



Placental CRH as a Signal of Pregnancy Adversity and Impact on Fetal Neurodevelopment

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Early life is a period of considerable plasticity and vulnerability and insults during that period can disrupt the homeostatic equilibrium of the developing organism, resulting in adverse developmental programming and enhanced susceptibility to disease. Fetal exposure to prenatal stress can impede optimum brain development and deranged mother's hypothalamic-pituitary-adrenal axis (HPA axis) stress responses can alter the neurodevelopmental trajectories of the offspring. Corticotropin-releasing hormone (CRH) and glucocorticoids, regulate fetal neurogenesis and while CRH exerts neuroprotective actions, increased levels of stress hormones have been associated with fetal brain structural alterations such as reduced cortical volume, impoverishment of neuronal density in the limbic brain areas and alterations in neuronal circuitry, synaptic plasticity, neurotransmission and G-protein coupled receptor (GPCR) signalling. Emerging evidence highlight the role of epigenetic changes in fetal brain programming, as stress-induced methylation of genes encoding molecules that are implicated in HPA axis and major neurodevelopmental processes. These serve as molecular memories and have been associated with long term modifications of the offspring's stress regulatory system and increased susceptibility to psychosomatic disorders later in life. This review summarises our current understanding on the roles of CRH and other mediators of stress responses on fetal neurodevelopment.

Keywords: CRH, fetal, neurodevelopment, maternal stress during pregnancy, CRH receptor-1, inflammation

INTRODUCTION

Early life, especially the first 1000 days from conception to age 2, is considered as one of the most critical periods of development (1), where the foundations of optimum health, growth, and neurodevelopment across the lifespan are established. During pregnancy, the mother must activate and coordinate multiple and diverse homeostatic mechanisms to support the growing fetus, especially neurodevelopment and achieve a favourable outcome. Optimal function of this adaptation process (2), involving mediators of the neuroendocrine stress response, promotes brain plasticity and resilience. In cases of excessive prenatal adversity, optimal homeostatic equilibrium might be severely impaired (3) and (mal)adaptive responses might contribute to development of prolonged pathogenic mechanisms. Pathological signals

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Altered developmental trajectories of the fetus, can lead to acute consequences or long-term outcomes such as enhanced susceptibility to adult disease (the fetal origin of adult disease) (5). A plethora of studies identified correlations between disrupted fetomaternal symbiosis and fetal programming sometimes leading to pathological birth phenotypes including abnormal immune function (6, 7) and increased metabolic risk of the offspring (8–10). Prenatal stress has also been associated with disrupted brain programming and function, since the perinatal period is a critical period of neurogenesis where the fetal brain can be remodelled or reprogrammed (11). Such exposures to adverse early life experiences that disrupt fetal neurodevelopment are associated with offspring's increased risk for various psychopathologies (12–15).

In this review we describe current knowledge and emerging evidence about the key players involved in maternal stress responses on fetal neurodevelopment, focusing on the two distinct but interacting mediators of hypothalamic-pituitaryadrenal axis (HPA axis) responses: corticotropin-releasing hormone (CRH) and its G-protein coupled receptors (GPCRs), and adrenal glucocorticoids (GCs).

ADAPTATION TO MATERNAL STRESS, RESILIENCE, AND NEURODEVELOPMENT

In human, the period between 20 and 32 weeks after conception is characterized by rapid brain development in particular neural migration and synaptogenesis and a high rate of fetal neuronal proliferation (16); this is associated with development of the fundamental anatomical structures for the initial functioning of early neural circuits *in utero* (17). Brain neurogenesis is remarkably complex and the fetal brain tissue can be particularly plastic and vulnerable to a hostile intrauterine environment. During pregnancy, coordinated actions of hormones produced by the mother, placenta and fetus, regulate fetal neurodevelopmental processes and fine-tune brain formation (18). Members from at least three families of GPCRs, (rhodopsin, secretin and adhesion) have been identified as crucial for mediating these actions (18–20) (**Table 1**).

Fetal exposure to maternal stressors and enhanced allostatic load, can disrupt optimum brain development. Mapping pathologic pathways implicated as central mediators of the effects of early life stress in brain function, identified altered HPA axis responses and perturbed glucocorticoid signalling as well as various modifications in the fetal brain function, including impaired GPCR signalling, alterations in neurotransmission and disturbed functionality of neuronal circuits (21–26). Thus, as the phenotype responds to the intrauterine environment, these adaptations can sometimes result in long term consequences and increased risk for disease in later life (27). See Howland et al. (28) for a recent review.

