

Characterizing Brain Cortical Plasticity and Network Dynamics Across the Age-Span in Health and Disease with TMS-EEG and TMS-fMRI

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Abstract Brain plasticity can be conceptualized as nature's invention to overcome limitations of the genome and adapt to a rapidly changing environment. As such, plasticity is an intrinsic property of the brain across the lifespan. However, mechanisms of plasticity may vary with age. The combination of transcranial magnetic stimulation (TMS) with electroencephalography (EEG) or functional magnetic resonance imaging (fMRI) enables clinicians and researchers to directly study local and network cortical plasticity, in humans in vivo, and characterize their changes across the age-span. Parallel, translational studies in animals can provide mechanistic insights. Here, we argue that, for each individual, the efficiency of neuronal plasticity declines throughout the age-span and may do so more

or less prominently depending on variable 'starting-points' and different 'slopes of change' defined by genetic, biological, and environmental factors. Furthermore, aberrant, excessive, insufficient, or mistimed plasticity may represent the proximal pathogenic cause of neurodevelopmental and neurodegenerative disorders such as autism spectrum disorders or Alzheimer's disease.

Keywords Cortical brain plasticity · Transcranial magnetic stimulation · Electroencephalography · Functional magnetic resonance imaging · Lifespan

Plasticity as an Intrinsic Property of Human Brain

The world we live in changes rapidly. Afferent inputs and efferent demands to the brain shift quicker than the time needed to implement genetic or even epigenetic changes. Brain plasticity can be conceptualized as nature's invention to overcome limitations of the genome and adapt to the rapidly changing environment (Fig. 1; Pascual-Leone et al. 2005). As such, plasticity represents an intrinsic property of the nervous system retained throughout life that enables modification of function and structure in response to environmental demands via the strengthening, weakening, pruning, or adding of synaptic connections and by promoting neurogenesis. This means that the brain does not remain static but, instead, continues to change as the obligatory consequence of each sensory input, motor act, association, reward signal, action plan, and awareness (Pascual-Leone et al. 2005).

However, it is worth remembering William James who defined plasticity is the "possession of a structure weak enough to yield to an influence, but strong enough not to yield all at once". Excess plasticity can be equally

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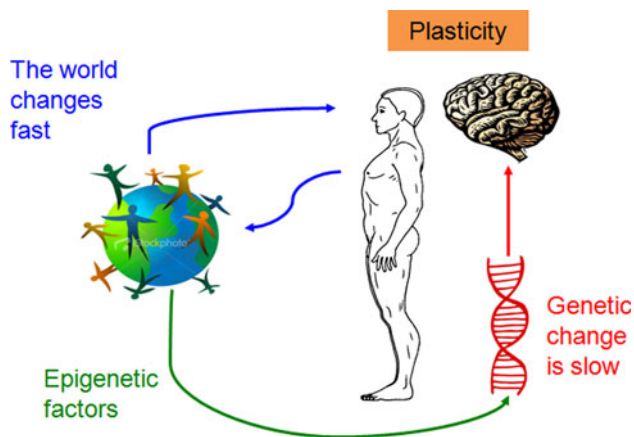


Fig. 1 Schematic representation of the concept of plasticity. Brain plasticity allows for rapid adaptation to environmental changes that occur quicker than genetic or epigenetic response times

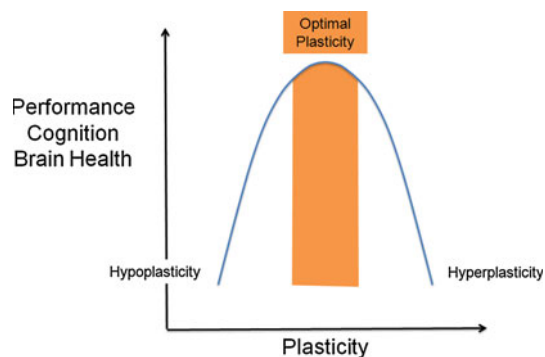


Fig. 2 A bell-curve response to plasticity levels suggests that both too little and too much plastic response can hinder cognitive performance and overall brain health. “Optimal amounts of plasticity” will necessarily be different for different individuals, varying across brain regions and networks and changing across the lifespan

deleterious as insufficient plasticity (Fig. 2). Plasticity is essential to the establishment and maintenance of brain circuitry, it can be beneficial for the individual enabling acquisition of new skills and adaptation after an injury, but it can also account for the symptoms of disease. Normal plasticity mechanisms can serve to compound the pathological consequences of a specific genetic mutation or sustained environmental insult, and aberrant plasticity mechanisms can act on a previously normal brain to induce pathological manifestations of disease. Early altered or mistimed plasticity may set the stage for otherwise innocuous processes to become pathogenic (Gogolla et al. 2009). A deficit in plasticity will render the brain unable to adjust to changing demands. On the other hand, if the brain is too plastic, structural connections may become unstable and functional systems necessary for cognition and behavior may be compromised.

The brain is highly interconnected. Therefore, plasticity plays out across the multiple levels of nervous system

complexity, from cellular through microcircuits to circuits and large-scale networks. Activity in lower levels may influence activity in higher levels, and vice versa. Changes in local plasticity can be compensated for by circuit and network adaptations in such a way that behavior may not deteriorate (in fact, some behaviors may even be paradoxically improved). Alternatively, local changes may be compounded by further maladaptive circuit and network dynamics giving rise to disability and the symptoms of disease. Thus, changes in local plasticity may constitute the first in a chain of events culminating in ‘circuitopathies’ in which symptoms are the consequence of dysfunctions of neural circuits and networks. If so, measures of cortical plasticity may provide very early local and network biomarkers of neuropsychiatric disease.

