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IONIC MODULATORS OF STEM CELL STATE

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Ionic Modulators of Stem Cell State
THESIS FOR DOCTORAL DEGREE (Ph.D.)

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To new beginnings

ABSTRACT

Tissue generation during development and maintenance throughout life relies on the proliferation and sequential specification of a group of cells. Stem cells are defined by the properties of self-renewal (division to produce daughter cells equipotent to the mother) and differentiation (division where at least one daughter is of a more restricted potential). In adult systems, two additional states - quiescence, a dormant state of infrequent proliferation, and activation, a state of increased proliferation, are described. Regulation of these states is a key determinant of health and fitness on a tissue and organism level, as it ensures proper development and regeneration. The aim of this thesis is to investigate how another key system at the cellular level – regulation of ion availability – can modulate cell states of embryonic and adult stem cells.

In **paper I** the effect of lithium chloride (LiCl) on juvenile mouse neural stem progenitor cells (NSPCs) from the subgranular zone (SGZ) of the hippocampus was investigated. Under maintenance conditions, treatment with LiCl increased NSPC proliferation, reducing the fraction of cells in G0/G1. Pre-treatment of NSPCs with LiCl prior to ionizing radiation (IR) exposure reduced DNA damage response activation, and attenuated the IR-induced G1 block, restoring proliferation, although cell death was not reduced.

In **paper II** the effect of ZD7288, a specific blocker of hyperpolarization-activated cyclic nucleotide-gated (HCN) channels, on mouse embryonic stem cell (ESCs) was examined. The blocker attenuated proliferation by extending G1 and S phases. This did not compromise pluripotency, but facilitated spontaneous serum-induced differentiation while reducing the efficiency of directed differentiation towards the neuronal lineage.

In **paper III** expression of HCN family channels and effects of their inhibition in adult NSPCs were described. *Hcn2* and *Hcn3* are expressed throughout the NSPC hierarchy, but only functional in S and G2/M phases. HCN inhibition or knockdown attenuated proliferation due to a reversible G0/G1 accumulation which was accompanied by alterations in activation marker expression, metabolism, and the molecular clock network. A small molecular agonist of Rev-erb- α , a clock component, recapitulated the proliferative effects. HCN inhibition-induced G0/G1 block was shown to have a protective effect during IR exposure of juvenile mice, reducing apoptosis and maintaining proliferation.

In conclusion, lithium, which is proposed to inhibit a number of enzymes by replacing magnesium as a cofactor, and HCN currents, which are involved in regulation of the electrochemical state of the cell, were shown to modulate stem cell state. This suggests that further investigation of these and other ionic modulators is warranted, both for therapy development and in the interests of basic science.

LIST OF SCIENTIFIC PAPERS

- I. Zanni G*, Di Martino E*, **Omelyanenko A**, Andäng M, Delle U, Elmroth K, Blomgren K. Lithium increases proliferation of hippocampal neural stem/progenitor cells and rescues irradiation-induced cell cycle arrest in vitro. *Oncotarget*, 2015, 6(35):37083-97
- II. **Omelyanenko A**, Sekyrova P, Andäng M. ZD7288, a blocker of the HCN channel family, increases doubling time of mouse embryonic stem cells and modulates differentiation outcomes in a context-dependent manner. *SpringerPlus*, 2016, 5:41
- III. Johard H*, **Omelyanenko A***, Gao F*, Zilberter M, Youssef R, Harisankar A, Trantirek L, Walfridsson J, Linnarsson S, Lundkvist G, Harkany T, Blomgren K, Andäng M. Hyperpolarization-activated cyclic nucleotide-gated channels modulate active proliferation and metabolism, and maintain molecular clock oscillations in adult mouse neural stem progenitor cells. *Manuscript*

*denotes equal contribution

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LIST OF ABBREVIATIONS

Akt	Protein kinase B
ALP	Alkaline phosphatase
APC/C	Anaphase-promoting complex
Ara-C	Arabinofuranosyl cytidine
ASCL1	Achaete-scute homolog 1
ATM	Ataxia telangiectasia mutated
ATP	Adenosine triphosphate
ATR	Ataxia telangiectasia mutated Rad3-related
BMAL1	Aryl hydrocarbon receptor nuclear translocator-like protein 1
BMP4	Bone morphogenic protein 4
BMPT1	Bisphosphate 3'-nucleotidase 1
BrdU	5-bromo-2'-deoxyuridine
Ca ²⁺	Calcium
cAMP	Cyclic adenosine monophosphate
Cdc20	Cell-division cycle protein 20
CDK	Cyclin-dependent kinase
CHD7	Chromodomain-helicase-DNA-binding protein 7
Chk1	Checkpoint kinase 1
CKI	Cell-cycle dependent kinase inhibitory protein
CNS	Central nervous system
Cry1	Cryptochrome 1
CsCl	Cesium chloride

DNA	Deoxyribonucleic acid
DNMT3a	DNA (cytosine-5)-methyltransferase 3A
E3.5	Embryonic day 3.5
EdU	5-Ethynyl-2'-deoxyuridine
EGF	Epidermal growth factor
EGFR	Epidermal growth factor receptor
ESC	Embryonic stem cell
FBS	Fetal bovine serum
FGF	Fibroblast growth factor
GABA	γ -aminobutyric acid
GFAP	Glial fibrillary acidic protein
giCSC	Human glioma-initiating cancer stem cell
GO	Gene Ontology
GSK3	Glycogen synthase kinase 3
HCN	Hyperpolarization-activated cyclic nucleotide-gated channel
hfNSC	Human fetal neural stem cell
HLA	Human leukocyte antigen
HSC	Hematopoietic stem cell
IMPase	Inositol monophosphatase
IP3	Inositol-1,4,5 triphosphate
IPPase	Inositol polyphosphate 1-phosphatase
iPSC	Induced pluripotent stem cell
IR	Ionizing radiation

K ⁺	Potassium
K _i	Constant of inhibition
LiCl	Lithium chloride
LIF	Leukemia inhibitory factor
Mad2	Mitotic arrest deficient 2
MAP2	Microtubule-associated protein 2
MEK	Mitogen-activated protein kinase
Mg ²⁺	Magnesium
Msi1	RNA-binding protein Musashi homolog 1
Na ⁺	Sodium
NANOG	Nanog
NCX	Sodium-calcium exchanger
NFIX	Nuclear factor 1 X-type
Nr1d1	Nuclear receptor subfamily 1 group D member 1
NSPC	Neural stem progenitor cell
OCT3/4	Octamer-binding transcription factor 4
OLIG2	Oligodendrocyte transcription factor 2
Pax6	Paired box protein Pax-6
PER2	Period 2
PIP2	Phosphatidylinositol 4,5-bisphosphate
PKC	Protein kinase C
qRT-PCR	Quantitative real-time polymerase chain reaction
Rb	Retinoblastoma protein

REST	RE1-Silencing Transcription factor
RNA	Ribonucleic acid
SGZ	Subgranular zone
shRNA	Short hairpin RNA
siRNA	Short silencing RNA
SMIT1	Sodium/myo-inositol cotransporter 1
SOX2	SRY (sex determining region Y)-box 2
SSEA1	Stage-specific embryonic antigen 1
SVZ	Subventricular zone
TEA	Tetraethylammonium
TLX	Nuclear receptor subfamily 2 group E member 1
γ -H2AX	Phosphorylated Histone H2A

1 INTRODUCTION

1.1 THE STEM CELL PARADIGM

We all start life as a single cell and end it as a multitude. From a single fertilised egg the many cell types found in the adult body emerge. This phenomenon represents a fundamental difference between uni- and multicellular organisms, yet the molecular mechanisms deciding cellular birth and death are remarkably similar from yeast to human. While the lifecycle of even a unicellular eukaryote, such as yeast, already allows access to a number of cell fates (division, mating, apoptosis), the increased complexity of multicellular life requires the introduction of additional fate states, at least in some cells. Today's answer to this conceptual challenge is introduction of a category of cells with greater developmental potential than mature somatic cells – the so called stem cells. These are cells possessing the two properties essential for making development (embryonic stem cells) and maintenance (adult stem cells) of multicellular life as we observe it possible: self-renewal and differentiation¹.

Self-renewal is the property of dividing to give rise to two daughter cells of which at least one has the same developmental potential as the original cell. This ensures the possibility of maintaining the stem cells population (in case of asymmetric division) while increasing cell number, or even expanding the stem cell pool (in case of symmetric division to two cells equipotent to the mother cell). Although not every stem cell will undergo self-renewal every time, it is a cardinal property of stem cells that under the right conditions they may do so. This property is essentially no different from regular division of adult cells – a fibroblast that will always be a fibroblast may divide to give rise to two fibroblasts of the same potential, which also can divide to give rise to fibroblasts. What makes self-renewal different from simple division is its tie to the second cardinal property of stem cells: differentiation.

Differentiation is the property of having the potential to give rise to cells with a developmental potential distinct from, and more restricted than that of the original cell. This allows first the creation of the germ layers and distinct lineages, and finally the generation of terminally differentiated cells, either capable of division or that have exited the cell cycle. These cells can then develop the distinct cell physiology that allows them to be best suited for their role in the tissue and the organism.

1.1.1 Stem cells in regeneration: quiescent, active, and differentiated

While embryonic stem cell lineages are generally a transient fate state, in organs with substantial cell turnover, life-long self-renewal and differentiation of a progenitor population

is the primary source of new cells. This allows mature cells to be functionalized without compromising tissue turn over, since when the cells reach the end of their life cycle they are replaced through differentiation of newly divided cells higher up along the differentiation hierarchy. Studies of the blood system in adult mice were instrumental both in the initial definition and development of the stem cell paradigm. Many credit James Till and Earnest McCullough with the “discovery” of stem cells during their work on hematopoiesis². Based on their findings that the blood system of a bone marrow-ablated host could be fully reconstituted, even in the long term, by some cells from the bone marrow of a donor, both the defining properties and the functional definition of stem cells in the blood system were formulated. Serving as a prototype in stem cell biology, the hematopoietic system is still one of the best understood and certainly most clinically relevant adult stem cell systems.