HPA Axis Activation and Pathological Offspring Phenotypes

The HPA responses during pregnancy are characterized by another major source of neuropeptide secretion, the placenta, which synthesises placental CRH (pCRH) as early as 7 weeks of gestation. Placental CRH exhibits distinct responses to glucocorticoids and during gestation there is a bi-directional release of pCRH into the maternal and fetal compartments (29, 30). Acting via type 1 (CRH-R1) and type 2 (CRH-R2) receptors, CRH coordinating homeostatic challenges (31) that might be crucial in fetal maturation and providing nutritional signals that ultimately control pace of fetal development. It is thus reasonable to assume, that hormonal imbalances associated with severe or prolonged stress response, could potentially affect optimal outcomes (18). For example, it has been suggested that prenatal maternal stress signals associated with elevated levels of CRH, influence fetal growth in directions that determine gestation outcomes and alter birth phenotypes (32). Results from the ELGAN (Extremely Low Gestational Age Newborns) study suggest that extremes of pCRH expression identify risk for adverse fetal developmental outcomes: low CRH mRNA concentrations are associated with placenta inflammation and predict ventriculomegaly whereas high CRH mRNA concentrations predict motor dysfunctions (33). In addition, pCRH through cortisol stimulation might induce a state of insulin resistance and increased glucose in maternal circulation available to the fetus.

CRH as a Neuroprotective Signal and Molecular Mechanisms

A potential neuroprotector role has been suggested for CRH by promoting neurogenesis, differentiation and survival of neuronal cells (34, 35), however, abnormal intrauterine exposure to excess CRH, can affect fetal neurodevelopment and result in brain alterations resulting in cognitive and emotional deficits that persist in later life (16, 36–39). Studies on early human embryos suggest that CRH can promote survival of the neural progenitor cells (NPCs) and serve as an endogenous neuroprotector. These actions involve CRH-R1 and downstream activation of multiple kinases including PKA and CREB activation (40) as well as MAPK and PI3K signalling pathways. The latter control apoptosis of NPCs, through inactivation of proapoptotic signals such as Glycogen Synthase Kinase (GSK-3 β) that prevents degradation of β -catenin, augmenting neurogenesis (41). CRH

TABLE 1	GPCRs in	volved in	CNS	development	
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Receptor family	Ligand example	Type of receptor	Actions on neuronal tissue		
Secretin GPCRs	CRH	CRH-R1	Neurogenesis, differentiation of neuronal cells, development of neuronal circuits (21, 22)		
Rhodopsin GPCRs	Serotonin	5-HT _{1A} , 5-HT _{2A} , 5-HT _{2C}	Development of neuronal circuits Modulation of memory, emotions and cognition (20)		
Adhesion GPCRs	Oxytocin	OT-R	Myelination, anti-inflammatory actions, neuroprotection (18) Architecture and wiring of cortical and subcortical brain areas (19)		

direct neuroprotective actions can oppose the neurotoxic effects of excess glucocorticoids on neuronal progenitors (34). Additionally, CRH serves as a key modulator in adult neurogenesis and genetic disruption of CRH/CRH-R impairs hippocampal neurogenesis. Exposure of hippocampal neural stem cells (NSCs) to CRH increases proliferation, survival, and differentiation *via* transcriptional processes involving upregulation of Notch3, a crucial regulator of adult tissue NSC quiescence and maintenance (35).

CRH control of neurogenesis involves regulators of neuronal connectivity and synaptic plasticity such as brain derived neurotrophic factor (BDNF). Hypothalamic CRH is positively regulated by BDNF via a mechanism that involves the cAMP response-element binding protein (CREB) coactivator CRTC2. This transcriptional regulator serves as a bidirectional switch for BDNF and glucocorticoids to control expression of CRH (42). Studies in CRH-overexpressing mice also identified a positive feedback loop between CRH and BDNF that enhances BDNF release. This leads to improved neuroprotective outcomes under acute excitotoxic stress, with reduced neurodegeneration and neuroinflammation of the hippocampus (43). Stress-induced epigenetic modifications, resulting in BDNF methylation and decreased expression in prefrontal cortex, amygdala and hippocampus of prenatally stressed rats (44, 45), disrupt the optimum neurodevelopmental processes, with similar effects also reported in humans (46).

