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# Learning and stroke recovery: parallelism of biological substrates

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## Abstract

Stroke is a debilitating disease. Current effective therapies for stroke recovery are limited to neurorehabilitation. Most stroke recovery occurs in a limited and early time window. Many of the mechanisms of spontaneous recovery after stroke parallel mechanisms of normal learning and memory. While various efforts are in place to identify potential drug targets, an emerging approach is to understand biological correlates between learning and stroke recovery. This review assesses parallels between biological changes at the molecular, structural and functional levels during learning and recovery after stroke, with a focus on drug and cellular targets for therapeutics.

### Keywords

Stroke; learning and memory; neurorehabilitation; dendritic spines; axonal sprouting; gene systems; cortical maps; functional motor recovery

## Introduction

Stroke causes lasting neurological impairments, making it the leading cause of adult disability. With advancements in early interventions, such as with recanalization and thrombolysis, the death toll resulting from a stroke has declined by 16.7 percent[1] from 2006 to 2016. While these efforts increase survival, the proportion of stroke survivors with motor and cognitive impairments is on the rise. Currently, neurorehabilitation is the only therapy for stroke recovery. Neurorehabilitative therapies encompass repetitive training[2], specific muscle training[3,4], constraint-induced movement[5], robot-assisted gait training[6] and other forms of reproducible behavioral activity, such as with virtual reality modalities[7,8]. The idea is that practice makes perfect. Repetitive and task-specific training have been hypothesized to induce forms of plasticity that are beneficial for stroke recovery. However, our understanding of the biological substrates that effect processes beneficial to recovery, or what these processes might be, are limited. Imaging studies in patients are limited to changes in gross anatomical features or macroscopic changes in functional connectivity across brain regions. While these outcomes can be loosely correlated to recovery, relevant

changes at the synapse, neuronal circuit, or in gene expression underlying recovery remain elusive. Pre-clinical models, such as rodent models of stroke, have furthered our understanding of these changes, allowing us to dissect mechanisms that lead to motor recovery after stroke. This review addresses biological substrates that have been defined, with an emphasis on parallels between mechanisms underlying normal learning processes in the brain and those of spontaneous biological recovery after stroke.

## Are learning-induced changes recapitulated during motor recovery?

The brain undergoes a remarkable degree of plasticity in an age and environmentally dependent manner. Both learning and stroke are marked by temporal periods of heightened plasticity. For learning, this period is present during development and shaped by sensory processing, and has been shown to be critical for language acquisition and development of other cognitive domains[9,10]. The most sensitive period of the brain to changes in circuitry and physiology from alterations in outside (sensory) input has been termed the critical period[9,10]. Following stroke, a critical period exists in the subacute phase of stroke in which the most significant change occurs in behavioral recovery and brain circuitry and physiology. This has been demonstrated with the use of neuromodulatory drugs and activity-based therapies[11–13].

Evidence from neurorehabilitative training and associative motor recovery suggests that learning processes may shape plasticity responses to output functional motor recovery. In other words, a learning process introduced after stroke, or by the stroke itself, gives rise to functional changes normally associated with learning, which act as substrates for motor recovery after stroke. While a direct causal link between learning-induced biological changes and recovery after stroke is yet to be experimentally determined, several lines of evidence show that biological changes within neuronal populations during learning and recovery after stroke share similar structural changes and gene transcriptional programs[14,15]. For example, induction of learning-induced biological changes, such as a genetic modulation underlying learning, may be evident during motor recovery after stroke. This review defines such processes that are active during learning as well as recovery after stroke, and hence identifies mechanisms that underlie motor recovery after stroke. Most descriptions on learning in this review pertain to learning of a motor skill, a modality most relevant to neurorehabilitation.

## Dendritic spine plasticity during learning and stroke recovery

Cortical plasticity, first described as instability in response to cortical stimulation[16], forms the basis of adaptation of the cortex to changes in the environment. Learning is an integral part of adapting to change. Although many studies have shown changes in cortical plasticity during learning, more recent studies have shown the involvement of the motor cortex in storing information that pertains to learned motor movements. A biological readout of this plasticity-related process has been through the quantification of sub-micron volume protrusions on dendrites that carry chemical information for synapses. These protrusions, termed dendritic spines, exhibit changes in numbers through formation of new spines or deletion of old spines, and can cluster on dendrites during learning processes (Figure 1).

