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# Studies on the role of cofilin signaling in hemin induced microglial activation

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A Thesis

entitled

Studies on the role of Cofilin signaling in Hemin induced Microglial activation

by

Muhammad Shahdaat Bin Sayeed

Submitted to the Graduate Faculty as partial fulfillment of the requirements for the

Master of Science Degree in Pharmaceutical Sciences

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May 2016

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An Abstract of  
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Intracerebral hemorrhage (ICH) is the most severe form of stroke and further exacerbates brain damage due to secondary injury following the initial burst. This secondary damage is partly due to the toxic effects of hemin, an endogenous breakdown product of hemoglobin. Microglia are CNS resident macrophages and the first to respond to any kind of insult (sterile or nonsterile) in the brain parenchyma. Cofilin is an actin depolymerizing factor and it has been primarily attributed to the control of actin dynamics.

Recently it was shown that cofilin is involved in mediating neuronal cell death in ischemic conditions. Cofilin is also involved in microglial activation induced by bacterial lipopolysaccharide. There are not many studies available that have clearly demonstrated the deleterious effects of extremely high doses of hemin released during ICH and its influence on microglia. Furthermore, the role of cofilin in this respect is still unknown. Therefore, investigations were conducted to study the effects of hemin on microglial activation and subsequent inflammation. The end-points measured were; NO and iNOS expression; cell survival; cytokine production (TNF- $\alpha$ , IL1- $\beta$ ); migration; hemin degrading

protein, HO-1 expression; Nrf2 expression; endoplasmic reticulum (ER) stress markers (Wfs-1, XBP-1, spliced XBP-1) and calcium ( $\text{Ca}^{2+}$ ) signaling.

In our studies, we found that hemin induces cofilin expression in microglia and increases NO production dose dependently. There were increased levels of iNOS, TNF- $\alpha$ , HO-1, Nrf2, Wfs-1, XBP-1 and spliced XBP-1 due to hemin treatment and were mediated by cofilin. However, hemin treatment did not affect IL-1  $\beta$  production. Acute hemin treatment did not evoke  $\text{Ca}^{2+}$  signaling but long-term treatment of hemin resulted in microglial failure to respond to acetylcholine evoked  $\text{Ca}^{2+}$  signaling. Knockdown of cofilin by siRNA also reduced acetylcholine evoked  $\text{Ca}^{2+}$  signaling. These studies demonstrate that cofilin signaling is important in ICH-induced brain injury and cofilin inhibition reduced hemin induced inflammation, oxidative stress, ER stress, microglial migration, and ability to evoke  $\text{Ca}^{2+}$  signaling and thus its inhibition can become novel drug therapeutics.

Keywords: Intracerebral Hemorrhage, Microglia, Hemin, Cofilin, Cell signaling

For the People of the United States of America

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## List of Abbreviations

μl	.....	Microliter
μM	.....	Micromolar
ABCG2	.....	ATP-binding cassette sub-family G member 2
ADF	.....	Actin depolymerizing factor
Aip1	.....	Actin-interacting protein-1
ATP	.....	Adenosine tri phosphate
BBB	.....	Blood–brain barrier
BMP	.....	Bone morphogenic protein
BSA	.....	Bovine Serum Albumin
Ca <sup>2+</sup>	.....	Calcium ion
CaMK	.....	Ca <sup>2+</sup> /calmodulin-dependent protein kinase
CAP1	.....	Cyclase-associated protein-1
cDNA	.....	Complementary DNA
CNS	.....	Central nervous system
CO	.....	Carbon Monoxide
DAMP	.....	Danger associated molecular patterns
DNA	.....	Deoxyribonucleic Acid
DTT	.....	Dithiothreitol
ELISA	.....	Enzyme-linked immunosorbent assay
ER	.....	Endoplasmic reticulum
ERK	.....	Extracellular-signal-regulated kinases
E-selectin	.....	Endothelial leukocyte adhesion molecules
FBS	.....	Fetal bovine serum
FLVCR1	.....	Feline leukemia virus subgroup C receptor
GAPDH	.....	Glyceraldehyde 3-phosphate dehydrogenase
GPx	.....	Glutathione peroxidase
GSH	.....	Glutathione
GSK3β	.....	Glycogen synthase kinase 3β
H	.....	Hour
H <sub>2</sub> O <sub>2</sub>	.....	Hydrogen peroxide
Hb	.....	Hemoglobin
HEPES	.....	4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid
HMGB1	.....	High mobility group box 1
HO	.....	Hemoxygenase
Hp	.....	Haptoglobin

HRP.....	Horseradish peroxidase
HS .....	Horse serum
HSP32 .....	Heat shock protein 32
Hx.....	Hemopexin
ICAM-1 .....	Intracellular adhesion molecules
ICH.....	Intracerebral hemorrhage
IL-10 .....	Interleukin-10
IL-1 $\beta$ .....	Interleukin 1- beta
IL-2 .....	Interleukin-2
INF $\gamma$ .....	Interfereon gamma
iNOS .....	Induced nitric oxide synthase
KCl.....	Potassium Chloride
LIMK .....	LIM kinase
LPS.....	Lipopolysaccharide
LRP1 .....	Low density lipoprotein receptor
MAPK.....	p38 mitogen-activated protein kinase
MAPKAPK-2/MK2 .....	MAPK)-activated protein kinase-2
MgCl <sub>2</sub> .....	Magnesium chloride
mM.....	Millimolar
MMP .....	Matrix metalloprotein mM
MPTP .....	Mitochondrial permeability transition pore
MTT .....	3-[4,5-dimethyl- thiazol-2-yl]-2,5-diphenyltetrazolium bromide
Na.....	Sodium
NaF.....	Sodium Flouride
nM.....	Nanomolar
NMDA .....	N-methyl-D-aspartate
NO.....	Nitrite oxide
NOS.....	Nitrite Oxide Synthase
NOX.....	NADPH oxidase
NP 40 .....	Nonidet P-40
Nrf2 .....	Nuclear factor (erythroid-derived 2)-like 2
PC-12 .....	Pheochromocytoma cell with embryonic origin from neural crest
PCFT .....	Proton-coupled folate transporter
PGE <sub>2</sub> .....	Prostaglandin E 2
PI3K .....	Phosphoinositide 3-kinase
PKD.....	Protein kinase D
PMN.....	Polymorphonuclear cells
PMSF .....	Phenylmethylsulfonyl fluoride
PP1/PP2A.....	Phosphoprotein phosphatases
PPAR $\gamma$ .....	Peroxisome proliferator activated receptor- $\gamma$
PtdIns (3,4) P <sub>2</sub> .....	Phosphatidylinositol (3,4)-bisphosphate
PVDF .....	Polyvinylidene fluoride
RNA .....	Ribonucleic Acid
RNS.....	Reactive nitrase species
ROCK .....	Rho-associated kinase
ROS.....	Reactive oxygen species

RT-qPCR.....	Real time quantitative polymerase chain reaction
SDS .....	Sodium dodecyl sulfate
SIM-A9 .....	Spontaneously Immortalized Microglia
siRNA .....	Small interfering RNA
SOD.....	Superoxide dismutase
SSH .....	Slingshot
TGF- $\beta$ .....	Transforming growth factor- $\beta$
TLR.....	Toll-like receptor
TNF- $\alpha$ .....	Tumor necrosis factor- $\alpha$
Trem2.....	Triggering receptor expressed on myeloid cells 2
UPR.....	Uncoupled Protein Response
VCAM-1 .....	Vascular cell adhesion molecule1
VEGF .....	Vascular endothelial growth factor
WB .....	Western blotting
Wfs-1.....	Wolframin ER transmembrane glycoprotein
XBP1 .....	X-box binding protein-1



# **Chapter 1**

## **Introduction**

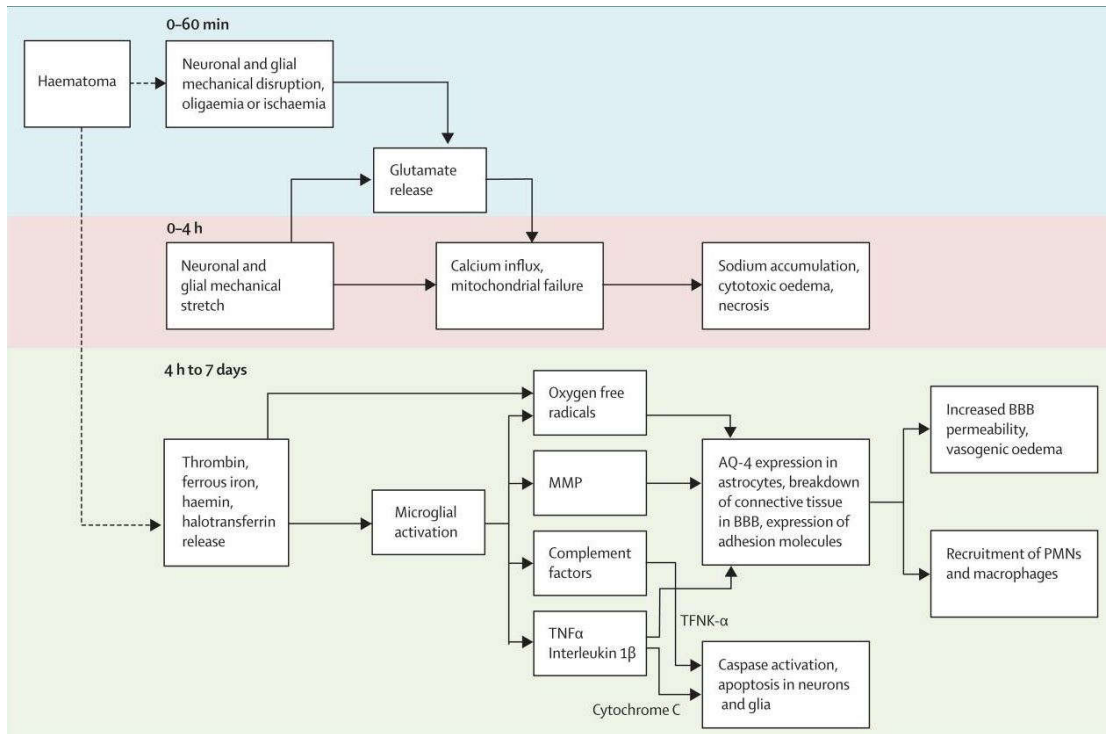
### **1.1 Intracerebral Hemorrhage**

Intracerebral hemorrhage (ICH), a subtype of stroke, is a devastating form of cerebrovascular disorder, with an annual incidence of 10–30 per 100,000, and about 2 million cases occurring worldwide. The very high morbidity and mortality rate reaches up to 40% within a month following ICH. Those who survive are left with severe neurological deficits (Mozaffarian, Benjamin et al. 2015). Six months after the onset of ICH, only 20% of ICH survivors are able to live independently (Kuramatsu, Huttner et al. 2013). The rate of ICH is expected to increase as a result of the aging population, but there are only few effective medical and surgical treatment strategies available currently. The mortality rates after one month of ICH have not decreased in the past three decades (van Asch, Luitse et al. 2010).

#### **1.1.1 Pathophysiology**

ICH causes immediate physical damage to the brain cellular architecture adjacent to the hemorrhage. After the first ictus of bleeding, about 20–40% of patients experience hematoma enlargement during the first day (Keep, Hua et al. 2012). The precise

mechanism for the hematoma growth is not known yet, and there are not much alternatives to stop this initial insult. However, the secondary brain injury after ICH is caused by a complicated cascade of events where therapeutic intervention could help to prevent the further damage (**Figure 1-1**). Immediately after ICH, the activation of hemostatic cascade occurs as an initial reaction to limit bleeding. Thrombin is produced in the brain immediately after hemorrhage affecting endothelial cells, astrocytes, neurons, and microglia (Hua, Keep et al. 2007, Keep, Hua et al. 2012), thereby contributing to early blood–brain barrier (BBB) disruption and edema formation (Keep, Zhou et al. 2014). The release of hemoglobin (Hb) and subsequent formation of hemin and iron from the hematoma is considered another prime contributor to the ICH-induced brain injury (Wagner, Sharp et al. 2003, Hua, Keep et al. 2007). Erythrocyte lysis occurs within 24 hour (h) after ICH, peaking at day 1–7, and hemoglobin- and iron-mediated brain injuries occur along with lysis of extravagated erythrocytes (Qureshi, Mendelow et al. 2009, Keep, Zhou et al. 2014).



**Figure 1-1: Cascade of neural injury initiated by intracerebral hemorrhage.** The steps in the first 4 h are related to the direct effect of the hematoma; later steps are due to the products released from the hematoma. BBB=blood-brain barrier. MMP=matrix metalloproteinase. TNF=tumor necrosis factor. PMN=polymorphonuclear cells (Adapted from (Qureshi, Mendelow et al. 2009) with permission from Elsevier) .

### 1.1.2 Oxidative stress and intracerebral hemorrhage

Oxidative stress is one of the major contributors of ICH, metabolites of Hb-heme-iron axis are key contributors to oxidative brain damage after ICH, although other factors, such as neuroinflammation and prooxidases play the primary role (Hu, Tao et al. 2016). The oxidative products of macromolecules significantly increase, whereas antioxidant enzymes, such as superoxide dismutase (SOD), glutathione peroxidase (GPx) and catalase,

correspondingly decrease as a result of ICH (Nakamura, Keep et al. 2005, Nakamura, Keep et al. 2006). Mitochondrial dysfunction has been reported to be responsible for the decreased oxygen metabolites after ICH (Kim-Han, Kopp et al. 2006). The mechanism of excessive ROS formation by mitochondria after ICH is not clear, but this has been attributed to the mitochondrial permeability transition pore (MPTP) since the inhibition of MPTP can attenuate ROS production (Ma, Chen et al. 2014).