Fetal Neurodevelopment in States of Maternal Stress and Excess CRH

Placental CRH can cross the immature fetal blood-brain barrier where it may alter the rate of maturation of developing neuronal structures. Differentiated cortical neurons in the fetal brain express CRH-R as early as 13 weeks' gestation. Studies in humans and animals linking pCRH with early life behavioural outcomes, show a positive correlation between maternal stress and aberrant neurodevelopmental function likely related to stunting of normal neuronal growth. See Lautarescu et al. (47) for a recent review. Reduced cortical volume in the frontal and temporal lobe, in the face of elevated levels of pCRH during gestation, have been associated with both cognitive and emotional deficits in pre-adolescent children (36). Moreover, in rodents models of disease, stress-induced dendritic remodelling has been linked with increased anxiety and depressive like behaviours and with memory impairment (48). In human studies, fetal exposure to accelerated pCRH trajectories during mid-gestation was associated with child internalizing symptoms at 5 years of age (16) and prenatal exposure to high pCRH can affect infant temperament. Full term infants of mothers with lower CRH levels at 25weeks of gestation showed lower levels of distress in infancy compared with infants exposed to elevated levels of the stress hormone (49).

Placental CRH levels have been positively correlated with thinning of selective brain regions during gestation. The impact of such developmental effects on maturation of neurons and brain circuits is long-lasting as it is evident into early life: children exposed to elevated levels of pCRH prenatally exhibit significant thinning in the whole cortical mantle at age 7. The timing of the exposure to altered pCRH levels also determines region specific changes; prefrontal thinning is associated with elevated pCRH levels at early gestation whereas temporal thinning is associated with pCRH levels later in gestation (36). Experimental animal models demonstrated CRH-induced alterations in dendritic brunching, specifically, decreased branching of cortical neurons (37); altered synaptic plasticity, impaired myelin formation and decreased dendritic spine density in the hippocampal region of the offspring (50, 51). Some evidence link maternal anxiety to reduced fetal amygdala volume during the late second and third trimester of pregnancy and alterations in fetal cortical gyrification of the frontal and temporal lobes in brains of human fetuses of stressed mothers (52).

The hippocampus and the HPA axis are functionally interconnected, therefore stress alterations to the HPA axis could mediate changes in the developing hippocampus (4). Recent studies demonstrated that maternal adversity involves brain CRH-R1 activation and regulation of neuronal connectivity and developmental trajectories of the immature hippocampus (38); this leads to structural remodelling of hippocampal CA3 neurons with significant reduction of complexity of apical dendrites and spine density (53–56).

CRH-Glucocorticoid Interplay

In addition to CRH, glucocorticoids (GC), the end-product of HPA axis, control a distinct HPA-driven regulatory pathway in fetal brain neurogenesis and neural cell proliferation (57, 58). GCs rise over the course of pregnancy to further enhance pCRH release in a distinct pCRH-adrenal GC positive feedback loop (59, 60). This loop also involves placental inflammatory pathways such as RelB and NF-kB2, molecules of the noncanonical NF-kB pathway (61–63). While GCs are key mediators of regulation of fetal growth and maturation of fetal tissues and organs (64, 65), excessive GCs during pregnancy has been associated with adverse fetal outcomes including intrauterine growth restriction (IUGR), cardiovascular disease, metabolic disorders and altered HPA set point of the neonate (18, 66).