The concept that the cortex acts as a seat of learning stemmed from observations on increased dendritic spine density in the visual cortex after animals were exposed to enriched conditions[17]. The plasticity of the cortex during skilled movement such as a unilateral reaching task was first demonstrated in the 1970's, where training of one limb led to structural changes in cortical layer 5 dendrites that were otherwise absent in the hemisphere contralateral to the trained limb[18]. Cortical layer V is the output layer of motor cortex to the spinal cord and brain stem movement centers. Since then, with the introduction of better imaging technology, the use of a similar training task has led to in-depth characterization of changes in synapse number, strength and structural changes during motor training. For example, repetitive motor training has been associated with the formation of new dendritic spines on layer 5 dendrites during the acquisition phase of skill learning[19]. Similarly, clustering of dendritic spines with repeated learning is indicative of strengthened synapses associated with the task[19], while destabilization or deletion of spines leads to a loss of the motor memory associated with the task[20,21]. The direct link between the association of newly formed spines with a new motor skill was recently demonstrated with an elegant use of tools where selective optogenetic silencing of task-associated dendritic spines led to a complete erasure of motor memory connected with the task[22]. Spatial characterization of spine dynamics has shown that task-associated spines are compartmentalized to specific dendritic branches[22]. These studies highlight the complexity with which motor learning affects neuronal circuits to the level of synapses on specific branches of neurons, presumably those directly involved in the task. Compartment-specific spine changes have also been observed in layers 2/3 where spines gained in superficial dendrites are lost as training ceases but after the skill has been gained, whereas dendrites in deeper layers retain spines gained during learning even in the absence of training[23]. This leads to the question of whether the disruption of neuronal circuits dedicated towards a specific motor task such as during a stroke can be remodeled onto different branches in adjacent spared tissue.

Synaptic dysfunction and spine loss are characteristic of neurodegenerative diseases. Paradoxically, spared tissue adjacent to the infarct in stroke shows a remarkable degree of spine plasticity. For example, dendritic branches from layer 5 neurons in peri-infarct cortex show increased spine formation two weeks after stroke, following the initial loss[24,25]. This increased plasticity has been partly attributed to increased blood flow, suggesting that a maintenance of tissue perfusion can augment spine formation or retention of spines during tissue stress. Despite the increased turnover of spines in peri-infarct cortex, this amount of plasticity is seemingly insufficient to produce motor recovery. Whether these responses are maladaptive or whether newly formed spines lack sufficient synaptic activation remains an area of further investigation. However, unlike the formation of new spines, preservation of dendritic spines within the first week after stroke is associated with early motor recovery[15]. The exact cellular mechanisms that lead to spine preservation and subsequent motor recovery are unknown, but presumably preserved spines could be better positioned to receive synaptic input from existing or new synaptic partners.

Similar to learning-induced changes in dendritic plasticity, rehabilitative training after brain injury forces numerous changes in dendritic branching and spine density in the contralesional hemisphere[26] and in motor and premotor areas on the same side of the stroke[27]. Dendritic complexity and increased spine density occur in task-specific neurons

in peri-lesion tissue following skilled rehabilitative training[28]. This broadens the possibility that spontaneous dendritic changes in peri-infarct neurons can be functionally stabilized through repetitive training. A similar output can be produced through frequent intermittent activation of synapses, such as through optogenetic stimulation[29]. For example, through the use of combinatorial tools to asses stimulation-induced synaptic function in peri-infarct cortex, optogenetic stimulation of thalamocortical axons projecting to peri-infarct somatosensory cortex causes formation of new and stable axonal boutons[30]. Importantly, these newly formed axonal boutons are responsive to forelimb-evoked calcium transients —indicating that the new synapses stabilized by this activity are functional. These data have been some of the first to directly link optogenetic stimulation with the formation of functional spines during stroke recovery. Given our limited understanding on how synaptic activation of spines through rehabilitation or optogenetic stimulation can augment stroke recovery, the identification of molecular effectors that drive similar changes opens a window for therapeutic potential.

Taken together, learning forces numerous changes in dendritic spines that are causally linked to a learned motor task. Stroke induces a transient loss of spines, whereas neurorehabilitative training increases spine density (Figure1). Changes in spine plasticity are associated with reorganization of molecular pathways. Perturbation of molecular pathways involved in learning and memory influences spine density after stroke. For example, downregulation of a learning and memory receptor, CCR5 (discussed later) prevents initial spine loss after stroke[15]. This points to the fact that molecular effectors of learning and synaptic plasticity prevent loss of spines and are associated with motor recovery.