Hb is released from erythrocyte protein into the extracellular space via complement-mediated cell lysis in the hours after ICH. This release is considered as the potent mediator of oxidative stress-induced injury (Regan and Panter 1993, Hua, Xi et al. 2000). Hb leads to increased reactive oxygen species (ROS), reactive nitrate species (RNS) (Yang, Chen et al. 2013). Heme from methemoglobin, and its oxidized product, hemin, trigger oxidative damage in brain tissue around the hematoma (Hu, Tao et al. 2016). Iron overload in the brain starts within 24 h and peaks at 7 days after hemorrhage, remains at least for a month, and consequently produces toxicity based on the Fenton reaction (Chaudhary, Gemmete et al. 2013). In experimental ICH models, it was confirmed that oxidative deoxyribonucleic acid (DNA) damage caused by free radicals occurs after ICH, and it leads to neuronal death (Nakamura, Keep et al. 2005). Neuroinflammation has been linked to increased oxidative stress after ICH (Matsuo, Kihara et al. 1995). Inhibition of NADPH oxidase (NOX) and nitric oxide synthase (NOS) has been found to improve ICH condition (Matveev 1966, Zia, Csiszar et al. 2009). The role of antioxidant system such as hemoxygenase (HO) is very complex and found to affect ICH in different ways based on context and site of action (Chen-Roetling, Lu et al. 2015). It has been found that in the free radical scavenging systems, SOD is severely impaired after ICH and therefore contribute to further increased

ROS, RNS induced brain damage (Wu, Hua et al. 2002). ROS can activate the Keap1/Nrf2/ARE pathway to counteract oxidative damage after ICH as an adaptive response mechanism but due to excessive oxidative stress the cells eventually fail to protect themselves (Shang, Yang et al. 2013).

### **1.1.3 Inflammation and intracerebral hemorrhage**

Inflammation is the key contributor of ICH-induced secondary brain injury. Inflammation progresses in response to various stimuli released after ICH (Zhou, Wang et al. 2014). Hematoma components initiate inflammatory signaling, with the production of thrombin playing a prominent role via activation of microglia and subsequently releasing proinflammatory cytokines and chemokines to attract peripheral inflammatory mediators (Keep, Hua et al. 2012). Hb, heme and iron released after red blood cell lysis aggravate ICH-induced inflammatory injury. Bilirubin, a Hb breakdown product, can also induce neutrophil entrance into brain but does not activate microglia (Loftspring, Johnson et al. 2011). Many preclinical studies have shown that targeting such signaling can affect ICH-induced brain injury. Inflammation is involved not only in early ICH-induced brain injury, but also in tissue repair (Egashira, Hua et al. 2015). Danger associated molecular patterns (DAMPs) such as the high mobility group box 1 protein (HMGB1) which is released from damaged or dead cells, trigger inflammation in the late stage of ICH (Lei, Lin et al. 2013). Preclinical studies have identified inflammatory signaling pathways that are involved in microglial activation, leukocyte infiltration, toll-like receptor (TLR) activation, and DAMP regulation in ICH. Recent advances in understanding the pathogenesis of ICH-induced inflammatory injury have facilitated the identification of

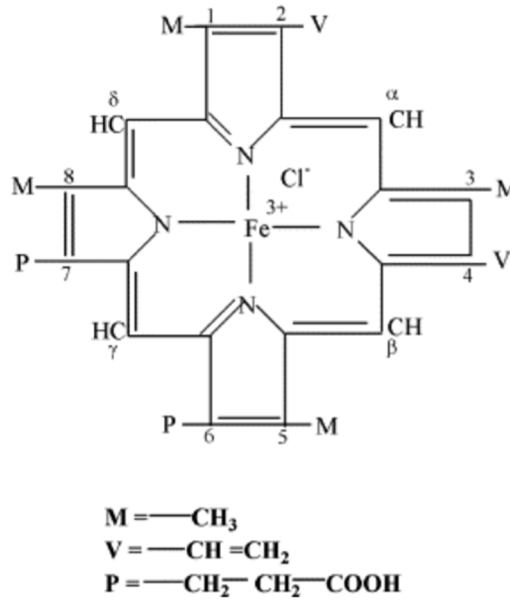
several novel therapeutic targets for the treatment of ICH such as thrombin inhibitors, desferrioxamine, minocycline, statins and therapeutic hypothermia; but none of these agents has proven to be effective in clinical settings (Belur, Chang et al. 2013).

## **1.2 Hemin**

Hemin is an erythrocyte derived endogenous chemical substance that is released due to breakdown of erythrocyte during different pathological conditions.

### **1.2.1 Chemistry of hemin**

Hemin is an iron–protoporphyrin complex consisting of four substituted pyrrole rings linked by –CH group (**Figure 1-2**). When the iron atom is in the ferrous state, the complex is called ferroprotoporphyrin or heme and the molecule is electrically neutral. When the iron atom is in the ferric state, the complex is called ferriprotoporphyrin or hemin, and the molecule carries a unit positive charge (and is consequently associated with an anion, chloride). Hemin is hydrophobic in nature and contains methyl groups in positions 1, 3, 5 and 8; vinyl groups in positions 2 and 4; propionic acid groups in positions 6 and 7 (Kumar and Bandyopadhyay 2005).



**Figure 1-2: Chemical structure of Hemin ( $Fe^{3+}$ -protoporphyrin).** Adapted from Kumar and Bandyopadhyay, 2005 with permission from Elsevier.

### 1.2.2 Catabolism and toxicity of hemin

Hemin is catabolized by the microsomal hemeoxygenase (HO) system consisting of HO and NADPH–cytochrome P450 reductase. This system catalyzes the oxidation of heme to biliverdin, carbon monoxide (CO) and ferrous iron, in equimolar amounts. This reaction requires the presence of oxygen and NADPH. Once biliverdin is formed, it is rapidly metabolized to bilirubin by biliverdin reductase (Liu and Ortiz de Montellano 2000, Wagener, Volk et al. 2003, Li and Stocker 2009). There are two isoforms of HO that are found in the CNS: HO-1 and HO-2. The former is also known as heat shock protein 32 (HSP32) and is highly expressed by microglia and astrocytes but is weakly expressed by neurons (Geddes, Pettigrew et al. 1996, Turner, Bergeron et al. 1998, Scapagnini, D'Agata et al. 2002). HO-1 expression can be induced by a wide range of conditions including

hemorrhage (Syapin 2008). On the other hand, HO-2 is primarily expressed by neurons and is not inducible, even by injections of whole or lysed blood into the subarachnoid space (Matz, Turner et al. 1996). It is generally considered that HO activity improves cellular viability only in instances where the amounts of heme to be metabolized are within the capacity of the cells to sequester the iron that is released. Conversely, HO activity becomes detrimental whenever excessive amounts of free iron are produced (Robinson, Dang et al. 2009) and no longer be stored in ferritin; this unbound iron is redox-active and produces toxic hydroxyl radicals.

Heme can also be cleared by forming a non-toxic heme-complex (e.g., hemopexin (Hx), albumin, heme-binding protein, etc.) or by its degradation (e.g., glutathione (GSH), xanthine oxidase, NADPH-cytochrome P-450 reductase, etc.) or by scavenging free redox-active iron (ferritin) released after heme catabolism (Kumar and Bandyopadhyay 2005). Hx-mediated transport of heme is an active process, requiring cellular energy,  $Ca^{2+}$  and a minimum temperature for heme uptake and also depends on the ability of the cells to respond adequately (Smith and Morgan 1981). Haptoglobin (Hp), albumin, lipocalin alpha 1-microglobulin, heme binding proteins have been suggested but detailed mechanism is yet to be discovered (Van Vlierberghe, Langlois et al. 2004, Kumar and Bandyopadhyay 2005).

Heme in the aqueous phase frequently aggregates in the membrane and promotes oxidation, which leads to the enhancement of permeability and membrane disorder. Oxidation of membrane components may promote cell lysis and death (Vincent 1989, Schmitt, Frezzatti et al. 1993). Free heme is a pro-inflammatory molecule, activates neutrophil responses. Heme increases the expression of interleukin-2 (IL-2) receptors in



human peripheral blood mononuclear cells and causes induction of mitogenicity, cytotoxicity and cytokine production (Tsuji, Wang et al. 1993). Hemin induces neutrophil chemotaxis, cytoskeleton reorganization as well as increased expression of chemokine interleukin-8 *in vitro* (Graca-Souza, Arruda et al. 2002). Hemin causes substantial hemolysis by affecting red cell membrane, stimulating potassium loss and swelling (Chou and Fitch 1981). This neurotoxic effect of hemin is iron-dependent, oxidative, and predominantly necrotic (Goldstein, Teng et al. 2003). In endothelial cells, hemin stimulates the expression of intracellular adhesion molecules (ICAM-1), vascular cell adhesion molecule1 (VCAM-1) and endothelial leukocyte adhesion molecules (E-selectin), probably through hemin-mediated generation of intracellular ROS and its subsequent signaling pathways (Shono, Ono et al. 1996).

Hemin catalyzes the degradation of proteins to small peptide fragments (Aft and Mueller 1984), oxidizes low- and high-density lipoproteins and forms conjugated dienes, thiobarbituric acid reacting substances (Camejo, Halberg et al. 1998), causes nicking and subsequent degradation of DNA (Kumar and Bandyopadhyay 2005). Even though hemin has been suggested to cause mitochondrial DNA damage (Suliman, Carraway et al. 2002), degradation product of hemin has been proposed to protect mitochondria (Mullebner, Moldzio et al. 2015). CO formed via degradation of hemin has been suggested to offer neuroprotection from hippocampal cell damage (Han, Yi et al. 2015) but bilirubin has been shown to induce endoplasmic reticulum (ER) stress in hepatocytes (Mullebner, Moldzio et al. 2015). Iron released from hemin breakdown is thought to damages lipids, proteins, and DNA through the generation of ROS (Rodriguez, Kemp et al. 2003).

### 1.2.3 Hemin and intracerebral hemorrhage

There is approximately 2.5 mM hemoglobin in blood (Robinson, Dang et al. 2009) and during ICH deposits of approximately 10 mM suspension of hemin is found in brain parenchyma (Letarte, Lieberman et al. 1993). To put this value into perspective, cell culture experiments have shown that 3–30  $\mu$ M hemin is sufficient to kill 60–70% of cultured neurons and astrocytes within 4–14 h (Regan, Wang et al. 2001, Regan, Chen et al. 2004). Although initially sequestered within erythrocytes, some is released and accounts for the high micromolar concentration of hemin that is present in one week old intracranial hematomas (Robinson, Dang et al. 2009). The exact mechanisms responsible for this hemin accumulation to cytotoxic levels have not been defined clearly. Obviously brain tissue is a poor microenvironment for erythrocytes survival. Whether their lysis are due to complement activation (Hua, Xi et al. 2000), heme-mediated oxidative stress (Baysal, Monteiro et al. 1990) or other mechanisms is still poorly defined (Wagner and Dwyer 2004, Jaremko, Chen-Roetling et al. 2010).

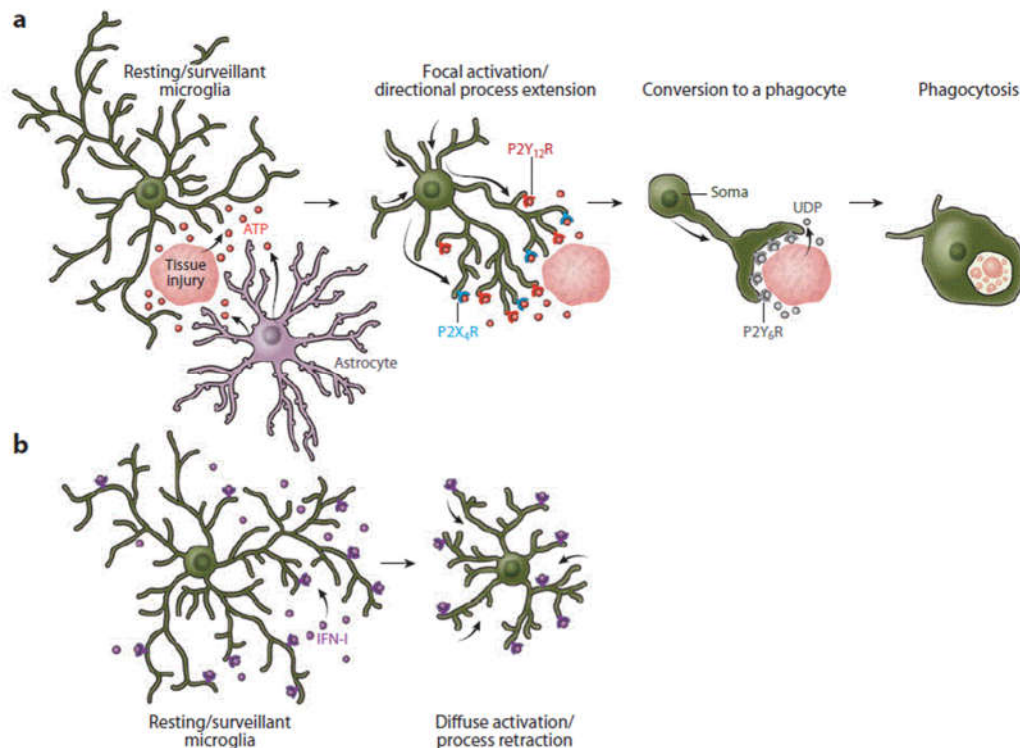
At the physiological temperatures, extracellular Hb is oxidized spontaneously to ferric methemoglobin, releasing a superoxide ion in the reaction (Chen-Roetling, Lu et al. 2015) and then it transfers them to higher-affinity lipid and protein binding sites (Bunn and Jandl 1968). Due to brain microenvironment's low concentration of the Hx, there is likelihood of increased hemin uptake by vulnerable cell populations such as neurons and astrocytes. Hb in the hematoma is scavenged by microglia and infiltrating macrophages, which phagocytose intact erythrocytes as well as take up Hb and Hb-haptoglobin (Hp) complexes (Kristiansen, Graversen et al. 2001, Schaer, Schaer et al. 2006). After being taken up by these phagocytic cells, the fate of hemoglobin-bound heme remains unclear. It

is likely that some percentage is ejected into the extracellular space by different exporters such as ATP-binding cassette sub-family G member 2 (ABCG2) and Feline leukemia virus subgroup C receptor (FLVCR1) where it is expected to take part in free radical reactions (Yang, Philips et al. 2010, Chen-Roetling, Cai et al. 2014). The accumulation of hemin to toxic concentrations within the hematoma and its remediation provides an obvious therapeutic target for ICH.

