adipose-derived mesenchymal stem cells (ADMSCs) isolation and characterization

ADMSCs were isolated, as described previously (Ahmadian Kia et al., 2011). In brief, adipose tissue was harvested from wildtype C57BL/6 mice and washed with PBS. Thereafter, the tissue was mechanically chopped before digestion with 0.1 % collagenase I (Gibco, Darmstadt, Germany) for 1 h at 37°C with intermittent shaking. The digested tissue was washed with high glucose Dulbecco's modified Eagle's medium (DMEM; Merck Group, Darmstadt, Germany) containing 10 % fetal bovine serum (FBS; Merck Group, Darmstadt, Germany) and then centrifuged at 1,000 rpm for 5 min to remove mature adipocytes. The cell pellet was resuspended in DMEM supplemented with 10 % FBS and 1% penicillin/streptomycin (PS; Gibco, Darmstadt, Germany) in a 37°C incubator with 5 % CO₂. The adherent cells were passaged with 0.25 % trypsin containing 0.02 % EDTA (Sigma-Aldrich, St. Louis, MO, USA) for 3–4 passages.

Cytometric evaluation of ADMSC surface profiles was carried out at passage 2. After two days of culture, the cells were washed with PBS and fixed in 4 % paraformaldehyde for 30 min. ADMSCs were identified with fluorescein isothiocyanate (FITC)-conjugated antibodies against CD29, CD34, CD45, CD90 (BD Biosciences, San Jose, CA, USA) and with phycoerythrin (PE)-conjugated antibodies against CD105 (Ebioscience, San Diego, CA, USA). The percentage of adhering cells that stained for CD29, CD34, CD45, CD90, and CD105, respectively, was quantified by FlowJO software (version 8.0). The results are shown in Figure S1.

2.3 | Preparation of primary cortical neurons

For the preparation of cortical neurons, pregnant C57BL/6 mice were killed by CO₂ inhalation at embryonic day 17. Embryos were dissected, and tissue pieces were trypsinized and dissociated using a fire-polished Pasteur pipette. Cells were seeded on poly-L-ornithine/laminin (Sigma-Aldrich, St. Louis, MO, USA)-coated 6 or 24 well plates at a density of 100,000/cm² containing neuroblast medium (Gibco, Darmstadt, Germany) with additional transferrin (Sigma-Aldrich, St. Louis, MO, USA), penicillin/streptomycin (PS; Gibco, Darmstadt, Germany), L-glutamine (Seromed, Dollnstein, Germany), and B27 supplement (Gibco, Darmstadt, Germany). Cells were used for subsequent experiments after 4 days of cell culture.

2.4 | Standard and oxygen-glucose-deprivation (OGD) model

For the standard (normoxia) condition, cells were maintained under an ambient atmosphere in an incubator at 37°C with 5 % CO₂. For the induction of OGD, primary neurons were incubated at 37°C in a hypoxic chamber (Toepfler Laborsysteme GmbH, Göppingen, Germany) (0.5 % O₂, 5 % CO₂) in Sterofundin medium (Braun, Melsungen, Germany) containing 1 mM mannitol (Serag-Wiessner, Naila, Germany) for different durations of time. Afterwards, the cells were re-incubated under standard cell culture conditions for 24 h (reoxygenation).

2.5 | ADMSC–primary neuron coculture and EV treatment

ADMSCs were plated onto 6-well (2 × 10⁴ cells/insert) transwell (3 μm pore size; Costar, Maryland, USA). After 24 h, the primary neurons were added to the plates, and the cells were cultured under the indicated (normoxia or hypoxia) conditions. To illustrate whether the effects of ADMSC-EVs were dose-dependent, three different ADMSC-EV concentrations were tested: ADMSC-EVs low (EVs equivalent to 2 × 10³ ADMSCs, 0.01 μg), ADMSC-EVs medium (EVs equivalent to 2 × 10⁴ ADMSCs, 0.1 μg), ADMSC-EVs high (EVs equivalent to 2 × 10⁵ ADMSCs, 1 μg). The EVs were added to primary neurons at the beginning of the OGD and at the beginning of the reoxygenation.

2.6 | ADMSC-EV assays – enrichment, characterization, purification, uptake and inhibition

Details can be found in Supplementary Materials and Methods S1.

2.7 | Autophagic flux

For autophagic flux assessments in primary neurons, cells were cultured on coverslips for 3 days and transfected with RFP-GFP-LC3B using the Autophagy Tandem Sensor RFP-GFP-LC3B Kit (Thermo Fisher Scientific, Waltham, Massachusetts, USA). After 24 h, the cells were cultured under the indicated experimental conditions (i.e., normoxia or OGD condition, with or without ADMSC-EVs) and fixed with 4 % paraformaldehyde (PFA). Images were obtained with a fluorescence microscope, and the number of autophagosomes and autolysosomes were quantified in each cell (20-25 cells per experimental group) in 3 independent experiments. For assessments of autophagic flux by LC-3 II, Bafilomycin A1 (Sigma-Aldrich, St. Louis, MO, USA) was used 3 h before harvesting. Western blots were performed, thereafter.

2.8 | Knockdown and overexpression

MiR-25b-3p mimicking oligonucleotides, anti-miR-25-3p oligonucleotides, siRNAs against Hrs and their control oligonucleotides (RiboBio, Guangzhou, China) were transfected into primary neurons or ADMSCs by using Turbofect Transfection reagents (Thermo Fisher Scientific, Waltham, Massachusetts, USA) according to the manufacturer's instructions. After 24 h of transfection, the cells were used for subsequent experiments.

2.9 | Real-time qRT-PCR

Total RNA was isolated using Trizol reagent (Invitrogen, Waltham, California, USA) according to the manufacturer's instructions. The KAPA SYBR[®] FAST One-Step Kit for LightCycler[®]480 (Merck Group, Darmstadt, Germany) was used to perform qRT-PCR as the manufacturer's instructions request, and the PCR primers were purchased from Eurofins Genomics (Ebersberg, Germany). MicroRNA expression was normalized to U6 and let-7a expression and calculated via the standard $2^{-\Delta\Delta Ct}$ method.

2.10 | RNase A and TritonX-100-treatment

To degrade RNA, 500 μ l of ADMSC-EVs (derived from 10×10^6 ADMSCs) were separated into four groups and each group was incubated in the presence of 20 μ g/ml RNase A (Invitrogen, Waltham, California, USA) and/or with 1% TritonX-100 (Sigma-Aldrich, St. Louis, MO, USA) for 30 min at 37°C under gentle agitation. RNase digestion was stopped by addition 1.25 μ l/ml RNasin (Sigma-Aldrich, St. Louis, MO, USA) and RNA-isolation was performed as previous described. All treated EVs were used immediately or were stored at -80 °C.

2.11 | Cell survival

Cell viability was measured by a colorimetric assay by using the MTT (thiazolyl blue tetrazolium, Sigma-Aldrich, St. Louis, MO, USA) viability assay according to the protocol (Venkataramani et al., 2018). Cell viability data are presented as relative changes in percent compared to untreated controls. Furthermore, cell death rate was also determined via fluorescence microscopy by using a LIVE/DEAD Viability kit (Lonza, Basel, Switzerland) as directed by the manufacturer's instructions; living cells were identified with calcein AM (4 μ M, green fluorescence), and dead cells were identified with ethidium homodimer-1 (2 μ M, red fluorescence). Three independent experiments were conducted and ≥ 200 cells were evaluated for each condition.

2.12 | Assessment of infarct volumes

One day after MCAO, the infarct volume was assessed by 2,3,5-triphenyl tetrazolium chloride (TTC) staining in 2-mm thick coronal brain slices. The infarct proportion was calculated by the formula: corrected percentage of infarct volume = (contralateral hemispheric volume – ipsilateral non-infarcted volume) / contralateral hemispheric volume x 100.

2.13 | Analysis of post-stroke motor coordination deficits

Mice were trained on days one and two before the induction of stroke to ensure proper test behaviour. The actual tests for analysis of motor coordination were performed at four, seven, ten and fourteen days using the tight rope test and the corner turn test, as previously described (Doepfner, Kaltwasser, Bähr, & Hermann, 2014). The details can be found in Supplementary Table S2.

2.14 | Protein extracts and Western Blot analyses

Details can be found in Supplementary Materials and Methods S2.

2.15 | Immunofluorescence staining

Brain injury as indicated by neuronal density was evaluated in 16 μm cryostat sections stained with a rabbit monoclonal anti-NeuN antibody (1:300; Merck Group, Darmstadt, Germany), which was detected by a donkey anti-rabbit Alexa Fluor 488 antibody (Invitrogen, Waltham, California, USA). Quantitative analyses for immunohistochemical stainings were performed defining five regions of interests (ROIs) at anterior-posterior +0.14 mm, medial-lateral ± 1.15 to +2.25 mm, and dorsal-ventral -2.25 to +3.25 mm. The number of NeuN⁺ cells was investigated by manual cell counting. The analysis was performed with a fluorescence microscope (Zeiss, Jena, Germany). Five sections per mouse were analyzed, and the mean neuronal densities were determined for all ROIs.

2.16 | Statistical analysis

All statistical analyses were performed using GraphPad Prism Software (GraphPad Software Inc., San Diego, CA, USA). For comparison of two groups, the two-tailed independent Student's *t*-test was used. For comparison of three or more groups, a one-way analysis of variance (ANOVA) followed by the Tukey's post-hoc-test was performed. The power calculations were performed using G*Power software. Unless otherwise stated, data are presented as mean values with SD. A *P* value of <0.05 was considered statistically significant.