## Molecular substrates that drive learning and stroke recovery

Both stroke and learning-induced gene expression changes are associated with brain excitability[31], synaptogenesis[23,32,33] and network connectivity[34] (Figure 2). Over the decades, many studies have shed light on molecular pathways that underlie memory formation and consolidation[35]. GABA and AMPA receptor trafficking play key roles in modulation of cortical excitability and synaptic strength during memory formation both in development and adulthood[36–39]. Dampening GABA signaling outside the synapse (extrasynaptic GABA signaling), by targeting the alpha5 subunit of the GABA receptor, which mediates inhibition in learning and memory process[39], promotes motor recovery after stroke[40]. Similarly, increasing AMPA receptor signaling also promotes motor recovery[11] (for a detailed review, see reference 31). Synaptogenic mechanisms are critical for memory formation, such as axon guidance molecules that mediate neuronal contact and synaptogenesis[41]. Axon guidance molecules are active during development, where signaling by binding of an axonal guidance receptor on the surface of a cell to its extracellular ligand meditates the migration of the cell or the axon across a region that is enriched in expression of the ligand. Neurons and glia express axon guidance molecules during development and adulthood[42,43]. In the adult, axon guidance molecules have roles in neurotransmission[44]. For example, an axon guidance cell-surface receptor-Eph and ligand Ephrin signal in astrocytes and neurons to regulate activity-dependent release of glutamate and subsequent activation of AMPA receptors during LTP and memory storage[44]. These molecules are also differentially expressed after injury such as stroke[45]. Ephrin-mediated neuro-glial signaling is active after

stroke. Ephrin A5 is highly upregulated in astrocytes following a stroke, where it interacts with neuronal EphA4. Blocking EphA4 in peri-infarct cortex during periods of reduced cortical excitability leads to functional motor recovery and robust axonal sprouting[45] (for a detailed review, see reference 46). A recent finding using a knockout model in the mice of Ephrin A5 reports that this molecule may not lead to functional recovery[47]. The report by de Boer et al uses a knockout model in mice of Ephrin A5 expression and reports no enhancement of functional recovery. The methods of determination of recovery and the data for a lack of effect are not shown. There is an important and critical limitation to this study, and to this overall approach, in the stroke field. Knockdown of a molecule for the entire brain development and life of a mouse, termed "constitutive knockout", produces compensation in other molecular systems, which assume the role of the knocked out molecular system[48,49]. Indeed, constitutive knockout for the life of an animal of one axonal growth inhibitor, leads to even compensatory upregulation of other axonal growth inhibitors[50]. In order to prove that a molecular system has, or does not have, a role in a disease like stroke, the brain cannot have the system knocked out for entire life of the animal (and its brain development), but must have the system knocked out at a discrete interval after the stroke.

In these and other studies of functional recovery after stroke, a key issue is alignment between the outcome measures for recovery in the animal model, often the mouse, to those in the human and the establishment of thresholds for meaningful functional gains. Humans have impairments in neurological function as a result of a stroke, which of course affect specific behavioral domains, such as motor, language, attention and sensory domains. In humans, behavioral measures of stroke recovery that do not test impairment, but instead rely on more global measures of disability, are not sensitive to stroke recovery[51,52]. This limitation in clinical trials with disability outcome measures stems directly from the non-specific—non-domain (motor, language, sensory, etc.)--measures of disability, such as the modified Rankin Scale or the Barthel Index. In mouse models of stroke, behavioral recovery measures that similarly aggregate many functional domains into global scales either do not detect recovery, or over-detect recovery. One such lumped test is the modified Neurological Severity Scale (mNSS), which finds positive behavioral recovery results in nearly every test of a candidate therapy in stroke[53]. Similarly, just as in human outcome measures in stroke, some commonly used mouse behavioral measures are affected by many different brain functions and brain areas. For example, the Rotorod test, which measures how long a mouse or rat can stay atop a moving, motorized rod, is actually sensitive to spinocerebellar tract damage and repair[54], and yet is commonly applied to outcome measures of stroke in the cortex or striatum. Thus, a key issue in this field relates to the use of adequate and aligned behavioral outcome measures in both human and rodent in stroke, which do not test stroke impairments at the general level of disability and which do not lump many functional domains of the nervous system into one common measure. In the studies described in for CREB and CCR5 (below), outcome measures in the mouse are foot placement in gait and individuated use of the forelimb in exploratory behaviors, designed to follow common human behaviors and to narrowly focus on motor control impairments. In these studies, the level of improvement in functional recovery is above the meaningful clinically important

difference established in human motor recovery in stroke[55,56], which can be extrapolated from the human to the mouse using effect size.