In the present article we discuss concepts of plasticity as they might evolve across the age-span and contribute to normal development, life-long cognitive abilities, and the manifestation of developmental or neurodegenerative disorders. We argue that, in all individuals, the efficacy of the mechanisms of plasticity changes over the lifespan, but that it does so from variable starting-points and with variable slopes depending on a number of genetic factors, environmental factors, and their complex interaction (Fig. 3). Empirical determination of each individual’s slope of changing brain plasticity across the lifespan is possible with real-time integration of transcranial magnetic stimulation (TMS) with electroencephalography (EEG) and functional magnetic resonance imaging (fMRI) (Freitas et al. 2011a, b). Such methods might ultimately provide early predictors of individual risk for age-related cognitive decline, diagnostic biomarkers for neurodevelopmental and neurodegenerative disorders, and enable plasticity-based interventions to optimize outcomes for each individual.

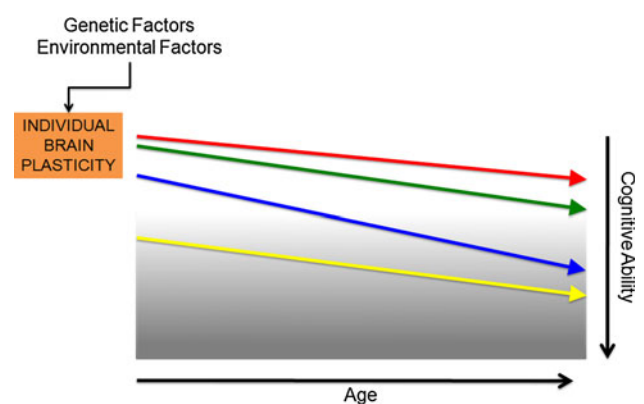


Fig. 3 Schematic representation of individual plasticity across the lifespan. Although mechanisms of plasticity show a downward trend over the course of a typical lifetime, this trend will manifest differently according to initial “baseline” levels, genetic factors, and environmental influences. Therefore, one may conceptualize each individual has a unique “slope of plasticity” across the lifespan

Plasticity Changes Across the Lifespan: Animal Models and Indirect Evidence in Humans

Animal studies building on pioneering work from Barnes in the late 1970s (Barnes 1979) have demonstrated an age-associated decline in synaptic plasticity that correlates with neurocognitive impairments (Rosenzweig and Barnes 2003). For example, once induced, hippocampal long-term potentiation (LTP) decays faster in older rats, and this appears to be associated with a greater degree of forgetfulness (Barnes and McNaughton 1980; Kelly et al. 2006). Moreover, deficits in the balance between LTP and long-term depression (LTD) result in impaired learning and memory (Larson et al. 1986; Roman et al. 1987; Bliss et al. 2003).

However, despite these and other findings in animal models, evidence in humans has remained largely indirect. Brain imaging studies with structural MRI, fMRI, diffusion tensor imaging (DTI), and positron emission tomography (PET) can provide evidence supporting the claim that plasticity changes across the lifespan. But, for the most part, these methods assess the *consequences* of plasticity and fail to directly assess the *mechanisms of plasticity* themselves.

Structural neuroimaging can assess the integrity of brain structures and provide indirect measures of plasticity phenomena. For example, cross-sectional studies have consistently identified age-associated morphometric brain changes encompassing regional cortical thinning, volumetric subcortical reductions, and ventricular enlargement (e.g., Fjell et al. 2009; Walhovd et al. 2005, 2009). Longitudinal studies have demonstrated annual atrophy rates for brain volume, hippocampus and entorhinal cortex (e.g., Scahill et al. 2003; Fotenos et al. 2005), and atrophy in cortical brain regions over different periods of time (Raz et al. 2005; Driscoll et al. 2009). Cortical thickness decreases over the lifespan are estimated at 0.5% a year (Thompson et al. 2007). These changes affect different neural systems differently: motor and visual cortices show regional thinning, whereas non-limbic temporal regions and parietal areas are relatively spared in normal aging (Salat et al. 2004; Raz et al. 2004). Furthermore, DTI can address structural changes in white matter structure (myelination) and connectivity. For example, DTI has demonstrated that white-matter connections, largely in fronto-striatal areas, have reduced myelination as age increases (Salat et al. 2005). It is assumed that such structural, morphometric MRI and DTI measures must be associated with cognitive and behavioral consequences (Fjell and Walhovd 2010).

There are changes in cognitive task performance associated with aging and, as a corollary to this, changes in the neural activation resulting from these cognitive tasks. It has been widely established that normal human aging in humans is associated with decrements in cognitive performance

across several domains including processing speed, working memory, episodic memory, attentional control, inhibitory control, and executive function (e.g., Gazzaley and D'Esposito 2007; Reuter-Lorenz and Park 2010; Park and Reuter-Lorenz 2009). In parallel, a host of neuroimaging investigations have described altered patterns of brain activation in healthy elders as they perform cognitive tasks. One such pattern is that of a reduction in prefrontal hemispheric asymmetry in elderly individuals, referred to as the HAROLD model (hemispheric asymmetry reduction in older adults) (Cabeza et al. 2002). According to the HAROLD model, the older brain displays less localizable and more bilateral activation during certain cognitive tasks. A second pattern is a shift in evoked neural activity from posterior to anterior cortex, a model referred to by Davis et al. as PASA (posterior-anterior shift in aging) (2008). The PASA model posits that the aging brain is more likely to recruit prefrontal, rather than occipito-temporal, cortex in the service of task execution.