Recent studies investigating impact of natural disasters as prenatal stressors associated with abnormal offspring HPA function and development (67), identified raised cortisol as a mediator of an angiogenic phenotype and a crucial role for GCs in altering placental transcriptome, especially reduced expression of GR-regulated endocrine genes expressed in syncytiotrophoblast. Fetal glucocorticoid exposure is partially regulated by the enzyme 11β-hydroxysteroid dehydrogenase type 2 (11β-HSD2), which is abundantly expressed in the placenta and other GC -target tissues catalysing the unidirectional conversion of cortisol to its inactive metabolite cortisone (68), thereby controlling fetal exposure to maternal cortisol. Placental activity and expression levels of 11β-HSD2 have been linked with fetal programming, and downregulation or deficiency of placental 11β-HSD2 have been associated with unfavourable birth outcomes such as significant restriction in fetal growth and low-birth weight (69, 70). Fetal brain development and limbic brain areas (e.g., amygdala, hippocampus, hypothalamus) are particularly vulnerable to

overexposure to high GCs levels. Maternal cortisol levels during pregnancy can predict amygdala volume in childhood and have been associated with temperament of the offspring (71, 72). Moreover, repeated antenatal corticosteroid administration has been linked with lower density of hippocampal neurons of neonates (73), cortical thinning (74) and reduced brain maturation (75) findings that are consistent with animal studies (76–80).

Moreover, a significant rise in GCs levels in response to disease or severe or prolonged stress can impair beneficial effects of CRH on neuronal brain tissue. As numerous studies have linked CRH to neuro-damaging effects in neuronal tissue of prenatally stressed offspring, a key question remains whether CRH is causally linked to structural brain alterations or whether it is an indirect indicator of raised GCs and neurodevelopmental negative outcomes.

PLACENTAL STRESS SIGNALS AND FETAL BRAIN NEUROTRANSMITTERS AND GPCRS

In addition to its traditional roles, placenta is recognised as a functional organ supporting fetal central nervous system (CNS) development through adaptive responses to the maternal environment. Recent gene expression and network analysis in murine studies demonstrated that the placenta transcriptome is tightly interconnected with the fetal brain and inhibition of neurotrophin signalling has been identified as a potential mediator of this crosstalk. A pattern of coordinated regulation suggests an extensive network of genes encoding specific receptors and ligands predicted to regulate functional interactions between the placenta and brain (81, 82). Prenatal adverse conditions that activate placenta responses also induce changes in fetal brain neurotransmitter circuits. For example, placental inflammation and raised proinflammatory cytokines (IL6 and $IL1\beta$) has been shown to alter the neural expression of dopamine D1 and D2 receptors in brain (83). Other studies have shown that maternal inflammation in midpregnancy results in an upregulation of tryptophan conversion to serotonin (5-HT) within the placenta, leading to exposure of the fetal forebrain to increased concentrations of this biogenic amine and to specific alterations of 5-HT-dependent neurogenic processes (25, 84). 5-HT receptors in the brain are expressed in neurons and glial cells and are involved in many neurodevelopmental events (e.g., neuronal formation, connectivity and synaptic formation) identifying a possible link between the neurodevelopmental complications of the offspring upon changes in the serotonergic system. Altered 5-HT levels can disrupt the expected thalamocortical and intracortical microcircuitry and modify CRH activation via the hypothalamic 5-HT1A and 5-HT2A receptors and via the 5-HT2C receptors at the hypothalamic paraventricular nucleus (PVN), resulting in HPA axis dysregulation and altered basal activity (21). The γ aminobutyric acid (GABA) system is also sensitive in prenatal environmental insults and the latter can lead to changes in GABAergic gene expression of presynaptic GABAergic genes and GABA receptor. The function of GABA receptors also appears

sensitive to such insults: while in the adult brain GABA neurotransmission serves as an inhibitory network, in the fetal and early postnatal brain, GABA signalling is primarily excitatory. Studies revealed that offspring born to immune-challenged mothers, exhibit altered gene expression of genes encoding the two cotransporters involved in the excitatory-to-inhibitory GABA switch, leading to an increased NKCC1:KCC2 ratio and thus experience a delay in the developmental switch of GABA signalling. This might represent a link between early life environmental hits and behavioural changes in adult life (22–24, 50).