### **1.3 Microglia**

Microglia are the central nervous system (CNS) resident innate immune cells constituting approximately 5–10% of the brain cell population depending on the brain region. They are the first responders in a defense network that covers the entire CNS parenchyma. They respond to immunogenic stimulus by migrating towards injured regions and releasing a cascade of cytokines and recent studies have shown their involvement in the synapse formation (Tremblay, Stevens et al. 2011, Sierra, Tremblay et al. 2014). Mice microglia originate from a unique primitive myeloid progenitors in the yolk sac (Ginhoux, Greter et al. 2010). These progenitors migrate and colonize the brain during fetal development before brain vasculature arborization is complete and before the BBB is fully formed. In the adult brain, microglia are confined by the fully developed BBB and become an autonomous, long-lived cell population retaining the ability to divide and self-renew throughout life (Ajami, Bennett et al. 2007). Under resting physiological conditions, microglia have a small soma and numerous long and thin processes extending several times the length of the soma into the microenvironment. This unique cytostructure allows them to cover the entire parenchyma in a non-overlapping, mosaic fashion in the normal adult

brain. *In vivo* imaging has shown that microglial processes are highly motile even under ‘resting’ physiological conditions (Nimmerjahn, Kirchhoff et al. 2005). Microglia exhibit at least four functional behaviors: surveillance, neuroprotection, phagocytosis, and toxicity (Chen and Trapp 2016). Acute insults and chronic disease states induce hypertrophy of their cell bodies, asymmetrical distribution of their processes and increased expression of activation molecules. Microglial morphology has intricately been related to their differential functions against different types of insults (**Figure 1-3**) but quantitative relation has not been fully elucidated yet (Nayak, Roth et al. 2014).



**Figure 1-3: Microglia activation following CNS injury.** Illustration of (a) focal and (b) diffuse microglia activation resulting from sterile brain injury and viral infection, respectively. (Adapted from Nayak et al., 2014 with permission from Annual Reviews).

### **1.3.1 Microglia and intracerebral hemorrhage**

Microglial activation is a very early event after ICH (Zhao, Grotta et al. 2009, Taylor and Sansing 2013) and ‘activated microglia’ produce many pro- and anti-inflammatory mediators. The response of microglia is highly dependent on the age of the subject with more prominence in aged than in young (Gong, Hua et al. 2004). Pro-inflammatory M1 phenotype of microglia produce interleukin-1 beta (IL-1 $\beta$ ), TNF- $\alpha$ , and ROS (Cherry, Olschowka et al. 2014). Such factors are known to activate other microglia and astrocytes, induce BBB disruption and cause the expression of adhesion molecules on the luminal surface of brain endothelial cells (Obermeier, Daneman et al. 2013), an important step for leukocyte migration into the brain. The M1 phenotype damage neurons by releasing oxidative metabolites and proteases (Kigerl, Gensel et al. 2009), induce a rapid switch towards M1 polarization (Kroner, Greenhalgh et al. 2014) and thus further aggravating the pathological conditions. In contrast, microglial cells with a M2 phenotype are primarily anti-inflammatory and secrete interleukin-10 (IL-10), transforming growth factor- $\beta$  (TGF- $\beta$ ), and arginase-1 which reduces NO production, phagocytic activity and is also involved in tissue repair (Zhao, Grotta et al. 2009, Cherry, Olschowka et al. 2014). M2 phase has also been suggested to produce pro-angiogenic factors, such as vascular endothelial growth factor (VEGF), and IL-8, and are also thought as a potential contributor for angiogenesis after ICH (Welser, Li et al. 2010). Therefore, there has been considerable interest in targeting microglia activation as a very

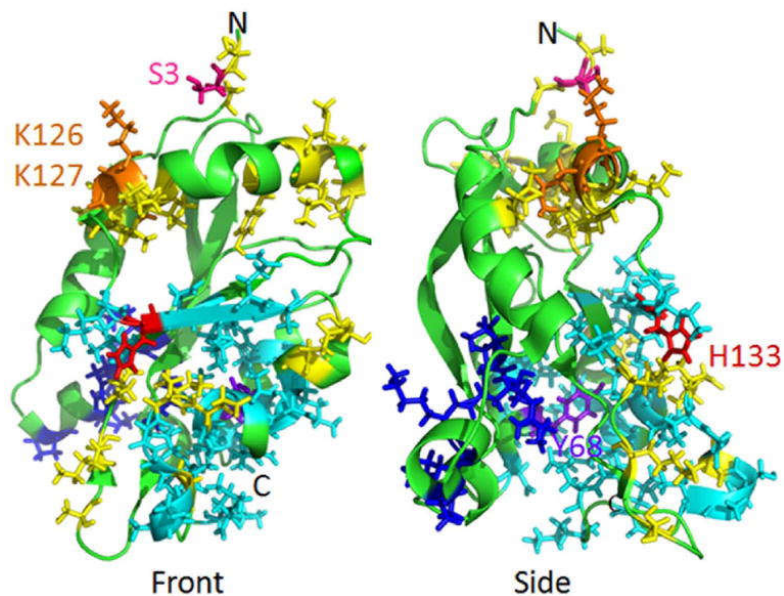
early event in the inflammatory cascade or altering microglia phenotypes to M2 like phenotype to reduce ICH-injury.

After ICH, microglia strongly express HO-1, the inducible isoform of HO (Wang and Dore 2007), that converts heme to iron, CO and biliverdin (Maines 1988). In order to prevent excessive ROS production, iron is temporarily stored within the phagocytes by iron-binding proteins, such as ferritin (Aronowski and Zhao 2011). This clean-up process persists 3–4 weeks after ICH onset (Zhao, Sun et al. 2007). Limiting extracellular hemoglobin degradation by early inhibition of HO-1 effectively ameliorated hemoglobin-induced brain edema in rats (Huang, Xi et al. 2002). However, in the later phase, HO-1 inhibition may interfere with phagocyte activity and the clean-up process after ICH (Aronowski and Zhao 2011). ICH induced extravasated erythrocytes are also cleared from brain parenchyma by microglia and thus microglia prevent further aggravation of secondary injury following ICH (Xi, Keep et al. 2006). Recent studies have shown that modulating phagocytic signal on erythrocytes could be a potential therapeutic target (Burger, de Korte et al. 2012, Zhou, Xie et al. 2014). It is generally accepted that microglia can have beneficial as well as detrimental effects during and after ICH and the effects of microglia inhibition could be planned time-dependently (Taylor and Sansing 2013). An alternate therapeutic approach might be to modify microglial activation to promote potential beneficial effects such as phagocytosis and tissue repair. In order to promote M2 phagocytic phenotype, peroxisome proliferator activated receptor- $\gamma$  (PPAR $\gamma$ ) agonists have shown benefit in animal ICH models and a PPAR $\gamma$  agonist is now in clinical trial for ICH (Gonzales, Shah et al. 2013). Further investigation into the

interaction of microglia and the mediators of ICH induced pathological cascade might lead to a better therapeutic treatment.

#### 1.4 Cofilin

Cofilin was purified and identified as a protein with molecular weight of 20-kDa that induces depolymerization of actin filaments in an extract from avian or porcine brain (Nishida, Maekawa et al. 1984, Bamburg, McGough et al. 1999). The cofilin family in mammals is composed of non-muscle type, cofilin-1 (also known as n-cofilin), muscle type, cofilin-2 (also known as m-cofilin) and actin depolymerizing factor (ADF) (also known as destrin) (Ono 2007, Poukkula, Kremneva et al. 2011).



**Figure 1-4: Human Cofilin Structure.** The residues shown in yellow are the ones those are in the actin-contact regions (right side of the molecule in the side view), those in cyan are in the Phosphatidylinositol (3,4)-bisphosphate (PtdIns (3,4) P2) binding region, and

those in dark blue are in the nuclear localization sequence. Also shown here is the Ser3 regulatory phosphorylation site, the two Lys residues (K126 and K127) to which Ser3 binds when it is phosphorylated, His133 that confers pH-sensitivity to the PtdIns (3,4)P<sub>2</sub>-binding, and Tyr68 that serves as a Src substrate for phosphorylation prior to ubiquitination and degradation of cofilin (Adapted from (Bernstein and Bamburg 2010) with permission from Elsevier).

The following section will use cofilin to mean cofilin-1. Aside from this canonical cofilin role in actin cytoskeleton modification and thus in cell motility, migration, and cytokinesis, cofilin has also been linked to modulate the intracellular Ca<sup>2+</sup> signaling, inflammation and different neurodegenerative diseases (Nusco, Chun et al. 2006, Schonhofen, de Medeiros et al. 2014, Ohashi 2015)

#### **1.4.1 Cofilin and actin dynamics**

Cofilin is distributed throughout the cytoplasm of the cell. It translocates to the cortical regions of the active cells in which the actin cytoskeleton is highly dynamic. It binds to ADP-bound F-actin and destabilizes and severs the actin filaments, resulting in depolymerization of the short actin filaments and such modulation depends on the ratio of cofilin to actin. At low cofilin/actin ratios, cofilin acts to sever actin filaments, whereas at high cofilin to actin ratio, cofilin nucleates actin assembly and stabilizes F-actin (Pollard and Borisy 2003). It enhances both actin filament disassembly and actin polymerization by severing and by supplying actin monomers, which results in higher dynamics of the actin cytoskeleton. Another suggested proposition of cofilin mediated actin severing is through



generation of short filaments and new barbed ends for actin assembly (Oser and Condeelis 2009). It promotes the generation of actin monomers, nuclei, and newly barbed ends for the assembly of actin structures such as lamellipodia, filopodia, and stress fibers, as well as the turnover of these structures during the cellular morphology changes. On the other hand, the inactivation of cofilin in response to extracellular signals decelerates the actin turnover and contributes to the assembly of actin structures (Ohashi, Fujiwara et al. 2011, Ohashi 2015).

#### **1.4.2 Regulation of cofilin activity**

Different regulatory kinases and phosphatases modulate cofilin activity through phosphorylation/dephosphorylation processes. Cofilin is inactivated by phosphorylation at Ser 3 by LIM kinase and testicular (TES) kinase and reactivated through dephosphorylation by the phosphatases like slingshot (SSH), chronophin, and phosphoprotein phosphatases (PP1/PP2A). Oxidative stress, proteins that modulate cofilin-actin interactions, phosphatidylinositol (4, 5) biphosphate, and pH are also found to regulate cofilin activities. Kinases and phosphatases are activated in response to different intracellular and extracellular signals like ROS, elevation in intracellular  $Ca^{2+}$ , and drop in intracellular ATP, resulting in cofilin induced actin dynamic modulation (dos Remedios, Chhabra et al. 2003, Bamburg and Bernstein 2010, Bernstein and Bamburg 2010).

#### **1.4.3 Cofilin and intracerebral hemorrhage**

There is not any direct experimental evidence showing the relation between cofilin and ICH but increasing evidence have suggested that the regulators of cofilin are affected

by ICH pathophysiology (Alhadidi, Bin Sayeed et al. 2016). Studies have shown that both the excess cofilin activity as well as defect in cofilin expression inhibits its proper functions in the cells. The dynamic imbalance of cofilin activity is one of the major pathophysiological features of ICH (Rust 2015). ICH leads to phosphatase activation in response to ROS mediated elevation in intracellular  $Ca^{2+}$  and drop in intracellular ATP, resulting in cofilin over-activation and actin dynamic disruption (Alhadidi, Bin Sayeed et al. 2016). ICH induced oxidative stress has been proposed to be the mechanism of both cofilin over-activation and excitotoxicity. Increased ROS affect cysteine residue in cofilin molecules resulting in intramolecular disulfide bond formation. Such structural modification disrupts cofilin-actin interaction and frees cofilin to translocate into the mitochondria, where it opens the permeability transition pore and initiates cytochrome C release, eventually leading to apoptosis. Inhibition of both cofilin expression with cofilin siRNA and cofilin oxidation prevented oxidant-induced apoptosis, suggesting the central role of oxidative stress in triggering cofilin-induced apoptosis (Zhang, Wang et al. 2004, Klamt, Zdanov et al. 2009). Studies have shown that oxidation of cofilin leads to oxidant-induced apoptosis (Klamt, Zdanov et al. 2009) as well as caspase-independent necrotic-like programmed cell death (Wabnitz, Goursot et al. 2010).