In addition to life-span changes in task-related brain activation patterns, resting-state fMRI is revealing age-related differences in the functional connectivity across large-scale brain networks. One such large-scale brain functional network, the default mode network (DMN) has been shown to undergo notable modifications with advancing age in health and disease (Buckner et al. 2008). Older individuals reportedly exhibit significantly lower DMN activity in the posterior cingulate as well as a tendency toward lower activity in all other DMN regions as compared to younger subjects (Koch et al. 2010). Functional connectivity within the DMN also seems to be reduced in older adults (Grady et al. 2010). During performance of a working memory task, the pattern of deactivation of the DMN also seems to be affected by aging, with older individuals not only showing decreased connectivity but also decreased ability to suppress low-frequency oscillations of the DMN (Sambataro et al. 2010). Age-specific changes in activation and connectivity are also seen in the task-positive network (TPN), though the functional significance of this remains uncertain (Grady et al. 2010; Sambataro et al. 2010). During memory encoding and recognition, age-related changes appear to occur mainly in the long-range connections with widespread reductions associated with aging in the fronto-temporal and temporo-parietal regions, and a few age-related increases in the posterior parietal regions (Wang et al. 2010). During developmental years, children and young adults appear to have similar patterns of functionally connected regions, but with differences in the size of functionally connected regions as well as in the strength of functional connectivity between brain regions (Jolles et al. 2010).

The apparent tendency for the aged brain to activate bifrontal cortex more anterior (as opposed to posterior)

cortical regions during certain cognitive tasks as well as the observed changes in resting-state functional brain connectivity are often explained on the basis of compensatory strategies. It has been theorized that these changes represent plasticity-based modifications to compensate for deficits associated with normal aging. However, there is also increasing discussion that age-related changes in brain activation patterns may also reflect a break-down of optimized brain function, an age-related increase in neural noise that might ultimately be maladaptive. Further longitudinal studies are needed and, in addition, greater mechanistic understanding of the nature of age-related changes in brain activity and structure is warranted. In relation to plasticity, direct experimental measures of plasticity mechanisms in humans that might be translated to animal models are desirable.

Direct Empirical Measures of Mechanisms of Plasticity in Humans

Transcranial Magnetic Stimulation (TMS)

TMS is a non-invasive procedure used to create electric currents in discrete brain regions (Kobayashi and Pascual-Leone 2003; Wagner et al. 2007; Horvath et al. 2011).

TMS is based on Faraday's principle of electromagnetic induction and features application of rapidly changing magnetic field pulses to the scalp via a copper wire coil connected to a magnetic stimulator. These brief pulsed magnetic fields painlessly pass through the skull and can create electric currents of sufficient magnitude in discrete brain regions to depolarize neurons. When applied to the motor cortex, this depolarization results in a series of descending (direct and indirect) cortico-spinal waves that can sum-up at the spinal segmental level, depolarize alpha motor neurons, and lead to the contraction of contralateral muscles. This contraction, known as a motor evoked potential (MEP), can be measured using electromyography (EMG). Applied to non-motor cortical regions, TMS evokes a local field potential that can be recorded with EEG, and represents a measure of cortical reactivity to TMS (Fig. 4). In addition, beyond the targeted cortical brain region, TMS exerts transynaptic distributed network effects that can be revealed combining TMS with neuroimaging.

Trains of repeated TMS pulses (rTMS) at various stimulation frequencies and patterns can induce a lasting modification of activity in the targeted brain region which can outlast the effects of the stimulation itself. Such lasting changes presumably represent alterations in plasticity mechanisms. One rTMS protocol, known as theta burst

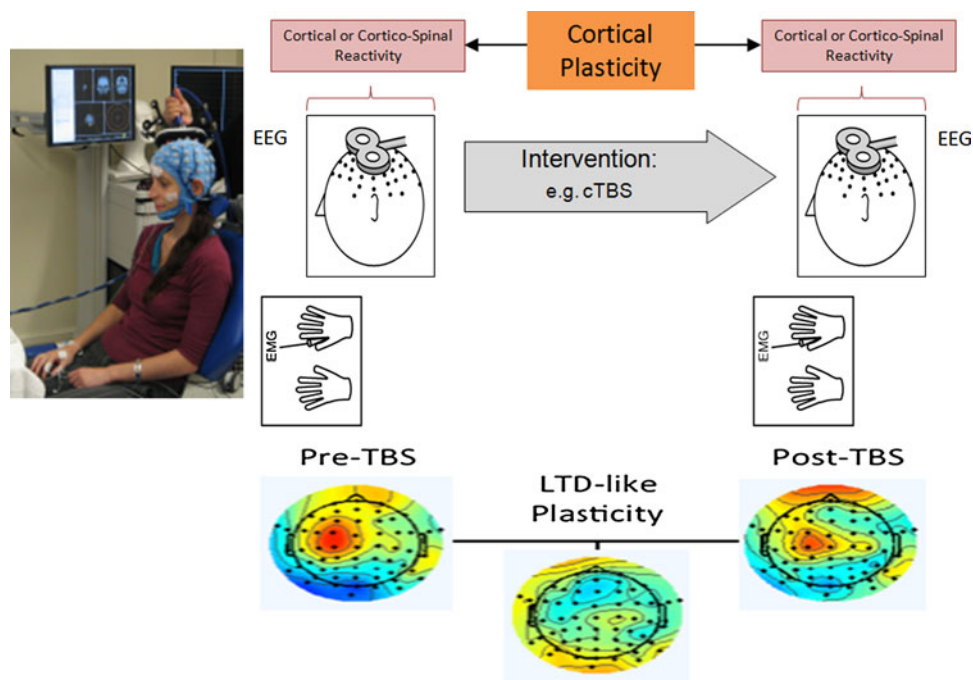


Fig. 4 TMS-based measures of cortical reactivity and plasticity. As recorded using either electromyography (EMG) or electroencephalography (EEG), brain responses to TMS can be measured as motor evoked potentials (when TMS is applied to motor cortex) or localized evoked field potentials. Such responses reflect cortical or cortico-spinal reactivity. These measures can be obtained before and

following a given intervention, for example controlled modulation of cortical excitability by repetitive TMS in the form of theta burst stimulation (TBS). The TBS-induced change in EMG or EEG responses to TMS provides a measure of local cortical plasticity. An example of TMS-EEG measures before and after continuous TBS illustrates the measurement of LTD-like plasticity

stimulation (TBS), mimics paradigms used to induce LTP and LTD in animal models (Huang et al. 2005). TBS consists of bursts of 3 pulses at 50 Hz repeated at intervals of 200 ms. When applied to the motor cortex, this protocol can induce robust and consistent responses across subjects, and lead to enhancement or depression of MEP amplitudes depending on stimulation parameters. After TBS is applied to the motor cortex in an intermittent fashion (iTBS), TMS-evoked potentials show increased amplitude. Conversely, after TBS is applied continuously (cTBS), TMS-evoked potentials show a marked decrease in amplitude.