INFLAMMATION AND ACTIVATION OF THE HPA AXIS

The HPA axis responds to a wide variety of maternal signals that disrupt fetomaternal equilibrium. In particular, maternal immune activation (MIA) during pregnancy by pathogen-derived stimuli, autoinflammatory conditions or environmental irritants (85) might be an important contributor to fetal or early life neurodevelopmental disorders such as spectrum autism disorders and schizophrenia (86-88). For a latest review see Depino A., 2018 (89). The pathophysiological processes implicated in the association between MIA and adverse fetal brain programming, involve not only the hyperactivation of the maternal stress system but also inflammatory processes, elevations in mother's circulating cytokine levels (e.g. IL-6) and oxidative stress in the maternal and fetal tissues as well as, sometimes, disruption of placental optimal functions due to inflammatory conditions (90, 91). Some prenatal factors that can potentially support fetal resilience to effects of MIA, include high maternal levels of vitamin D, iron and zinc, availability of omega-3 fatty acids and efficient anti-inflammatory and antioxidant response systems. On the other hand, maternal hypoferremia and anaemia, gestational diabetes mellitus, maternal stress during pregnancy, dysbiosis of the maternal gut microbiota and maternal history of cannabinoid exposure (90) can increase the susceptibility of the offspring response to MIA. As inflammatory mediators can pass through fetal blood-brain barrier they can cause neuroinflammation leading to neuronal loss, white matter abnormalities and impaired synaptic development and neurotransmission (92). In addition, studies in rodents suggest that MIA during pregnancy can also result in profound changes in protein synthesis of the fetal brain (involving translation initiation factors and other regulators of protein synthesis) disrupting the neurodevelopment of the fetus (85). MIA due to environmental insults by endocrine disrupting chemicals (EDCS) can also affect the developing embryo by altering synaptic connectivity, neurotransmitter or neuropeptide expression, and neuronal differentiation (93).

MIA appears to induce dysregulation of the bi-directional gutbrain axis (GBA) in particular the gut microbiome and HPA axis. Pro-inflammatory cytokines can modulate the offspring's HPA axis activity, in particular CRH-R1 expression shown in rodent environmental 'two-hit' insult models (94), or modify its neuroimmune function and gut microbial colonization (95). Many studies highlight the role of gut microbiome in health (96) and disease (97). An increasing body of research indicates an association between intestinal microbes and brain function, as intestinal microbes can modulate anxiety-like behaviour and cause endocrine abnormalities in the HPA axis (98). See Morais et al. (99) for a recent review. During pregnancy, maternal microbiome composition, influenced by prenatal conditions (such as MIA), can dynamically affect the offspring and potentially program susceptibility to psychiatric disorders later in life (86, 100, 101). The latter finding is of crucial importance for fetal neurodevelopment as maternal gut microbiome promotes fetal thalamocortical axonogenesis, probably via signalling by microbially modulated metabolites to neurons in the developing brain (102). Recent findings suggest a novel interplay between maternal gut microbes and CCL2 in mediating fetal brain inflammation via raised IL-6 and placental serotonin metabolism in mediating the programming effects of prenatal stress leading to aberrant sociability and anxiety-like behaviour in adult offspring (25). Moreover, murine studies suggest that early prenatal stress disrupts maternal-to-offspring microbiota transmission and has lasting effects on metabolism, physiology, cognition, and behaviour in male offspring. Maternal vaginal microbiota appears to contribute to the long-lasting effects of prenatal stress on offspring gut and reprogramming of the developing hypothalamus associated with neurodevelopmental disorders (103).

EPIGENETIC MECHANISMS LINKING STRESS AND NEURODEVELOPMENT

Fetal neurodevelopment is associated with considerable epigenetic changes targeting a wide range of genes and molecules with diverse biological roles (104). Recent studies exploring activity of HPA axis in maternal prenatal adversity, demonstrated enhanced methylation of the promoter region of NR3C1, leading to transcription silencing (105, 106). This results in decreased GCs negative feedback and a concomitant increase in both basal and stress-activated (107) HPA activity and elevated CRH and cortisol levels in response to stressors (108, 109). Disruption of key epigenetic processes during critical periods of brain development and the modifications observed in the stress regulatory mechanism, can increase an individual's vulnerability to psychopathology later in life (110). For example, human post-mortem hippocampal tissue of suicide victims showed reduced GC receptor expression in the hippocampus of those with a history of childhood abuse (111). In previous studies, NR3C1 methylation has also been associated with internalizing psychopathology in children and adolescents (112, 113). Other perceived stressors, such as socioeconomic status, health conditions, and lifestyle can also influence NR3C1 gene regulation, revealing the complexity of environmental impacts on epigenetic modifications (114). During human hippocampal neurogenesis, exposure to GCs can lead to lasting DNA methylation changes and subsequent alterations in gene transcription (115). Similar findings have been described in guinea pigs with the intriguing finding that hippocampal changes in gene transcription and DNA methylation persist across three generations of the juvenile female offspring (116).