ICH induced ionic imbalance that in turn inhibits the uptake of glutamate by glutamate transporters and even causes the reversal of the transporters, induces over-accumulation of extracellular glutamate (Rossi, Oshima et al. 2000). Studies have shown that cofilin activation mediate Bax translocation to the mitochondria during excitotoxic neuronal death (Posadas, Perez-Martinez et al. 2012) and both N-methyl-D-aspartate (NMDA) and non-NMDA receptors mediate glutamate stimulation-induced cofilin

dephosphorylation and cofilin-actin rod formation (Chen, Jiang et al. 2012). Following ICH, glial cells release a wide array of pro-inflammatory cytokines, chemokines, and free radicals. Therefore, inhibiting the inflammatory response can reduce injury and improve neurological outcomes. But inhibiting cofilin might not reduce inflammation fully resulting in low grade inflammatory response during the chronic phase which in turn can lead to repair, regeneration, and functional recovery (Zhou, Wang et al. 2014). Cofilin has also been suggested to play a major role in ICH glial cell migration and proliferation (Li, Briehner et al. 2007, Jonsson, Gurniak et al. 2012). Cofilin knockdown in microglia inhibited NADPH oxidase activity and ROS formation (Rasmussen, Pedersen et al. 2010). Therefore, inhibition of the cofilin might prevent progression of over-activation induced inflammation and might also reduce the levels of the injury.

### **1.5 Aim and hypothesis**

The primary goal of this investigation is to study the role of cofilin in different microglial functions as well as underlying molecular mechanism induced by high doses of hemin exposure to mimic the condition of ICH.

**Aim 1:** Evaluate the role of cofilin in hemin induced microglia inflammation and survival. This can be done by exposing microglia to high doses of hemin followed by studying the levels of NO, nitric oxide synthase (iNOS), cytokines, and cell viability. To study the role of cofilin, microglia were transfected with cofilin siRNA followed by similar experimental studies. Microglia are the main source of NO in ICH (Zhou, Wang et al.

2014); increased iNOS and higher cytokines levels are the markers of pathogenesis of ICH (Kim, Im et al. 2009, Lin, Yin et al. 2012, Ndisang and Jadhav 2013) and extremely high doses of hemin presumably affect the viability of all cell types of brain parenchyma in ICH (Egashira, Hua et al. 2015). Therefore, these investigations are highly relevant to the context of ICH.

**Aim 2:** Investigate the role of cofilin in hemin induced microglial migration. This can be accomplished by exposing microglia to hemin followed by streaking a scratch in the culture media and then counting the number of cells migrated to the area of scratch. Similar experimental paradigm would be followed after cofilin siRNA transfection in microglia for studying the role of cofilin. Migration of microglia towards the site of injury is one of the primary functions of microglia (Kettenmann, Hanisch et al. 2011) and it has been suggested to be mediated by cofilin (Bernstein and Bamburg 2010). Hemin at a very high concentration as in the case of ICH is hypothesized to affect migration of microglia.

**Aim 3:** Illustrate the role of cofilin in microglial antioxidant defense system. By incubating microglia with hemin followed by the protein extraction and expression by Western blotting of hemin degrading enzyme, HO-1 (Chen-Roetling, Lu et al. 2015) and one of the upstream regulators of HO-1, nuclear factor (erythroid-derived 2)-like 2 (Nrf2) (Gonzalez-Reyes, Guzman-Beltran et al. 2013) a broad overview of antioxidant defense system of microglia due to hemin exposure could be studied. Microglia transfection by cofilin siRNA followed by the same experimental paradigm would show the role of cofilin in microglial antioxidant defense system relevant to hemin exposure. Studies have shown the cytoprotective role of HO-1 and Nrf2 in different cellular insults (Gozzelino, Jeney et al. 2010, Paine, Eiz-Vesper et al. 2010, Li, Wang et al. 2014) and therefore, the

investigation on the role of cofilin in hemin induced microglial cytoprotective antioxidant system is highly relevant to those in case of ICH.

**Aim 4:** Elucidate the role of cofilin in hemin induced ER stress and  $Ca^{2+}$  signaling in microglia. Incubation of microglia with hemin followed by ribonucleic acid (RNA) extraction, complementary DNA (cDNA) synthesis and then quantifying ER stress makers such as Wolframin ER transmembrane glycoprotein (Wfs-1), X-box binding protein-1 (XBP-1), sliced XBP-1 with reverse transcriptase quantitative polymerase chain reaction (RT-qPCR) would provide the extent of ER stress due to hemin exposure. Cofilin siRNA transfection of microglia followed by similar experimental paradigm would give the role of cofilin in hemin induced ER stress markers. Studies have shown that hemin affect cytostructure of the cell and increase ER stress in endothelial cell (Mullebner, Moldzio et al. 2015). Therefore, investigation on the role of cofilin in hemin induced ER stress in microglia is highly relevant to ICH. ER is one the primary sources of intracellular  $Ca^{2+}$  that plays a critical role in the process of microglial transformation and migration (Sharma and Ping 2014, Tsai, Seki et al. 2014).  $Ca^{2+}$  signaling measurement after acute exposure of hemin as well as after long term exposure of hemin would provide answer to the questions whether hemin evokes  $Ca^{2+}$  signaling and whether hemin exposure affect the efficacy of microglial ability to respond to acetylcholine evoked  $Ca^{2+}$  signaling respectively. Experimentation with cofilin siRNA transfected microglia would offer the role of cofilin in evoking  $Ca^{2+}$  signaling.

## Chapter 2

### Materials and Methods

#### 2.1 Cell culture

Spontaneously Immortalized Microglial Cells (SIM-A9 Cells) were generous gift from Dr. Kumi Nakamoto-Combs at the University of North Dakota and were grown in Dulbecco's Modified Eagle Medium (DMEM): Nutrient Mixture F-12 (HyClone, Thermo scientific, West Palm Beach, FL, USA) supplemented with 5% horse serum (HS), 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin.

Pheochromocytoma cell with embryonic origin from the neural crest (PC-12) cells (ATCC, Manassas, VA, USA) were grown in DMEM supplemented with 10% horse serum (HS), 5% fetal bovine serum (FBS) and 1% penicillin-streptomycin. To study the impact of cofilin knockdown in hemin activated microglia on the survival of PC-12 cells, a conditioned medium from microglial cells was added to the PC-12 cells and cells were incubated for 24 h.

#### 2.2 Microglial transfection by siRNA

SIGENOME SMARTpool mouse cofilin1 siRNA with the following sequences AGACAAGGACUGCCGCUAU, GGUGGCAGCGCCGUCAUUU, GUUCGCAAGUCUUCAACAC, and GAGAAUGCACCCCUCAAGA was ordered from

GE Healthcare Dharmacon, Inc. In addition, scrambled siRNA was used as a control. SIM-A9 cells were plated at the density of  $1.25 \times 10^5$  cells/ml. The cells were transfected with scrambled/cofilin1 siRNA at 50 nanomolar (nM) concentrations according to previously optimized conditions in our lab. Xfect transfection reagent (Clontech Laboratories, Inc) was used to perform the transfection experiment according to the manufacturer's instructions. Briefly, in two separate sets of microtubes, siRNA and Xfect transfection reagent were diluted using Xfect buffer. Then, the transfection reagent was added to the siRNA and was kept at room temperature for 20 min to ensure complex formation. After that, transfection mix was added to the cell culture and then incubated back for 48 h before starting hemin treatment.

### **2.3 Hemin induced microglial activation and nitrite oxide production**

Hemin at 75, 100, 200, 500 micromolar ( $\mu\text{M}$ ) concentration were used initially to activate microglia and then 100  $\mu\text{M}$  were selected for subsequent experiments due to consistent result in NO production. SIM-A9 cells were plated in 24-well plate and then transfected with scram/cofilin siRNA for 48 h prior to hemin activation. Cell culture medium from the respective wells was mixed with equal volume of Griess reagent (Sigma-Aldrich) in 96-well plate at room temperature. The amount of NO released into cell culture medium was quantified calorimetrically at 540 nm according to the manufacturer instructions.

### **2.4 ELISA assay**

SIM-A9 cells were plated in 24-well plate, transfected with scrambled/cofilin siRNA and then 100  $\mu\text{M}$  hemin was added to the cells. After that, the levels of TNF- $\alpha$  and IL-1 $\beta$  released into cell culture medium were quantified using commercially available

enzyme-linked immunosorbent assay (ELISA) kits (eBioscience, San Diego, CA, USA) according to the manufacturer protocol.

### **2.5 MTT-cell survival assay**

Cell viability assays were determined using the 3-[4,5-dimethyl- thiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) assay. In this assay, the viability of siRNA transfected SIM-A9 cells treated with hemin as well as PC-12 cells, treated with microglia conditioned medium, were measured. MTT assay protocol involves incubation of the cells (SIM-A9 or PC-12 cells) with MTT reagent (Promega Corporation, Madison, WI, USA) for 3 h in 5% CO<sub>2</sub> at 37°C. After that, the whole medium was discarded and DMSO was added to dissolve formazan crystals. Viable cells were quantified by measuring the absorbance at 570 nm.

### **2.6 Scratch migration assay**

SIM-A9 cells were cultured in 6-cm plate and then transfected with scrambled/cofilin siRNA for 48 h. Hemin at a dose of 100 µM were added to the cells for 20 h. By using a sterile 200 µl pipette tip, a scratch was made in the middle of each well followed by washing with PBS to remove damaged cell remnants. Then similar media as used earlier were added to the cells. At different time intervals (3 h, 12 h, 24 h of scratch and 23 h, 32 h and 44 h of hemin treatment), cells migrated to the scratched area were imaged and then the number of cells were counted by using ImageJ software (NIH, USA).

### **2.7 Subcellular fractionation and western blotting**

To study protein expression levels by Western blotting (WB), SIM-A9 cells were plated in 6-cm plate and then stimulated with 100 µM hemin for 24 h before cell lysis. To



obtain subcellular fractions, SIM-A9 cells were harvested using ice- cold lysis buffer [250 mM sucrose, 1.5 magnesium chloride (MgCl<sub>2</sub>), 10 mM potassium chloride (KCl), 20 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES) (pH 7.5), 1 mM dithiothreitol (DTT), 1 mM phenylmethylsulfonyl fluoride (PMSF), 50 mM sodium fluoride (NaF), 10 mM Na vanadate, 20 mM Na pyrophosphate and protease inhibitor cocktail (Thermo Scientific)]. Cell homogenates were kept on ice for 15 min and then Nonidet P-40 (NP-40) was added at 0.5% for 2 min extra. The resultant supernatant was used for studying protein expression. Bradford reagent (Bio-Rad Laboratories, CA, USA) were used for measuring protein concentrations, and samples were analyzed by loading equivalent amounts of protein (20-30 µg) onto 10–15% SDS-polyacrylamide gels. Proteins separated on the gels were transferred onto a pretreated polyvinylidene fluoride (PVDF) membrane and were blocked with 3% bovine serum albumin (BSA) for 1 h to prevent nonspecific binding. The membrane was then incubated overnight at 4°C with the following primary antibodies: rabbit anti-cofilin, rabbit anti- $\alpha$ -tubulin, rabbit anti-GAPDH, rabbit anti-HO-1, rabbit anti- Nrf2, and rabbit anti- iNOS (1:2000; Cell Signaling Technology, Danvers, MA, USA). Following the incubation, the blots were washed and incubated with horseradish peroxidase (HRP) conjugated goat anti-rabbit secondary antibody (1:2000; Jackson ImmunoResearch) for 1 h at room temperature. Anti-rabbit  $\alpha$ -Tubulin, rabbit anti-GAPDH were used as a loading control for cytosolic proteins. The images were analyzed using Bio-Rad ChemiDoc™ XRS Image Lab Software.

## **2.8 RT-qPCR**

SIM-A9 cells were transfected with scramble/cofilin-1 siRNA for 48 h followed by hemin treatment for 24 h. Total RNA was extracted from SIM-A9 cells with the treatments

as mentioned earlier by using TRIzol method (Life technologies) according to the manufacturer's protocol. Then 1 µg total RNA was transcribed into cDNA templates in a 20 µl a reaction mix of 5X iScript reaction mix, iScript reverse transcriptase, and nuclease-free water (Bio Rad, Hercules, CA). PCR was performed in a 10 µl reaction, containing 5 µl cDNA (1/5 diluted), 1X SYBR Green PCR Master Mix (Applied Biosystems) and 100 nM of each primer. Ct values (cycle threshold) were used to calculate the amount of amplified PCR product relative to 18S (the endogenous control). The relative amount of mRNA was calculated using the  $\Delta\Delta$ CT method. Minus-reverse transcriptase samples were used as negative controls to test for DNA contamination. The experiments were repeated thrice with duplication. For ER stress, Wfs-1, XBP1 and spliced XBP1 were used whose primers are presented in **Table 2-1**.