TBS is safe if appropriate safety guidelines are followed (Oberman et al. 2011) and various lines of evidence support the notion that such TMS protocols can indeed provide a measure of synaptic plasticity in humans (Cárdenas-Morales et al. 2010). Physiologic and pharmacologic studies of TBS in humans show involvement of glutamatergic and GABAergic mediators (Huang et al. 2008; Stagg et al. 2009), and the effects and their time-course are consistent with LTP- or LTD-phenomena. Furthermore, Tokay et al. (2009) demonstrated that high-frequency magnetic stimulation induces LTP in rat hippocampal slices and that stimulation-induced LTP was prevented by the presence of a selective N-methyl-D-aspartate receptor (NMDAR) blocker. These findings further suggested that presynaptic involvement of the induced LTP may be excluded based on the lack of changes in paired-pulse ratio and the afferent fiber volleys. However, studies to date are limited and further understanding of the mechanistic effects of TMS is of obvious and crucial importance.

Combining TMS with EEG and fMRI

Real-time integration of single- and paired-pulse TMS with EEG can provide information about local cortical reactivity and distributed network dynamics in health and in disease (Thut et al. 2005; Thut and Pascual-Leone 2010). Contrasting such measures before and serial following various rTMS protocols (for example TBS) can provide insights into LTP- and LTD-like cortical plasticity and network dynamics in humans in vivo across cortical region.

Using EEG as the outcome measure enables the implementation of a variety of sophisticated techniques to identify and characterize connectivity networks between different brain regions (Sameshima and Baccala 1999; Kaminski et al. 2001; Stam and van Dijk 2002; Kramer et al. 2009). Graph theoretic techniques can be utilized to derive an understanding of some of the important properties of these networks (Bullmore and Sporns 2009). Several studies have shown that these network architectures vary during normal aging (Micheloyannis et al. 2009; Boersma

et al. 2011), and may be abnormal in patients with a variety of neuropsychiatric conditions including schizophrenia (Micheloyannis et al. 2006; Medkour et al. 2010), depression (Fingelkurts et al. 2007; Leistedt et al. 2009), ASD (Murias et al. 2007), epilepsy (Bettus et al. 2008; van Dellen et al. 2009; Douw et al. 2010), TBI (Cao and Slobounov 2010; Sponheim et al. 2011), and AD (Stam et al. 2007; Jelles et al. 2008; de Haan et al. 2009). The combination of such EEG approaches with TMS allows clinicians and researchers to fully characterize dynamic properties of local and distributed brain cortical activity by offering a behaviorally independent input of quantifiable and parametrically scalable magnitude applicable across ages, individuals, and neural states.

Esser et al. (2006) demonstrated that after trains of high-frequency (5 Hz) rTMS, EEG responses to single TMS pulses—TMS-evoked potentials (TEP)—are significantly potentiated. Conversely, amplitude of TEPs is significantly decreased after low-frequency (0.6 Hz) rTMS (Van der Werf and Paus 2006). These findings are consistent with the differential effects of high and low frequency rTMS on cortical excitability and plasticity (Valero-Cabre et al. 2007). A number of studies have shown that the cortical responses to various sensory stimuli are also altered by different rTMS protocols; for example, Ishikawa et al. (2007) demonstrated that the amplitude of somato-sensory evoked potentials is altered after TBS, with the effects persisting for up to 50 min. Repetitive TMS protocols also induce quantifiable changes in various EEG metrics of intrinsic cortical activity, such as changes in EEG power in various frequency bands (Okamura et al. 2001; Schutter et al. 2001; Griskova et al. 2007, 2009; Brignani et al. 2008; Fuggetta et al. 2009), or synchrony and coherence between different cortical regions (Jing and Takigawa 2000; Strens et al. 2002; Oliviero et al. 2003; Schindler et al. 2008). Julkunen et al. (2008, 2011) assessed navigated TMS-evoked EEG responses in patients with AD, and showed prominent changes in functional connectivity (and reactivity) in AD subjects. In particular, TMS-evoked response at 30–50 ms decreased significantly in AD patients over widespread brain regions, thus suggesting dysfunction of a large-scale sensorimotor network or less network synchronization in patients with AD.

TMS-EEG local cortical and network plasticity measures can be further elucidated by obtaining simultaneous neuroimaging with fMRI. Functional MRI allows for the investigation of whole brain response to and recovery from patterned stimulation. For example, combining TMS with resting-state fMRI allows clinicians and researchers to experimentally manipulate local cortical responses with patterned stimulation and observing change in network responses (Halko et al. 2010).

Direct Evidence of Changing Plasticity Mechanisms in Humans Across the Life-Span

Direct evidence of brain plasticity changes in humans throughout the lifespan in health and disease can be measured via real-time combination of TMS with EMG, EEG, and/or fMRI (Fig. 4). Moreover, measures of TMS before and serially following cTBS or iTBS can provide measurements of LTP- or LTD-like plasticity.