Returning to molecular mechanisms of stroke recovery in systems involved in the normal synaptic plasticity of learning and memory, recent evidence suggests that biological changes during learning processes and stroke recovery share common molecular pathways outside of the realm of immediate neurotransmitter receptor signaling of glutamate via AMPA receptors or GABA via extrasynaptic GABA receptors. Some examples include neurotrophin signaling via BDNF[57], immediate early genes such as Arc[58] and growth inhibitors such as Nogo[59]. While these act as striking examples, our review will focus on the two most recently characterized molecules that have robust effects in both systems-- learning and motor recovery.

### CREB

The nuclear transcription factor cAMP response element binding protein (CREB) is a key modulator of learning and memory processes and has been investigated in different model systems that range from aplysia to rodents[60,61]. CREB is involved in memory formation, storage and retrieval[60]. At a finer level, CREB expression determines which neurons integrate into a network that stores information pertaining to an event[62,63]. CREB also regulates learning-induced structural plasticity, such as formation of new spines potentiated through learning[64]. It is important to note that CREB is involved in learning-dependent structural changes but does not otherwise alter the normal structure of the CNS. In terms of function, CREB downregulation impairs long-term potentiation (LTP)[64]. The dependency on CREB for learning-induced structural changes indicates that CREB induces a plasticity state in neurons.

Based on its role as a 'plasticity molecule,' several studies have shown that CREB upregulation promotes axonal regeneration *in vitro* as well as in *in vivo* models of CNS injury[65,66]. In the context of stroke, CREB upregulation in peri-infarct cortex has substantial effects on motor recovery[14]. In tasks that test fine motor control of the forelimb, animals with CREB overexpression show enhanced performance such as reduced foot faults, less bias towards use of the impaired limb and improved fine motor control for handling and grabbing, one month after a stroke. Overexpression of CREB through viral transduction of a small population of neurons adjacent to the infarct is sufficient to induce recovery, following a stroke to either the motor cortex or in a larger stroke model inclusive of the motor cortex and striatum. Interestingly, silencing CREB-overexpressing neurons with chemogenetic tools, such as with an inhibitory DREADD (hM4Di)[67], reverses its effects on recovery, whereas lifting inhibition in CREB-induced neurons reinstates recovery. In other words, CREB induction drives motor recovery, whereas silencing neurons that selectively upregulate CREB silences the effects on motor recovery. A possibility is that CREB activation within neurons attributes a property of functional dominance, wherein a motor network becomes dependent or is driven by pools of neurons with higher levels of CREB expression. This attribute of functional dominance is present even in the absence of stroke. In healthy animals, neurons with CREB overexpressing neurons in motor cortex, when silenced, show motor deficits similar to animals that have received a stroke to the

motor cortex[14]. A decline in motor performance suggests a selective integration of CREB-overexpressing neurons to a motor network and that this feature is based on a state of increased excitability.

Motor recovery resulting from CREB overexpression after stroke results in widespread gene expression changes[14]. In healthy animals, CREB overexpression alone causes differential regulation of approximately 200 genes. Being a ubiquitous transcription factor, CREB affects various signaling pathways, including pathways distinct to post-stroke signaling. Prominently, these include fibroblast growth factor (FGF) and suppressor of cytokine signalling-2 (SOCS2) signaling, known to regulate neuronal development and homeostasis,[68,69] and have been shown to influence neural repair and axon regeneration. In addition to stroke, a large body of evidence supports a pro-growth role for FGF in other systems of injury[68–72]. Interestingly, both SOCS2 and FGF have been implicated in signaling during neurogenesis, a process that is upregulated after stroke and can cause functional motor recovery through integration of new neurons into functional cortical circuits in an activity-dependent manner[73]. These gene expression changes indicate that CREB directly influences recovery after stroke by enabling neurons to engage in larger functional connections in the brain tissue adjacent to the stroke.