The impact of stress-induced methylation on CRH gene expression is not well understood. Recent studies identified positive association between prenatal maternal stress and methylation of the transcription factor binding site of CRH gene in the neonatal cord blood, maternal and placenta blood samples. Methylation in several NR3C1 and CRH CpG sites, predicted a negative correlation with birth weight (117). Maternal pCRH concentration correlates with cord blood cells DNA methylation, especially methylation of the leptin (LEP) gene promoter, and these epigenetic alterations are present into mid-childhood. As higher LEP methylation has been associated with lower BMI in childhood, these results might suggest an underlying link between pCRH and metabolic fetal programming (118). Studies on rodents reveal that chronic stress can induce epigenetic alterations that can mould the central stress response and ultimately affect gene expression and CRH transcriptional and translational activities in many brain areas, in a sex specific manner (119). Although maternal interaction is a major determinant of hypothalamic CRH expression in early life (28), data from in utero epigenetic studies are not available but required to explore the impact of prenatal interaction on later-life stress responsiveness. DNA methylation is not the only epigenetic alteration that can be detected; early life stress can also lead to histone modifications, that can modify chromatin architecture, alter transcription, and ultimately affect gene expression of candidate genes during early brain development (120-122). Chronic maternal stress can also generate major alterations in the antioxidant levels, and in the cellular pathways implicated in neurodevelopmental processes and DNA damage; a recent study demonstrated that maternal chronic stress downregulates levels of β -catenin and BDNF and upregulates GSK-3β, resulting in compromised neurogenesis in the prenatally stressed offspring (41, 123). Long-term alterations on signalling pathways interfering with the inflammatory/immune response and metabolism in the prenatally stressed offspring, have been reported especially with the neuroinflammation signalling pathway, the NFkB and p38 MAPK signalling (124). Epigenetic changes may also interfere with GPCR signalling and changes in the methylation status of genes encoding G proteins or GPCRs may result in inability of G protein signalling initiation or even in abruption of the GPCR signalling transduction. For example, in male neonates, there is a positive correlation between pregnancy anxiety and fetal methylation of the GABBR1 gene, that encodes the G protein coupled receptor subunit GABA-B1 (26). Likewise, maternal emotional stress and cortisol levels during pregnancy are associated with fetal DNA methylation of GNASXL, the extra large isoform of Gas protein involved in networks that control fetal growth and development (125). For a recent review see Cao-Lei et al. (126).

CONCLUSIONS

The placenta is pivotal in the development of the fetal brain and extensive molecular networks and pathways functionally link the two tissues. Placental adaptive inflammation and epigenetic responses to the maternal environment under the influence of prenatal stress, activate mechanisms that exert adverse roles in fetal neurodevelopment (**Figure 1**). The underlying biological



inflammatory mediators ultimately drives fetal neuroinflammation and IL-6 elevation in the fetal brain.

mechanisms only recently began to unravel, especially the roles of altered HPA responses involving placental CRH and glucocorticoids. However, the precise mechanisms employed by the fetus to protect itself from an unfavourable intrauterine environment have not yet been fully elucidated. The fetus plays an active role in its own development and efforts to establish successful pregnancy outcomes, and via developmental programming, alters the birth phenotype to adjust better to the postnatal life. This "fight response", is controlled by the HPA at multiple levels with pCRH exerting a major influence by integrating the homeostatic mechanisms that will ultimately promote adaptation to maternal adversity. Nonetheless, in cases of extremely unfavourable intrauterine conditions, the fetus can also choose to 'escape' from a hostile maternal environment, triggering the "flight response" via HPA axis activation, that will ultimately initiate parturition and lead to preterm birth. This hypothesis places pCRH in a central role in controlling the placental "clock" determining the length of pregnancy and the

onset of labour (32, 39, 127–129) but at extreme situations at the expense of optimal fetal development. The impact of HPA and pCRH activation on neurodevelopment is crucial and dissecting these actions could provide novel mechanistic (and potentially actionable) insights especially for understanding susceptibility to psychosomatic disorders later in life.

AUTHOR CONTRIBUTIONS

All authors contributed to the article and approved the submitted version.

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