Preliminary experimental evidence reveals a decline in corticomotor plasticity across the human lifespan (Freitas et al. 2011a, b). In a cross-sectional study of 36 healthy volunteers throughout the adult age-span ranging from 19 to 81 years, we found the duration and magnitude of corticospinal excitability modulation by cTBS was inversely and significantly correlated with age (Fig. 5). These data provide direct experimental evidence that, in humans, LTD-like plasticity becomes increasingly less efficient with advancing age. Such decreasing plasticity in the motor cortex with advancing age may be associated with the decrement of hand motor function (e.g., longer reaction time) observed during normal aging in both men and women (e.g., Carmeli et al. 2003) and to the age-related deficits in motor learning (e.g., Brown et al. 2009). TMS studies using EEG instead of EMG measures will enable assessment of non-motor cortical regions and provide direct experimental assessments of age-related changes in mechanisms of plasticity.

Advantage of Translation of Method to Animals

Through the use of TMS in controlled animal models of healthy and diseased states, the underlying mechanisms

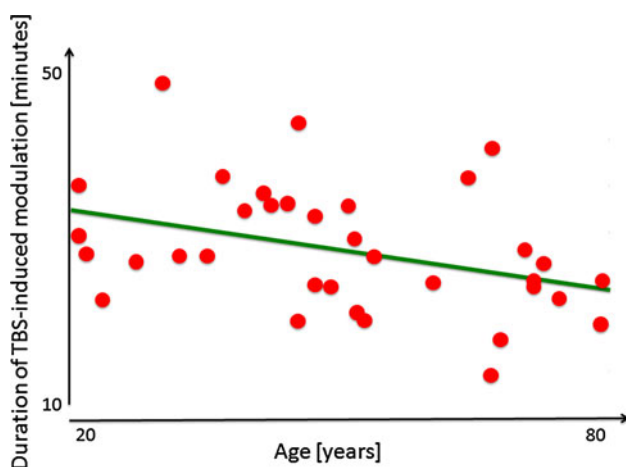


Fig. 5 Modified from Freitas et al. (2011a, b). Correlation between age and duration of the modulation of cortico-spinal responses to TMS after cTBS in cognitively intact, healthy adults ranging in age from 19 to 81 years. This measure of plasticity shows a significant decrease with age

and the role of TMS in evaluating neural plasticity may be more thoroughly defined. Animal models allow for more precise pharmacologic manipulation, improving our ability to correlate specific neurochemical changes with TMS effects. In addition, the use of rodent models provides for the potential to evaluate the role of TMS under specific genetic backgrounds, i.e., transgenic mice. In turn, TMS may also improve animal studies by allowing a noninvasive quantitative neurophysiologic measure that can be obtained in real-time and evaluated serially in longitudinal studies within the same subjects. In translational animal research, such as in neuroprotective stroke research, failure to bring the successes of animal studies to clinical fruition may be, in part, the result of different outcome markers between animal and human investigations (Gladstone et al. 2002). TMS measures of neuronal excitability and plasticity may serve as a unifying outcome measure amongst animal and human research to better facilitate clinical translation.

The use of TMS in animal studies has been met with some difficulty. Namely, conventional TMS coils, designed for human application, generate relatively broad volumes of electrical current which may activate multiple cortical and subcortical regions when applied to the rat. As such, the focality of TMS in animals is brought into question and obscures the origin and interpretation of resultant TMS metrics. Furthermore, TMS animal studies require anesthesia to both minimize animal discomfort and suppress spontaneous muscle activation. The presence of neurochemically-active anesthetics may further obscure TMS findings (Luft et al. 2001; Rotenberg et al. 2010).

Yet, rat TMS studies have made important advancements in recent years. TMS in rats and mice has been shown to elicit stable motor evoked potentials (MEPs) with similar recruitment curves as seen in humans (Rotenberg et al. 2010). In addition, when placed over the scalp in an eccentric orientation, TMS is able to evoke MEPs lateralized to the contralateral limb. These findings suggest a resolution of rodent TMS to at least one hemisphere. Further, the longer latency of TMS-evoked MEPs when compared with MEPs from cervical electrical stimulation, as well as the similarities in latency and morphology with MEPs from focal cortical electrical stimulation, favors a presumed cortical origin of TMS (Luft et al. 2001, 2002; Rotenberg et al. 2010; Schlag et al. 2001; Zandieh et al. 2003). Studies have also begun to demonstrate the role of animal TMS as a marker of neuronal plasticity. Luft and colleagues demonstrated the effect of motor learning, simulated by somato-sensory afferent stimulation, on the enhancement of cortical motor excitability (Luft et al. 2002). More recent research has translated paired-pulse TMS techniques to murine models, providing measures of cortical inhibition as an indicator of neuronal plasticity

(Vahabzadeh-Hagh et al. 2010). TMS-EEG studies in animal models can also parallel those conducted in humans (Ives et al. 2006). Although many variables remain, rodent TMS research is proving to be a viable model through which many TMS questions may be further investigated.

Individual Differences in Plasticity

TMS-based experimental methods to examine cortical plasticity mechanisms in humans enable longitudinal studies that are critically needed to further examine brain changes across the lifespan. Longitudinal studies are particularly critical as individual differences in brain plasticity are likely quite large, and understanding the factors that condition such individual differences will offer unique and novel targets to promote individual brain health and well-being across the lifespan.