## CCR5

C-C Chemokine Receptor-5, CCR5, is a G-protein coupled receptor with functions in adaptive immune signaling[74]. CCR5 is expressed in immune cells such as macrophages, monocytes, T-cells and NK cells. In particular, CCR5 signaling is important for determining localization of CD8 T-cells within lymphoid tissue and subsequent recruitment within inflammatory sites[74]. CCR5 is also expressed in the CNS, such as in astrocytes, microglia and endothelial cells. Although CCR5 signaling is associated with immune responses, a definitive role for CCR5 in the adult brain was only recently characterized through an extensive screen for potential learning and memory candidate genes[75]. In a reverse genetic screen of 148 transgenic mouse lines, mice with CCR5 knockdown showed improved performance in various learning and memory tasks, whereas CCR5 overexpression significantly impaired learning and memory. The study clearly constituted a role for CCR5 as a suppressor of cortical plasticity. Although chemokine receptors are not usually envisioned as modulators of synaptic plasticity, a growing body of evidence supports such a role through interactions with neurotransmitters and transcription factors[76]. In the context of learning and memory, CCR5 knockdown elevates CREB and MAPK signaling[75]. Furthermore, binding of CCR5 to its ligand RANTES (for 'regulated upon activation normal T cell expressed and secreted') biases CCR5 signaling towards activation of the $G_{\alpha i}$ pathway that inhibits cAMP production[77]. cAMP production facilities synapse formation during learning and memory[60]. Production of cAMP following CNS injury promotes a regenerative state[78]. Given the link between CCR5 and cAMP production, dampening CCR5 could lead to increased cAMP levels required for memory formation. Collectively, G-protein coupled receptor signaling along with interactions with transcription factors such as CREB enable CCR5 to commune plasticity associated-signaling events that range from cAMP generation to synaptogenesis; events that are hallmarks for learning and memory.

We have recently shown that dampening CCR5 signaling within neurons induces substantial motor recovery after stroke[15]. Knocking down CCR5 function after stroke induces motor recovery in motor tasks that allow quantification of fine motor control of the forelimb. Animals with CCR5 knockdown show improved navigation on a wired grid with fewer foot faults and less bias towards use of the impaired forelimb. CCR5 signaling is active in neurons and glia after stroke[79,15]. Interestingly, CCR5 is not normally expressed in cortical neurons, but is selectively upregulated following a stroke[15]. Knockdown of CCR5 in pre-motor cortex following a stroke to the motor cortex induces early and sustained motor recovery. This effect is associated with structural changes differentially regulated by CCR5 knockdown, such as the growth of unique projections to the contralateral pre-motor cortex and preservation of dendritic spines. CCR5 knockdown induces pro-regenerative signaling pathways through upregulation of CREB and dual leucine zipper kinase (DLK) expression. This data indicates a link between the recovery-promoting molecular systems of CCR5 and CREB. DLK, also known as MAP3K12, acts as a prominent axonal regeneration signal in other systems of CNS injury[80]. DLK activates various transcriptional profiles that pertain to either regeneration or apoptosis. The activation of these opposing pathways is dependent on targets downstream of DLK[80]. Following a stroke, diminishing CCR5 signaling positively modulates DLK signaling towards switching on a regeneration and/or repair program[15]. In fact, DLK upregulation is critical for this process. Downregulating DLK function diminishes motor recovery induced through CCR5 knockdown, showing that DLK acts as a signaling hub through which neuronal knockdown of CCR5 restores motor function[15].

A goal of molecular medicine has been to identify pharmaceutical targets for disease that have translational potential. CCR5 fits this description. Maraviroc, an FDA-approved CCR5 antagonist for HIV therapy, when delivered after acute or chronic stroke induces motor recovery, similar to viral genetic manipulations of CCR5[15]. Similarly, stroke patients that carry CCR5 32, a mutation that inactivates CCR5, show improved stroke outcomes compared to non-carriers[15]. The effects of CCR5 in rodent models, its druggable potential and clinical significance in patients makes CCR5 a highly promising target for stroke. Collectively, the role of CCR5 as a transcriptional substrate for memory and learning process parallels its role as a potent genetic target for stroke recovery through induction of shared mechanisms of plasticity.

## Learning-associative axonal remodeling and motor recovery

A consequence of gene expression changes during learning result in changes in synaptic strength or remodeling of synaptic inputs. For example, thalamocortical inputs are received by corticospinal neurons in the cortex that control distal and proximal hand function. During motor learning of a skilled reach task, thalamocortical projections onto neurons that project to the distal forelimb involved in grasping show greater firing release probability and amplitude of the signaling events, compared to the proximal limb[81]. Here, thalamocortical projections show a bias in increased connectivity towards neurons involved in learning. Axonal remodeling that involve growth of new projections have been reported, but these responses as a result of learning are limited, unlike after injury. In Pavlovian eye blink conditioning, a form of motor learning required for timed eyelid closure, skilled performers show structural changes in the cerebellum, such as increased mossy fiber collaterals and *de*

*novo* axonal connections from the basal pontine nucleus to various regions in the cerebellar nuclei[82]. In a more relevant motor learning model, where macaques were trained for tool use, new axonal projections form from higher visual centers to the intraparietal regions in the cortex[83], suggesting that learning is associated with changes in number and growth of axonal projections. An appreciable body of evidence in humans through longitudinal MRI imaging during learning points towards similar observations on gross structural changes in gray and white matter[84,85].