**Table 2-1: Primers sequences**

<b>Genes</b>	<b>Primer Forward Sequence (5'-3')</b>	<b>Primer Reverse Sequence (5'-3')</b>
<b>Wfs-1</b>	CGCCTCGTCAGCAGTGAAT	GGAACAGGTTGGTGGGAATG
<b>XBP-1</b>	TGGCCGGGTCTGCTGAGTCCG	GTCCATGGGAAGATGTTCTGG
<b>Spliced XBP-1</b>	CTGAGTCCGAATCAGGTGCAG	GTCCATGGGAAGATGTTCTGG
<b>18S</b>	TTCGAACGTCTGCCCTATCAA	ATGGTAGGCACGGCGACTA

## **2.9 Calcium signaling**

SIM-A9 cells were transfected with scramble/cofilin siRNA for 48 h followed by hemin treatment for 24 h. Then the cells were incubated with fura-2 AM for 1 h at 37°C. Fluorescence images were obtained using a polychrome IV monochromator-based high-speed digital imaging system (TILL Photonics, Gräfelfing, Germany) ported to a fiber optic guide and epifluorescence condenser and coupled to a Nikon TE2000 microscope equipped with DIC optics. Cytosolic Ca<sup>2+</sup> concentration in fura-2 AM-loaded SIM-A9 cells was

measured by alternatively illuminating the SIM-A9 cells with dual wavelength light (340 and 380 nm) focused onto the image plane via a DM400 dichroic mirror and Nikon SuperFluor  $\times 40$  oil-immersion objective, and fluorescence obtained through a  $525 \pm 25$  nm band-pass filter (Chroma Technologies, Brattleboro, VT, USA). Alternating transmitted light and fluorescent images were acquired using a high-speed Uniblitz VS35 optical shutter (Vincent Associates, Rochester, NY, USA) placed in the tungsten lamp illumination path in experiments. Pairs of transmitted light and fluorescence images (30 and 50 ms exposure, respectively) were collected at 2 Hz. Images were generated by subtracting each transmitted light frame from its preceding frame as described previously (Chen, Warner et al. 2005).

### **2.10 Statistical Analysis**

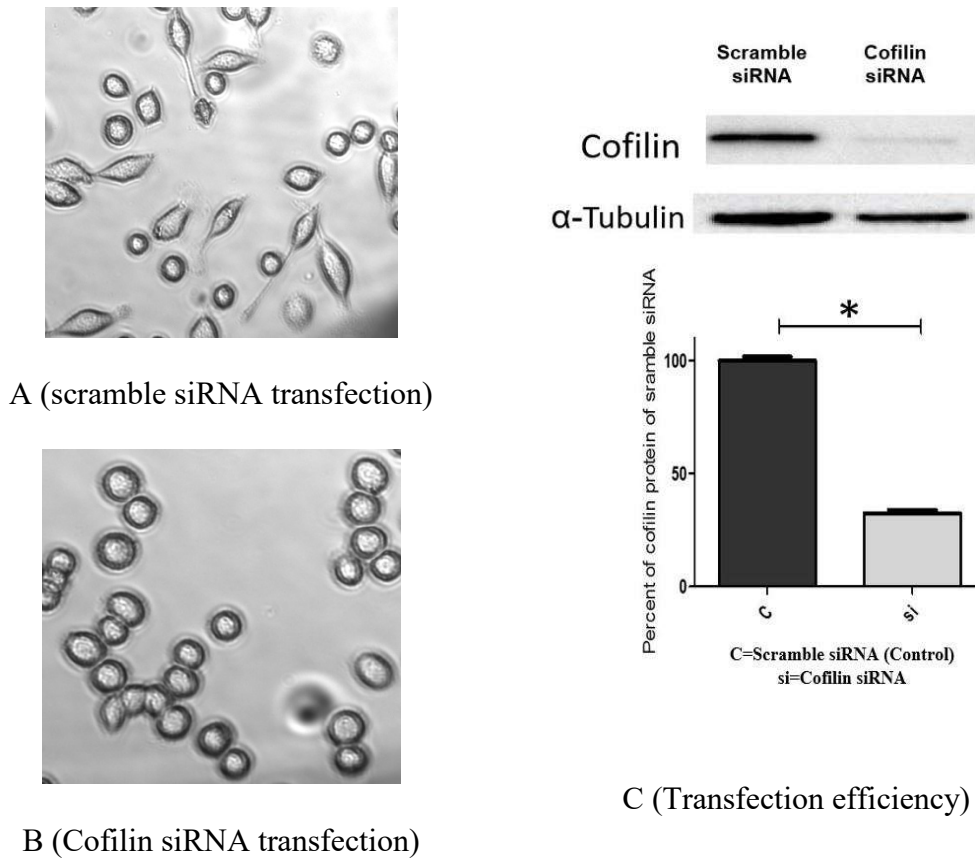
The experimental results were expressed as the mean  $\pm$  standard error of mean (SEM) and are accompanied by the number of observations (independent preparations of cultured cells). Data were analyzed by ANOVA followed by Bonferroni post Hoc using Graph Pad Prism (GraphPad Software, San Diego, CA) and IBM SPSS Statistics 22. Furthermore, student's unpaired t-test was used to determine significant differences between two groups. A value of  $p < 0.05$  is considered to be statistically significant.

## Chapter 3

### Results

#### 3.1 Cofilin knockdown changed microglial cells morphology

SIM-A9 cells assume ramified morphology in cell culture DMEM/F12 media and most of the cells transfected with scramble siRNA assumed spindle shape. Some assume globular shape indicating microglial functional state to stress conditions (**Figure 3-1A**). On the other hand, SIM-A9 cells transfected with cofilin siRNA assumed globular morphology (**Figure 3-1B**). There was no signs of pointed region in any corner of the cell whereas in scramble siRNA transfected SIM-A9 cells, pointed region in one or more than one regions was observed. Based on our optimized transfection experiment there was more than 80% reduction of cofilin protein expression as observed by the WB protein analysis (**Figure 3-1C**).

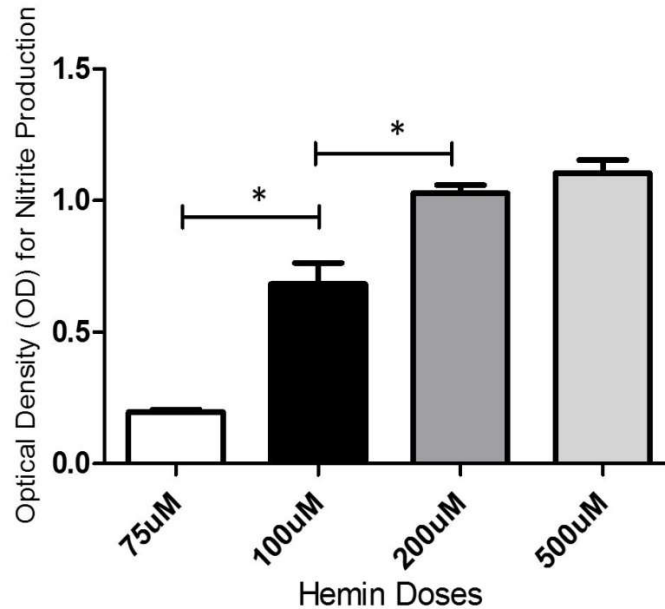


**Figure 3-1: SIMA9 cell morphology and cofilin expression after transfection by scramble (A) and cofilin siRNA (B).** Images were taken at 20X bright field and protein expression was performed by Western blot analysis (C). The Western blot results were expressed as mean  $\pm$  SEM of the ratio of cofilin and  $\alpha$ -tubulin. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$ ).

### 3.2 Hemin dose adjustment and nitrite Oxide production

Previously, different doses of hemin were used to induce microglia activation (Huang, Xi et al. 2002, Lin, Yin et al. 2012, Hanafy 2013). In order to mimic ICH conditions, higher dose of hemin such as 75  $\mu$ M, 100  $\mu$ M, 200  $\mu$ M and 500  $\mu$ M were

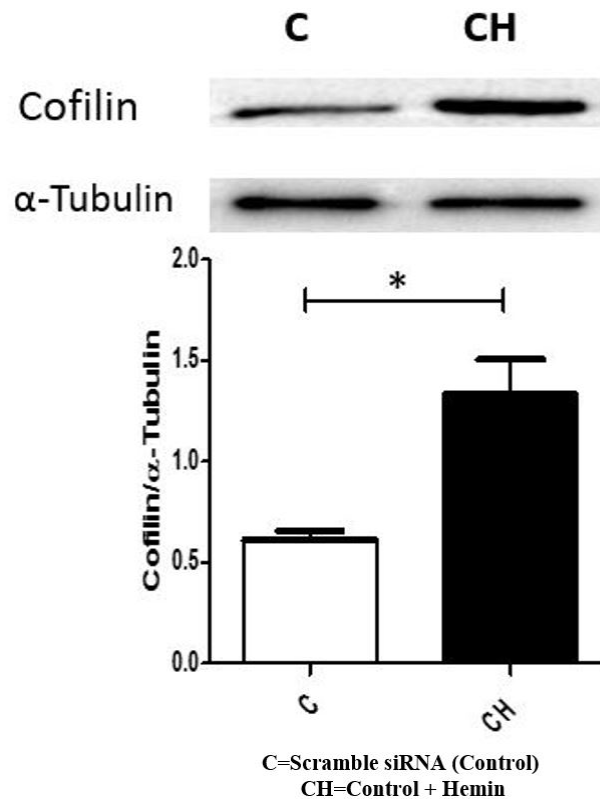
initially selected. It was found that there was a dose dependent increase of NO production (Figure 3-2) but due to interference with NO detection at higher hemin dose exposure levels (200 and 500  $\mu$ M), 100  $\mu$ M hemin was selected for subsequent hemin treatments.



**Figure 3-2: Dose dependent hemin treatment-induced nitrite oxide (NO) production.** There was gradual increase in the production of NO production due to increase in hemin dose of 75  $\mu$ M, 100  $\mu$ M, 200  $\mu$ M. But further increase of hemin dose after 200  $\mu$ M lead to interference with optical detection of NO production and therefore giving rise to inconsistency results. The results were expressed as the mean  $\pm$  SEM of OD; 3 independent experiment in triplicate. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant.

### 3.3 Hemin induces increased cofilin expression

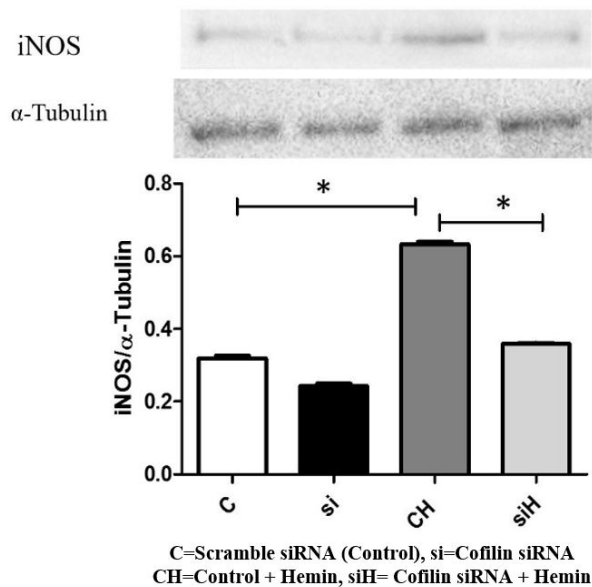
Our lab has previously shown that lipopolysaccharide (LPS) increases the expression of cofilin (unpublished results). Similar conditions were applied in case of hemin treatment in microglia in order to investigate the effect of hemin treatment on cofilin expression and it was found that there was significant increase in cofilin expression ( $p < 0.05$ ) after hemin treatment (**Figure 3-3**).



**Figure 3-3. Hemin treatment-induced increased expression of cofilin.** The Western blot results were expressed as mean  $\pm$  SEM of the ratio of cofilin and  $\alpha$ -tubulin. Independent t test was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$ ).

### 3.4 Cofilin mediates hemin treatment-induced iNOS production

There was a significant difference in microglial iNOS levels among different treatment groups with  $F(3, 8) = 680.4, p < 0.0001$ . Bonferroni post Hoc test indicated that there was no significant difference in iNOS expression levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$ ). However, scramble siRNA transfected microglia treated with hemin showed significantly higher levels of iNOS ( $p = 0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment group (control). On the other hand, significantly reduced ( $p = 0.000$ ) iNOS levels were observed in cofilin siRNA transfected microglia treated with hemin group as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that microglia iNOS expression increase due to long term exposure to hemin treatment is mediated by cofilin (**Figure 3-4**).



**Figure 3-4: Cofilin mediates hemin treatment-induced increase in iNOS production.** The Western blot results were expressed as mean  $\pm$  SEM of the ratio of



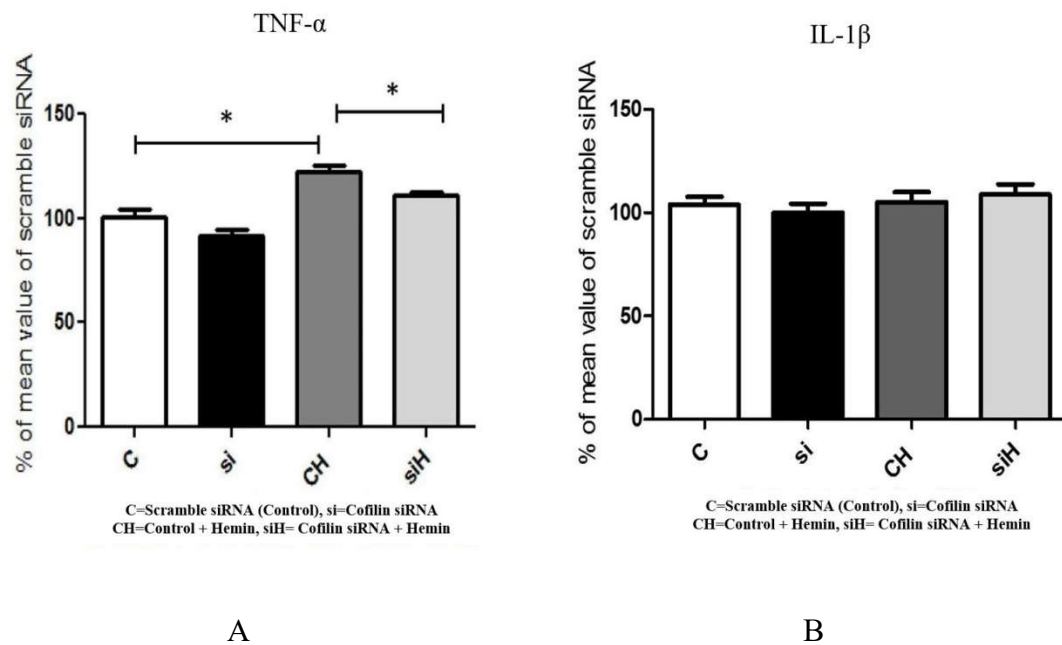
iNOS and  $\alpha$ -tubulin. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$ ).

