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Special issue: Neurobiology of temperament, personality and psychopathology: what's next?

Title: Gut microbiome-brain axis and inflammation in temperament, personality and psychopathology

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Abstract

Full understanding of temperament, personality and psychopathology must consider biological mechanisms beyond the brain. Over 4000 species of commensurate microbiota inhabit our bodies and influence almost all aspects of human physiological function. Evidence supports bidirectional relationships between the gut microbiome and brain function. For example, gut environment directly influences limbic function via the vagal nerve, modulating affect and stress-responsivity. In turn, states of distress affect the ecology of the gut, physiologically and through behavioural alteration (diet, social interaction). Furthermore, the gut microbiome modulates the release of inflammatory molecules and hormones, indirectly affecting brain structure and function. Thus, development of gut microbiome from gestation, through birth, during childhood, adulthood and into old age is associated with temperament, personality and psychological wellbeing, including sexual differentiation in psychological function during puberty, and vulnerability to developmental, psychiatric and neurological disorders. Moreover, nutrients known to affect gut function and inflammation (e.g., fatty acids) are associated with temperament and personality in clinical and nonclinical groups. These relationships may reflect the influence of psychological traits on the microbiome by determining how an organism explores the environment, seeks reward, its social interaction and food preferences. However, it might also reflect an effect of microbiome status on psychological function. There is a need for further systematic multidisciplinary studies that integrate psychology, neurosciences, immunology and microbiology to determine the direction of these relationships and fully understand the biological basis of temperament, personality and psychopathology.

Disciplines covered: neuroanatomy, psychiatry, endocrinology, psychology, personality, immunology, gastroenterology; neuroimaging, neurotransmitter function

Introduction

Substantial advances in understanding brain networks underpinning temperament, personality¹ and psychopathology have been made over the last few decades [1,2]. However, recent evidence suggests that a full understanding of what makes us 'us', may require consideration of biological mechanisms beyond the brain [3,4]. Over 4000 commensurate microbiota species inhabit our bodies, influencing almost all aspects of human physiological function. Indeed, only 1% of the combined human and microbial DNA in our bodies is human. Given temperament and personality affect perceived threat (Neuroticism), social (Extraversion) and environmental (Openness) interaction, the resulting physiological and behavioural changes may influence the gut microbiome (i.e., the gut microbial community, including genetic components, microbial biodiversity, and their resulting functionality) [5]. This, in part, contributes to the remarkable individual differences in gut microbiome diversity and composition, which influences physical and psychological health [3,6]. In turn, the gut microbiota might also influence temperament and personality, via bidirectional communication with the brain [3,7], involving both direct (cranial nerve X) and indirect (e.g., via immune and endocrine systems) pathways [8]. Thus, the gut may represent a missing link between psychological traits and health. The current manuscript reviews evidence in this field that is rapidly gaining ground in neuroscience, psychology and mental health [9].

Gut microbiome development

The gut contains 10^{13} microorganisms with over 1000 unique bacterial species, comprising four major phyla: Firmicutes (e.g., *Lactobacillus*, *Lachnospira*, *Veillonella*, *Eubacterium*, *Roseburia*, *Ruminococcus*), Bacteroidetes (e.g., *Bacteroides*, *Parabacteroides*), Actinobacteria (e.g., *Bifidobacterium*) and Proteobacteria (several gram-negative pathogens e.g., *Escherichia-Shigella*, Gammaproteobacteria). Infants receive the majority of their initial microbiome via the mother during, and possibly prior to, birth [10]. Several factors - mode of birth (vaginal, C-section), duration of gestation, antibiotic exposure, nutrition, genetics, social and environmental interaction, trauma/stressor exposure (to name a few; [11,12]) - influence an intense period of infant microbiome development, that is intertwined with early brain [13] and immune system development [14,15]. The gut microbiome stabilises by 3 years to resemble the young adult composition [8,10], at which time it is predominantly determined by horizontal transmission (e.g., social contact [7]), but follows various age-related milestones throughout life [16].