Till and McCullough also reported equipotent cells that were long- or short-term repopulating based on the length of time during which they were able to sustain hematopoiesis. The long-term repopulating cells were found to be label retaining, indicating low division frequency. This combination of high potency and low division frequency represents the state now commonly referred to as “quiescence”. This concept is applicable in many adult stem cell systems and the quiescent and activated (proliferative) states are associated with expression or repression of particular genes. Well studied in the blood system, these genes, and even hematopoietic stem cell (HSC) properties, vary depending on the timing and location of hematopoiesis, indicating that stem cell state is an emergent property of intrinsic factors and environmental influence.

Environmental factors (secreted and physical) that a stem cell experiences are often collectively called “the niche”. The niche is known to play an important role in regulating all aspects of stem cell behaviour. Both secreted and physical niche factors have been reported to regulate stem cell pool size³, and inherent proliferative properties. As an example, long-term repopulating HSCs in the liver, the site of fetal hematopoiesis in mammals, are more proliferative and express different surface markers than they do in the bone marrow, the site of adult hematopoiesis⁴. Interestingly, the location of the HSC niche varies not only through development but also among species: from kidney in fish, to bone marrow in birds. Perhaps most intriguingly, the HSC niche seasonally migrates between the bone marrow and the liver in some frogs^{4,5}. This diversity may reflect different functional requirements for the niche (e. g. accessibility of external cues, protection from damage), understanding which may be instrumental to mobilizing endogenous or using adult stem cells for therapy.

1.2 ADULT NEUROGENESIS

The brain and the central nervous system (CNS) were long believed to exclusively be populated by non-proliferative cells, as there is no obvious ongoing turnover or need for regeneration, unlike in the blood system⁶. This was questioned extensively in the 1960's, when Altman and coworkers published a number of papers showing proliferation in the adult brain and providing evidence of neurogenesis in the mouse and other mammals⁷⁻⁹. Evidence of adult neurogenesis in humans only became available in 1998, when Eriksson and coworkers published their work identifying dividing cells in the brains of cancer patients¹⁰. At present, the two neurogenic niches most extensively studied in mammalian brains are the subgranular zone (SGZ) of the hippocampus, and the subventricular zone (SVZ) (see Figure 1) of the lateral ventricle, with ongoing controversy regarding the existence of additional niches¹¹. Although proliferative activity is more abundant in the SVZ in the mouse¹², both niches are populated by cells which undergo division and give rise to new neurons. The SGZ produces granule neurons and astrocytes for the dentate gyrus, whereas SVZ gives rise to oligodendrocytes and to neuroblasts that migrate along the rostral migratory stream, becoming olfactory bulb interneurons^{13,14}. The dynamics of neurogenesis have also been investigated, and the number of dividing cells as well as newly born neurons was shown to decrease with age both in mice^{15,16} and humans¹⁷.

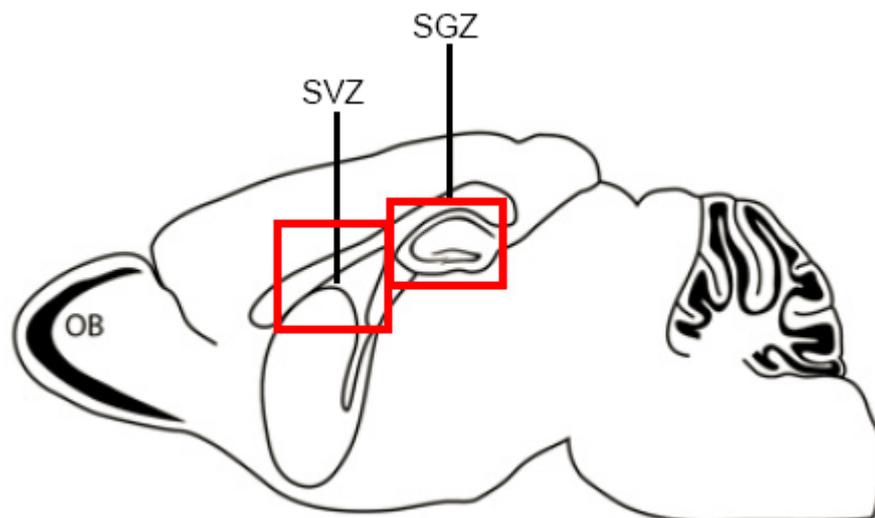


Figure 1: Neurogenic zones in the mouse brain. Modified from *Genes and Development*, 26 (10), Hsieh J, Orchestrating transcriptional control of adult neurogenesis, 1010-1021 (2012) under the Creative Commons license.

Unlike in other adult tissues, the main role of the proliferative progenitor compartment in the brain appears to be facilitation of plasticity, rather than tissue maintenance and replacement of exhausted cells. On a cellular level new-born neurons in the adult brain possess electrical

properties unique from those of mature neurons and can thus code information differently¹⁸. On a tissue level, newly generated neurons form new synapses, modifying existing neural circuitry¹⁹. When looking at the level of cognitive function and behaviour in the mouse, neurogenesis contributes to both short- and long-term memory. Olfactory bulb neurogenesis is believed to be important for short-term memory and olfactory learning^{20,21}, while neurogenesis in the hippocampus is important for long-term memory formation and pattern recognition^{20,22}. Adult neurogenesis has also been suggested to be involved in mood regulation and hippocampal neurogenesis was shown to be necessary for function of a number of antidepressants^{23,24}. Intriguingly, neurogenesis was also shown to be increased in response to damage, such as due to epileptic seizures²⁵ and stroke²⁶, in line with a more conventional role of tissue repair and regeneration for the neurogenic pool.

With the homeostatic and potentially injury repair functions of adult neurogenesis in mind, a number of clinical interventions are possible. Transplantation or activation of endogenous neural precursors into an injured brain is a potential strategy. This has been tested in Parkinsonian patients who received fetal cell transplants²⁷. Additionally, modulation of proliferation to affect mood disorders may in fact be current practice already, as a number of antidepressant therapies are hypothesized to function through inducing proliferation^{28,29}. In patients experiencing loss of neurogenic potential an alternative clinical strategy is preservation or enhancement of the proliferative pool to improve cognitive outcomes. This is particularly relevant for when the injury is a side effect of therapy, such as what is seen in treatment of pediatric brain cancers³⁰. Adults who had suffered from brain cancer as children were found to have reduced neurocognitive abilities, with degree of impairment proportional to dose of radiation received³¹. Given that postnatal neurogenesis is highest at younger ages, it has been proposed that irreversible damage to the proliferating progenitor pool may underlie the impairments observed later in life³². Strategies to address this have included stimulating neurogenesis after the event³³, as well as neuroprotective neoadjuvant treatments³⁴, including the well-known mood stabilizer, lithium³⁵.

1.2.1 Stem cell hierarchy in the adult mouse brain

Neural cells originate from the ectodermal layer of the embryo, which forms the neurotube made up by neuroepithelial cells, a symmetrically and rapidly dividing population of early neuronal progenitors. Neuroepithelial cells are further specified to become radial glia cells, which continue to divide symmetrically, self-renewing until the onset of neurogenesis. As neurogenesis commences radial glia cells start to divide asymmetrically giving rise to

intermediate progenitor cells that go on to further differentiate to neurons. Following neurogenesis, gliogenesis is initiated, and when that nears its close, radial glia cells start to divide symmetrically, exhausting their pool^{36,37}. Neurogenesis in gyrencephalic species (e.g. sheep, primates) involves additional intermediate cell types, necessary to build up the many cortical layers that are not found in the most commonly studied lissencephalic animals (mouse) discussed here³⁸.

It is not entirely clear which neurogenic cells in the embryonic developmental cascade give rise to adult neural stem progenitor cells (NSPCs). It has been suggested that SGZ NSPCs are generated late during gestation in the ventral hippocampus and migrate to their eventual location by birth³⁹, whereas SVZ NSPCs are generated *in situ* and simply remain quiescent until the end of neurogenesis⁴⁰. Postnatally, the neurogenic cascade in the SVZ (see Figure 2A) starts with the mostly quiescent, radial glia-like type B cells, located close to the ventricular wall projecting a cilium into the ventricle. These cells express the GFAP and stem cell marker SOX2 and persist over a long time. The transit-amplifying type C cells occupy the same niche, but are further away from the ventricle, proliferate much more, and give rise to type A cells, which are the migrating neuroblasts destined to become neurons⁴¹. In the

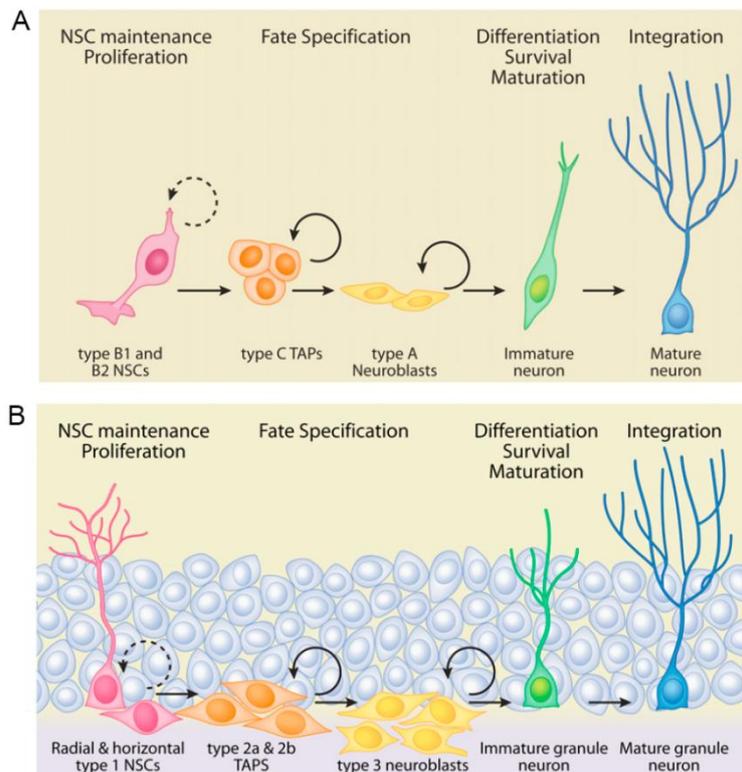


Figure 2: (A) In the SVZ NSPC hierarchy starts with type B cells, transit amplifying type C cells, and type A neuroblasts. (B) in the SGZ radial type I are followed by transit amplifying type II cells and type 3 neuroblasts. Modified from Genes and Development, 26 (10), Hsieh J, Orchestrating transcriptional control of adult neurogenesis, 1010-1021 (2012) under the Creative Commons license.