3 | RESULTS

3.1 | Purification, Isolation, and Characterization of ADMSC-EVs

We first enriched ADMSC-EVs from conditioned medium of ADMSCs using two well established methods (UC and PEG). Complete cell culture medium has been processed in the same PEG-based manner as conditioned cell culture medium (CCM). The subsequent characterization of such enriched ADMSC-EVs included transmission electron microscopy (TEM), nanosight tracking analysis (NTA), and Western blotting. In line with previous reports (Zheng et al., 2020), Western blotting analysis on the expression patterns of so called EV surface markers revealed CD9, CD63, Alix, and TSG101 but not albumin, Histone, and TOMM20 abundance in these ADMSC-EVs (Figure 1a). The data showed no difference between the UC and the PEG method. The results from the NTA revealed the majority of ADMSC-EVs to be about 100 nm in size in both groups (Figure 1b). TEM analysis showed no significant morphological differences between UC-EVs and PEG-EVs, as both preparations contained smaller and larger vesicles (Figure 1c).

Because PEG protocols may be incapable of discriminating EVs from non-EV nanoparticles and protein (Patel et al., 2019), our next goal was to set up another separation assay to determine which are the biologically relevant nanoparticles of the aforementioned samples, i.e., UC and PEG samples. Density gradients like iodixanol are a classic means to separate membrane-enclosed vesicles according to their floatation speed and equilibrium density (Colombo, Raposo, & Théry, 2014). We first used a self-forming iodixanol (OptiPrep) gradient to subfractionate EVs isolated by UC or PEG (Figure 1d). After subfractionation, ten fractions were recovered and analysed for the presence of protein markers like CD9, CD63, and Alix in both UC and PEG fractions. As shown in Figure 1e, the samples recovered floated mostly in fraction seven in both UC and PEG groups. Consistent with the Western blot results, NTA revealed a vast majority of vesicles to be in fraction seven, regardless of the enrichment method chosen (Figure 1f). These vesicles found in fraction seven typically had a diameter of 30–150 nm (Figure 1g). Thus, floatation into iodixanol gradients allows separation of subtypes of EVs displaying different buoyant densities and sizes, with small EVs (sEVs, 50 to 200 nm diameter) strongly enriched in the seventh fraction regardless of the isolation method.

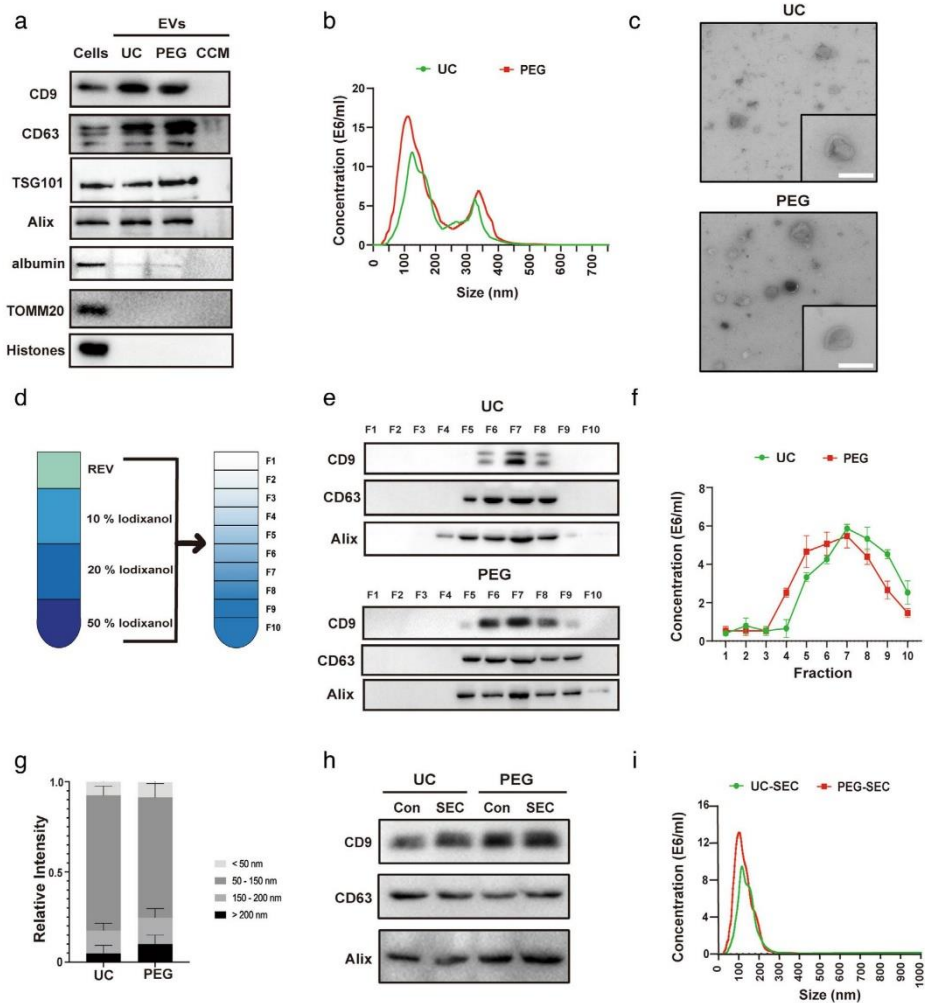


FIGURE 1 Characterization and purification of ADMSC-EVs. Adipose-derived mesenchymal stem cells (ADMSCs) were cultured under standard cell culture conditions, and conditioned cell medium (CCM) was obtained after passage three. CCM was used for the enrichment of extracellular vesicles (EVs) using either the differential centrifugation (i.e. ultracentrifugation, UC) or the polyethylene glycol (PEG) method. (a) Western blot analysis of EVs against exosomal markers of CD9, CD63, TSG101 and Alix, with albumin, TOMM20 and Histones being used as negative markers. Western blots were performed on total cell lysates (cells), EV lysates (UC and PEG) and the CCM. (b) Nanoparticle tracking analysis (NTA) from enriched EVs (UC and PEG) depicting size distribution patterns. (c) Representative transmission electron microscopy (TEM) analysis from EVs enriched by either UC or the PEG method. Scale bar, 100 nm. (d) Resuspended EVs isolated by UC or PEG were allowed to float into an overlaid iodixanol gradient to purify and isolate the small extracellular vesicle (sEV) population. (e) The iodixanol cushion gradient fractions for UC and PEG were analysed by Western blotting (fraction 1–10) using exosome markers. Equivalent volumes of each fraction were loaded per lane. Representative images were shown for CD9, CD63 and Alix which were enriched in fraction seven. (f) NTA was used to assess EV concentrations for each fraction (fraction 1–10). The two-tailed independent Student's *t*-test was used. (g) Representative size distribution of EVs isolated by UC or PEG from their corresponding fraction seven gradients. (h) Western blotting against EV markers (CD9, CD63, and Alix) were performed on ADMSC-EVs isolated by UC or PEG after (or without, Con) size exclusion chromatography (SEC) purification. (i) The size analysis of SEC purified EVs (UC and PEG) was again done by means of NTA

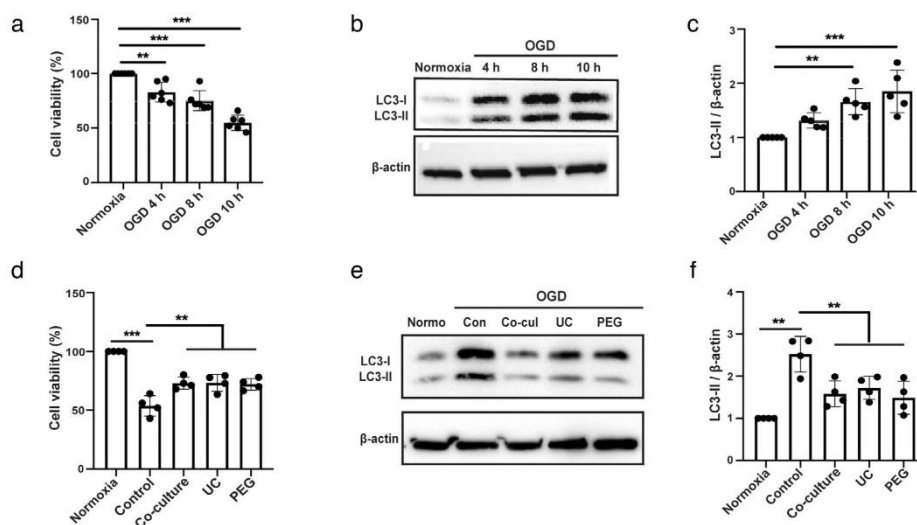


FIGURE 2 ADMSC-EVs protect neurons from oxygen-glucose deprivation (OGD) injury through autophagy regulation. Neuroprotective effects of ADMSC-EVs in cultivated mouse primary cortical neurons were detected by cell viability assays. Primary neurons were incubated for 4 days as mentioned in the materials and methods part. (a) Cell viability was analysed in primary neurons exposed to 4, 8 and 10 h of OGD followed by 24 h of reoxygenation using the MTT assay ($n = 3$). Cells incubated under standard cell culture conditions ('Normoxia') were defined as 100 % cell survival. (b-c) Both qualitative and quantitative analysis of autophagy levels as indicated by the abundance of the autophagy associated protein LC3-II under the aforementioned time points ($n = 3$). Cells incubated under standard cell culture conditions ('Normoxia') were used as negative control. (d) The neuroprotective effect of ADMSC-EVs in cultured primary neurons was evaluated by the MTT assay. After 4 days of cell culture, cells were exposed to 10 h of OGD followed by 24 h of reoxygenation, as mentioned by "OGD" for the later assays. Cell viability was analysed in neurons incubated with either PBS (Control), ADMSCs (Co-culture) or ADMSC-EVs (EVs) isolated by UC or PEG after induction of OGD followed by reoxygenation. PBS, ADMSCs and EVs were given at the beginning of both hypoxia and reoxygenation. Cells incubated under standard cell culture conditions (Normoxia) were defined as 100 % cell survival ($n = 3$). (e-f) LC3 levels were evaluated by Western blotting in OGD exposed primary neurons treated with ADMSCs (Co-culture) or ADMSC-EVs isolated by UC or PEG. ADMSCs and EVs were given at the beginning of both hypoxia and reoxygenation. Neurons treated with PBS under OGD conditions served as positive control (Control). Cells incubated under standard cell culture conditions (Normoxia) were used as negative control ($n = 3$). A representative Western blot is shown in (e), whereas the quantitative analysis for LC3-II is shown in (f). One-way ANOVA followed by the Tukey's post-hoc-test was used. Data are shown as mean \pm SD. Data are statistically different from each other with * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$

In subsequent experiments, we collected layer seven as the purified sEVs (iodixanol) for further experiments. Size exclusion chromatography (SEC) is considered to be one of the best methods for isolating and purifying EVs from different matrices (Stranska et al., 2018). Then, EVs isolated by UC or PEG were applied to the SEC column and collected after filtration. UC-SEC and PEG-SEC were characterized based on the size distribution and the presence of the EV-enriched proteins. Western blot analysis revealed that both UC-SEC and PEG-SEC were enriched for EV markers (e.g., CD9, CD63, and Alix), showing no difference between filtered and untreated ones (Figure 1h). Figure 1i shows a representative size distribution profile of UC-SEC and PEG-SEC based on NTA, which reflects a similar size distribution pattern of UC-SEC and PEG-SEC with the highest peak at approximately 100 nm.