Early infant microbiome is characterised by an abundance of *Bifidobacterium* (due to a predominantly milk-based diet). As the child explores other foods (e.g., starches) and environments, *Bacteroides* (and other genera) increase, and the microbiome becomes more diverse. Whilst transition to a *Bacteroides*-dominated community at 12 months **has been associated with** better cognitive development at 24 months [17], other genera and specific

¹ Whilst overlap between the terms temperament and personality is acknowledged, the authors view temperament as emerging earlier in ontogenetic development and continuing throughout life, whilst more complex personality traits emerge with the development of executive function and the social brain. Thus, currently, we refer to temperament in sections on infancy and childhood, whilst in sections on adulthood we refer to personality.

species remain important determinants of health (e.g., *Bifidobacterium*, *Lactobacillus*) and illness (e.g., several Proteobacteria). In adults, distinct entero-types (e.g., characterised by a predominance of *Bacteroides*, *Prevotella* or *Ruminococcus*) have been proposed that depend in part on the hosts' long-term diet (and thus, culture [16,18] and personality, e.g., openness) and show different patterns with regard to emotion processing, brain function and vulnerability to psychopathology [19].

Microbiome and early life temperament

Regarding infant (18–27 months) temperament [20], microbiome phylogenetic diversity is associated with Surgency/Extraversion [21]. In boys only, Surgency/Extraversion has been associated with overall diversity and relative abundances of certain Firmicutes (*Dialister*, Ruminococcaceae) and Bacteroidetes (Rikenellaceae, *Parabacteroides*). This latter finding was in part associated with age and diet, while the relationship between phylogenetic diversity and Surgency (Extraversion) seemed unaffected by these variables. Although the directionality of these associations is unclear, they may reflect a tendency towards high levels of positive affect, engagement with the environment, and activity [21]. In another study an abundance of *Bifidobacterium* (and *Streptococcus*) assessed at 2.5 months was associated with later Surgency assessed at 6 months [22], whilst higher *Bifidobacterium* and Enterobacteriaceae (and lower *Bacteroides*) was associated with Regulation.

Gut-microbiome-brain research is vulnerable to inter-study inconsistency on many levels, not in the least due to methodological differences in measurement of microbiota and psychological function [23]. Indeed, distinct species in the same genus differentially associate with psychological function. For example, impulsivity in childhood is positively associated with *Bacteriodes xylanisolvans*, but inversely with *Bacteriodes fragiles* [24]. Thus, associations based on phyla and genera, might be less consistent than those that group microbiota based on functional properties, for example, production of short chain fatty acids (e.g., butyrate), bile and/or pro/anti-inflammatory molecules, which influence physiology and behaviour [25].

Individual differences in study populations (sex, diet, age, enterotype) might also lead to apparently inconsistent findings. For example, girls (18–27 months) show a different pattern to boys with inverse associations between phylogenetic diversity (Shannon Diversity Index) and effortful control, and positive associations between Beta diversity and fear reactivity [21]. This is in line with animal studies showing that associations between the gut and behaviour are sex dependent [26,27], and suggest that in humans, such differentiation begins prior to puberty.

As diet diversifies and the infant brain develops, the role of the gut in regulating negative emotions may vary. Thus, in new-borns (25 days old), *Bifidobacterium* enrichment is associated with negative emotionality (*Bifidobacterium pseudocatenulatum*) and emotion regulation/orienting (*Bifidobacterium pseudocatenulatum* and *Bifidobacterium catenulatum*). However, in infants, an abundance of *Bifidobacterium* (along with lower *Clostridium*) at 2.5 months has been associated with reduced “fear bias” (i.e., attention toward fearful versus happy/neutral faces) at 8 months [28]. Similarly, in a Chinese cohort (12 months olds), *Bifidobacterium* was related with Soothability [29], whereas *Hungatella* (also

associated with constipation and eczema in children [30]) was associated with reduced Cuddliness. Indeed, several other families and species have been associated with the fear response, such as Rikenellaceae (18–27 months old girls [21]), *Parabacteroides distasonis*, *Bilophila unclassified*, and *Roseburia intestinalis* (5–7 year olds [24]), and Lachnospiraceae and *Bacteroides* (in 5–11 year olds **were associated with** activation of frontal networks in response to fear [11]). Whilst these studies highlight the possibility that gut microbiome composition from infancy directs development of temperament, clearly further **hypothesis-driven** work using standardised protocols is needed in this sparsely investigated field to replicate findings and resolve inconsistencies, control for potential confounding variables (e.g., maternal age, womb environment, diet, virulence etc; [31,32]), and clearly delineate the direction of relationships [23]. **Such work is likely to require corroborating human and animal investigations, including dietary intervention and elimination studies, as demonstrated in the case of Autism** [33].