SGZ (see Figure 2B), type I cells represent the main progenitor population and, similar to radial glia during embryonic neurogenesis, have projections spanning the entire granule cell layer. Like type B cells in the SVZ, they express GFAP as well as SOX2. There are also type

II cells that are less similar to radial glia than type I cells, due to lack of GFAP expression and a shortened morphology, but that still express SOX2. There is significant debate regarding their identity in relation to type I cells, with some arguing that they are more restricted progeny, while others believe the two cell types represent interconvertible pools^{19,42}. Finally, the SGZ also contain intermediate progenitors that undergo symmetrical division to give rise to neural cells without further proliferative potential⁴³. Currently, identification of the various NSPC types relies on their position and morphology *in vivo*, as well as expression of a number of characteristic markers, of which SOX2, GFAP, Prominin1, and Nestin are associated with the most potent cell type (type B and type I).

SVZ and SGZ NSPCs are usually isolated for *in vitro* culture as neurospheres, by dissecting out the relevant anatomical brain region, dissociating the tissue and growing the resulting cell in a defined media supplemented with fibroblast growth factor (FGF) and epidermal growth factor (EGF), which have been shown to support NSPC expansion. Bulk cultures from both neurogenic regions have been shown to exhibit stem cell properties, as they are self-renewing and multipotent, giving rise to neurons, astrocytes and oligodendrocytes *in vitro*^{44,45}. *In vitro* differentiation of clonal SVZ-derived cells initially indicated bipotentiality only, with only neuronal and glial progeny observed⁴⁶, and more recently, it has been suggested that clonally SVZ NSPC can only give rise to neurons⁴⁷. This is in line with *in vivo* reports of single lineage differentiation of SVZ precursors, which also suggested their rapid exhaustion once proliferation is initiated⁴⁸. Similar controversy applies to the SGZ, where there are reports of both exhaustion⁴⁹ and cycling between active proliferation and quiescence⁵⁰. These disparate findings leading to the use of the term “stem progenitor cell”, as they bring up the questions of whether NSPCs are *bona fide* stem cells very tightly controlled by their niche, or represent separate progenitor populations with limited stem cell potential. However, work showing that *in vitro* cultured SGZ NSPCs give rise to SVZ-specific cell types *in vivo*⁵¹ when transplanted into the SVZ argues for the former.

1.2.2 Cell cycle in NSPCs

Cell cycle control in NSPCs involves the canonical cell cycle machinery and follows the usual order of G1-S-G2-M, but with a prominent G0, especially *in vivo*. While G0 is the quiescent phase when cells are neither undergoing nor preparing for division, G1 and G2 are growth phases when the cells are preparing for DNA synthesis and for cell division, respectively. The entire process is regulated by sequential activation of cyclin-dependent

kinases (CDKs) and transcription of specific genes, and involves three checkpoints, at G1/S transition, in G2, and in M phase (see Figure 3).

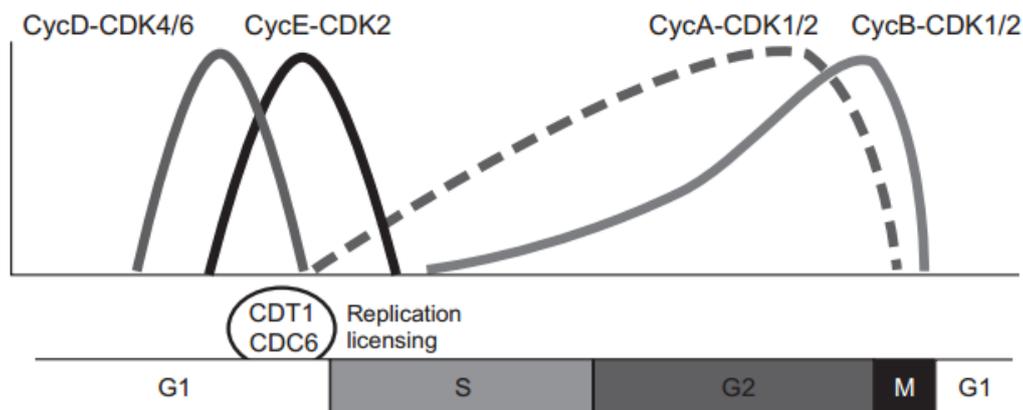


Figure 3: Cyclical activity of cyclins and CDKs regulates progression of somatic cells through the cell cycle. Reprinted from Current topics in developmental biology, 104, Tsubouchi T, Fisher A. Reprogramming and the Pluripotent Stem Cell Cycle. 223-241 (2013) with permission from Elsevier.

Cyclins D (D1/D2/D3) are regulated by mitogen signalling and together with CDK4 and 6 regulate G1 progression by phosphorylating the retinoblastoma protein (Rb), releasing it from binding E2F. Once sufficient E2F is available, cyclins E and A are transcribed and the cell cycle proceeds when cyclin E-CDK2 phosphorylation of Rb reaches critical levels, passing the so called “restriction point” and committing the cell to division. CDK2 then cooperates with cyclin A to phosphorylate DNA synthesis machinery as the S-phase progresses. With the end of S-phase, CDK2 is replaced with CDK1, which cooperates with cyclin A and the G2/M cyclin B to phosphorylate G2/M targets. Once the M-phase is complete and the cell has divided, the cell cycle machinery is reset to the start and cyclin D starts to build up in response to mitogen signalling, if cycling is continuous. The activity of CDKs is regulated not only by the availability of cyclins, but also by their subcellular localization, degradation, and inhibition by cyclin-dependent kinase inhibitor proteins (CKIs). CKIs include INK4 proteins (p15, p16, p18, and p19) that regulate CDK4 and 6 activity and G1 progression, and the Cip/Kip family (p21,p27,p57) that regulate CDK activity in both G1 and G2, and DNA synthesis directly (in the case of p21)^{52,53}.

Expression of CKIs is most notably regulated by p53, which is upregulated in response to DNA damage and is the main effector in the G1 DNA damage checkpoint. In response to DNA damage during G1, p53 upregulates CKI expression, preventing G1/S progression and inducing G1 arrest. It can also play a role in G2 DNA damage response, where p53-induced p21 expression can inhibit CDK1. G2/M progression is abrogated if DNA damage is present

even in the absence of p53, due to the action of checkpoint kinases 1 and 2 (Chk1 and Chk2). Cell cycle stalling in response to DNA damage allows repair or induction of apoptosis to safeguard genomic integrity in the daughter cells, and is regulated by the ataxia telangiectasia mutated (ATM) and Rad3-related (ATR) proteins. The final checkpoint is the mitotic spindle checkpoint in the M-phase, which ensures correct chromosomal segregation during division. If the mitotic spindle and chromosomes are not correctly assembled, the cells stall due to mitotic arrest deficient 2 (Mad2)-mediated inhibition of cell-division cycle protein 20 (Cdc20), resulting in inhibition of the anaphase-promoting complex (APC/C), which would otherwise be driving the cell cycle forward by ubiquitination and degradation of cell cycle machinery^{52,54}.

1.2.3 Quiescence and activation in NSPCs

The more intriguing and unique aspect of NSPC cell cycle regulation is the regulation of G0/G1 balance. Early studies examining neurogenesis in the adult brain using anti-mitotic drug identified that the majority of NSPCs *in vivo* are quiescent in the G0 cell cycle phase⁴¹. Recent reports are suggesting that although not mitotically active, quiescence is not a passive state and presence of signalling is necessary for its maintenance⁵⁵. WNT signalling has been shown to regulate quiescence⁵⁶, as does γ -aminobutyric acid (GABA) signalling through the GABA_A receptor, the absence of which resulted in NSPC reactivation following irradiation injury^{57,58}. Prominin1, a hallmark of NSPCs, is expressed in primary cilia, known for acting as signalling centres, and is also characteristic of quiescence⁵⁹. In fact, Prominin1 expression coupled with lack of epidermal growth factor receptor (EGFR) expression is commonly used to identify quiescent cells⁶⁰, with the absence of EGFR highlighting their low dependence on mitogens.

In line with this, quiescent cells have been shown to have altered metabolic requirements⁶¹, be resistant to mitochondrial perturbations⁶² and have gene expression signatures consistent with glycolytic metabolism⁶³. A number of cell-intrinsic parameters, such as expression of lineage-related transcription factors (SOX2, REST⁶⁴, OLIG2⁶⁵, ASCL1⁶⁶), the nuclear receptor TLX⁶⁷, circadian regulator PER2⁶⁸, epigenetic modifiers (NFIX⁶⁹, DNMT3a⁷⁰), and cell cycle machinery (CHD7⁷¹, p27⁷²) have been linked to regulation of the stem cell pool and the balance between quiescence and activation. Niche factors such as growth factor signaling through FGF2 and bone morphogenic protein 4 (BMP4)⁷³ and vascular contact⁷⁴ have also been implicated in maintenance of quiescence and activation. Unfortunately, the mechanisms

of interconversion between the two pools and, in particular of quiescence induction, are still poorly understood, highlighting the need for further study.

1.3 EMBRYONIC STEM CELLS

Mouse embryonic stem cells (ESCs) were first isolated and cultured in 1981 from the inner cell mass of the preimplantation blastocyst at embryonic day 3.5 (E3.5), or the epiblast at E4.5, and represent cells of the late pre-implantation blastocyst⁷⁵ (see Figure 4). A similar cell type can be derived from early human pre-implantation blastocysts that are the by-product of *in vitro* fertilization, although these cells represent a developmentally distinct cell type. In normal development, ESCs give rise to the three embryonic germ layers – mesoderm, endoderm, and ectoderm, making all the adult tissues of embryonic origin. *In vitro*, they can be maintained indefinitely (as they express telomerase and are not subject to telomere shortening) without losing their stem cell properties, provided they are cultured under permissive conditions.