3.2 | ADMSC-EVs protect neurons from oxygen-glucose-deprivation (OGD) injury

EVs enriched from bone marrow-derived MSCs induce neuroprotection and enhance neurological recovery in preclinical stroke models, as previously described by us and others (Bang & Kim, 2019; Doepfner et al., 2015; Xin et al., 2013). The present study, however, analysed the underlying mechanisms of ADMSC-EV-induced neuroprotection using both in vitro and in vivo stroke models. Primary neurons exposed to OGD followed by 24 h of reoxygenation displayed a significant extent of cell injury, depending on the duration of the OGD (Figure 2a). Although recent evidence suggests a role for autophagy to be involved in cell injury due to hypoxia or ischemia, published data appears to be contradictory with regard to autophagy being beneficial or detrimental (Chen et al., 2014; Wei, Wang, & Miao, 2012). We, therefore, analysed autophagy levels using the expression of the autophagy-

associated protein LC3-II in our OGD model. LC3-II levels significantly correlated with the extent of cell injury, i.e., LC3-II was significantly more abundant in neurons exposed to 10 h of OGD than in neurons exposed to 4 h or 8 h of OGD (Figure 2b-c). Further, *in vitro* experiments were performed using the 10 h OGD experimental paradigm only. Treatment of primary neurons exposed to OGD with ADMSCs or with EVs isolated from UC or PEG yielded a significant reduction of cell injury when compared to controls (Figure 2d). No differences between the various methods used were observed. Interestingly, ADMSC-EV treatment was not inferior to ADMSC treatment with regard to survival rates of neurons exposed to OGD (Figure 2d).

3.3 | ADMSC-derived EVs inhibit autophagic flux and protect primary neurons from OGD-injury through regulation of autophagy

In order to study whether or not ADMSC-derived EVs enhance the resistance of primary neurons against OGD by regulating autophagic activity, autophagy levels were assessed using the expression of autophagy-associated protein LC3-II. When primary neurons were exposed to OGD and co-cultured with ADMSCs or treated with ADMSC-derived EVs enriched from UC or PEG, the stroke-induced increased protein abundance of LC3-II was reversed (Figure 2e-f). We also compared the effects of the concentrates from iodixanol separated layers one, three, and seven on alleviating cell damage after iodixanol density gradient ultracentrifugation (DGUC). Not surprisingly, the seventh layer of concentrates, but not the first or the third layer which occupied a typical sEV diameter, has a significant effect on enhancing cell viability. This effect is the same as the unstratified one (Figure S2A). Consistent with the results from DGUC, SEC filtered EVs have the same effects as the unfiltered EVs with regard to reducing cell damage after OGD, regardless of the enrichment method chosen (Suppl. Fig. S2B). More importantly, all of the manipulated EVs (iodixanol and SEC) can down-regulate neuronal autophagy levels after OGD, and the effects were not inferior to the unpurified one (Figure S2C-D). Combined with the aforementioned results, we showed that sEVs are the key compounds mediating reduced levels of autophagy and neuroprotection. Importantly, we confirmed that sEVs are the key compounds in both EV-isolation methods (UC and PEG), and that these two methods display no significant difference between each other. We therefore used the PEG method to enrich ADMSC-EVs for the remainder of the study.

To confirm the physiological properties of ADMSC-EVs in the aforementioned OGD model, the ADMSC-EV uptake in primary neurons was studied using DiI staining. Indeed, primary neurons displayed an intracellular uptake of EVs after OGD (Figure S2E). We then asked the question whether or not EV-induced effects are dose-dependent. Hence, three different doses (2×10^3 , 2×10^4 , and 2×10^5 cell equivalents) of ADMSC-EVs were used in the OGD model. ADMSC-EVs significantly repressed the LC3-II expression (Figure S2F-G) and rescued cell injury (Figure S2H) in a dose-dependent manner. Although we do not have clear evidence, it appears to be feasible that high dosages (2×10^5) might lead to excessively down-regulated autophagy levels, resulting in the loss of further protection from EVs to neurons after OGD. On the contrary, low dosages (2×10^3) of EVs result in an insufficient concentration of either cargo proteins or non-coding RNAs, thus failing to induce neuroprotection. However, the medium dosages (2×10^4) of ADMSC-EVs showed beneficial effects regarding cell survival. Further experiments were therefore performed choosing the medium dosage.

In addition, we also measured the cell survival after co-culturing neurons with ADMSC-EVs that were pretreated with either RNase A, the detergent Triton X-100 (to disrupt the lipid bilayer of vesicles), both of them, or with the untreated EVs after OGD as shown in Figure S2I. When EVs were pretreated with RNase A, but not with Triton X-100, and co-cultured with primary neurons, there was no significant difference between RNase A treated and the untreated group in terms of cell viability after OGD. Pretreating EVs with Triton X-100 alone resulted in some decrease in neuroprotection. Conversely, pretreating EVs with Triton X-100 and RNase A led to near-total elimination of the protective properties of EVs. These results support the hypothesis that EVs are likely to be the therapeutically efficacious substance in our OGD model.

Although LC3-II, which is specifically recruited to the autophagosome membrane (Mizushima & Yoshimori, 2007), is a good marker for analyzing autophagy, the static nature of such measurements is hard to interpret. As a matter of fact, an increase in LC3-II levels can either be a consequence of enhanced autophagosome formation or due to a lack of autophagosome fusion with lysosomes (du Toit, Hofmeyr, Gniadek, & Loos, 2018). To distinguish between these two states of autophagy processing, we evaluated the effect of autophagy by treating primary neurons with bafilomycin A1 (BafA1), which is a potent V-ATPase inhibitor that impedes autophagosome-lysosome fusion. If the breakdown of autophagosomes is blocked, the absolute LC3-II accumulation (BafA1 treated minus the untreated one) indicates how many new autophagosomes are being produced over a particular time. The magnitude of BafA1-induced LC3-II accumulation under OGD conditions was significantly lower in primary neurons cultured with either ADMSCs or EVs when compared to controls (Figure 3a-b). Furthermore, we employed a tandem fluorescence RFP-GFP-LC3B reporter system to monitor the autophagic flux in primary neurons. Incubation of primary neurons exposed to OGD with ADMSC-EVs yielded a reduced number of both autophagosomes and autolysosomes (Figure 3c-d).

To further reveal the relationship between ADMSC-EVs and autophagy, the autophagy activator rapamycin and the inhibitor 3-MA were used in primary neurons exposed to OGD (Figure 3e). The combined treatment of neurons with different concentrations of rapamycin and ADMSC-EVs resulted in a reversal of the former EV-induced neuroprotection, indicating that ADMSC-EVs apparently act in an opposite way than rapamycin. On the contrary, inhibition of autophagy using 3-MA signif-