### **3.5 Cofilin knockdown inhibits hemin treatment-induced microglial TNF- $\alpha$ production but not IL-1 $\beta$**

In this investigation, there was a significant difference in microglia derived TNF- $\alpha$  levels among different treatment groups with  $F(3, 32) = 28.14$ ,  $p=0.000$ . Bonferroni post Hoc test indicated that there was no significant difference in TNF- $\alpha$  levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$ ). However, scramble siRNA transfected microglia treated with hemin showed significantly higher levels of TNF- $\alpha$  ( $p=0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment group (control). On the other hand, significantly reduced ( $p=0.000$ ) TNF- $\alpha$  levels were observed in cofilin siRNA transfected microglia treated with hemin and as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that increased TNF- $\alpha$  released from microglia is due to long term exposure to hemin treatment which is mediated by cofilin (**Figure 3-5A**).

On the other hand, there was no significant difference in microglia derived IL-1 $\beta$  levels among different treatment groups with  $F(3, 32) = 0.6468$ ,  $p= 0.5907$ . Bonferroni post Hoc test indicated that there was no significant difference in IL-1 $\beta$  levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$ ). Difference was also not observed in scramble siRNA transfected microglia treated with hemin derived IL-1 $\beta$  ( $p=1.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment

group (control). Similarly, there was no significant difference observed in IL-1 $\beta$  level ( $p=0.067$ ) between cofilin siRNA transfected microglia treated with hemin and scramble siRNA transfected microglia treated with hemin. These results indicate that increased IL-1 $\beta$  released from microglia is not affected due to long term exposure to hemin treatment (Figure 3-5B).



**Figure 3-5: Hemin induced cytokine production.** The ELISA results for TNF- $\alpha$  (A) and IL-1 $\beta$  (B) were expressed as mean  $\pm$  SEM of the percent of the mean optical density (OD) of the control group. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$  in triplicate).

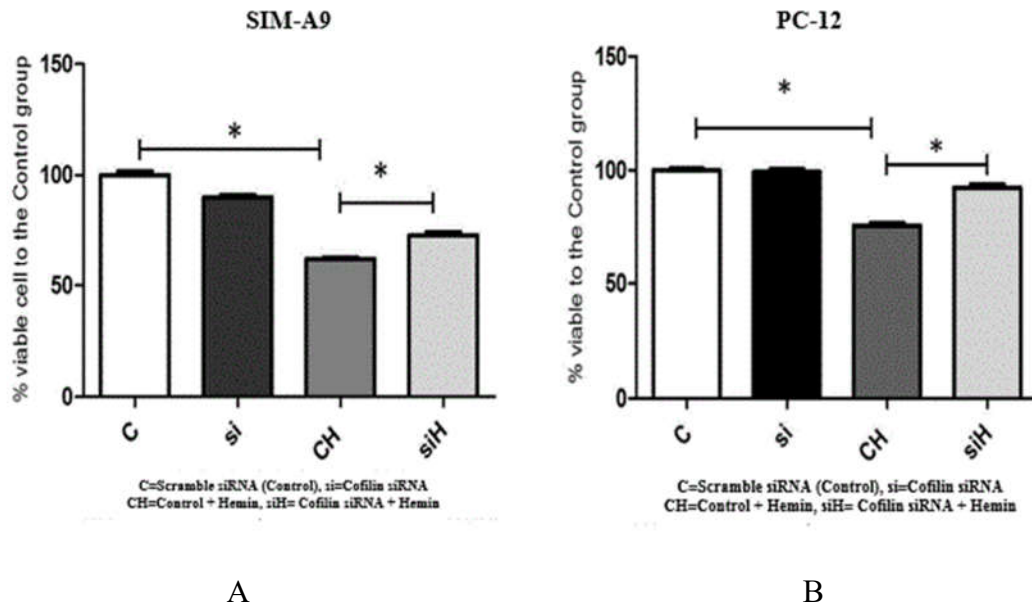
### **3.6 Cofilin knockdown reduces hemin treatment-induced microglial cell death as well as protects undifferentiated PC-12 cells from neurotoxicity from conditioned medium of hemin activated microglia**

There was a significant difference in microglial viability between different treatment groups with  $F(3, 32) = 81.05, p=0.000$ . Bonferroni post Hoc test indicated that there was no significant difference in cell viability between scramble siRNA and cofilin siRNA transfected microglia ( $p>0.05$ ). However, scramble siRNA transfected microglia treated with hemin showed significantly lower cell viability ( $p=0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment group (control). On the other hand, significantly higher cell viability ( $p=0.000$ ) were observed in cofilin siRNA transfected microglia treated with hemin and as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that decreased microglial viability due to long term exposure to hemin treatment is mediated by cofilin (**Figure 3-6A**).

To test the neurotoxic potential of the activated microglia, conditioned medium of transfected and hemin treated microglia was added to undifferentiated PC-12 cells and incubated for 24 h. The viability of treated PC-12 cells was measured using MTT assay.

There was a significant difference in PC-12 viability between different treatment groups with  $F(3, 32) = 86.11, p=0.000$ . Bonferroni post Hoc test indicated that there was no significant difference in cell viability between scramble siRNA and cofilin siRNA transfected group ( $p>0.05$ ). However, scramble siRNA transfected group treated with hemin showed significantly lower cell viability ( $p=0.000$ ) as compared to scramble siRNA transfected group without hemin treatment group (control). On the other hand, significantly higher cell viability ( $p=0.000$ ) was observed in cofilin siRNA transfected group treated

with hemin and as compared to scramble siRNA transfected group treated with hemin. These results indicate that decreased PC-12 viability due to long term exposure to hemin treatment is mediated by cofilin (**Figure 3-6B**).



**Figure 3-6: Cell viability assay after; hemin treatment in SIM-A9 cells (A) and exposure of undifferentiated PC-12 cells to conditioned media from SIM-A9 cells (B).** The MTT assay results were expressed as mean  $\pm$  SEM of the percent of the viable cells to the control group. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$  in triplicate).

### 3.7 Cofilin knockdown inhibits microglial cell migration

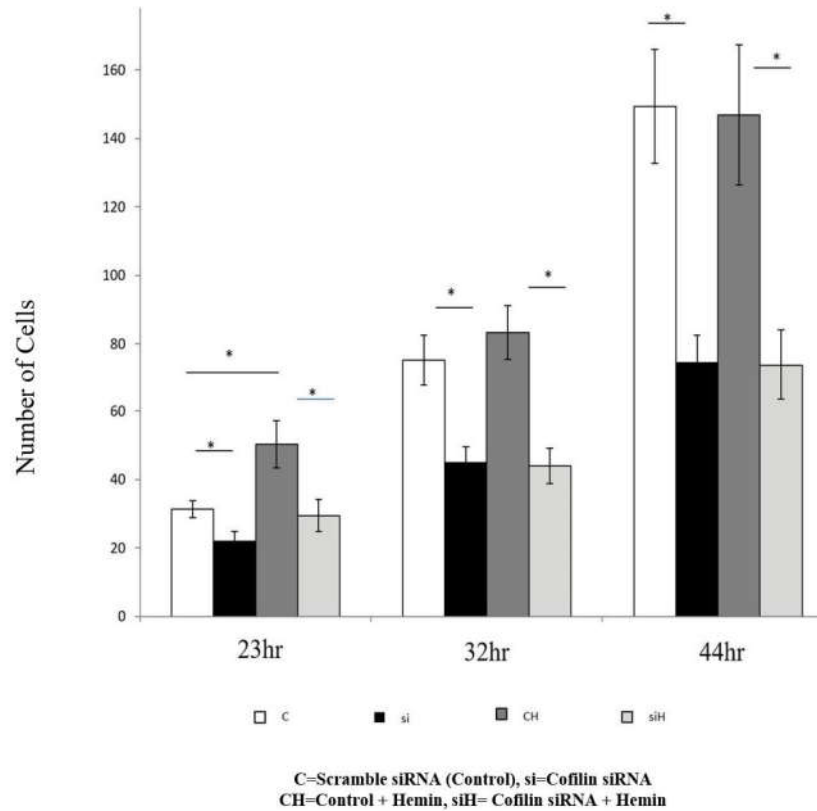
Scratch assay was conducted to test the effect of hemin as well as cofilin knockdown on microglial cells migration (**Figure 3-7**). Only at 3 h of scratch and 23 h of

hemin treatment there was significant reduction of cell migration ( $p < 0.05$ ) but in all time points the number of microglia migrated to the scratched area was significantly less in cofilin siRNA transfected cells (**Figure 3-7**). Different time periods were considered to reduce the impact of cell proliferation in the scratched area.

At 3 h of scratch and 23 h of hemin treatment, there was significant difference among different treatment groups with  $F(3, 44) = 6.944$ ,  $p = 0.001$ . Bonferroni post Hoc test indicated that there was a significant difference in migration between scramble siRNA and cofilin siRNA transfected microglia ( $p = 0.031$ ). Similar results were obtained ( $p = 0.025$ ) in case of scramble siRNA transfected microglia treated with hemin as compared to scramble siRNA transfected microglia without hemin treatment group (control). There was significantly reduced number of microglia ( $p = 0.000$ ) in cofilin siRNA transfected microglia treated with hemin group as compared to scramble siRNA transfected microglia treated with hemin. This indicated that microglial migration is reduced due to both hemin treatment as well as cofilin knockdown at 3 h after scratch and 23 h after hemin treatment.

At 12 h of scratch and 32 h of hemin treatment, there was significant difference among different treatment groups with  $F(3, 44) = 10.152$ ,  $p = 0.000$ . Bonferroni post Hoc test indicates that there was significant difference in migration between scramble siRNA and cofilin siRNA transfected microglia ( $p = 0.010$ ). However, such difference was not observed ( $p = 1.00$ ) in case of scramble siRNA transfected microglia treated with hemin and as compared to scramble siRNA transfected microglia without hemin treatment group (control). There was significantly reduced number of microglia ( $p = 0.000$ ) in cofilin siRNA transfected microglia treated with hemin group as compared to scramble siRNA transfected microglia treated with hemin. This indicated that microglial migration at 12 h of scratch

and 32 h of hemin treatments is reduced due to cofilin knockdown but hemin was not found to affect migration of microglia neither in scramble siRNA group nor in cofilin siRNA group.



**Figure 3-7: Cofilin mediated microglial migration after hemin treatment.** The number of microglia migrated to the area of scratch in the arbitrary dimension of 330X800 or 800X330 depending of the orientation of the scratch. The cells were counted manually with the help of Image J software (NIH, US). The results were expressed in mean  $\pm$  SEM of the number of the cells migrated to the randomly chosen area with fixed dimension within scratch area. One-way ANOVA followed by Bonferroni post hoc was

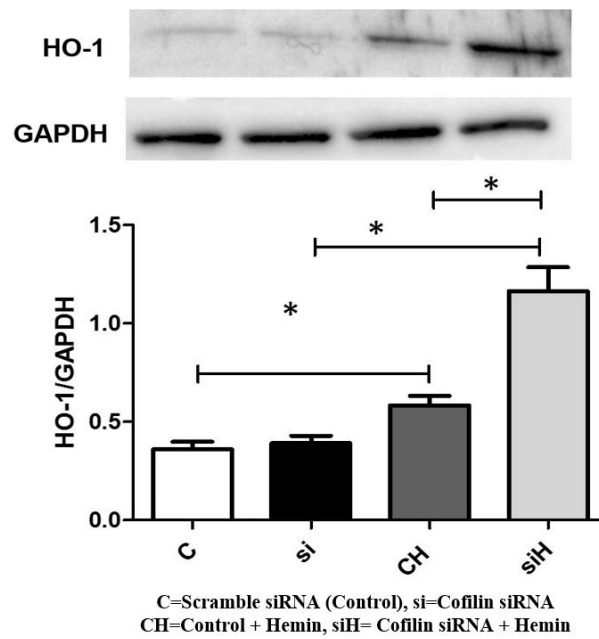
performed and the asterisk (\*) indicates statistically significant differences and the value  $p < 0.05$  was considered significant ( $n=3$ ).

At 24 h of scratch and 44 h of hemin treatment of scratch, there was significant difference among different treatment groups with  $F(3, 44) = 8.428$ ,  $p=0.000$ . Bonferroni post Hoc test indicates that there was significant difference in migration between scramble siRNA and cofilin siRNA transfected microglia ( $p=0.005$ ). However, such difference was not observed ( $p=1.00$ ) in case of scramble siRNA transfected microglia treated with hemin as compared to scramble siRNA transfected microglia without hemin treatment group (control). There was significantly reduced number of microglia ( $p=0.000$ ) in cofilin siRNA transfected microglia treated with hemin and as compared to scramble siRNA transfected microglia treated with hemin. This indicated that microglial migration at 24 h of scratch and 44 h of hemin treatments is reduced due to cofilin knockdown but hemin was not found to affect migration of microglia neither in scramble siRNA group nor in cofilin siRNA group.