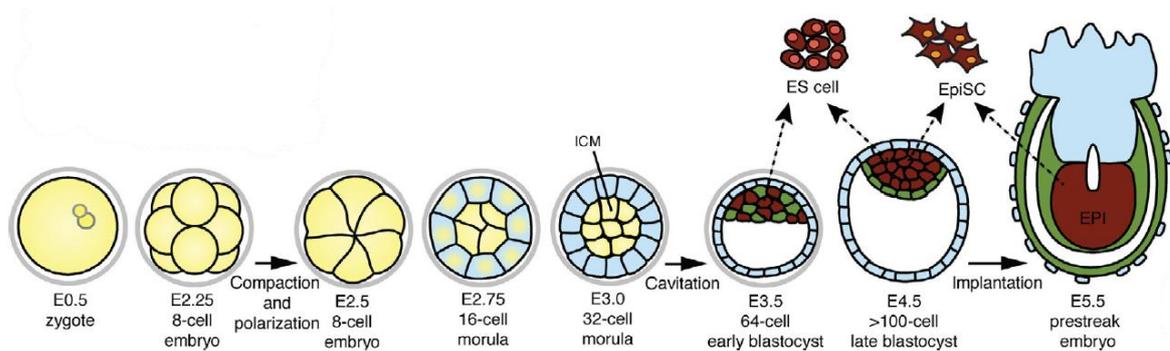


Figure 4: Early development of the mouse. Embryonic stem cells are derived from and correspond to E3.5 or E4.5 the pre-implantation blastocyst. Reprinted from Current topics in developmental biology, 107, Posfai E, Tam OH, Rossant J. Mechanisms of pluripotency in vivo and in vitro. 1-37 (2014) with permission from Elsevier.

The ability of ESCs to be cultured long term is remarkable not only because they are not transformed, but also because unlike adult stem cells, which persist for many years *in vivo*, they usually represent a very transient state in mouse development. An exception exists in diapause – a facultative stalling of mouse embryo development at an about 130 cell stage induced by metabolic restriction, such as lactation of the mother⁷⁶. At this time, metabolic processes such as glycolysis, in the cells of the blastocyst are downregulated⁷⁷, and autophagy has been proposed to take place in some cells to generate nutrients for the others⁷⁸. The process is proposed to be regulated by microRNAs⁷⁹ and results in G0/G1 stalling⁸⁰ and major alterations in cell metabolism⁷⁶, resembling quiescence of adult stem cells. Recent work comparing transcriptional profiles of diapaused and developing blastocysts has in fact

identified E4.5 as the active stage most similar to that seen in the diapaused epiblast⁸¹, suggesting that *in vitro* cultured ESCs should also be capable of reaching the diapaused state, but the conditions necessary for reaching it are only just beginning to be described.

ESC are fascinating for many reasons, not the least of which being their extensive proliferative capacity and ability to generate most mature cell types, given the proper inductive cues. As with many cell types, identification of conditions permissive of efficient *in vitro* maintenance and expansion is critical for allowing both basic and applied research using ESCs. Generation of knockout mice, which has become a cornerstone of biological research, has been based on ESC technology for many years now and is one example of the usefulness of ESC as tools. This process has been significantly facilitated by the development of defined culture methods, which use small molecule inhibitors to generate uniform populations of undifferentiated ESCs⁸². However, the final goal remains to develop this cell type for regenerative medicine. A decade ago a breakthrough in stem cell biology was made by the team of Shinya Yamanaka, who developed the technology for reverting mature cells to the ESC stage by induction of a set of key genes linked to ESC identity⁸³, drawing this goal closer. However, clinical use of ESC-based technology will still require, at the very least, efficient and well-defined protocols for stem cell differentiation as well as culture, and adaptation of them to human cells.

1.3.1 Stemness and differentiation in embryonic stem cells

Embryonic stem cell state is characterized by self-renewal and the ability to differentiate to all somatic cell types (pluripotency). As such, the gold-standard functional test to determine stem cell capacity in ESCs is aggregation with or injection into a developing blastocyst, which should result in chimeric embryos with ESC contribution to all germ layers. The key to ESC identity lies in the expression of the core pluripotency transcription factors: OCT3/4, SOX2, and NANOG. Expression of these genes endows ESCs with their characteristic properties: clonogenicity (ability of a single cell to give rise to a colony) and pluripotency (ability to give rise to cells of all embryonic germ layers). Insufficient expression of these genes, on the other hand, results in ESC differentiation even under maintenance conditions, and in the case of OCT3/4, even overexpression perturbs pluripotency⁸⁴. These transcription factors function by regulating their own and each other's expression, and cooperatively bind peripheral targets that are also involved in pluripotency maintenance, making the network even more robust. They also bind the promoters and repress the expression of differentiation inducers, such as GATA6 which drives the first *in vivo* differentiation event – the formation

of the primitive endoderm⁸⁵. Conclusive proof of this is the ability of the core pluripotency factors to induce the ESC state in mature cells when overexpressed during the process of reprogramming, giving rise to induced pluripotent stem cells (iPSCs)⁸³. This technology has been extended to human cells⁸⁶, and has initiated discussions of comprehensive human leukocyte antigen (HLA)-based donor cell line biobanking initiatives that would provide nation-wide coverage and near-universal access to future stem-cell based therapies with minimal immunological complications⁸⁷.

Efficient maintenance of the pluripotent state *in vitro* was traditionally achieved by culture in media supplemented with leukemia inhibitory factor (LIF)⁸⁸ and serum (with BMP4 believed to be the key molecule)⁸⁹, that have more recently been replaced by two small molecule inhibitors: one targeting mitogen-activated protein kinase (MEK) and the other targeting glycogen synthase kinase 3 (GSK3)⁸². The role of MEK inhibition in this culture protocol is to inhibit FGF signalling, which is responsible for initiation of differentiation already at the level of the inner cell mass, where it is essential for giving rise to the differentiated hypoblast along with the pluripotent epiblast⁹⁰. FGF signalling is endogenous to ESCs, which produce FGF4 and express the FGF receptor, and the knockout of *Fgf4* results in impaired differentiation to both ectoderm and mesoderm lineages⁹¹, suggesting that it plays a role in creating the heterogeneity necessary to allow exit from pluripotency and induction of lineage specification. The role of GSK3 inhibition is less clear, although the majority of its effect is believed to be related to activation of WNT signalling. Activation of WNT signalling through genetic perturbation of GSK3 phosphorylation-mediated β -catenin degradation showed a dose dependent inhibition of differentiation⁹², although other effects of GSK3 inhibition, such as increase in colony forming ability and c-Myc expression were not recapitulated⁸².

Major progress in defining protocols for ESC differentiation has also been made. Whereas differentiation in general is easily achieved through withdrawal of maintenance factors (LIF), induction of particular cell identity is often a lengthy process of multistage differentiation, in many cases closely mimicking development and using complex components, such as stromal cells, which are not compatible with eventual clinical use⁹³. Significant progress is being made, and the development of a clinically-compatible differentiation protocol for retinal epithelium⁹⁴ has resulted in one of the first ever ESC-derived cell products entering clinical trials⁹⁵. However, many differentiation protocols still rely on modulation of kinase or receptor activity with often costly peptides and proteins, often of non-human origin, although efforts are being made to address this⁹⁶.

1.3.2 The unique ESC cell cycle and self-renewal

Given the speed of proliferation necessary to accommodate the growth seen in the first days of development, it is not surprising that the architecture of ESC cell cycle is drastically different from that of mature cells. *In vivo*, cell cycle length at the developmental stages corresponding to ESCs has been estimated at 9.1-11.5 hours⁹⁷, while ESC doubling times under standard culture conditions vary between 8 and 12 hours^{98,99}, although they have been reported to increase to over 30 hours in serum-free culture⁹⁹. In comparison, many mature cells do not proliferate, and generation times for commonly cultured mature cells, such as mouse embryonic fibroblasts, are reported to be around 19 hours¹⁰⁰. The observed shorter generation time in ESCs is due to shortening of both G1 and G2 growth phases, with ~60% of cell cycle spent in S-phase, as opposed to ~16-20% of cycling time devoted to S-phase in mature cell types⁹⁸.

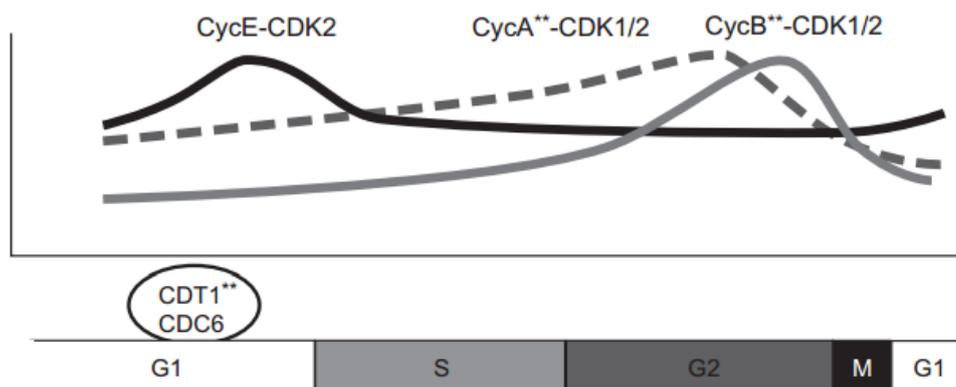


Figure 5: ESCs have a unique cell cycle characterized by constitutive activity of cycling and CDKs. Reprinted from Current topics in developmental biology, 104, Tsubouchi T, Fisher A. Reprogramming and the Pluripotent Stem Cell Cycle. 223-241 (2013) with permission from Elsevier.

This is possible due to radically altered activity of cell-cycle machinery and absence of checkpoints. In somatic cells, progression through the cell cycle is tightly regulated to ensure conditions are right for the next phase of the cell cycle to occur. As described above (see Figure 3), this is coordinated by successive activation of CDKs by their cyclin partners, which are sequentially expressed and promptly ubiquitinated and degraded, ensuring kinase activity is restricted to the proper cell cycle phase⁵². In ESCs this oscillatory behaviour is lacking (see Figure 5), with constitutive CDK2, cyclin A and cyclin E activity⁹⁸, geminin expression, which blocks DNA replication under normal conditions¹⁰¹, and attenuated APC/C oscillations¹⁰². The usual checkpoints that are present in the cell cycle of somatic cells are also absent. The restriction point is absent from ESC, where Rb is in a phosphorylated state throughout the cell cycle¹⁰³. Consistent with this, serum starvation does not result in cell cycle

arrest¹⁰⁴ indicating that mitogen signals are dispensable for ESC proliferation. p16 fails to inhibit the CDK6-CycD3 complex¹⁰⁵ further indicating a lack of a G1/S checkpoint. Induction of checkpoints associated with DNA damage and resulting in p53-induced cell cycle arrest, leads instead to loss of pluripotency, as translocation of p53 to the nucleus downregulates Nanog expression in ESCs¹⁰⁶. Similarly, activation of the mitotic checkpoint, which is functional in ESCs, does not result in apoptosis as in somatic cells¹⁰⁷, a G2/M accumulation and impaired apoptosis were also seen upon CDK1 inhibition in human ESCs¹⁰⁸.