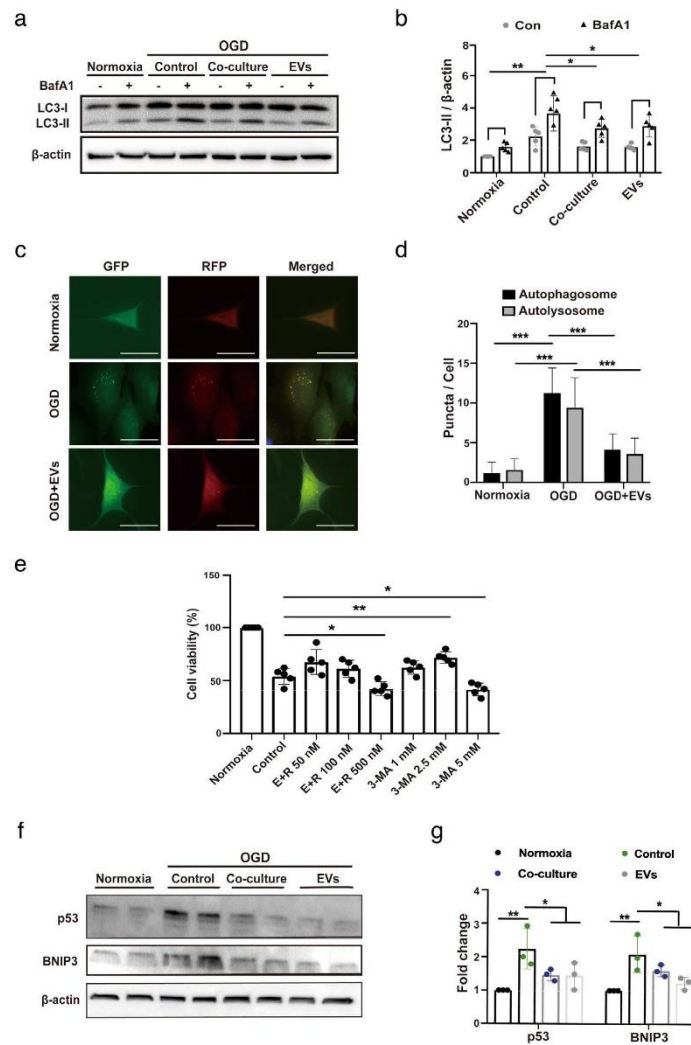


FIGURE 3 ADMSC-EVs inhibit autophagic flux and increase cell viability through p53 and BNIP3 signalling. (A-B) Assessment of the autophagic flux was done using bafilomycin A1 (BafA1) in the aforementioned groups (Normoxia, Control, Co-culture and EVs). BafA1 was added 3 h before harvesting the cells. LC3 levels were evaluated again by Western blotting in the presence of DMSO or BafA1 ($n = 3$). Quantitative analysis of LC3-II blotting is shown in (b). (c-d) Autophagosomes (yellow) and autolysosomes (red) were detected in OGD-exposed primary neurons that express mRFP-GFP-LC3. The neurons were treated with PBS or ADMSC-EVs. PBS and EVs were given at the beginning of hypoxia and reoxygenation. Cells incubated under standard cell culture conditions (Normoxia) were used as negative control. Scale bar, 10 μ m. The number of autophagosomes and autolysosomes in each cell (20-30 cells per group) was quantified in (d) ($n = 3$). (e) The impact of autophagy on neuronal survival after OGD was evaluated using different concentrations of the autophagy inhibitor 3-MA in comparison to neurons treated with the solvent DMSO using the MTT assay. ADMSC-EVs together with different concentrations of the autophagy stimulator rapamycin (E+R; $n = 3$) were also used on primary neurons exposed to OGD. All the drugs were given twice, at the beginning of hypoxia and reoxygenation. Cells incubated under standard cell culture conditions (Normoxia) were defined as 100 % cell survival. (f-g) Both p53 and BNIP3 were evaluated by Western blotting in OGD exposed primary neurons treated with PBS, ADMSCs (Co-culture) or ADMSC-EVs (EVs). Neurons treated with PBS under OGD conditions served as positive control (Control). Cells incubated under standard cell culture conditions (Normoxia) were used as negative control ($n = 3$). EVs were given at the beginning of hypoxia and reoxygenation. The quantitative analysis of p53 and BNIP3 Western blotting is shown in (g). One-way ANOVA followed by the Tukey's post-hoc-test, data are given as mean \pm SD. Data are statistically different from each other with * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$

icantly increased neuron viability in a concentration dependent manner (Figure 3e). The protective impact of 3-MA, however, was lost when 3-MA was applied in a concentration of 5 mM, suggesting that over-suppression of autophagy may be detrimental for primary neurons. Rather, a moderate decrease in autophagy confers significant protection against OGD-induced cell death in primary neurons.

3.4 | ADMSC-derived EVs regulate autophagy by signalling pathways involving p53 and B-cell lymphoma 2-interacting protein 3 (BNIP3)

To understand the mechanisms by which ADMSC-derived EVs regulate autophagy, we investigated autophagy-related signalling cascades. Autophagy is known to be regulated by at least four signalling pathways, among which are p53 and BNIP3. Interestingly, protein levels of p53 and BNIP3 were significantly increased in OGD-exposed primary neurons when compared to standard cell culture conditions (Figure 3f-g). Incubation of neurons with either ADMSCs or with ADMSC-EVs significantly decreased the OGD-induced upregulation of both p53 and BNIP3 (Figure 3f-g), indicating that ADMSC-EVs possibly modulate autophagy through p53 and BNIP3 signalling.

3.5 | Inhibition of exosome biogenesis reverses ADMSC-EV-induced effects in neurons exposed to OGD

As mentioned before, exosomes form an important subgroup of EVs. Two mechanisms of exosome formation have been described by ceramide synthesis (Trajkovic et al., 2008) and by the endosomal sorting complex required for transport (ESCRT) machinery (Hanson & Cashikar, 2012), respectively. To determine the role of exosomes in the regulation of autophagy and neuroprotection in primary neurons exposed to OGD, we inhibited ceramide synthesis with the neutral sphingomyelinase-targeting inhibitor GW4869 and inhibited ESCRT-mediated exosome biogenesis by knocking down Hrs, a member of the ESCRT-0 complex (Tamai et al., 2010). As expected, both Hrs-KD and GW4869 pre-treatment led to a large decrease in exosome secretion shown by NTA analysis (Figure 4a), showing no difference between the two methods. Moreover, the effects of ADMSC-EVs (EVs) on autophagy regulation (Figure 4b-c) and neuroprotection (Figure 4d) were significantly blocked when ADMSCs were pretreated with the inhibitor of exosomal secretion GW4869 or Hrs-KD. Then, we applied GW4869 in the coculture model, the effects of ADMSC-EVs (EVs) on neuroprotection (Figure 4d) and autophagy regulation (Figure 4e-f) were significantly blocked when ADMSCs were pretreated with the inhibitor of exosomal secretion GW4869.

Importantly, the incubation with ADMSC-EVs down-regulated the abundance of p53 and BNIP3 in neurons exposed to OGD, whereas incubation of neurons with EVs obtained from such pretreated ADMSCs failed to inhibit p53 and BNIP3 protein levels (Figure 4g-h). Thus, exosomes forming a subgroup of ADMSC-EVs appear to be critical in mediating the anti-autophagic activity of ADMSCs in primary neurons exposed to OGD.

3.6 | The miR-25-3p is abundant in ADMSC-EVs reducing the autophagic flux in hypoxic primary neurons

To further identify the compounds of EVs that are responsible for ADMSC-EV-induced regulation of autophagy, we first explored miRNAs that directly target the p53/BNIP3 autophagy signalling pathway. Out of more than 10 miRNAs known to target p53 (Liu, Zhang, Zhao, & Feng, 2017), we selected 6 candidates miRNA (miR-25-3p, miR-98, miR-125a-5p, miR125b-5p, miR-214-3p, and miR-30) that are more closely related to neurons for further screening. In order to eliminate the bias due to the selection of internal controls, we have selected two internal controls, U6 and let-7a, for relative quantification of miRNAs (Li et al., 2015). The results from qRT-PCR analyses indicated that miR-25-3p was one of the most abundant miRNAs in ADMSC-EVs (Figure 5a). In order to show in which physicochemical state miR-25 can be found in our conditions, we treated EVs with either RNase A, the detergent Triton X-100 (to disrupt the lipid bilayer of vesicles), both of them, or with the solvent alone (Control). We then measured the levels of miR-25 by RT-qPCR. When EVs were treated with RNase A, but not with Triton X-100, the PCR signal for the miR-25 remained within the margin of error compared to the signal for solvent treated one. This suggests that RNAs captured by the column were located inside vesicles and were therefore protected by the lipid bilayer from RNase A digestion (Figure 5b). Treating EVs with Triton X-100 alone resulted in some degradation of RNA, supposedly because disruption of the vesicle membrane exposes the RNA to residual RNase present in the environment. Conversely, treating with Triton X-100 and RNase A led to near-total digestion of the RNAs (Figure 5b), confirming that their initial resistance to digestion was due to sequestration within vesicles, which proved miR-25 to be inside of ADMSC-EVs. Nevertheless, the intracellular concentration of miR-25-3p in primary neurons significantly decreased when these cells were exposed to OGD (Figure 5c). On the contrary, treatment of primary neurons with ADMSC-EVs significantly increased the concentration of miR-25-3p (Figure 5c).