Important factors to consider that likely contribute to differences in mechanisms of plasticity include genetic and epigenetic mechanisms (e.g., polymorphisms, genetic expression), hormonal factors (e.g., gender, menstrual cycle), impact of morbidities (e.g. diabetes, cancer, or infections), and lifetime experiences (e.g., traumatic brain injury, exposure to toxins, stress, sleep deprivation, substance abuse, poor cognitive reserve, poor nutrition, sedentariness, etc.). Therefore, dissimilar ‘starting points’ for different individuals and distinct lifelong ‘slopes of change’ in plasticity might be postulated (Fig. 3). Beyond that, Fig. 6 schematically illustrates how different genetic and environmental factors may impact each individual’s brain plasticity ‘slope’ throughout the age-span, inducing distributed neural network responses that may prove

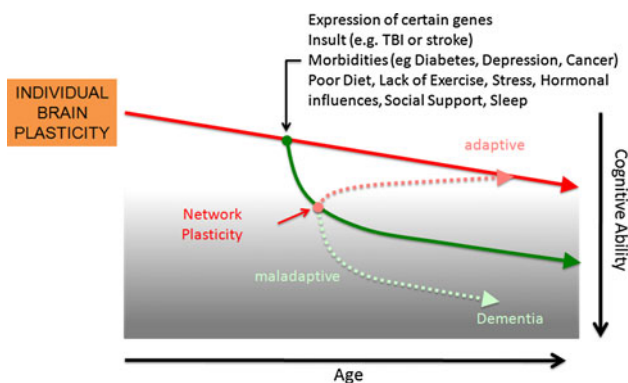


Fig. 6 Schematic representation of the influence of genetic or environmental impacts on brain plasticity. Alteration of local plasticity will trigger secondary adaptive responses across diffuse neural networks that may prove ultimately adaptive or maladaptive for the individual. Depending upon the amount and scope of such secondary responses, initial insult effects may be alleviated or heightened

adaptive or maladaptive, and might determine age-related cognitive abilities and even the risk of late-life dementia.

The Role of Genetic Differences

A number of genetic factors have been identified that relate to regulation of human brain plasticity (Pearson-Fuhrhop et al. 2009). For example, the brain derived neurotrophic factor (*BDNF*) gene plays an important role in neuroplasticity. The *BDNF* gene contains a functional single-nucleotide polymorphism (rs6265) that results in the substitution of valine to methionine (Val66Met), leading to reduced mature *BDNF* expression. The *BDNF* Val66Met polymorphism has been shown to differentially modulate human cortical plasticity and the response to training (Kleim et al. 2006), brain stimulation (Cheeran et al. 2008) and motor learning (Fritsch et al. 2010). Thus, the *BDNF* example illustrates the fact that individuals with a certain genetic predisposition may show a different response to interventions that modulate brain plasticity. Furthermore, the *BDNF* Val66Met polymorphism has also been shown to distinctively influence both brain structure and function by exerting a more global, rather than local, effect. For instance, brain allometry shows that *BDNF* 66Met carriers display larger total lobar volumes, while regional differences might only reflect the proportion of the volumes occupied by different brain regions relative to the total lobar volumes (Toro et al. 2009). Similarly, the *BDNF* Val66Met polymorphism appears to differently influence whole-brain—but not regional-activation patterns (McHughen et al. 2010). The functional relevance of such effects remains unclear. Yet, it is conceivable that the presence of the Val66Met polymorphism may have distinct repercussions on the mechanisms of plasticity during normal development and aging. Accordingly, individuals carrying this polymorphism may have distinct starting points and plasticity slopes throughout the age-span. Such differences may, eventually, contribute to late-life neurodegeneration and dementia. Indeed, it has been suggested that the presence of the *BDNF* Val66Met polymorphism renders higher susceptibility to AD, although conflicting results have been found (e.g., Desai et al. 2005; Matsushita et al. 2005; Saarela et al. 2006; Tsai et al. 2006; Fehér et al. 2009). Recently, the low-activity Met66 allele was shown to be an additional risk factor for rapid disease progression during the preclinical period of AD (Hashimoto et al. 2009) and may also constitute a risk factor for the development of psychotic symptoms in AD (Pivac et al. 2010).

Another common genetic mutation that illustrates the impact of genetic polymorphisms on distinct aspects of brain plasticity is the apolipoprotein E (*APOE*) susceptibility gene located on chromosome 19 and its $\epsilon 4$ allele, which is thought to raise the risk of AD (e.g., Corder et al.

1993) in a dose-dependent manner (Bertram et al. 2007; Bertram and Tanzi 2009). Wolk and Dickerson (2010) have shown that the presence of *APOE-ε4* differentially influences large-scale brain networks and this contributes to the clinical phenotype of AD. Thus, it seems that the presence of *APOE-ε4* may influence network plasticity and the slope of plasticity throughout the lifespan.

In addition, mutations in a number of other genes involved in synaptic plasticity have been shown to confer an increased risk for certain mental disorders that are associated with changes in cortical local and network plasticity. Examples can be found in candidate genes for autism spectrum disorders (ASD), including *Neurologin 3* and *4*, which are implicated in synaptogenesis (Jamain et al. 2003), *CNTNAP2*, which contributes to neural migration (Durand et al. 2007), *SHANK3*, which encodes a protein involved in dendritic development (Durand et al. 2007), and, more recently, *c3orf58*, *NHE9*, and *PCDH10*, also critically involved in synaptic development and plasticity (Morrow et al. 2008).

The interface between genes and environment may translate into differential genetic expression between individuals and also throughout the lifespan. For instance, Lu et al. (2004) used transcriptional profiling to identify a set of genes whose expression decreases with increasing age; among the genes most affected were those involved with synaptic plasticity, including the gene coding for a subunit of the Glu R1 AMPA receptor and genes coding for subunits of the NMDA and GABA_A receptors. Additionally, these authors found that genes involved with the synaptic calcium signaling system (which is critical for LTP) were downregulated with advancing age, including *calmodulin 1* and *CAM kinase IIα*. Moreover, as argued by some (Bray 2008), individual differences in gene expression are likely to be responsible for much of the human diversity.