The role of this altered cell cycle architecture in pluripotency maintenance is debated, but the prevailing view is that a link exists. G1 length has been most discussed in this context, as differentiation is accompanied by G1 lengthening¹⁰⁹, while reprogramming to a pluripotent state both requires and results in G1 shortening^{110,111}. Additionally, findings that differentiation factors are preferentially expressed in G1¹¹², and the G1/S inhibitor p27 plays a role in repressing *Sox2* during differentiation¹¹³ further link cell cycle control in G1 and pluripotency maintenance. In complete contradiction, it has also been reported that extension of G1 by exogenous expression of p21 and p27 did not compromise pluripotency under maintenance or differentiation conditions¹¹⁴, although the same authors later reported improved differentiation when G1 lengthening was induced by culture in low serum conditions¹¹⁵.

1.4 THE ROLE OF IONS AND THEIR REGULATION IN EUKARYOTIC CELLS

Whether *in vivo* or *in vitro*, cells exist in an aqueous environment which allows diffusion of nutrients, signalling molecules, and wastes to and from the cell, thus sustaining cellular metabolism. While the cell membrane is permeable to water, it is not permeable to many of the molecules and ions found in the interstitial fluid. In fact, concentrations of many ions and metabolites inside and outside of the cell are unequal, often differing by multiple orders of magnitude¹¹⁶. Sodium (Na⁺) and potassium (K⁺) are the most differentially distributed intracellular ions, each with an over 10-fold difference in cytoplasmic and external concentrations. This gradient is used to drive nutrient transport¹¹⁷, regulate cell volume^{118,119}, and in signalling, by allowing rapid changes in the cell's electrical properties. Whether directly or through their downstream targets, Na⁺ and K⁺ currents influence multiple aspects of cellular physiology including cell division¹²⁰ and differentiation¹²¹.

In addition to Na⁺ and K⁺, concentrations of other metal ions are also tightly controlled in the cell. Calcium (Ca²⁺) can directly influence protein function by inducing conformational

changes upon binding, and is used as a messenger and as a cofactor in processes such as vesicular transport¹²². It has been well established as a regulator of differentiation¹²³, apoptosis¹²⁴, and neuronal firing¹²⁵. The temporal dynamics of fluctuations in Ca^{2+} concentration, and not just the absolute concentration, have also been shown to have a regulator role.

Somewhat less well-understood are the regulatory functions of metals such as magnesium (Mg^{2+}), iron (Fe^{3+}), zinc (Zn^{2+}), copper (Cu^{2+}), and manganese (Mn^{2+}). Their concentrations are tightly regulated in the cell and they are known to be essential cofactors for essential cellular processes, such as adenosine triphosphate (ATP) utilization (Mg^{2+} ¹²⁶), ATP production in the mitochondria (Fe^{3+} ¹²⁷), detoxification (Cu^{2+} , Zn^{2+} , and Mn^{2+} ¹²⁸), and transcriptional regulation (Zn^{2+} ¹²⁹). In fact, presence of exogenous ions, such as lithium or lead, can have significant effects on development and cell physiology due to competitive inhibition of enzymes through replacements of their endogenous ionic cofactors. Despite their clear importance, study of these ions is complicated by difficulties in detection and lack of specificity for any eventual clinical applications. Although little is known about the regulation of these ions, a number of successful attempts at modulating intracellular ion concentrations to achieve cell fate effects by varying cell media compositions have been made^{130,131}, reflecting that the role of ions in cellular physiology is becoming increasingly appreciated.

1.4.1 Ion channels in proliferation and differentiation

Ion channel activity has been suggested to regulate proliferation already in the 1950's, when it was observed that non-proliferative cells had lower resting membrane potential than proliferative cells¹³² and membrane potential varied during cell cycle progression¹³³. This was further supported by the findings of Cone that hyperpolarization of dividing cells induced arrest¹³⁴, while depolarization of even post-mitotic neurons induced their proliferation¹³⁵. Since then ion channels, in particular K^+ channels, have been linked to cell cycle progression, both due to their cell-cycle phase-specific expression and function^{136,137}, and the ability to manipulate proliferation via ion channel inhibition¹³⁸⁻¹⁴⁰. Ca^{2+} channels¹⁴¹ and chloride channels¹⁴² have also been implicated in cell cycle progression in multiple cell types, as has the Na^+ - K^+ ATPase, which is proposed to be responsible for G1 hyperpolarization¹⁴³.

Ion channel control of stem cell proliferation has also been shown. While it may not be as surprising for NSPCs, which although not electrically active mature to be electrically active cells, it is even the case for ESCs. Regulation of ESC proliferation by ligand-gated ion

channels has previously been reported¹⁴⁴, with activation of the GABA_A receptor carrying a chloride current resulting in accumulation of cells in the S phase. Similarly, it induced quiescence in NSPCs¹⁴⁵. Block of L- and T-type Ca²⁺ channels was shown to attenuate ESC proliferation, with T-type channel block resulting in an accumulation of cells in G1 and G2/M cell cycle phases and differentiation¹⁴⁶. L-type channel block caused a reduction in BrdU incorporation and cell numbers¹⁴⁷, which is compatible with an induction of differentiation reported for these channels¹⁴⁸. Similarly, block of tetraethylammonium (TEA)-sensitive K⁺ channels induced an accumulation of ESCs in G0/G1 and compromised pluripotency¹⁴⁹, while block of the hyperpolarization-activated cyclic nucleotide-gated (HCN) K⁺ channel family resulted in S-phase accumulation of ESCs with uncharacterized effects on stem cell fate¹⁵⁰.

Recent publications have suggested that modulation of cellular electric properties can be an instructive factor in stem cell differentiation: optogenetic stimulation improved neural and neuronal differentiation of ESCs¹⁵¹, activation of Ca²⁺-activated potassium channels resulted in enrichment in cardiac pacemaker cells¹⁵², and block of L-type Ca²⁺ channels, known to have an instructive effects on NSPC proliferation and differentiation¹⁵³, was shown to restrict ESC differentiation to non-mesodermal lineages¹⁴⁸. Curiously, when L-type Ca²⁺ channels were blocked with a different compound, differentiation to cardiomyocytes, a mesodermal lineage, was improved¹⁵⁴, underlining the importance of further characterization of ion current effects on stem cell fate.

1.4.2 Lithium

Lithium is a metal that is already well established in the clinic. First introduced by John Cade for the treatment of manic patients in 1949¹⁵⁵, it has been a cornerstone of therapy and the target of extensive study, as deciphering its mechanism of action could be key to understanding psychosis. Its effects are however not limited to behaviour. Patients treated with lithium and responding to treatment were shown to have increased grey matter volume in the prefrontal cortex, which was not present in the non-responders¹⁵⁶. In line with this, lithium enhanced proliferation of hippocampal neural progenitor cells in the mouse both under homeostatic^{157,158} and radiation injury conditions³⁵. Outside the brain, lithium treatment is known to increase neutrophil and reticulocyte counts¹⁵⁹, and has been shown to affect osteoblastic differentiation *in vitro*¹⁶⁰. Interestingly, while reportedly only resulting in a mild increase in heart defects in humans^{161,162}, lithium can severely dysregulate embryonic

development, resulting in excessive endomesoderm formation in *Xenopus laevis*¹⁶³, and defective patterning in zebrafish and mouse embryos^{164,165}.

Despite many known effects and many proposed modes of action with experimental support, there is no consensus on which molecular pathway is responsible for mediating lithium action. The most discussed pathways are the inositol and GSK3 β pathways. Lithium is reported to inhibit inositol monophosphatase (IMPase) with an inhibitory constant (concentration achieving 50% inhibition, K_i) of 0.8mM¹⁶⁶ and inositol polyphosphate 1-phosphatase (IPPase) with $K_i=0.3\text{mM}$ ¹⁶⁷, two enzymes critical in inositol metabolism, by replacing their cofactor Mg^{2+} . This results in reduced availability of the second messenger inositol-1,4,5 triphosphate (IP3) which is a major mediator of extracellular signals and lies upstream of protein kinase C (PKC). Alternatively, lithium has been reported to inhibit the transcription of the sodium/myo-inositol cotransporter 1 (SMIT1), a Na^+ coupled inositol transporter^{168,169}, also resulting in lower inositol availability. The fact that levels of inositol *in vivo* are not affected at therapeutic lithium concentrations argues against this as the major pathway through which lithium acts. In the GSK3 β pathway, lithium inhibits GSK3 β , also by competing with magnesium¹⁷⁰, resulting in accumulation of β -catenin and dysregulation of WNT signalling, which is a key regulator of multiple cellular processes, including patterning during development and cell fate. Conflicting reports regarding behaviour of GSK3 β knockout mice and the fact that lithium-induced mouse embryogenesis defects were not mediated by WNT signalling argues against GSK3 β -WNT axis as the major effector of lithium action.

Some less commonly discussed targets of lithium have also been described, many in metabolic pathways. Bisphosphate 3'-nucleotidase 1 (BMPT1), an enzyme in the sulfate metabolism pathway which compromised protein synthesis when knocked out¹⁷¹, is inhibited by lithium with K_i of 0.153 mM¹⁷². In the glycolysis and glycogenesis pathway, fructose 1,6-bisphosphatase¹⁷³ has been suggested to be inhibited, while glucose 1,6-biphosphate synthase¹⁷¹ enzymatic activity was greatly slowed down by lithium. In line with this, earlier work showed that rate constants for reactions catalyzed by lithium-bound phosphoglucomutase was 4×10^8 times slower than when the enzyme is bound to its usual cofactor magnesium¹⁷⁴. In cell systems, treatment with lithium has been linked to inhibited glycogen synthesis in astrocytes¹⁷⁵, and long term lithium treatment *in vivo* resulted in reduced glycogen content in rat livers¹⁷⁶ and perturbed brain metabolite concentrations when examined over the 8 hours post injection¹⁷⁷.