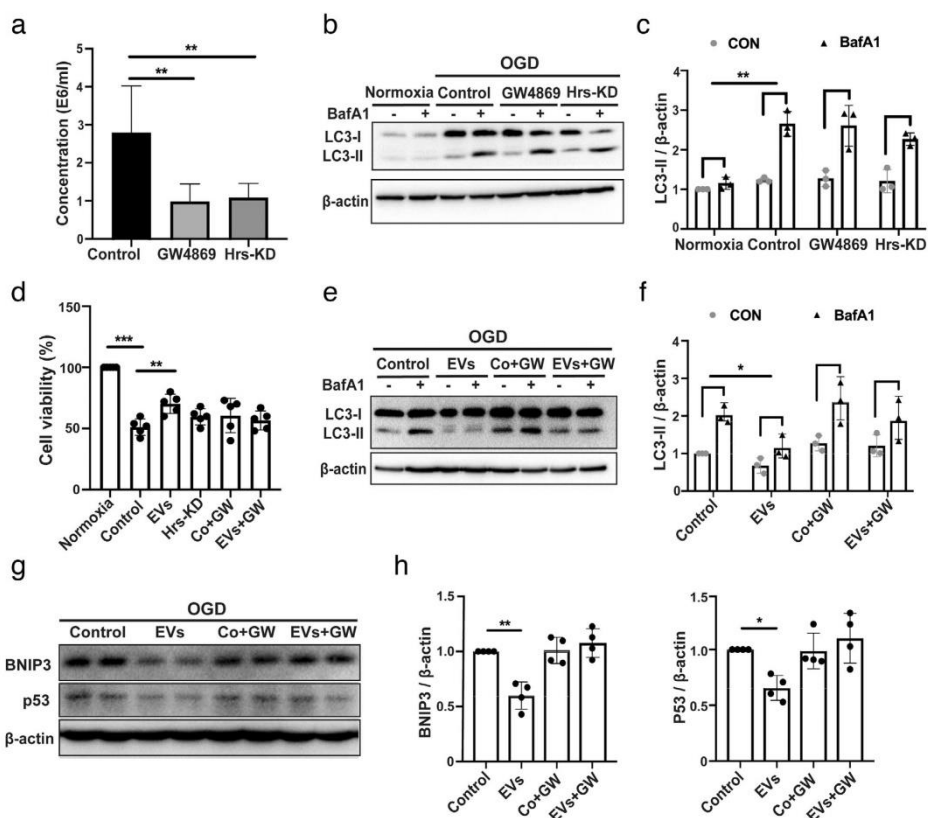


FIGURE 4 The regulation of the autophagic flux by ADMSC-EVs depends on exosomes. (a) ADMSC-derived EVs obtained from ADMSCs pre-treated with DMSO, GW4869 or Hrs-siRNA (knockdown, Hrs-KD) were isolated by the PEG method. The quantification of EV numbers is shown in (A) ($n = 5$). (b-c) LC3 levels were detected by Western blotting followed by densitometric analysis in primary neurons exposed to oxygen-glucose-deprivation (OGD). OGD-exposed neurons were either incubated with EVs isolated from ADMSCs pre-treated with the exosome secretion inhibitor GW4869 (GW4869) or with ADMSC-derived EVs obtained from ADMSCs that were pre-transfected with Hrs-siRNA (Hrs-KD). All experimental conditions were performed with or without the autophagic flux inhibitor BafA1 ($n = 3$). (d) Cell viability was examined in OGD-exposed primary neurons that were treated with PBS, ADMSC-EVs (EVs), ADMSC-derived EVs obtained from ADMSCs that were pre-transfected with Hrs-siRNA (Hrs-KD), ADMSCs treated with the exosome secretion inhibitor GW4869 (Co+GW) or ADMSC-EVs isolated from conditioned medium containing GW4869 (EVs+GW; $n = 3$). Cells incubated under standard cell culture conditions (Normoxia) were defined as 100% cell survival. (e-f) OGD-exposed neurons were either incubated with ADMSC-derived EVs (EVs), ADMSCs pre-treated with the exosome secretion inhibitor GW4869 (Co+GW), or with ADMSC-derived EVs obtained from ADMSCs that were pretreated with GW4869 (EVs+GW). EVs, GW4869, EVs+GW were given at the beginning of both hypoxia and reoxygenation. All experimental conditions were performed with or without the autophagic flux inhibitor BafA1 ($n = 3$). BafA1 was added 3 h before harvesting the cells. (g-h) P53 and BNIP3 were quantified by Western blotting in OGD-exposed primary neurons co-incubated with PBS, ADMSC-EVs (EVs), ADMSCs with GW4869 (Co+GW) or with EVs isolated from ADMSCs pre-treated with GW4869 (EVs+GW; $n = 3$). Quantitative analysis of p53 and BNIP3 is shown (h). One-way ANOVA followed by the Tukey's post-hoc-test was used, data are shown as mean \pm SD. Data are statistically different from each other with * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$

To confirm whether or not miR-25-3p has a functional role in the aforementioned EV-induced regulation of autophagy under OGD conditions of neurons, EVs were obtained from ADMSCs that were pretreated with an anti-miR-25-3p oligonucleotide (ADMSC-EV^{anti-miR-25}) or with a scrambled construct as control (ADMSC-EV^{NC}). The cell viability after OGD was significantly higher in primary neurons cultured with ADMSC-EVs^{NC} (EVs^{NC}) when compared to PBS-treated controls (Figure 5d). However, these effects were reversed when the cells were treated with ADMSC-EVs^{anti-miR-25} (EVs^{anti-miR-25}, Figure 5d). The magnitude of BafA1-induced LC3-II accumulation under OGD conditions was also significantly lower in primary neurons cultured with ADMSC-EVs^{NC} when compared to PBS-treated cells (Figure 5e-f), whereas these results were again reversed when the neu-

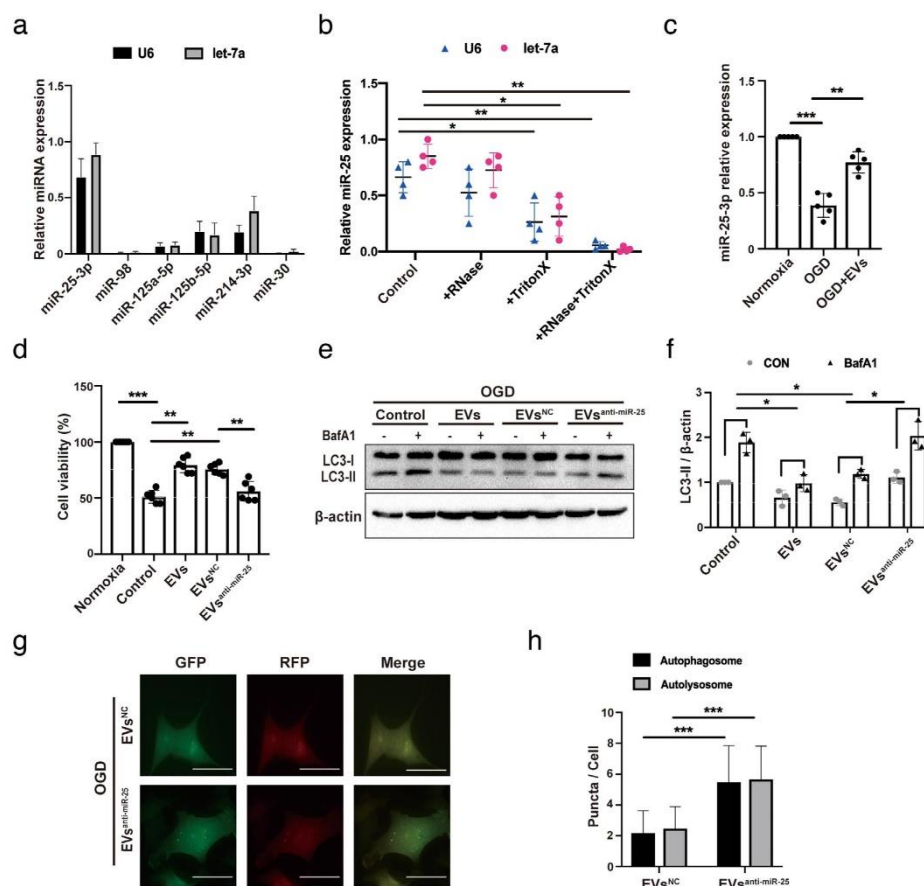


FIGURE 5 ADMSCs regulate autophagy and induce neuroprotection by miR-25-3p. (a) Real time Quantitative Polymerase chain reaction (qRT-PCR) quantification of miRNA concentrations in EVs obtained from ADMSCs as stated in the materials and methods section. (b) Levels of miR-25-3p were detected in ADMSC-EVs that were pretreated either RNase A (+RNase), the detergent Triton X-100 (+Triton X), both of them (+RNase+Triton X), or with the solvent alone (Control) ($n = 3$). (c) Levels of cellular miR-25-3p were measured in primary neurons treated with PBS or ADMSC-EVs in the OGD model by qRT-PCR. PBS and EVs were given at the beginning of hypoxia and reoxygenation. Data refer to neurons cultured under standard cell culture conditions (Normoxia). (d) Cell viability was examined in primary neurons exposed to OGD that were treated with PBS, EVs obtained from normal ADMSCs (EVs), EVs isolated from ADMSCs that were pretreated with anti-miR-25-3p (EVs^{anti-miR-25}) or with EVs isolated from ADMSCs that were pretreated with scramble (EVs^{NC}; $n = 3$). All of them were given at the beginning of hypoxia and reoxygenation. Cells incubated under standard cell culture conditions (Normoxia) were defined as 100 % cell survival. (e-f) Western blot identification for LC3 in primary neurons after incubation with different EVs, i.e., EVs isolated from normal ADMSCs (EVs), EVs obtained from ADMSCs that were pretreated with anti-miR-25-3p (EVs^{anti-miR-25}) or with EVs isolated from ADMSCs that were pretreated with scramble (EVs^{NC}). OGD experiments were performed with or without BafA1 under each condition ($n = 3$). Quantitative analysis of LC3-II is shown in (f). (g-h) Autophagosomes and autolysosomes were detected in EVs^{anti-miR-25}-treated primary neurons or in EVs^{NC}-treated neurons that expressed mRFP-GFP-LC3 under OGD conditions. The number of autophagosomes and autolysosomes in each cell (20-30 cells per group) was quantified ($n = 3$). EVs^{anti-miR-25} and EVs^{NC} were given twice as mentioned before. Scale bar, 10 μ m. The two-tailed independent Student's t -test was used in H. One-way ANOVA followed by the Tukey's post-hoc-test was used except for H, data are shown as mean \pm SD. Data are statistically different from each other with * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$

rons were incubated with ADMSC-EV^{anti-miR-25} (Figure 4e-f). Likewise, using a luciferase system, a significant increase in the number of autophagosomes and autolysosomes was counted in neurons incubated with ADMSC-EVs^{anti-miR-25} (EVs^{anti-miR-25}) compared to ADMSC-EVs^{NC} (EVs^{NC}) (Figure 5g-h). Thereafter, the effects of miR-25-3p on the regulation of autophagy were further analysed, using a miR-25-3p mimic and its inhibitor directly in primary neurons that were exposed to OGD. Hypoxic neurons transfected with the miR-25-3p mimic recapitulated the inhibitive effects on both autophagy regulation (Figure S3A-B)

and autophagic flux (Figure S3C-D) when compared with the scramble group. Through the live/dead assay, we found that the transfection itself did not increase cell death compared with the untransfected group, suggesting that the autophagy regulating effect was due to miR-25a itself but not due to the protein loss after cell death (Figure S3E-F). Conversely, transfection of neurons with the miR-25 inhibitor yielded opposite results (Figure S3G-H). Being consistent with the autophagic flux, the levels of p53 and BNIP3 were decreased in response to the treatment with the miR-25-3p mimic (Figure S3I-J), while abundance of these proteins was enhanced when the neurons were treated with the miR-25-3p inhibitor (Figure S3 H-L). Finally, incubation of neurons exposed to hypoxia with the miR-25-3p mimic resulted in increased cell viability (Figure S3M), whereas inhibition of miR-25-3p decreased neuron viability (Figure S3N).