Axonal growth responses are active as early as 7 days after a stroke[72]. These responses are substantial and have been identified through various anatomical mapping studies[15,46,74,87,88]. The location, amount and the associated outcome of axonal sprouting vary depending on the size and location of stroke, molecular interventions applied and rehabilitative training. The peri-infarct cortex undergoes a remarkable degree of structural reorganization and growth, the result of time-dependent gene expression changes characteristic to this region. For example, following a focal cortical stroke, the peri-infarct cortex sends out new axonal branches to areas within motor, somatosensory and pre-motor cortices[15,46,88]. In addition to new axonal outgrowth in local areas, novel growth of bihemispheric projections are also induced[15]. One could argue that larger sprouting responses have been observed with larger infarcts. In models of large strokes in which more than a quarter of the cortical hemisphere is lost, responses from the contralateral hemisphere to denervated regions in the cervical spinal cord have been reported[89]. While new projections are formed as a result of the brain's endogenous response to reorganize after stroke, these axons follow an unconventional trajectory and populate regions of the cortex that are distinct prior to injury. For example, a focal stroke in the somatosensory cortex leads to a loss in connections to the thalamus; however, new axons that sprout take an alternative path into adjacent intracortical regions[87]. It is unclear if these responses contribute to spontaneous motor recovery, or if these responses are maladaptive.

Genetic manipulations to promote motor recovery or motor training from neurorehabilitation have also shown to induce axonal sprouting responses. Manipulations in EphrinA5, GDF10, or NgR1[45,88,90] signaling produce new and expansive axonal outgrowth from peri-infarct cortex to adjacent cortical areas, while blockade of CCR5[15] causes axons to sprout to cortical areas contralateral to the stroke site. These responses were unique in location and were absent in groups with stroke without treatment. Similarly, administration of Nogo-A antibody following a lesion to the primary motor cortex causes sprouting to callosal homotypic pre-motor areas in the contralateral cortex in macaques[91]. In addition to inter-intra-cortical connections, axonal sprouting resulting from such manipulations after stroke have shown to alter growth responses in descending pathways. Treatment with Nogo-A or inosine induces sprouting of corticospinal axons into the denervated cervical spinal cord [59,90,91,92]. Sprouting responses were further intensified when manipulation was paired with skill training[59,90]. It is worthy to mention that increased sprouting does not positively correlate with motor recovery. A clear demonstration of this is seen when Nogo-A antagonism simultaneously paired with skill training produces a robust sprouting response with null or negative effects on motor recovery[59]. Interestingly, motor recovery was induced with the introduction of a delay period between NogoA therapy and motor training, but with a less robust axonal sprouting response. These studies suggest that while stroke primes

post-stroke neurons to send out new projections, recovery requires complex target selection mechanisms. Non-selective axonal sprouting, or the formation of projections after stroke, may result in detrimental synaptic competition in off-target brain regions.

## Cortical map plasticity as a substrate for learning and motor recovery

Growth triggered after loss of axons at the stroke site can lead to changes in topographical brain regions activated during a motor response. These changes in the brain can be mapped using electrical or optical stimulation techniques by pairing the activation of a brain region with the elicited motor response. Characterization of cortical representations during motor behavior, such as a dexterous reach task, have shown the existence of maps with clear boundaries for wrist, digit and trunk movements[93]. The areas that represent these movements are enlarged during skill training—in other words, the part of the body that is trained shows an enlarged brain representation in its corresponding cortical map. Map reorganization is also dependent on age[94]. In the aged brain after stroke, where paucity for regenerative potential is pertinent, map reorganization has been shown to be less pronounced or absent.