The Role of Environmental and Other Personal Differences

Beyond genetic factors, and certainly contributing to genetic expression via epigenetic mechanisms or interactions with susceptible genes, environmental factors also contribute to shape differences in the efficacy and functional consequences of plasticity mechanisms across the lifespan. This includes such diverse factors as educational experience, family upbringing and other social interactions, cultural background, hormonal factors (e.g., gender), morbidities (e.g. diabetes, cancer, infections, or depression), nutritional factors, exposure to toxins, stress, sleep deprivation, substance abuse, physical exercise, as well as unexpected insults (e.g., traumatic brain injuries). Physical exercise is increasingly recognized as a powerful influence on plasticity across the lifespan (e.g. Voss et al. 2011).

Nutrition also has a significant influence on the aging process (Woo 2011), possibly due in part to an impact on brain plasticity slopes across the lifespan. The richness of one's environment (Volkers and Scherder 2011) as well as cultural influences (Park and Gutches 2002) impact the aging process, perhaps via modified cortical plasticity mechanisms.

Another important factor potentially contributing to inter-individual variability in efficacy of plasticity and its lifespan slope is intelligence. Indeed, brain plasticity and intellectual ability both seem to be partly driven by shared genes (Brans et al. 2010). In their longitudinal study of adult twins, Brans et al. (2010) showed that increased thickening in the medial temporal lobe and attenuated thinning of the frontal cortex, until age 35, were associated with higher IQ scores and further showed that genes implicated in cortical thickness overlapped with those involved in the level of intelligence. This may be related to early life experience and contribute to the construct of cognitive reserve.

Epidemiological studies suggest that early-life abuse and neglect increase susceptibility to psychopathology. Reviews of large-scale research studies conclude that there is clearly an excess of early traumatic events in individuals who experience psychosis, schizophrenia, and other psychiatric conditions in adulthood (Larkin and Read 2008; Read et al. 2005). Indeed, the quality of each individual's social environment can have profound influences on the development and activity of neural systems, with repercussions on a variety of behavioral and physiological responses (Curley et al. 2011). If so, changes in brain plasticity in individuals living in dysfunctional environments may be distinct from the changes of those in protective and supportive ones.

At the opposite end of the lifespan, one might consider the construct of cognitive reserve (Stern 2003, 2009), which allows cognitive function to be maintained—or minimally disrupted—in older age and can enable individuals to sustain a greater amount of neuropathological insults before they manifest signs and symptoms of cognitive decline. A number of lifestyle factors including education, work complexity, social network, and leisure activities seem to contribute to this reserve (Fratiglioni and Wang 2007; Scarmeas and Stern 2003). For instance, highly-educated individuals, even those with neuropathologic Alzheimer's disease (AD), seem to have a reduced risk of clinical manifestation of dementia (Roe et al. 2007), perhaps related to greater white matter integrity (Teipel et al. 2010). In mild AD patients with the same degree of cognitive deterioration, highly educated patients have been found to have more advanced pathological and functional brain changes (Kemppainen et al. 2008). This suggests that the clinical manifestation of advancing AD pathology is

delayed in individuals with higher educational attainment (Stern et al. 1992; Alexander et al. 1997), presumably due to more efficient compensatory, plasticity-based mechanisms against the underlying pathology. In the proposed model of plasticity across the lifestyle, adaptive network plasticity might represent the neurobiological substrate of cognitive reserve.

Link to Pathology

Acquired brain insults, such as traumatic brain injury, or certain systemic diseases like diabetes, depression, or cancer may impact the capacity for plasticity and the slopes of change in plasticity thereafter. For instance, former athletes with a history of sports concussion show altered cognitive and physiological responses more than three decades post-concussion (De Beaumont et al. 2009). These findings suggest that the efficiency of plasticity mechanisms may become impaired and the slope of decline in brain plasticity may be more pronounced in those having sustained a sports concussion. Indeed, recent evidence demonstrates reorganization of functional connectivity after acquired brain injury (Castellanos et al. 2010) that, while initially adaptive, may ultimately limit compensatory adaptations for age-related changes in brain plasticity mechanisms.

In addition to such instances, though, dissimilar starting points and different slopes of change in brain plasticity across the lifespan may represent the proximal cause for neuropsychiatric diseases, either during early developmental or aging years. Consistent with such notion, early findings suggest that the mechanisms of plasticity are aberrantly altered in neurodevelopmental and neurodegenerative disorders such as ASD and AD, respectively.

Autism Spectrum Disorders (ASD)

In a recent study, we have found compelling direct evidence for abnormal plasticity in individuals with fragile X and ASD (Oberman et al. 2010). As compared with neurotypical, age-, gender- and IQ-matched controls, adult subjects with ASD showed greater and longer-lasting modulation of MEPs following TBS (Fig. 7a). The return to baseline latency following TBS was on average between 80 and 90 min in the ASD group compared to 25–30 min in the control group. Additionally, the return to baseline latency was significantly correlated with performance on behavioral motor learning tasks. This finding was replicated in a second cohort of individuals and suggests that ASD is associated with a *hyperplastic* state.