3.7 | ADMSC-EVs induce sustained neuroprotection and promote neurological recovery after stroke in mice

In light of the aforementioned *in vitro* data on primary neurons, we investigated whether or not ADMSC-EV administration improves post-stroke neurological recovery by modulating autophagic activity. We first checked the EV biodistribution patterns under ischemic conditions. Consistent with previous results (Zagreen, Hermann, Opris, Zagreen, & Popa-Wagner, 2018), ADMSC-EVs indeed do reach the central nervous system, at least under ischemic stroke conditions (Figure 6a). Following a previously published protocol (Doepfner et al., 2015), EVs released by 2×10^6 ADMSCs diluted in 200 μ l of PBS were systemically administered immediately at the beginning of the reperfusion or at 12 h after reperfusion. Mice that received ADMSC-EVs immediately during the beginning of the reperfusion exhibited significantly smaller infarct volumes when compared to PBS controls (Figure 6b-c). Likewise, treatment of stroke mice with EVs at 12 h also yielded significantly reduced infarct volumes when compared to controls, with no significant difference between the two EV groups (Figure 6b-c). Along with such a reduction of acute brain injury, the behavioural test analyses revealed better test scores of mice treated with EVs at either time point (Figure 6d-e). Of note, this better test performance in both the corner turn and the tight rope test was long-lasting and stable until the end of the observation period of 14 days. We subsequently analysed neuronal survival in the ischemic striatum at 14 days after the stroke. In line with the reduction of neurological impairment, increased neuronal densities were found in mice treated with ADMSC-EVs of both injection time points (Figure 6f-g). Conclusively, ADMSC-derived EVs reduce post-stroke brain injury on both the histological and the functional level in experimental stroke models.

3.8 | ADMSC-EV administration after MCAO reduces autophagic flux through p53-BNIP3 signalling

To further confirm whether autophagy inhibition contributes to neuroprotection and functional improvement, we used 3-MA to inhibit autophagy after MCAO, and then assessed neuronal survival as well as neurological recovery. Treatment of mice with the inhibitor of autophagy, 3-MA, at 12 h post-stroke confirmed the aforementioned results of the behavioural tests, thus further supporting the hypothesis that inhibition of autophagy contributes to neurological recovery upon stroke induction (Figure 6d-e). Whereas delivery of 3-MA at 12 h post-stroke yielded better neurological recovery in these mice when compared to PBS controls, delivery of 3-MA immediately at the beginning of the reperfusion only partially enhanced neurological recovery (Figure 6d-e). Analysis of neuronal survival at 14 days post-stroke showed increased neuronal densities in mice treated with 3-MA at 12 h, whereas treatment immediately at the beginning of the reperfusion did not result in neuroprotection (Figure 6f-g). Again, treatment of stroke mice with 3-MA at 12 h significantly reduced the autophagic flux 24 h post-surgery as indicated when BafA1 was simultaneously used in these animals. The LC3-II levels were significantly lower in these mice as compared to the PBS group (Figure 6h-i). In contrast to this, treatment with 3-MA immediately during the beginning of the reperfusion had no impact on the autophagic flux. These results suggest that modulating post-stroke over-activated autophagic activity induces neuroprotection in mice after induction of MCAO.

To further investigate whether or not neuroprotection caused by ADMSC-EVs is associated with a regulation of autophagy, the temporal resolution patterns of autophagy levels in non-treated stroke mice were assessed. Western blotting against LC3-II revealed a peak of protein abundance at 24 h post-stroke when compared to sham animals (Figure S4A-B). Treatment of mice with ADMSC-EVs immediately at the beginning of the reperfusion (EVs 0 h; Figure S4C-D) or at 12 h (EVs 12 h; Figure S4E-F) significantly reduced the stroke-induced activation of autophagy as measured at 24 h post-stroke. These observations suggest that ADMSC-EVs significantly reduce the stroke-induced autophagic flux in mice.

We next detected the expression of p53 and BNIP3 *in vivo*. Consistent with our *in vitro* data, both p53 and BNIP3 protein levels were increased in the infarct region of MCAO mice. Application of ADMSC-EVs at the aforementioned two injection time points (0 h and 12 h) significantly reduced abundance of both p53 and BNIP3 in stroke mice (Figure S4G-H).

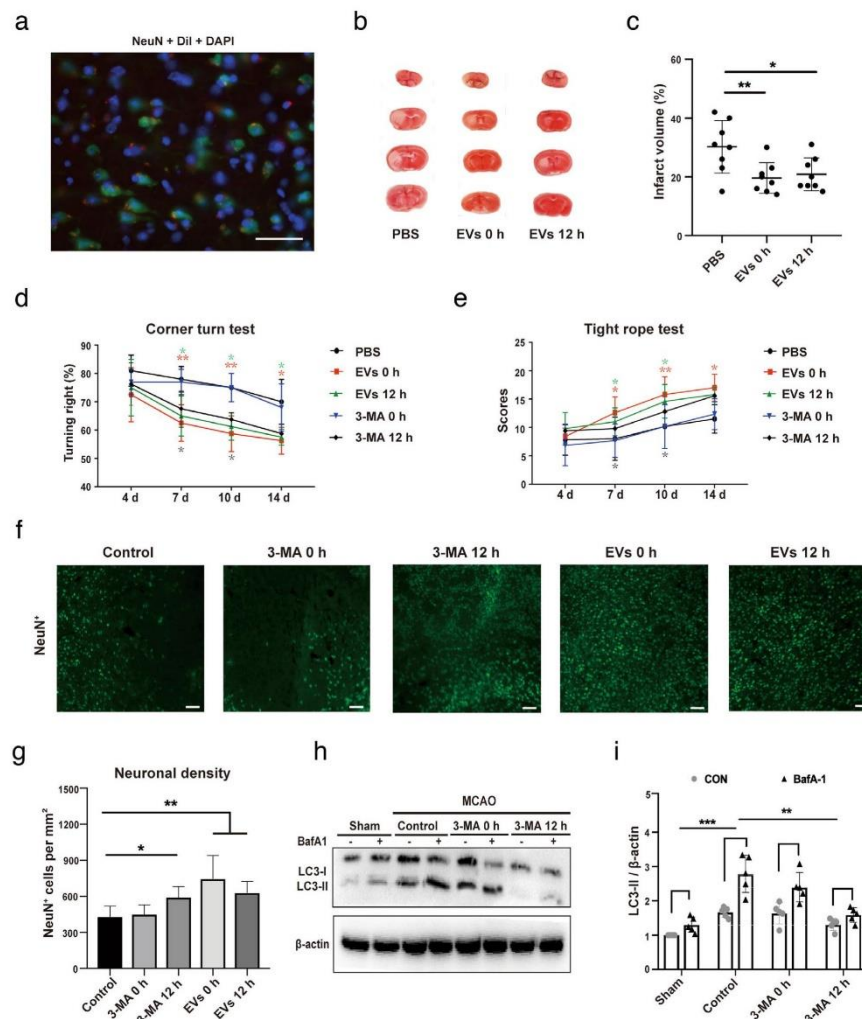


FIGURE 6 EV-induced regulation of autophagy reduces post-stroke brain injury and improves neurological recovery. (a) Representative immunofluorescence images displaying the biodistribution of ADMSC-EVs within the ischemic hemisphere. Dil (red spots), NeuN (green) and DAPI (blue). Scale bars, 25 μ m. (b-c) Neuroprotective effects of ADMSC-EVs in mice exposed to 1 h of middle cerebral artery occlusion (MCAO) followed by 24 h of reperfusion were evaluated by TTC staining. ADMSC-EVs were injected at the beginning of the reperfusion (EVs 0 h) or at 12 h after reperfusion (EVs 12 h). Mice treated with PBS served as control ($n = 8$ per group). Quantitative analysis of the infarct size is shown in (c). (d-e) Mice ($n = 10$ per condition) were exposed to 1 h of MCAO with subsequent reperfusion for 14 days during which the corner turn test (d) and the tight rope test (e) were performed. Mice received systemic delivery of ADMSC-EVs or of 3-MA (15 mg/kg) immediately at the beginning of the reperfusion (EVs 0 h) or 12 h after reperfusion (EVs 12 h). Control mice received PBS only. Motor coordination tests were done at 4, 7, 10 and 14 days after cerebral ischemia. Both EVs 0 h and EVs 12 h groups showed significant improvement in the tight rope test compared to the PBS control group. On day fourteen, only EVs 0 h significantly improved tight rope performance. In the corner turn test, the EVs 0 h and 12 h group showed improvement on day seven, ten and fourteen compared to the PBS control group, and the 3-MA 12 h group showed improvement on day seven and day ten compared to the PBS control group. (f-g) The neuronal density was measured in mice treated with PBS (Control), ADMSC-EVs or of 3-MA. EVs and 3-MA were systemically injected at the beginning of the reperfusion or 12 h after reperfusion. NeuN staining within the ischemic lesion site was done on day 14 ($n = 10$ per group). Scale bars, 50 μ m. (g-h) The autophagic flux was evaluated with BafA1 in MCAO mice that received 3-MA injection immediately at the beginning of (3-MA 0 h) or 12 h after reperfusion (3-MA 12 h). PBS was given to control animals ($n = 6$ per group). Quantitative analysis of LC3-II is shown (h). One-way ANOVA followed by the Tukey's post-hoc-test was used, data are shown as mean \pm SD. Data are statistically different from each other with $*P < 0.05$, $**P < 0.01$, and $***P < 0.001$