1.4.3 Hyperpolarization-activated cyclic nucleotide-gated (HCN) family channels

Electrochemical changes on a cellular level are essential for cardiac pacemaking, muscle contraction, and neural processes. This is accomplished by the concerted action of gated ion channels, which taking advantage of the electrochemical gradients of some ion species existing between the cytoplasm and interstitial fluid, facilitate rapid and drastic changes in the electrical properties of cells. The heart is perhaps the best known example of this. Opening of voltage-gated sodium channels causes rapid Na^+ influx and cell depolarization, which deactivates the channels and activates other voltage gated channels sensitive to depolarization. Influx of Ca^{2+} and efflux of K^+ through these channels restore membrane potential to below activating voltage of the Na^+ channels that initiated the event, giving time for Na^+ - Ca^{2+} exchanger (NCX) and the Na^+ - K^+ ATPase to restore the ion balance. The current that then drives partial depolarization allowing the entire event to happen again, is the “funny” current (I_f , also known as I_h) carried by HCN4, one of the members of the HCN channels family¹⁷⁸.

Activation at hyperpolarized potentials is a unique feature of this channel family, which consists of four channel isoforms, activating at different voltages and differentially modulated by cyclic adenosine monophosphate (cAMP)¹⁷⁹. The channels are permeable to Na^+ and K^+ and to a much lesser degree to Ca^{2+} , but generally conduct inward Na^+ currents due to the electrochemical gradient of the ion. The four isoforms have different tissue distribution, with HCN1 and HCN2, which have the fastest kinetics, expressed in the CNS and the heart, while HCN3 and HCN4 are expressed in selected regions of the heart and CNS, as well as in other organs, such as the lung, liver, and muscle¹⁸⁰. Interestingly, the channels assemble into tetramers, which can be composed of multiple isoforms, giving rise to a wide range of conductance and activation properties. Channel function is also modulated by trafficking¹⁸¹⁻¹⁸³, auxiliary subunits^{184,185}, phosphatidylinositol 4,5-bisphosphate (PIP2)¹⁸⁶, and G-protein coupled receptors¹⁸⁷.

In addition to cardiac rhythm generation, HCN function is implicated in many neuronal functions, ranging from mood regulation to regulation of circadian pacemaking and learning¹⁸⁸. In non-excitatory cells, a pro-apoptotic effect of HCN2 has been described¹⁸⁹, and the channels have been reported to regulate the proliferation of ESCs¹⁵⁰. The channels can be blocked by cesium chloride (CsCl), and due to significant interest in modulation of HCNs in epilepsy and cardiac arrhythmias, a number of small molecule inhibitors have been developed. ZD7288 is one of the most specific and commonly used HCN inhibitors^{190,191},

although this has recently been put into question¹⁹². An elegant way of targeting the brain while avoiding cardiac effects is being explored by developing drugs targeting the interaction of HCNs with their auxiliary subunit TRIP8b¹⁹³ which is not expressed in the heart.

2 AIMS

The aim of this thesis was to investigate the action of ionic modulators on stem cell state.

More specifically, the following three sub-aims were defined:

- I. For lithium chloride, an ionic modulator with well-investigated non-ionic downstream effector pathways, to investigate the proliferative and anti- apoptotic effects on *in vitro* cultured juvenile mouse neural stem progenitor cells in the context of ionizing radiation.
- II. For ZD7288, an inhibitor of the hyperpolarization-activated cyclic nucleotide-gated (HCN) channel family, to investigate the effects on stem cell state in a well-defined, robust, *in vitro* stem cell system (mouse embryonic stem cells).
- III. To investigate effects of HCN inhibition on the proliferative state of mouse neural stem progenitor cells and identify potential downstream effector pathways underlying cell state effects.

3 MATERIALS AND METHODS

Please refer to the materials and methods sections of the included papers for detailed descriptions of experimental procedures, techniques, and reagents used.

4 RESULTS AND DISCUSSION

4.1 PAPER I: LITHIUM INCREASES PROLIFERATION OF HIPPOCAMPAL NEURAL STEM PROGENITOR CELLS AND RESCUES IRRADIATION-INDUCED CELL CYCLE ARREST *IN VITRO*

Cancer therapy involving irradiation of brain tissue in juvenile patients is associated with reduced cognitive performance later in life. One of the mechanisms proposed to mediate this effect is irreparable damage to the neural stem progenitor cells (NSPCs) in the developing brain, which are then not able to sustain proper development. Lithium chloride (LiCl), an antipsychotic drug with a long history of use in the clinic has been suggested to be a promising candidate for neuroprotection. This paper investigates the effects of lithium on mouse NSPCs *in vitro*, focusing on the cell physiology and outcomes following drug treatment and ionizing radiation (IR) exposure. To reflect the clinically relevant population, juvenile (post-natal day 8) mice were used for NSPC preparation. The SGZ, which is the site responsible for cognitive development in humans and likely affected in cancer patients, was chosen as the anatomical site from which NSPCs were isolated.

4.1.1 Lithium has a concentration-dependent effect on NSPC proliferation

To characterise the effects of LiCl on NSPCs, cells were adapted to *in vitro* culture by neurosphere formation for 4 days, following which they were dissociated to a single cell suspension and exposed to 0, 1, or 3 mM LiCl in standard media.

Analysis of sphere size 24 and 48 hours following treatment initiation found increased sphere volume in 3 mM LiCl treated cultures as compared to control at both time points (increases of ≈ 3.7 fold and ≈ 3.2 fold, respectively), reflecting larger cell numbers. 1 mM LiCl treated cultures also showed a trend toward increased sphere volume. While this could be due to altered rates of cell death, levels of both apoptosis and necrosis were found to be similar in 3 mM LiCl treated and control cells. To investigate possible cell cycle effects, cell cycle distribution of treated and control cells was examined by flow cytometry analysis of DNA content at 12, 24, 48, 72, and 96 hours post treatment initiation. While there was no effect at 12 hours, the G1 fraction was reduced while the S fraction was increased in the cultures treated with 3 mM LiCl at all later time points. The G2/M fraction in LiCl treated cells was reduced compared to control at 24 hours, but increased at 48 hours following treatment. Similar trends were seen in 1 mM LiCl treated cultures. S-phase findings were confirmed by increased bromodeoxyuridin (BrdU) incorporation at 48 hours. Unaltered frequencies of cells

staining positive for glial (GFAP) and neuronal (MAP2) markers indicate that LiCl treatment was not associated with differentiation.

These findings support the previous reports that lithium increases proliferation of NSPCs, where GSK3 β inhibition causing activation of WNT signalling was implicated in the response. They are also in line with reports of lithium-induced proliferation in other cell types. The observed changes in the G1 fraction may reflect altered cell cycle dynamics and a shorter G1, activation of previously quiescent cells, or some combination of these two effects.

4.1.2 Lithium protects NSPCs from irradiation, recruits them into proliferation following IR-induced G1 arrest and reduces DNA damage, but does not affect apoptosis.

To investigate potential neuroprotective effects of LiCl, NSPCs grown *in vitro* were treated with 0, 1, or 3 mM LiCl for 12 hours, following which cultures were exposed to 0 or 3.5 Gy of IR and allowed to form spheres in the same medium. IR treated cells formed smaller spheres than sham irradiated cells, with effects of radiation clear at all time points analysed (up to 72 hours) and significantly more drastic in the cells that were not pre-treated with lithium. While sphere volume in non-treated irradiated cells was ≈ 7.5 fold smaller than in sham irradiated cells, the reduction was only ≈ 2 fold in the cells that were pre-treated with 3 mM LiCl. Similar effects were observed at 24 hours and with lower (1 mM) LiCl concentration.

BrdU incorporation rates indicated that proliferation after irradiation remained higher in the 3mM LiCl treated than untreated cells, although it was lower than in 3mM LiCl treated sham irradiated cells. Interestingly, the proliferative differences were more pronounced in the irradiated cultures due to very low proliferation in the non-LiCl treated group. In fact, irradiation induced a significant G1 and G2 accumulation in both pre-treated and non-treated culture at the expense of the S-phase fractions. LiCl-treated cells recovered quicker, with more cells entering S than in the control cultures already at 24 hours after irradiation. To test if reduced DNA damage could underlie the recovery, levels of γ -H2AX staining, a marker of DNA damage, were examined 30 min after irradiation in treated and control cells. 3 mM LiCl treated cells were found to have on average 13 % lower staining intensity, suggesting a more limited extent of DNA damage. Finally, effect of treatment on cell death was examined. Staining for Annexin-V, a marker of apoptosis, and quantifying the subG1 DNA fraction indicated that LiCl treatment did not affect apoptosis or necrotic cell death.

Although LiCl treatment did not prevent IR-induced cell death, as was suggested previously¹⁹⁴, it did reduce IR-induced DNA damage, or at least the extent of response. This did not abolish, but rather attenuated the G1 accumulation seen after irradiation in the LiCl pretreated cells, which returned to proliferation faster than the non-pretreated cells. It is possible that this is due to lower rates of DNA damage, faster DNA repair, or more worryingly, leaky DNA damage checkpoints. It is interesting that the G2/M fraction in the LiCl pretreated cells remained elevated for up to 72 hours following irradiation. A potential explanation of this could be differential activation of DNA damage pathways in cells at different stages of the cells cycle, or impaired clearing of γ -H2AX foci due to reduced protein kinase B (Akt) activation¹⁹⁵ following lithium effects on the inositol pathway.

In summary, we found that LiCl induced NSPC proliferation under homeostatic conditions, and attenuated IR-induced DNA damage response and G0/G1 stalling. Although treatment did not reduce levels of cell death, cell numbers in treated cultures were increased due to increased proliferation.