A substantial body of work has shown that targeted strokes in the motor-sensory areas lead to the displacement or formation of new motor maps in adjoining cortical zones[95–99]. Map displacement is dependent on the location of the infarct. Map displacement is more prevalent following strokes to the somatosensory cortex where remapping of sensory forelimb function is seen in the motor cortex, although this does not reciprocate in the event of a stroke to the motor cortex[95]. Maps topographically do not displace following a loss of motor cortex; instead changes are more localized to perilesional cortex such as through an increased state of excitability. These differences in map topography based on lesion location suggest that cortical substrates underlying these maps are segregated based on differences in their cortical output projections or perhaps the intrinsic circuit structure of the particular cortical region (motor vs somatosensory).

A causal relationship between map reorganization and recovery of motor function is yet to be established. While the initial phase of learning induces changes in map topography, these changes normalize over time, making it difficult to assess the importance of topographical changes[100]. It is possible that map changes are less dynamic following the initial phase of learning. An interesting feature is that map enlargement can be induced through raising cortical excitability by increasing CREB activation after stroke[14]. However, further studies are required to understand if map enlargement leads to motor recovery. The gross topography of a map lacks resolution to determine changes at the circuit level, such as changes in cortical microcircuits and circuit reorganization during learning and recovery after stroke.

## Conclusion

Biological mechanisms underlying learning and memory formation parallel mechanisms of stroke recovery that range from changes at the synapse to motor circuits. Studies on cortical plasticity have identified distinct cellular substrates and molecular systems shared

between learning, memory and motor recovery. Synaptic plasticity as viewed in changes in dendritic spine remodeling, axonal remodeling and cortical map function show similar patterns during motor learning and recovery after stroke. Molecular systems that promote neuronal excitability underlying memory formation can enhance recovery after stroke. In fact, these systems drive changes in synapse density and integrate neurons that normally engage in storage of motor memories into a network that accomplishes a motor task that underlies motor recovery. Molecular systems, such as CREB and CCR5, provide pharmacological targets for drugs that promote stroke recovery. However, questions remain in the understanding of links between synaptic plasticity mechanisms and recovery from stroke. While brain reorganization processes have been described at the anatomic levels, organization at the functional level within populations of neurons are areas of further investigation. Furthermore, how does learning such as during neurorehabilitation augment spatiotemporal activity within neuronal assemblies in the motor system to output motor recovery? Is there a state of maladaptiveness reflective at the population level that help understand adaptive and maladaptive structural plasticity? While existing studies discussed here have greatly furthered our understanding on changes in the motor system after a stroke and during functional motor recovery, future studies will dissect mechanistic processes to help understand complex computations within neuronal circuits that underlie recovery of motor function.