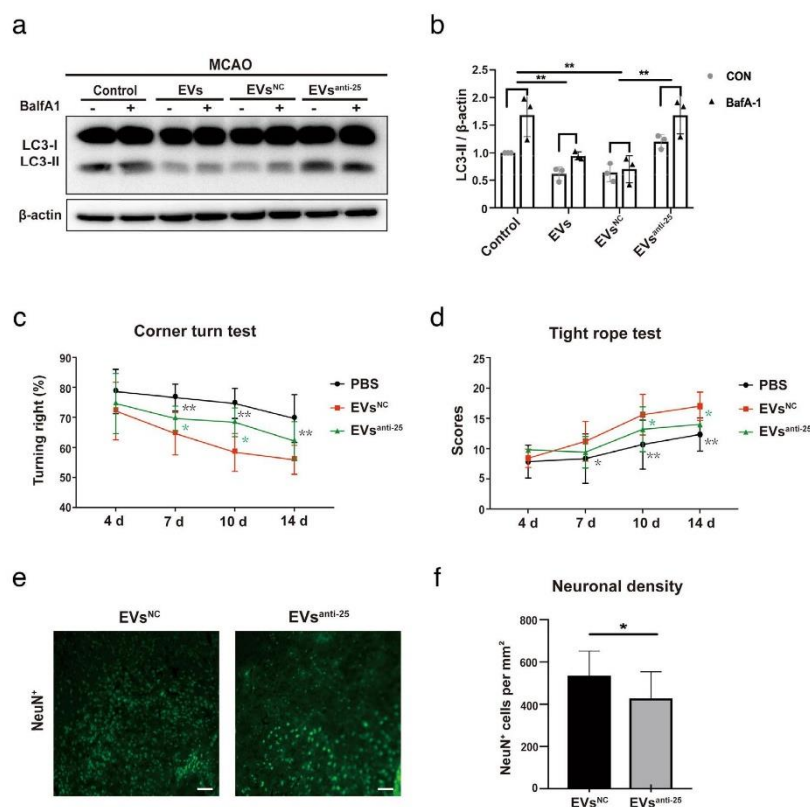


FIGURE 7 Loss of miR-25-3p in ADMSC-EVs diminishes post-stroke EV-induced regulation of autophagy and neuroprotection. (a-b) The autophagic flux was evaluated with BafA1 in MCAO (middle cerebral artery occlusion) mice that either received PBS or ADMSC-EVs 12 h after the induction of ischemic stroke, including ADMSC-EVs, ADMSC-EVs^{NC}, and ADMSC-EVs^{anti-miR-25} ($n = 6$ per group). BafA1 was injected 3 h before sacrifice. Quantitative analysis of LC3-II is shown in (b). (c-d) Motor coordination was evaluated by the tight rope test and by the corner turn test at four, seven, ten and fourteen days after cerebral ischemia to verify the effects of ADMSC-EVs^{NC} and ADMSC-EVs^{anti-miR-25} in MCAO animals compared to PBS group. In the corner turn test, the EVs^{NC} group showed improvement on day seven, ten and fourteen compared with PBS control group. The EVs^{NC} group showed improvement on day seven and day ten compared to the EVs^{anti-miR-25} group. EVs^{NC} group showed significant improvement in the tight rope test compared to the PBS control group. EVs^{NC} groups also showed significant better tight rope performance compared to EVs^{anti-miR-25} on day ten and fourteen. (e-f) The neuronal density in ADMSC-EVs^{NC} and ADMSC-EVs^{anti-miR-25} was measured on day fourteen as indicated by NeuN staining within the ischemic lesion site ($n = 10$ per group). The quantitative analysis of neuronal density is shown in (f). Scale bars, 50 μ m. The two-tailed independent Student's *t*-test was used in. One-way ANOVA followed by the Tukey's post-hoc-test was used except for F, data are shown as mean \pm SD. Data are statistically different from each other with * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$.

3.9 | Post-stroke autophagic regulation by ADMSC-EVs depends on miR-25-3p

In order to confirm that the decline in autophagic flux associated with ADMSC-EV administration is mediated by the EV-delivery of miR-25-3p, mice were injected with either PBS, ADMSC-EVs, ADMSC-EVs that had been pretreated with anti-miR25-3p (EVs^{anti-miR25}), or with ADMSC-EVs that had been pretreated with the control oligonucleotide (EVs^{NC}). The injection was done 12 h after MCAO surgery. After 24 h of reperfusion, the BafA1-induced LC3-II accumulation in the ischemic striatum was significantly lower in both the ADMSC-EV and the ADMSC-EVs^{NC} groups compared to the PBS group (Figure 7a-b). However, treatment with ADMSC-EVs^{anti-miR25} failed to show similar effects on the autophagic flux (Figure 7a-b).

Consistently, neurological recovery as assessed by the corner turn and the tight rope test was enhanced in mice treated with ADMSC-EVs^{NC} when compared to both controls and mice treated with ADMSC-EVs^{anti-miR25} (Figure 7c-d). Analysis of the neuronal density 14 days after MCAO revealed that mice treated with ADMSC-EVs^{NC} displayed reduced brain injury compared

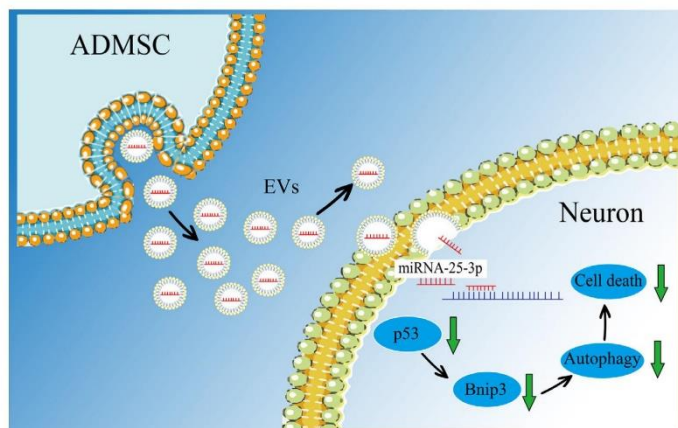


FIGURE 8 A schematic diagram showing the role of miR-25-3p derived from ADMSC-EVs in a preclinical stroke model. ADMSCs release EVs enriched with miR-25-3p, which are uptaken by neurons. In neurons, miR-25-3p induces degradation of the mRNA of p53, resulting in the downregulation of the p53 protein level and subsequent reduction of BNIP3. The inhibition of BNIP3, in turn, further reduces the levels of autophagy which exerts the neuroprotective effect

to animals treated with ADMSC-EVs^{anti-miR25} Figure 7e-f). Conclusively, ADMSC-derived EVs thus reduce brain injury on both the histological and the functional level. The anti-autophagic activity associated with ADMSC-EV administration after MCAO is at least partially mediated by the EV transfer of miR-25-3p.

4 | DISCUSSION

EVs from various tissue sources induce both neuroprotection and neuroregeneration in preclinical stroke models. The underlying mechanisms of such EV-induced beneficial effects under stroke conditions, however, remain elusive. Using both an in vitro and an in vivo stroke model, the present work identified a new mode of action by which EVs derived from ADMSCs mediate post-stroke neuroprotection and neurological recovery. The findings presented herein suggest that ADMSC-EVs are able to inhibit ischemia-induced autophagy. A key mechanism of ADMSCs-induced neuroprotection involves the EV-mediated transfer of miR-25-3p from ADMSCs to the neuron, where it interferes with the p53/BNIP3 signalling pathway (Figure 8).

Intercellular transfer of RNA by small EVs has opened a new avenue of research centred on these tiny cellular particles in diverse physiological systems. However, these small size particles have imposed numerous technical hurdles for their isolation and study. Related to EV isolation, the most popular method remains differential ultracentrifugation (UC). UC, however, is limited in scale and cumbersome (Lötvald et al., 2014). Other concentration methods have therefore to be adopted for concentrating these vesicles. The use of polyethylene glycol (PEG) has recently gained popularity due to its rapid and simple-to-use nature. Previous work from our own group systematically analysed the PEG precipitation approach in direct comparison with standard EV enrichment procedures on HEK293T cells (Ludwig et al., 2018). We revealed the PEG approach to be not inferior to standard EV enrichment procedures using HEK293T cells. In the present study, we also compared two different isolation methods, UC and PEG enrichment in the enrichment of ADMSC-EVs. The purification rates obtained by these methods were high for ADMSC-EVs, as indicated in the Western blot results. Especially, EV negative markers such as Albumin, TOMM20, and Histones were close to the detection threshold or not detectable at all in our UC and PEG method group. Likewise, the distribution and morphological patterns were similar between EVs enriched with either UC or PEG.

The PEG method has been reported to retain certain amount of protein contaminants (Patel et al., 2019), and these co-precipitated protein complexes may have related functions and affect the experimental outcome. To obtain what fraction is the active fraction mediating neuroprotection, it was necessary to combine several separation procedures and to follow a nonbiased approach (Freitas et al., 2019; Kowal et al., 2016). We showed that density gradient ultracentrifugation (DGUC) or size exclusion chromatography (SEC) can separate sEVs from other co-precipitates, which was indicated by strong signals of EV markers and typical particle diameters of 30–150 nm after purification. Moreover, the further purified sEV showed a similar performance on alleviating cell damage and reducing autophagy levels after ischemic stroke compared to unpurified EVs, suggesting that sEV are the therapeutically efficacious factors in our stroke models. In particular, both DGUC and SEC have a positive effect on the

determination of functional fractions for large-scale isolation of EVs from conditioned media by applying precipitation using PEG coupled with a DGUC or a SEC approach. Therefore, PEG offers an option to remedy the limitations of ultracentrifugation for the initial concentration step.