### **3.8 Cofilin knockdown increases hemin treatment-induced HO-1 and Nrf2 expression**

There was a significant difference in microglial HO-1 expression among different treatment groups with  $F(3, 8) = 28.35$ ,  $p < 0.0001$ . Bonferroni post Hoc test indicated that there was no significant difference in HO-1 expression levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$ ). However, scramble siRNA transfected microglia treated with hemin showed significantly higher levels of HO-1 expression ( $p=0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment

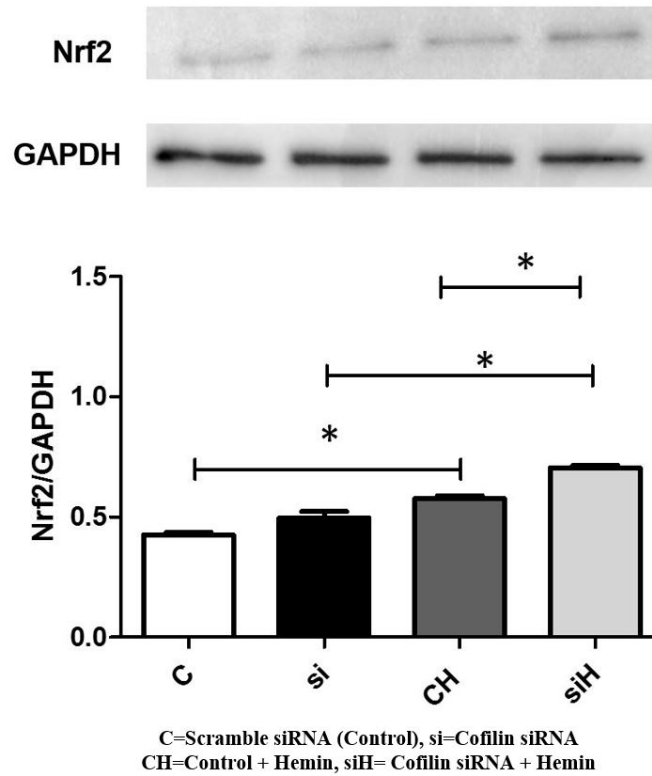
group (control). On the other hand, significantly higher ( $p=0.000$ ) HO-1 expression levels were observed in cofilin siRNA transfected microglia treated with hemin and as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that microglial HO-1 expression is increased due to long term exposure to hemin treatment and cofilin knockdown further increased HO-1 expression induced by hemin treatment (**Figure 3-8**).



**Figure 3-8: Cofilin mediates hemin treatment-induced hemoxygenase- 1 (HO-1) expression.** The Western blot results were expressed as mean  $\pm$  SEM of the ratio of HO-1 and glyceraldehyde 3-phosphate dehydrogenase (GAPDH). One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value  $p<0.05$  was considered significant ( $n=3$ ).



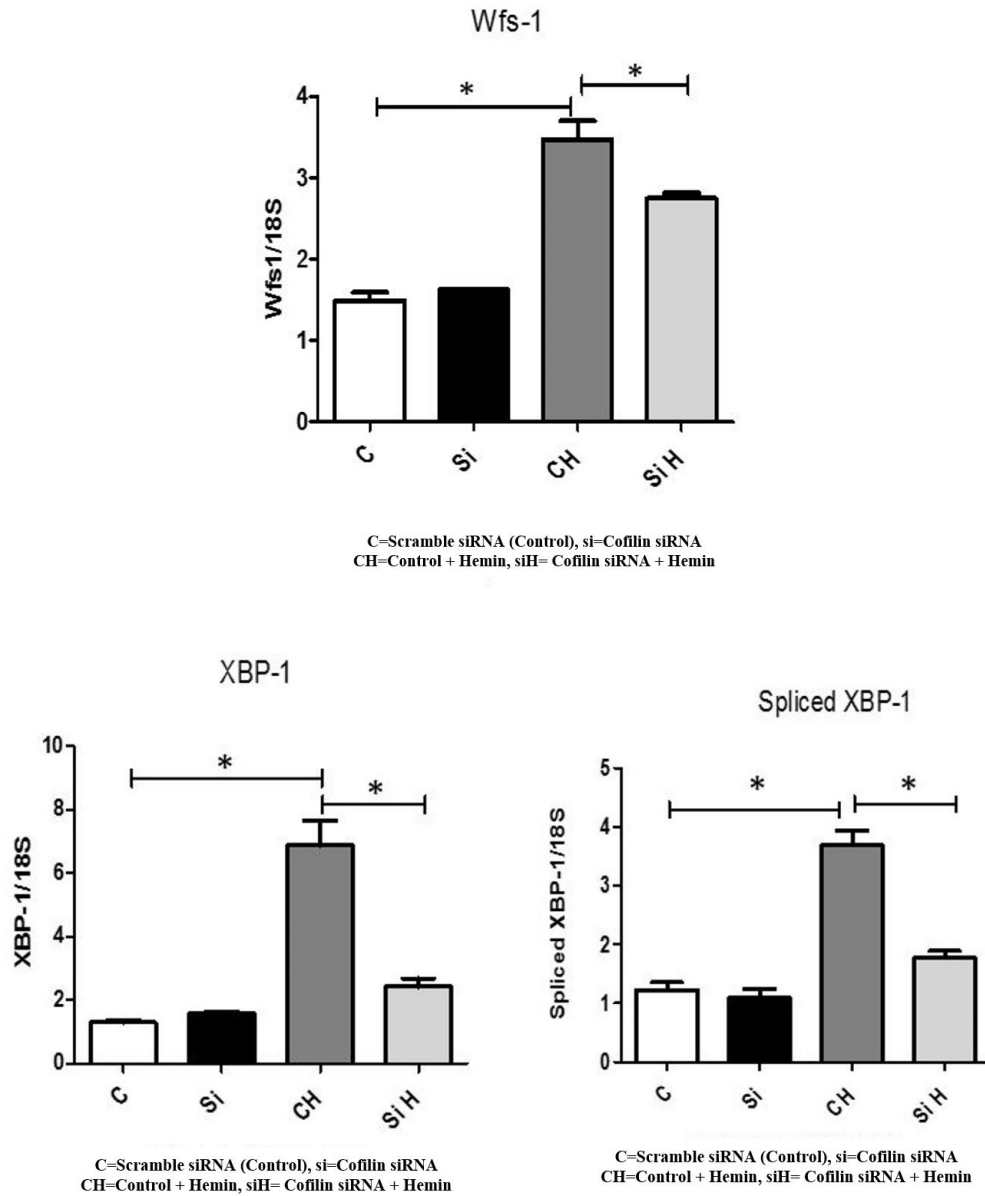
There was a significant difference in microglial Nrf2 expression among different treatment groups with  $F(3, 8) = 51.15, p < 0.0001$ . Bonferroni post Hoc test indicated that there was no significant difference in Nrf2 expression levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$ ). However, scramble siRNA transfected microglia treated with hemin showed significantly higher levels of Nrf2 expression ( $p = 0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment group (control). On the other hand, significantly higher ( $p = 0.000$ ) Nrf2 expression levels were observed in cofilin siRNA transfected microglia treated with hemin group as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that microglial Nrf2 expression is increased due to long term exposure to hemin treatment and cofilin knockdown further increased Nrf2 expression induced by hemin treatment (**Figure 3-9**).



**Figure 3-9: Cofilin mediates hemin treatment-induced nuclear factor (erythroid-derived 2)-like 2 (Nrf2) expression.** The Western blot results were expressed as mean  $\pm$  SEM of the ratio of Nrf2 and glyceraldehyde 3-phosphate dehydrogenase (GAPDH). One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant ( $n=3$ ).

### 3.9 Cofilin knockdown decreases hemin treatment-induced endoplasmic reticulum stress

There was significant difference in microglial Wfs-1, XBP-1, spliced XBP-1 expression with  $F(3, 20) = 852.94, p=0.000$ ;  $F(3, 20) = 38.50, p=0.000$ ;  $F(3, 20) = 48.61, p=0.000$  respectively (**Figure 3-10**).



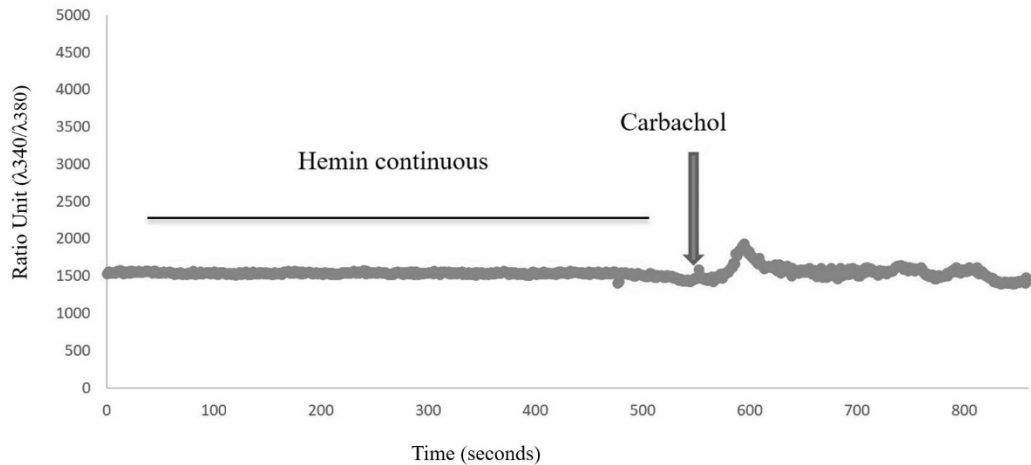
**Figure 3-10: Cofilin mediates hemin treatment-induced endoplasmic reticulum (ER) stress markers (Wfs-1, XBP-1, spliced XBP-1).** RT-qPCR results were expressed

in fold change as mean  $\pm$  SEM (n=3 with duplicate). One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and value of  $p < 0.05$  was considered significant.

Bonferroni post Hoc test indicated that there was no significant difference in Wfs-1, XBP-1, spliced XBP-1 expression levels between scramble siRNA and cofilin siRNA transfected microglia ( $p > 0.05$  for all). However, scramble siRNA transfected microglia treated with hemin showed significantly higher levels of Wfs-1, XBP-1, spliced XBP-1 expression ( $p = 0.000$  for all) as compared to scramble siRNA transfected microglia without hemin treatment group (control). On the other hand, significantly higher ( $p = 0.000$ ) Wfs-1, XBP-1, spliced XBP-1 expression levels were observed in scrambled siRNA transfected microglia treated with hemin group as compared to cofilin siRNA transfected microglia treated with hemin. These results indicate that microglial Wfs-1, XBP-1, spliced XBP-1 expressions are increased due to long term exposure to hemin treatment and cofilin mediate the process.

### **3.10 Hemin treatment alone does not evoke calcium signaling**

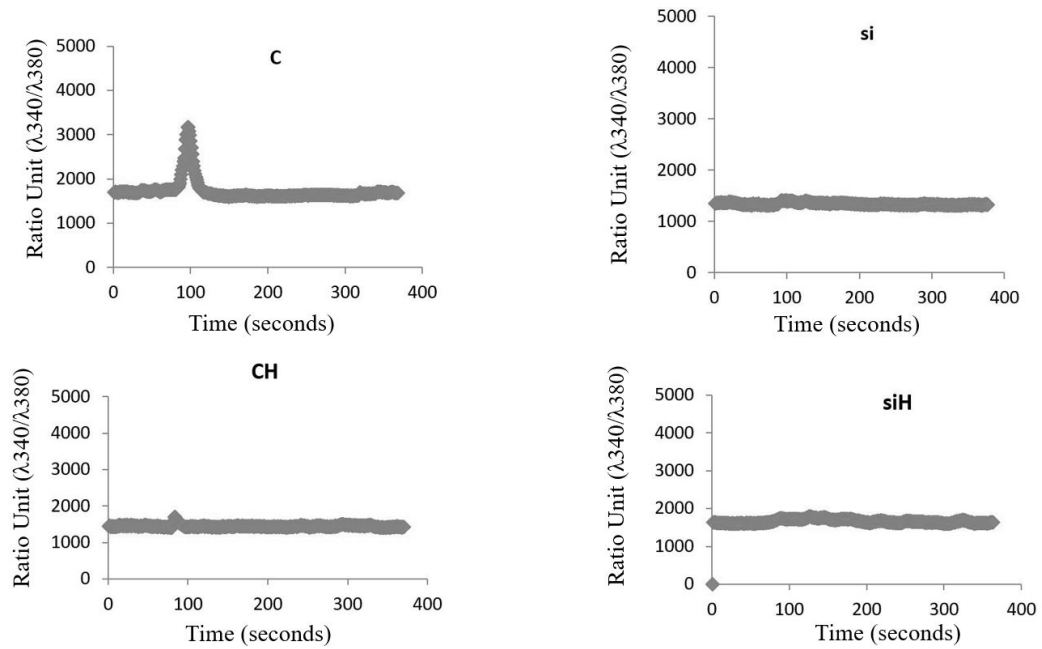
In order to study whether hemin evokes calcium signaling, different concentration of hemin was exposed to microglia and it was found that hemin alone cannot evoke  $\text{Ca}^{2+}$  signaling. However, due to short-term treatment of hemin, microglia do not lose their ability to respond to acetylcholine evoked calcium signaling induced by acetylcholine agonist carbachol (**Figure 3-11**).