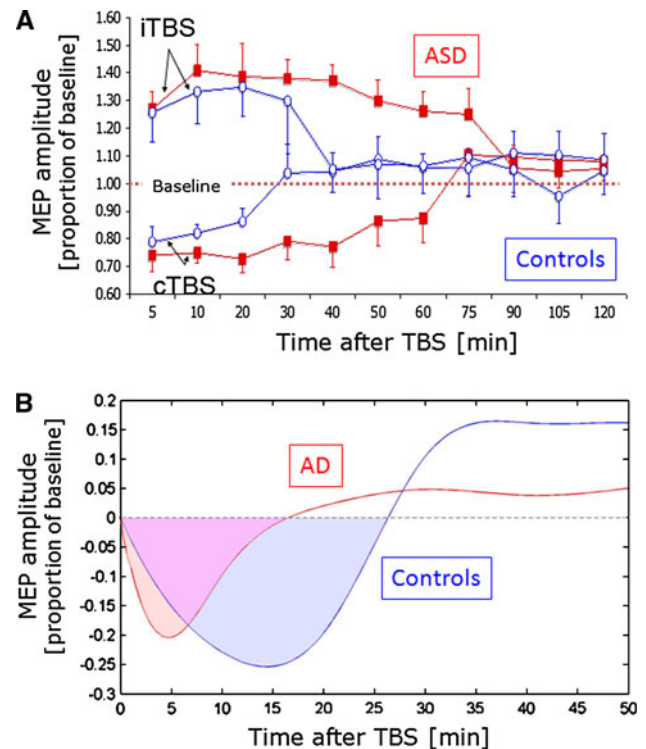
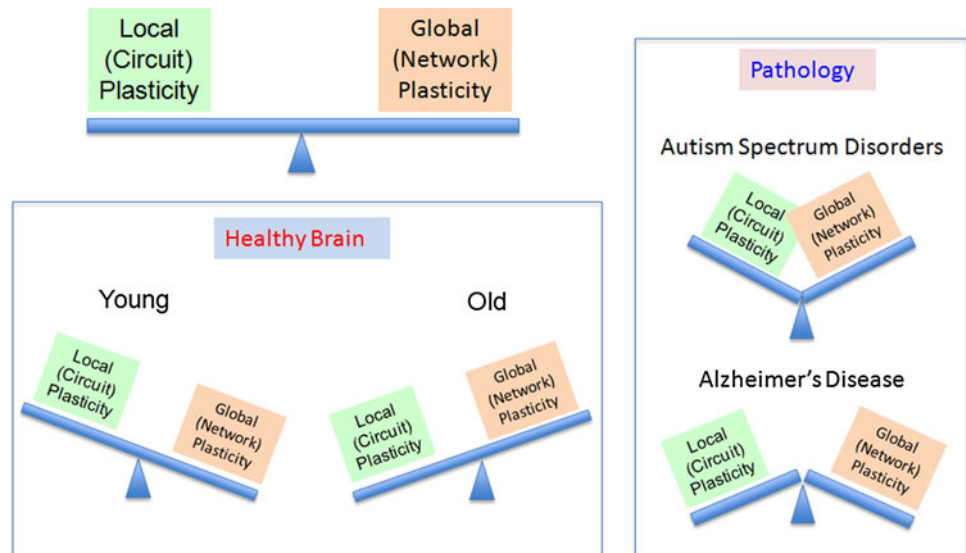


Fig. 7 Alterations of TMS-based measures of plasticity in nervous system pathology exemplified by autism spectrum disorder (ASD; **a**) and Alzheimer's diseases (AD; **b**). **a** As compared with age-, gender- and IQ-matched controls, subjects with ASD show stronger and longer-lasting modulatory responses to continuous theta burst stimulation suggesting abnormal, hyper-plastic mechanisms. This is true for both, LTP- and LTD-like plasticity. **b** As compared with age-, gender- and IQ-matched controls, subjects with mild AD show weaker and shorter-lasting modulatory responses to continuous theta burst stimulation suggesting abnormal, hypo-plastic mechanisms

Alzheimer's Disease (AD)

Recently, we collected proof-of-principle data supporting the existence of aberrant mechanisms of plasticity in patients in early stages of AD as compared with cognitively normal, age-matched elderly individuals. Results show that the duration and magnitude of the modulation of corticospinal excitability by cTBS is significantly shorter in individuals with early AD than in controls (Fig. 7b; unpublished data). This finding suggests that a *hypoplastic* state may underlay the cognitive and behavioral decline in AD. Our findings corroborate and extend those of others (for review, Freitas et al. 2011b). For example, Battaglia et al. (2007) studied neocortical (motor) LTP-like plasticity in AD and healthy individuals using a paired associative stimulation (PAS) protocol and found it to be significantly reduced in AD patients. Furthermore, Inghilleri et al. (2006) tested the effects of cortical motor modulation induced by suprathreshold high-frequency (5 Hz) rTMS and found the amplitude of MEPs progressively decreased

Fig. 8 Schematic representation of the hypothesized balance between local and network plasticity, its change across the lifespan, and its alterations by pathology



in patients while increasing in controls. This suggests impaired LTP-like plasticity.

Conclusions

The integrity of the neurophysiological mechanisms underlying brain plasticity play an important role throughout the lifespan—during developmental and aging years—in health and also in disease. In health, local cortical and network plasticity might keep a fine tuned balance which optimizes functionality. The mechanisms of plasticity, and thus the balance between local and network plasticity, change over the lifespan (Fig. 8). In the young brain, local cortical plasticity appears to be higher, and cross-sectional studies suggest that it decays with age. Longitudinal studies are critically needed as individual factors are presumed to be critical. On the other hand, in the older healthy brain, network dynamics appears to incorporate non-functionally connected regions and may provide a critical neurobiological substrate to cognitive reserve. Hypo- or hyperplastic mechanisms may set the stage for abnormal circuit development, network compensation, and, accordingly, pathological behavior. Mismatched balance between local cortical plasticity and network plasticity is also likely to be an important contributor to pathophysiology in developmental (e.g., ASD) and neurodegenerative diseases (e.g., AD). Likewise, a progressive dampening in the efficiency of plasticity mechanisms that may be induced by a number of genetic, lifestyle, and environmental factors may trigger for a cascade of events potentially leading to cognitive deterioration and even dementia in the absence of successful network compensatory strategies.

Empirically measuring plasticity may provide critical predictors and early biomarkers of disease. Additionally, in the setting of longitudinal studies, plasticity measures, as those proposed here, may allow for the assessment of the impact of genetic, biological, and environmental factors on changes in brain plasticity throughout the lifespan. Ultimately, modulating brain plasticity may minimize, delay or even prevent symptoms of brain disease. Such measures of cortical brain plasticity may thus serve to guide plasticity-based therapeutic interventions aimed at promoting brain health and cognitive function across the age-span.

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Conflict of interest None.

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