The pathophysiology of cerebral ischemia involves a plethora of different signalling cascades (Fricker, Tolkovsky, Borutaite, Coleman, & Brown, 2018). Among these different mechanisms, more recent data suggests a role of autophagy being an interesting target for stroke treatment (Descloux, Ginet, Clarke, Puyal, & Truttmann, 2015; Puyal, Ginet, Grishchuk, Truttmann, & Clarke, 2012; Wang et al., 2018). However, the precise role of autophagy under such ischemic stroke conditions is still under debate, i.e., autophagy might have both beneficial and detrimental effects. Some studies found pharmacological induction of autophagy by rapamycin to reduce apoptotic and necrotic cell death during hypoxia, whereas others reported inhibition of autophagy by 3-MA to enhance cell death (Carloni et al., 2010; Wang et al., 2012). Contrary to this, many groups reported that an increase in autophagy contributes to cell death. Using a model of excitotoxicity, the application of the glutamate receptor activator resulted in the death of cortical neurons, whereas delivery of the autophagy inhibitor 3-MA or the genetic knockdown of Atg7 and beclin-1 yielded neuroprotection (Ginet et al., 2014). Similar findings were supported by other models of cerebral ischemia (Koike et al., 2008; Xing et al., 2012).

In our study, autophagy was strongly activated upon induction of either cerebral ischemia or oxygen-glucose-deprivation (OGD). Activation of autophagy under such conditions represented a dynamic evolutionary process with a peak at 24 h after MCAO induction. Interestingly, down-regulation of autophagy using 3-MA at the onset of reperfusion did not induce any significant effect on brain injury, whereas 3-MA delivery at 12 h after reperfusion significantly improved both the neurological outcome and the neuronal density. *In vitro* studies confirmed the beneficial effect of inhibition of autophagy using 3-MA in a dose dependent manner. The lack of 3-MA when given immediately after the stroke in our experimental setting is most likely a consequence of the rather low half-life time of 3-MA with about 2–3 h (Sweet, Carda, & Small, 1981). As such, 3-MA is already metabolized before the peak of post-stroke autophagic activity is reached at 24 h. In light of both previous studies and the present work, it stands to reason that basal levels of autophagy are essential for proper neuronal function, with moderate increases of autophagic activity during mild conditions of ischemia, hypoxia, or nutrient deprivation still allowing neurons to maintain a sufficiently high energy supply (Kim et al., 2018). More severe noxious stimuli such as acute ischemic stroke, however, ensue an increase in autophagic flux that may culminate in autophagic cell death (Ginet et al., 2014; Liu & Levine, 2015). Consequently, a well-balanced inhibition of autophagy under stroke conditions seems to be in order (Shi et al., 2012).

Whereas pharmacological drugs such as 3-MA might only transiently affect post-stroke autophagy due to metabolism and short half-life time, stem cells like ADMSCs including their secretion products such as EVs are known to induce long-term effects under various circumstances (Lai, Yeo, Tan, & Lim, 2013; Tian et al., 2018). Indeed, the application of ADMSCs resulted in a pronounced downregulation of the autophagy marker LC3-II under both *in vitro* and *in vivo* conditions, contributing to both neuroprotection *in vitro* and *in vivo* as well as to neurological recovery *in vivo*. Interestingly, the inhibition of exosome secretion from ADMSCs resulted in a therapeutic loss of both such pretreated ADMSCs and their enriched EVs. The latter suggests a role for exosomes as a subgroup of the EV family that might be responsible for the biological effects observed in the study.

Inhibiting autophagy itself, however, does not exclusively predict the therapeutic impact of such a kind of therapy. Rather, the success of such a therapeutic intervention depends on the precise autophagic signaling pathway that is supposed to be inhibited. Multiple reports indicate that cerebral ischemia activates HIF-1 α which induces p53 or BNIP3 expression (Althaus et al., 2006; Li, Sun, Ni, Chen, & Guo, 2013). The induced p53 stabilization as a consequence of up-regulated HIF-1 α promotes further expression of BNIP3 (Wang et al., 2013; Xin et al., 2011). The latter contains a single Bcl-2 homology 3 (BH3) domain and belongs to the Bcl-2 family proteins that serve as an important target gene of HIF-1 α (Chinnadurai, Vijayalingam, & Gibson, 2008). BNIP3 can compete with beclin-1 for binding to Bcl-2. Released beclin-1, in turn, thus triggers autophagy and cell death (Glick, Barth, & Macleod, 2010; He & Klionsky, 2009). Herein, both p53 and BNIP3 levels were increased upon induction of hypoxia or cerebral ischemia, and ADMSC-EVs reversed the aforementioned increase of these proteins. The present study thus shows that autophagy inhibition plays a critical role in ADMSC-EV-based therapy of ischemic stroke, although other autophagy-related signalling pathways being regulated by ADMSC-EVs cannot be completely excluded.

The role of EVs in intracellular signalling, as well as their therapeutic potential for the treatment of ischemic stroke (Cunningham et al., 2018; Webb et al., 2018), has only recently become a prominent topic of research. Previous work from our own group and from other scientists have shown that EVs have both proangiogenic and anti-inflammatory properties (Dabrowska, Andrzejewska, Lukomska, & Janowski, 2019; Doepfner et al., 2015; Keshtkar, Azarpira, & Ghahremani, 2018; Xin et al., 2013). As indicated before, EVs do not only carry diverse sets of proteins (Doepfner et al., 2017; Doepfner et al., 2012; Doepfner et al., 2009; Doepfner, Kaltwasser, Fengyan, Hermann, & Bähr, 2013) but also contain non-coding RNA and DNA, among which microRNAs are of particular interest (Bertoli, Cava, & Castiglioni, 2015). The latter guides the binding of the RNA-induced silencing complex to regions of partial complementarity located mainly within 3' untranslated regions (UTRs) of target messenger RNA (mRNA) molecules, resulting in mRNA degradation and/or translational inhibition (Bartel, 2009). We have already demonstrated that miR-124, which is among the most abundant microRNAs in the adult mammalian brain (Cao, Pfaff, & Gage, 2007; Mishima, Mizuguchi, Kawahigashi, Takizawa, & Takizawa, 2007) affects a plethora of signaling molecules such as inhibition of deubiquitination of Usp14, significantly contributing to the reduction of post-stroke brain injury in rodents (Doepfner et al., 2013).

Moreover, it is well known that miRNAs impact the core autophagy pathway, including p53 (Frankel & Lund, 2012). Since EVs are able to transfer miRNA cargo to target cells (Abels & Breakefield, 2016), it is fair enough to hypothesize that ADMSC-EVs yield protection of neurons and brain tissue using this way of action. We herein demonstrated that ADMSC-EVs improve neurological recovery by impeding autophagy and the key anti-autophagic and cytoprotective compound of ADMSC-EV cargo may be miR-25-3p. The latter is one of several miRNAs found in EVs that target p53. Previous research on miR-25-3p has primarily focused on its biological function under conditions of cancer (Zeng et al., 2018; Zhang et al., 2019). To the best of our knowledge, our results from both in vitro and in vivo data suggested for the first time that miR-25-3p likely contributes to the pathogenesis of stroke. To test the hypothesis that miR-25-3p was responsible for ADMSC-EV-mediated neuroprotection, ADMSC-EVs that had been pretreated with anti-miR-25-3p were administered via femoral vein injection. As expected, the miR-25-3p knockdown abolished the therapeutic effects of ADMSC-EVs, as evidenced by the markedly decreased neuronal density and the lack of motor coordination recovery.

In conclusion, the findings presented herein provide novel insights into the mechanisms by which EVs induce therapeutic actions under experimental stroke settings. The data provides evidence that native ADMSC-EVs yield neuroprotection and enhanced neurological recovery by inhibiting ischemia-induced autophagy. Inhibition of autophagy, in turn, is mediated by EVs transferring miR-25-3p from ADMSCs to their target cells. The latter results in interference with the p53/BNIP3 signaling pathway. These novel observations on EVs derived from stem cells might lead to the development of new therapeutic targets and strategies for the treatment of ischemic stroke.

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AUTHOR CONTRIBUTION STATEMENT

Research and experiments were performed by Kuang, Zheng, Zhang, Ai, Venkataramani and Doeppner. Design and concept of the study were from Kuang and Doeppner. Kuang, Doeppner, Hermann, Kilic, Majid, and Bähr wrote the manuscript. Financial support was provided by Bähr and Doeppner.

CONFLICT OF INTEREST

The authors declare to have no conflict of interest of any kind.

ETHICAL APPROVAL

This article does not contain any studies with human participants performed by any of the authors. All animal experiments were performed with governmental approval according to the NIH guidelines for the care and use of laboratory animals. Both the STAIR criteria and the ARRIVE guidelines have been followed.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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8 Curriculum Vitae

My name is Kuang Yaoyun. I was born in Guiyang, China, on April 27, 1990. I studied clinical medicine at the Shenyang Medical College between 2008 and 2013. After graduating from college, I went on studying neuroscience and got my master's degree in 2016. During these 3 years of clinical internship and laboratory work, I found that physicians' work was just repeating routine work established by other people. But each patient is heterogeneous; the experience of the predecessor does not give you all the clinical guidance. There are still countless diseases that are largely untreatable, such as the most common neurological disease, ischemic stroke. Patients are often left with severe disabilities and suffer from poor quality of life after the onset of the disease. Unfortunately, the available neuroprotective drugs have not shown significant beneficial effects either. A large number of patients are unable to take care of themselves. Therefore, I became interested in discovering the new treatment of ischemic stroke, especially during the intervention in the subacute phase. So, I started my MD thesis in June 2018 under the supervision of Prof. Döppner at the Department of Neurology. We are dedicated to the study of mesenchymal stem cell-derived extracellular vesicles (EVs) for the treatment of ischemic stroke. From previous studies, numerous *in vitro* and *in vivo* works have revealed that EVs derived from MSCs have a neuroprotective role in ischemic stroke, but the exact mechanism remains unclear. Through my study, we found that ADMSCs are able to secrete large numbers of EVs that participate in intercellular signaling and regulating the functions of the recipient cells. By exploring the biological information contained in the EVs, the microRNAs inside it caught our interest. With inhibitory effects on the recipient cells at the transcriptional level, we have identified potential new targets for stroke intervention and provided new experimental evidence.