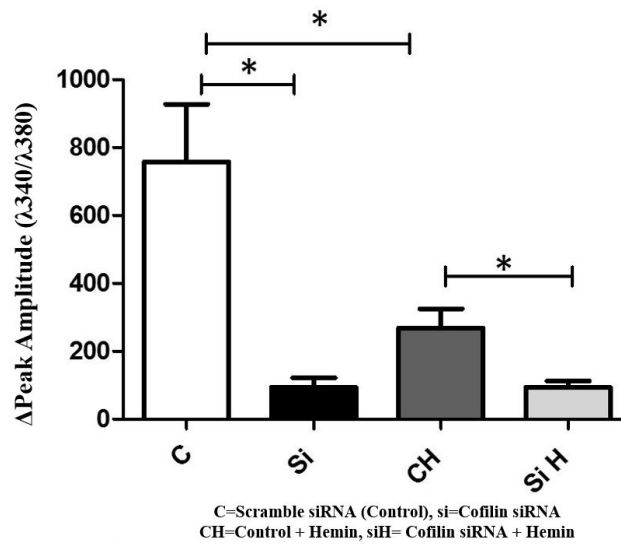
**Figure 3-11: Hemin treatment alone neither evoke calcium signaling nor affect acetyl choline evoked calcium signaling.** 100  $\mu\text{M}$  hemin was continually passed over microglia culture media for 550 seconds followed by 100  $\mu\text{M}$  carbachol (acetylcholine agonist) running over the media for next 250 seconds. The experiments were repeated trice with 10-15 cells stimulated in each independent experiment.

### **3.11 Long term exposure of hemin and cofilin knockdown decreases acetylcholine evoked calcium signaling in microglia**

For evoking  $\text{Ca}^{2+}$  signaling 100  $\mu\text{M}$  carbachol was used which was based on the previous investigations on microglia (Whittemore, Korotzer et al. 1993) and SIMA9 cells responded to acetylcholine evoked  $\text{Ca}^{2+}$  signaling (**Figure 3-11 and Figure 3-12A**)



A



B

**Figure 3-12: Acetylcholine evoked calcium signaling due to hemin treatment. (A)** Cofilin knockdown as well as hemin treatment decreased microglial ability to respond

to acetylcholine evoked calcium signaling. Locke solution was passed for 60 seconds over the microglia media followed by 100  $\mu$ M carbachol (acetylcholine agonist) passing for up to 300 seconds. Experiments were conducted in triplicate with 8-14 microglia in each experiments. (B) The results were expressed in fold change as mean  $\pm$  SEM. One-way ANOVA followed by Bonferroni post hoc was performed and the asterisk (\*) indicates statistically significant difference and the value of  $p < 0.05$  was considered significant.

There was a significant difference in acetyl choline evoked  $Ca^{2+}$  signaling among different treatment groups in microglia with  $F(3, 47) = 16.61, p = 0.000$ . Bonferroni post Hoc test indicated that there was statistically significant difference in  $Ca^{2+}$  signaling between scramble siRNA and cofilin siRNA transfected microglia ( $p = 0.000$ ). Scramble siRNA transfected microglia treated with hemin showed significantly lower levels of  $Ca^{2+}$  ( $p = 0.000$ ) as compared to scramble siRNA transfected microglia without hemin treatment group (control). Significantly lower ( $p = 0.000$ )  $Ca^{2+}$  signaling were observed in cofilin siRNA transfected microglia treated with hemin group as compared to scramble siRNA transfected microglia treated with hemin. These results indicate that that microglial ability to respond to acetylcholine evoked  $Ca^{2+}$  signaling is mediated by cofilin as well as hemin treatment for 24 h (**Figure 3-12B**).

## Chapter 4

### Discussion

In the present investigation, we found that hemin induces microglial inflammation which was observed to be mediated by cofilin. Two previously reported results are relevant in this context. One is on the report that hemin induced inflammation via MyD88/TRIF signaling pathway (Lin, Yin et al. 2012), and other in hemorrhage, that reported inflammation is not due to hemin but due to methemoglobin (Gram, Sveinsdottir et al. 2013). Our investigations are aligned with the former but contradict with the later. This apparent contradiction could be explained by the fact that microglial inflammatory responses *in vitro* is slightly different from those of *in vivo* and the studies on microglia *in vitro* tend to provide increased inflammatory response (Kettenmann, Hanisch et al. 2011). Extremely high dose of hemin induced increased oxidative stress might also lead to microglial inflammation (Hu, Tao et al. 2016). Similar explanation could be provided for hemin induced increased production of TNF- $\alpha$  production but further investigation is warranted in this regard. The observed decrease in TNF- $\alpha$  production in cofilin siRNA transfected microglia might be due to the conditions that lead to decreased cofilin-actin rod formation which is in compliance with previously reported role of cofilin in mediating cellular stress in several disease conditions (Cichon, Sun et al. 2012).



Hemin treatment reduced the viability of the cells which is opposite to the previous report (Mullebner, Moldzio et al. 2015). The explanation lies in the dose use. Previous study used low dose that increase cellular protective mechanism but extremely high dose as in ICH or in our study conditions give rise to cell death. Cofilin knockdown increased cell viability which could be due to decreased hemin take up due to globular microglial cell surface or decreased mitochondrial transition pore permeability induced cell death (Klamt, Zdanov et al. 2009, Posadas, Perez-Martinez et al. 2012). This is also in compliance with previous report with cellular stress induced by chemicals or oxygen glucose deprivation (Madineni, Alhadidi et al. 2016). Hemin treated conditioned media from microglia induced increased cell death of undifferentiated PC-12 cell could be attributed to the increased production of TNF- $\alpha$  or other toxic cytokines from microglia.

Time dependent hemin exposure effects on microglial migration could be due to differential microglial response. Acute hemin treatment affects microglial migration negatively but over time microglia recover by themselves to assume their normal migratory function. However, cofilin knockdown decreased microglial migration is due to dysregulation of cytoskeletal dynamics (Sidani, Wessels et al. 2007).

Since hemin is a known inducer of HO-1 and therefore it is likely that microglia expressed higher HO-1 levels due to hemin treatment. Hemin induced higher expression of Nrf2 by microglia implies that HO-1 might also be upregulated due to Nrf2 translocation to the nucleus and upregulation of antioxidant system (Andreadi, Howells et al. 2006, Li, Dong et al. 2014). Cofilin knockdown alone cannot induce HO-1 and Nrf2 expression but cofilin knockdown microglia treated with hemin have shown increased HO-1 and Nrf2 expression indicating that cofilin knockdown did not reduce microglial ability to degrade

hemin. We found that there was more HO-1 and Nrf2 expression in cofilin knockdown microglia in comparison to scramble siRNA transfected hemin treated microglia. One possible explainable could be that due to cofilin knockdown microglia lose the ability to change the morphology against stress which is not enough to upregulate HO-1 and Nrf2 but when stressor like hemin is used to challenge microglia, the higher expression of HO-1 and Nrf2 might be the work of preconditioning like effects of microglia preparing cells to protect via antioxidant defense mechanism (Cai, Cho et al. 2011).

Wfs-1 is a cellular ER stress marker that attenuates unfolded protein response (UPR) and protects cells from apoptosis (Fonseca, Fukuma et al. 2005, Ueda, Kawano et al. 2005, Zatyka, Da Silva Xavier et al. 2015). XBP1 and spliced XBP-1 are markers of ER stress and indicate adaptive response of the cells (Xu, Bailly-Maitre et al. 2005, van Schadewijk, van't Wout et al. 2012). The increased ER stress due to hemin treatment as observed in our study could be due to increased oxidative stress caused by hemin exposure. Increasing evidence suggest that protein folding and generation of ROS in the ER are closely linked events. Activation of the UPR on exposure to oxidative stress is an adaptive mechanism to preserve cell function and survival (Malhotra and Kaufman 2007). Even though increased ER stress have increased protective HO-1 expression (Liu, Peyton et al. 2005) but such protective mechanisms fail due to extremely high amounts of hemin. In such context, there is sudden increase of oxidative stress both in the cytoplasm as well as in the ER. Even though it was reported earlier that hemin acts as a generic and potent protein misfolding inhibitor (Liu, Carver et al. 2014), in our experiment condition microglia might have lost their ability to tolerate persistent oxidative stress and protein misfolding. Another possibility is that the antioxidant system like SOD, GPx and catalase

inside microglia might have lost their ability to rescue increased oxidative stress caused by hemin exposure. This failure to scavenge free radicle might increase in ER stress. Based on previous studies on different cell types, it can be said that hemin induced ER stress in microglia is due to iron and bilirubin formed from hemin degradation (Barateiro, Vaz et al. 2012, Mullebner, Moldzio et al. 2015). Since cofilin knockdown did not increase Wfs-1, XBP-1 and spliced XBP-1 expression indicating that actin dynamics dysregulation alone is not sufficient to induce ER stress in microglia. However, the cofilin knockdown microglia treated with hemin had reduced expression of Wfs-1, XBP-1 and spliced XBP-1 indicating that cofilin mediate hemin induced ER stress. Such effect could be due to microglial decreased ability to sense external environment or increased protective Nrf2 expression but further study is warranted to substantiate this finding (Glover-Cutter, Lin et al. 2013).

Hemin neither evokes  $\text{Ca}^{2+}$  signaling nor affect acetylcholine evoked  $\text{Ca}^{2+}$  due to short term exposure to microglia indicating that acute hemin exposure cannot induce  $\text{Ca}^{2+}$  dependent downstream effector functions of microglia or other machinery that evoke  $\text{Ca}^{2+}$  signaling. On the other hand, hemin exposure of 24 h reduced acetylcholine evoked  $\text{Ca}^{2+}$  signaling in microglia. This apparently opposite results implies that microglia respond differently based on the duration of hemin treatment (Moller 2002).

Since the mechanism as well as the rate of hemin uptake in microglia has not been determined yet, it cannot be said whether this differential function is due to the interaction of hemin with the surface receptors of microglia or due to interaction with intracellular component of microglia by hemin or degradation products of hemin such as iron, biliverdin bilirubin, and CO. Iron stimulates ryanodine receptor-mediated  $\text{Ca}^{2+}$  release through ROS

produced via the Fenton reaction leading to stimulation of the extracellular-signal-regulated kinases (ERK) signaling pathway (Hidalgo and Nunez 2007). In case of acute hemin exposure, the non-stimulation of  $\text{Ca}^{2+}$  signaling might be due to the fact that, within that time frame, hemin could not enter inside the cell in order to stimulate the cell. Or it could be the fact that hemin enters inside cell but the degradation of hemin does not occur for iron production in order to stimulate the  $\text{Ca}^{2+}$  signaling. Another possibility is that after degradation of hemin, the released iron is taken up by iron-storage protein, ferritin (Ponka, Beaumont et al. 1998) and therefore there is not sufficient iron to stimulate  $\text{Ca}^{2+}$  signaling. On the other hand, long term exposure to hemin causes microglia to take up more hemin, increased production of iron but over production of iron might lead to increased oxidative stress to the cells that is beyond the ability of microglia to cope up with. It might be possible that ferritin is no longer able to store iron and thus causes the cells to turn on their defense system resulting in reducing microglial ability to respond to stimuli. CO produced from hemin breakdown has shown to increase  $\text{Ca}^{2+}$  signaling (Wilkinson and Kemp 2011) and this could be due to increased generation of mitochondrial oxidative stress. In acute hemin treatment on microglia, no effect on  $\text{Ca}^{2+}$  signaling might also be due to the inability of microglia to generate CO adequately within the short period of time. On the other hand, long-term exposure of hemin might cause overproduction of CO resulting in increasing the microglial protective mechanism. Thus, affecting reduced physiological response of microglia to respond to external stimuli. Another possibility is that microglia might lose their ability to process sufficient oxygen which is a requirement for hemin breakdown (Tenhunen, Marver et al. 1968) resulting in their inability to evoke  $\text{Ca}^{2+}$  signaling adequately. Biliverdin and subsequent production of bilirubin from hemin breakdown

might also affect Ca<sup>2+</sup> signaling but the mechanism is poorly understood. This response might be similar to those of iron and CO.

Acetylcholine evoked Ca<sup>2+</sup> signaling in microglia has been suggested to be due to release of Ca<sup>2+</sup> from internal stores such as ER (Whittemore, Korotzer et al. 1993) but it is not clear how hemin interact in this process. Since other signaling pathways have not been investigated, the possibility that hemin or its degradation products affect other internal Ca<sup>2+</sup> signaling pathways cannot be ruled out. Further investigation is suggested to substantiate this finding.

Cofilin knockdown reduced the microglial ability to respond to acetylcholine evoked Ca<sup>2+</sup> signaling could be explained from the fact that the role of cofilin in actin cytoskeleton dynamic is the basis of several effector function (Uhleemann, Gertz et al. 2015). One of the important morphological modification due to cofilin knockdown is that microglia assume globular shape. This shape is different from ‘activated microglia’ or ‘phagocytic microglia’. It is assumed that the actin cytoskeleton dynamic is affected due to the loss of depolymerization of actin by cofilin. Cofilin knockdown results in reduced microglial ability to respond to external stimuli and thus might lose their ability to respond to acetylcholine evoked Ca<sup>2+</sup> signaling but how this interaction translated into near ‘non-respond cell to acetylcholine evoked microglia’ signaling is still not clear (Han, Stope et al. 2007). Therefore, further study is needed to unravel the detailed mechanism.

In this investigation, our hypothesis was that inhibition of cofilin might be a potential therapeutic target for the treatment of ICH. Previous studies in our lab have shown the beneficial effect of cofilin inhibition in ischemic stroke as well as preventing LPS induced microglial inflammation (Madineni, Alhadidi et al. 2016). As an extension of the

project, in this current study we have demonstrated that cofilin inhibition might be a potential therapy to prevent hemin induced microglial excitotoxicity, inflammation, oxidative stress and ER stress. However, the ability of microglia to migrate towards the injurious site is negatively affected therefore affecting negatively on the microglial phagocytosis of cellular debris. Investigation using animal models of hemorrhage will provide further insight into the possibility of cofilin inhibition as a therapeutic target for ICH.

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