4.2 PAPER II: ZD7288, A BLOCKER OF THE HCN CHANNEL FAMILY, INCREASES DOUBLING TIME OF MOUSE EMBRYONIC STEM CELLS AND MODULATES DIFFERENTIATION OUTCOMES IN A CONTEXT-DEPENDENT MANNER

The unique cell cycle properties of pluripotent stem cells have been linked to their identity and postulated to be essential for pluripotency maintenance. ZD7288, an inhibitor of the HCN channel family, has previously been reported to modulate cell cycle progression in mouse embryonic stem cells (ESCs), however the effects of this compound and the cell cycle perturbation it caused on cell fate were not described. This paper investigates the cell cycle dynamics and cell fate of mouse ESCs treated with ZD7288.

4.2.1 ZD7288, an HCN channel blocker, reduced ESC proliferation by increasing doubling time due to extended G1 and S phases

ESCs cultured in maintenance media with the addition of ZD7288 were found to be significantly fewer after 4 days of culture, with a reduction in cell number of over 50% as compared to control culture conditions, accompanied by only a disproportionately small reduction in viability, in line with the previously described action of the compound¹⁵⁰. Similarly, cells plated at subclonal densities and cultured in the presence of the compound formed significantly smaller colonies (27 vs 44 cells per colony in treated and control cells,

respectively). Reduced cell numbers could be explained by fewer proliferating cells or longer cell cycle. G0 is usually absent from ESCs, but to ensure that all cells were in active cell cycle, cells were stained for KI-67, a marker of active proliferation. As expected, no differences were observed, suggesting that the cell cycle was longer in ZD7288 treated cells. Based on these cell number observations, doubling time for treated cultures was calculated to be ≈ 37 hours, as compared to ≈ 28 hours for control cells, in line with previous reports⁹⁹.

To examine what accounted for the increased cell cycle length, we performed DNA content analysis using flow cytometry. This showed a significant increase in the fraction of cell in S-phase (37.6 % in control, 41.8 % in treated) and a corresponding decrease in G2/M phase (28.3 % in control, 23.4 % in treated) in the treated cultures. Taken together with the calculated doubling time, this corresponds to an increase in the length of G1 and S phases in response to ZD7288 treatment. This was in contrast to what had previously been reported. Previous report by Lau *et al*¹⁵⁰ indicated an accumulation in S-phase with a compensatory decrease in the G1 fraction, indicating an increase in the length of S-phase only. Interestingly, the cell cycle length in the previous publication was much shorter than in the present one, possibly due to difference in culture conditions. Lau *et al*¹⁵⁰ used serum-based, whereas this study used serum-free media, and it has previously been reported that ESC doubling time varied between 13 and 34 hours depending on media formulation, with serum-free medias resulting in longer doubling times.

4.2.2 DNA replication is altered in ZD7288 treated cells

Surprisingly, the number of 5-ethynyl-2'-deoxyuridine (EdU) incorporating cells, a marker of cells in S-phase, appeared to be lower in treated cultures when examined using flow cytometry. This was puzzling because the number of S-phase cells was increased in treated cultures based on DNA content quantification, prompting us to take a closer look at the at EdU incorporation pattern. EdU staining pattern was distinctly different in the treated cells, which formed a single wide peak, instead of the two peaks corresponding to positive and negative cells usually seen, and observed in the control.

Potential causes of this alteration could be intra-S phase stalling, reduced DNA replication, or reduced availability of the EdU nucleotide. To investigate the possibility of intra-S-phase stalling, cells were staged as “early” or “late” S-phase based on EdU incorporation pattern following visualization using confocal microscopy. The fraction of “early” phase cells was similar in both conditions (55.7 % in control and 53.3 % in treated culture), suggesting altered EdU availability or slower DNA replication, which would be in line with the observed

prolonged S-phase transit time. It is possible that the two proposed scenarios are linked, with reduced nucleoside availability hampering replication. Alternatively, ion concentrations are known to affect DNA tertiary and quaternary structure, altered sodium or potassium concentrations due to HCN inhibition could induce structural changes attenuating replication speed. Finally, DNA replication speed varies during S-phase¹⁹⁶, and the observed effect may be related to modulation of a mechanism involved in DNA replication speed regulation.

4.2.3 Treatment with ZD7288 decreases colony formation frequency but does not alter expression of the core pluripotency genes and effects cell differentiation in a context dependent manner

To test the effects of cell cycle perturbation on pluripotency, cells were plated at subclonal density to test clonogenicity and self-renewal. The number of colonies formed per well was much lower in the treated condition, and more colonies had rough edges, which could be a sign of differentiation. To test this, cells were stained for the various markers of pluripotency (SSEA1, OCT3/4, SOX2, NANOG, and ALP activity), with no significant changes detected. Similarly, no changes in pluripotency marker expression were detected on an RNA level. To functionally test maintenance of pluripotency in ZD7288 treated cultures, treated cells were re-plated and both colony formation rates and colony morphology were restored to control levels. A possible explanation for the observed effect is altered attachment/cytoskeletal dynamics, which could affect both the propensity of the cells to attach and form colonies and cellular morphology on colony borders.

To functionally test the effects of ZD7288 on the other cardinal stem cell property – differentiation, ESCs were induced to differentiate either spontaneously in fetal bovine serum (FBS) –containing media with LIF withdrawal, or toward the neural lineage as per defined serum-free protocol. Spontaneous differentiation was potentiated by ZD7288 addition, with more pronounced downregulation of the pluripotency markers and upregulation of most lineage markers. Surprisingly, neural differentiation did not benefit from ZD7288 addition, with similar levels of expression of most pluripotency factors in control and treated cells, and a less efficient induction of 2 of the 3 ectoderm markers tested. These findings suggest that the action of ZD7288 in differentiation is context dependent. This is rather similar to the difference in cell cycle effects between the previous and current reports. Intriguingly, one of the most striking differences is also similar – serum-containing media. Although the underlying mechanisms of these differences are unclear, this work adds to the weight of

evidence that extending G1 does not necessarily result in differentiation, at least not if the G1-S ratio is maintained¹⁹⁷.

In summary, we found that ZD7288, an inhibitor of the HCN channel family, reduced ESC proliferation by extending G1 and S residence time. This did not compromise cell identity, with treated cell maintaining pluripotency and the ability to differentiate to all lineages. Differentiation in FBS-containing media was facilitated by the compound, while directed differentiation to neuroectoderm in defined media was less efficient in its presence.

4.3 PAPER III: HYPERPOLARIZATION-ACTIVATED CYCLIC NUCLEOTIDE-GATED CHANNELS MODULATE ACTIVE PROLIFERATION AND METABOLISM, AND MAINTAIN MOLECULAR CLOCK OSCILLATIONS IN ADULT MOUSE NEURAL STEM PROGENITOR CELLS

Data from recent sequencing studies attempting to define genes specific to the activated and quiescent stem cell states have indicated that a number of ion channels are differentially expressed in the two pools, with members of the HCN family on the list. Taken together with the established fact that voltage gated potassium channels are involved in cell cycle regulation, and data from other stem cells systems, including Paper II above, this work set out to investigate a potential role for HCN family channels in adult mouse NSPC proliferation.

4.3.1 HCN2 and 3 are expressed throughout the NSPC hierarchy with their function restricted to the proliferative (S and G2/M) phases

Gene expression analysis with qRT-PCR indicated that *Hcn2* and *Hcn3* were expressed in both bulk primary SVZ NSPCs, and in all populations of NSPCs sorted according to GFAP::GFP intensity, believed to correspond to different levels in the NSPC differentiation hierarchy, including type B and C cells. This was promising, as HCN3 was the isoform previously reported to modulate ESC proliferation, confirmed in **paper II**, and was identified as part of the quiescent stem cell signature in data from Codega *et al*¹⁹⁸. While expression of HCN3 was not confirmed on a protein level due to technical limitations, HCN2 co-stained BrdU+/SOX2+ NSPCs in tissue sections from the SVZ and SGZ. To further verify channel expression, function was assessed with electrophysiology. The typical hyperpolarization-activated current (I_h), sensitive to the specific blocker ZD7288, could be recorded from some, but not all NSPCs.

To investigate the basis for non-uniform presence of I_h , cells were cell-cycle sorted based on DNA content. This showed that while no cells in the G0/G1 cell cycle fraction carried the

current, the current was present in many S, and the majority of G2/M cells. Correspondingly, HCN2 staining intensity was found to be higher in those cell cycle phases as ascertained by flow cytometry, and differential channel localization was observed in the three cell cycle phases by confocal microscopy. The association of HCN function with the proliferative phases further supported the hypothesis that the channel could be implicated in the regulation of proliferation.

4.3.2 Inhibition of HCN channel activity attenuated NSPC proliferation due to a reversible G0/G1 block *in vitro* and its effects extend *in vivo*

Culture of primary SVZ NSPCs in the presence of ZD7288, an HCN channel inhibitor, resulted in smaller neurosphere size, while block of HCN function with different concentrations of its inhibitors ZD7288, Zatebradine, or CsCl resulted in lower SVZ NSPC number as determined by a viability assay. To ensure the difference in cell numbers was not due to cell death, as HCN has been reported to regulate apoptosis¹⁸⁹, levels of activated caspase-3 and the subG1 fraction of cell cycle were quantified. No significant induction of cell death was observed. Moreover, EdU incorporation was significantly reduced (by 46%) in treated cells 24 hours after treatment initiation. The proliferative effects were confirmed on a genetic level with si- and shRNA. Rather conflictingly, siRNA showed that HCN2 but not HCN3 knockdown reduced cell numbers, while shRNA, showed that knockdown of either gene was sufficient to see a reduction in proliferating cells. This could be due to different potency of knockdown, but regardless, confirmed the expectation that HCN activity is involved in regulation of NSPC proliferation.

To further investigate the effect of HCN inhibition on the cell cycle, DNA content analysis of treated and untreated cells was performed, indicating an increase in G0/G1 cell cycle fraction accompanied by a reduction in the S and G2/M. To further identify whether it was G0 or G1 that was affected, cells were stained for KI-67, which showed an increase in cells no longer in active proliferation, as indicated by loss of KI-67 marker expression. This was further supported by increased survival of ZD7288 treated NSPCs upon long term (48 hours) exposure to the genotoxic agent arabinofuranosyl cytidine (Ara-C), and the full reversibility of the induced proliferative block upon washout. The reversibility of the proliferative effect distinguished the observed block as quiescence, as opposed to senescence.