## Acknowledgements

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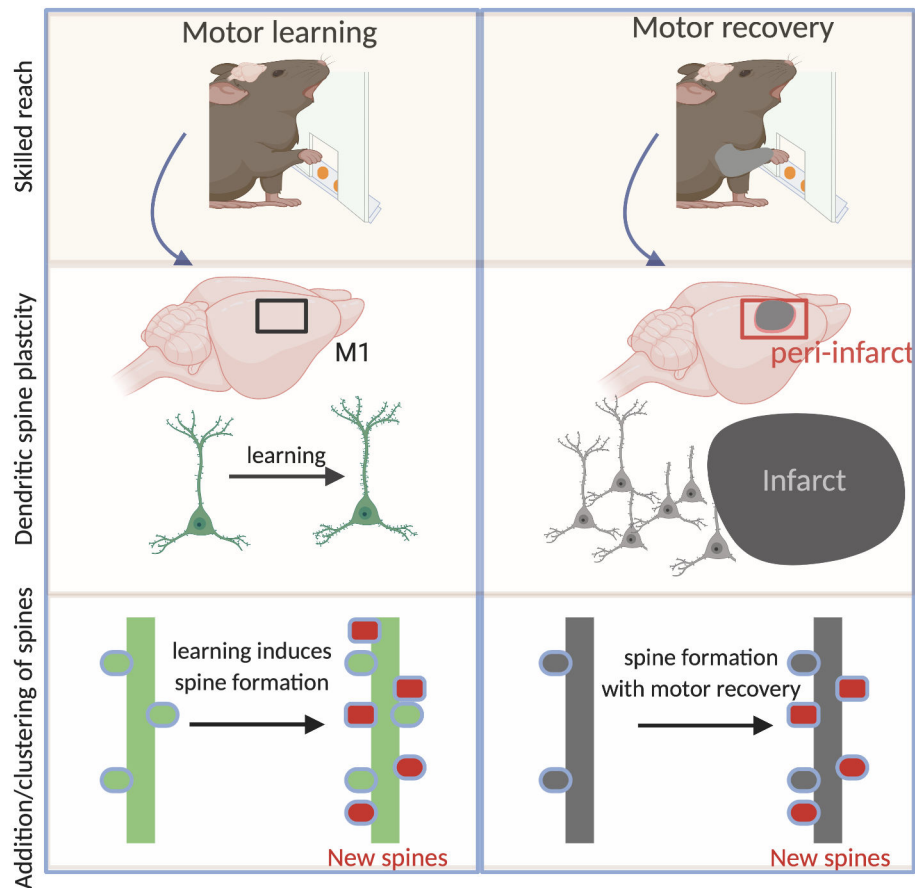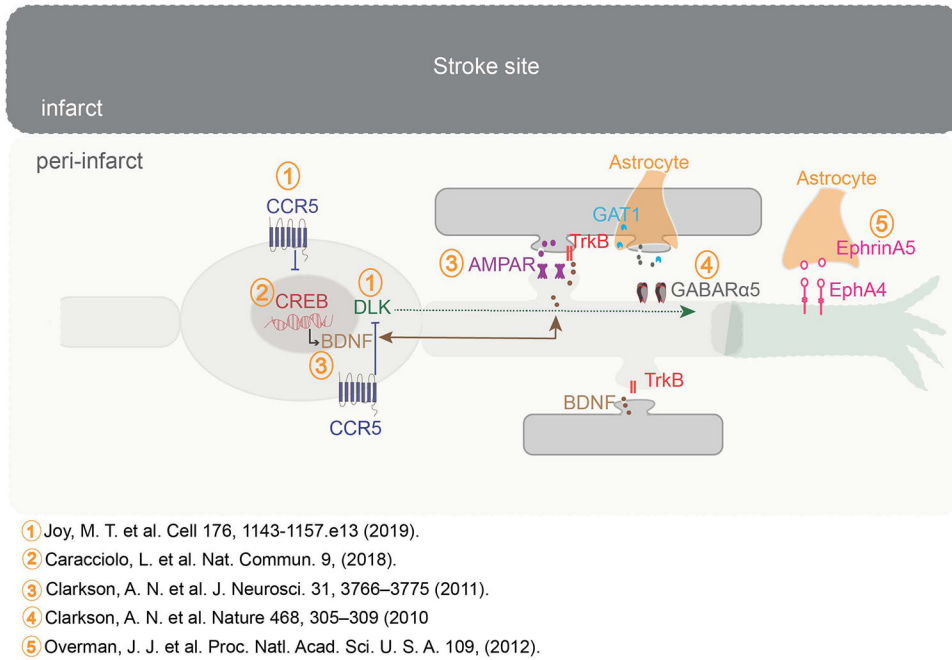
**Figure 1:**
Dendritic spine plasticity- a substrate for motor learning and motor recovery after stroke.
Left panel- Motor learning in rodents such as skilled reaching to grab food pellets forces
changes on post-synaptic structures on dendrites called spines. These changes occur in
motor cortex (M1). Changes include addition of new spines (red) and clustering of spines
that lead to strengthened synaptic inputs. Right panel- After stroke, there is loss of synaptic
connectivity, particularly near the infarct (peri-infarct tissue). Rehabilitation, spontaneous
motor recovery or genetic manipulations such as with CCR5 knockdown induces similar
changes on spines as reported with motor learning in the normal brain.

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**Figure 2:**

Molecular systems that underlie learning regulate recovery after stroke. The peri-infarct cortex acts a hub for molecular events that regulate recovery after stroke. Transcription factor CREB is critical for memory formation. Increasing CREB signaling after stroke improves recovery of motor function. CREB signaling can lead to signaling pathways that enhance BDNF production. BDNF activates Trk signaling and this pathway in turn is positively modulated by AMPA receptor signaling. Slow and persistent (tonic) GABA signaling through the alpha5 receptor at extra synaptic sites dampens stroke recovery. GABA signaling is persistent due to defects in the GABA transporter (GAT1) expressed by astrocytes that function to transport and clear GABA at extrasynaptic sites. Blocking this pathway enhances motor recovery. CREB signaling is impeded by expression of CCR5 early after stroke. Blocking CCR5 signaling increases CREB signaling and signaling via DLK that acts as a regeneration signal for axonal sprouting. Axonal sprouting is further facilitated through reduced signaling via Ephrin A5 expressed on reactive astrocytes and its receptor EphA4 on neurons, that otherwise serve to collapse arrangement of cytoskeletal proteins.