Importantly, similar effects on proliferation (reduction in proliferation by BrdU incorporation, phospho-histone 3 (a marker of M-phase) expression, and KI-67 expression) were seen *in vivo* in both SVZ and SGZ when animals were injected with 4mg/kg of ZD7288.

This was not accompanied by induction of apoptosis, with onset of action observed at 24 hours and complete attenuation of the effect at 72 hours, further supporting reversibility.

4.3.3 Transcriptome analysis identified alterations in expression of cell cycle regulators, metabolic enzymes, and members of the molecular clock in treated cells, while NSPC identify marker expression was not altered

Transcriptome analysis confirmed that treated cells were not undergoing differentiation, as expression of key NSPC regulatory genes *Sox2*, *Pax6* and *Msi1* was similar in treated and untreated cells, and expression of differentiation genes was not induced by treatment.

Markers of quiescence and activation, on the other hand, were altered. *Nestin* and *Ascl1*, both implicated in activation and stem cell proliferation, were downregulated on an RNA level, while western blot for EGFR indicated that the receptor was downregulated on a protein level. Inhibition of proliferation was further confirmed by reduced expression of many positive cell cycle regulators in treated cells, while some negative regulators of proliferation were upregulated. p21 and p27 canonical inducers of cell cycle arrest, were upregulated on an RNA, but not protein level, suggesting a novel mechanism for induction of quiescence in response to HCN inhibition.

Gene Ontology (GO) term enrichment analysis of differentially expressed genes identified metabolism as one of the major gene categories altered in treated cells. Enzymes involved in glycolysis were strongly downregulated. To complement this analysis and look at metabolism more functionally, we analyzed metabolite abundance. Levels of glycolysis intermediates, lactate, and amino acids were found to be significantly altered, confirming a metabolic perturbation as either the mediator or a consequence of the effects of HCN inhibition.

When a more narrow approach was taken GO enrichment analysis was run on genes shared between the quiescence signature published by Codega *et al*¹⁹⁸ and those altered in response to ZD7288. “Rhythmic processes”, containing many genes known to be components of the molecular clock network (*Per1*, *Per2*, *Cry1*, *Cry2*, *Nr1d1* and *Nr1d2* (Rev-Erb- α and β), *Timeless* and *Tipin*), was the most significantly enriched term. To test functional significance of the observed change, luminescence from Per2-driven luciferase was recorded in adherent NSPCs and SVZ explants with and without ZD7288 in the media. In line with observations of deregulation on the transcriptional level, clock oscillations were abolished upon HCN inhibition. In light of the previous reports of clock function regulating proliferation in other stem cell systems^{199,200}, we tested whether ZD7288 effect could be phenocopied by activation

of Rev-Erb- α , the most upregulated component of the clock network identified in our dataset. Treatment of NSPCs with the Rev-Erb- α agonist SR9009 resulted in reduced proliferation, suggesting its involvement in mediating the effect of HCN inhibition. Interestingly, hippocampus hyperproliferation was reported in Rev-Erb- α knockout animals²⁰¹, supporting its function as a regulator of proliferation in NSPCs. It has also been reported to be inhibited by lithium²⁰², representing an alternative pathway for proliferative increase seen in **paper I**. Although it is not clear if or how the observed changes in metabolism and the molecular clock are related, recent publications have reported metabolic perturbations in the kidneys of clock-null animals²⁰³.

4.3.4 Pre-treatment of juvenile mice with ZD7288 and a neuroprotective effect, reducing the deleterious effects of ionizing radiation.

Following up on work in **paper I**, neuroprotective potential of HCN blockade was tested in an animal model of ionizing radiation (IR)-induced NSPC damage. Juvenile animals (post-natal day 8) were treated with ZD7288 (8 mg/kg) 24 hours prior to exposure to 2Gy of IR. Animals were sacrificed 6 hours following exposure and levels of proliferation and cell death were examined in the neurogenic areas (SGZ and SVZ). The levels of apoptosis as seen by activated caspase-3 staining were significantly reduced, and proliferation as measured by BrdU incorporation was similar in treated and untreated animals. No cell cycle analysis was performed, as in **paper I**, to assess potential additional stalling of cells in G0/G1, but similar proliferation rates in both irradiated and non-irradiated animals suggest that proliferation was not affected by IR, possibly due to the fact that the most sensitive cells were in proliferative arrest due to ZD7288 treatment, and thus protected from IR damage and the apoptosis that would have followed. An alternative explanation of HCN inhibitor action could be related to calcium influx-induced apoptosis, in which HCN2 was previously implicated.

To test whether there may be a therapeutic window with HCN inhibition, and a similar protective effect would not be observed for cancer cells, 17 patient-derived cancer stem cell lines were tested for growth response following culture in media containing 0-49 μ M ZD2788. Across all lines there was no change in viability signal to any of the tested inhibitor concentrations, suggesting that HCN has no proliferative effect in cancer stem cells. As the cell line tested were of human origin, while all other work had been done on murine cells, to ensure the observed lack of effect was not due to species differences, proliferation kinetics in response to ZD7288 treatment were tested in human fetal neural stem cells (hfNSCs) and human glioma-initiating cancer stem cells (giCSCs). Confirming the viability test results, the

number of EdU positive giCSCs did not change in response to ZD7288, while proliferation of hfNSCs was inhibited resulting in fewer EdU incorporating cells, as observed earlier in mouse NSPCs. The differential response of cancer and tissue stem cells to HCN inhibition could reflect abnormal cell cycle regulation in cancer cells²⁰⁴, but also disrupted clock function, which has been reported previously²⁰⁵. Finally, alterations in metabolism, a hallmark of cancer²⁰⁶, is also a possible explanation of observed differences.

In summary, HCN2 and HCN3 are expressed in mouse NSPCs with function restricted to the S and G2/M phases. Their inhibition inhibited proliferation inducing a reversible G0/G1 accumulation accompanied by changes in markers of quiescence/activation and cell cycle, but not stem cell identity. This was accompanied by downregulation of glycolysis and changes in metabolite levels, as well as abolition of molecular clock oscillations due to upregulation of negative clock components. An agonist of the peripheral clock component Rev-erb- α could mimic the proliferative effects of HCN inhibition. This effect could be confirmed in both proliferative regions *in vivo*, and resulted in reduced cell death in juvenile animals pre-treated with ZD7288 prior to IR exposure.

5 CONCLUSION AND FUTURE DIRECTIONS

The work in this thesis showed that:

- I. Lithium induces proliferation of juvenile mouse NSPCs *in vitro*, and although it does not reduce apoptosis, it protects them from IR-induced DNA damage, and induces their return to proliferation following G1 block
- II. Treatment of mouse embryonic stem cells with the HCN blocker ZD7288 attenuates their proliferation by extending their cell cycle, specifically in the G1 and S phases. Treatment does not compromise pluripotency under maintenance conditions, but can facilitate differentiation in a context-dependent manner.
- III. HCN2 and HCN3 are expressed in NSPCs and functional in the S and G2/M phases. Block of their activity reduces proliferation *in vivo* and *in vitro* by inducing a reversible cell cycle block in the G0/G1 phase, which exhibits many features of quiescence: reducing glycolytic metabolism, downregulating markers of activation, and maintenance of NSPC identity. I_h current block also disrupts oscillations of the molecular clock, with modulation of the clock component Rev-erb- α having similar effects on proliferation.
- IV. HCN activity can protect NSPCs from genotoxic stress both *in vitro* and *in vivo*, and attenuates IR-induced apoptosis *in vivo*. HCN inhibition may have the potential to be developed into a neuroprotective therapy for pediatric cancer patients.

Additional work exploring the mechanisms behind these findings and investigating their potential translation to the clinic is needed. The work presented here makes a case for modulation of ion-dependent processes as an in-point for cell state regulation. Modulation of ion availability as a co-factor (in the case of lithium effects) or ion trafficking (in the case of HCN inhibition) have both been shown to be viable strategies for regulation, although the downstream mechanisms and exact targets of each intervention remain to be elucidated. An interesting area of overlap is metabolism, which is regulated in a circadian manner both on a whole-organism and on the cell level²⁰⁷, and is known to regulate stem cell dynamics, both in adult stem cell systems^{208,209} and in the embryo through diapause and reprogramming^{210,211}. It is however unclear how the metabolic and circadian regulatory systems relate to each other and proliferation – are these parallel processes that can exert proliferative control? Is one process subservient to the other? Are perturbations in one system just a side effect of the

deregulation of the other? The effect observed in ESCs in **paper II**, if mediated in a similar fashion to the NSPC effect in **paper III**, would put metabolism at the top of the hierarchy, and circadian perturbations as a side effect, because ESCs lack a circadian clock²¹². It is unclear how well this would generalize to somatic cells, as ESCs also lack conventional cell cycle control. To more appropriately address this question, it would be interesting to test how the effects of HCN inhibition observed in **paper III** would change in BMAL1-deficient mice, which lack circadian rhythmicity²¹³, but not the usual somatic cell cycle machinery. To more closely examine the interaction of HCN-inhibition, metabolism, and proliferation, it would be interesting to follow up the work in **paper II** on proliferative slow down of ESCs with further experiments looking at metabolic effects in that cell system. ESCs are a relatively homogenous and robust cell system, while still relevant in a developmental context, making them a good platform for research into fundamentals of cellular physiology.

This thesis also investigated two very different approaches to neuroprotection – temporary attenuation of proliferation (**paper III**) and a combination of DNA damage protection and induction of proliferation to overcome cell loss due to IR treatment (**paper I**). In addition to developing compounds and techniques for more precise delivery of active compounds to injury site, this work could be followed-up by investigations into how these two interventions may be combined to improve outcomes. Pre-treatment with an HCN inhibitor in combination with acute or follow-up treatment with lithium may be a worthwhile approach to neuroprotection, taking advantage of both strategies to overcome IR-induced damage. Since significant additional work would be needed to characterize such a regiment, lithium is likely a more promising first approach for clinical tests of neuroprotection due to its fairly well understood toxicity profile. However, long-term implications of lithium use should first be more thoroughly investigated. Deregulation of proliferation can lead to tumorigenicity, especially in combination with genotoxicity and stress, which NSPCs face following IR treatment. Ideally, development of therapy should be guided by mechanistic insights into the mode of action of both lithium and HCN inhibition, until then, the empirical approach should not be overlooked.

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