Microbiome and personality in adults

Regarding adult personality, Gammaproteobacteria (mostly pathogenic, including *Escherichia coli*) has been associated with Neuroticism (in line with inflammatory models of neuroticism), and indeed major depressive disorder [34,35], whilst Conscientiousness was characterised by lower Proteobacteria, and higher Lachnospiraceae (butyrate-producing bacteria related to anti-inflammatory mechanisms) [36]. Bacterial diversity has been associated with openness and agreeableness, which may reflect a greater tendency for exploration (trying new foods) and social contact. Using home kits for microbiome assays and an online questionnaire in a large cohort ($n > 600$)², a composite sociability measure (extraversion, social skills, communication) was associated with a microbiome profile similar to that seen in autism: positive relationship with several genera with anti-inflammatory properties (*Akkermansia*, *Lactococcus*, and *Oscillospira*) and negative association with others implicated in systemic inflammation (*Desulfovibrio*, *Sutterella*). Neurotic tendencies were negatively associated with *Corynebacterium* and *Streptococcus* [7]. In comparison, *Streptococcus salivarius* has been associated with both emotional reactivity and externalising behaviour in children; and *Akkermansia muciniphila* with depression [24]. In older adults, *Megamonas* (Firmicutes with anti-inflammatory action via propionate production) were negatively associated with conscientiousness, neuroticism, and openness, and positively with agreeableness, whilst *Fusobacterium* (opportunistic pathogens) were negatively associated with openness and extraversion [37].

Microbiome, limbic and executive function

The varied relationship between gut microbiota and negative emotionality or fear with age may reflect differential subregion development of limbic networks involved in the response to threat, including those centred around the frontal-medial temporal connections [38] and neuroendocrine systems (i.e., hypothalamus-pituitary-adrenal axis [8]). Whilst development of limbic regions (e.g., amygdala) begins soon after conception, progressive structural and

² This study controlled for several potential confounding variables (e.g., sex, age, body mass index, birth delivery mode, type of infant feeding method, antibiotic use etc).

functional segregation between sub-regions' cortical networks continues throughout childhood and adolescence [38,39]). Influence over limbic function by the gut microbiome is supported by several animal studies [8,40] and emerging neuroimaging findings in human infants [41], children [11], and adults [42–44]. For example, Alpha diversity³ is associated with amygdala-thalamic, anterior cingulate-insula and sensorimotor-inferior parietal connectivity in 1-year olds [41], and might also affect interhemispheric connectivity in newborns [32]. In adults, bacterial microbiota diversity (particularly *Prevotella* and *Bacteroides* abundance) has been associated with insula connectivity with several brain regions [45].

Regarding executive ability, **a role has been proposed for *Bacteroides fragilis***, which protects against pathogen induced inflammation, and is positively associated with cognitive control, and inversely with impulsivity and sadness in children [24]. This species also showed weaker inverse associations with aggressive behaviour, emotional reactivity, externalizing behaviour, and incidents of family turmoil. In that study [24], impulsivity was positively associated with *Eubacterium siraeum* and *Bacterioides xylanisolvens*, whilst inhibitory control was inversely associated with *Eubacterium rectale* (a butyrate producer). In response to fear-based stimuli, *Bacterioides* enrichment was associated with a very specific activation of the medial prefrontal cortex (PFC), whilst Lachnospiraceae enrichment was related to activation in the left lateral PFC, medial PFC, precuneus/cerebellum, and deactivation in the post central gyrus, possibly reflecting frontal control over fear. Concordantly, Lee et al. [19] show that abundance of a novel Lachnospiraceae genus (PAC001043_g taxa) was associated with higher positive affect and lower negative affect. *Lachnospiraceae* is also implicated in interoception, the production of serotonin, adaptive response to stress and pain (along with other family members of the Clostridiales order, e.g., *Ruminococcaceae*) [46], and deficiency may underpin depression [47]. Inflammatory molecules [48], neuropeptides and hormones (i.e., cortisol, ovarian, oxytocin [49]) are regulated by the gut [8] and influence the limbic and executive systems [50,51].

Inflammation

As well as being the principal armament of our physiological defence system, adaptive inflammation collaborates with the microbiome to affect registration of environmental threats and opportunities, biasing perception to promote harm avoidance and facilitate recovery from illness [25]. Indeed, Gassen and Hill [25] suggest a specific adaptive role for inflammation in 'social' perception (punishment and reward) and associated personality traits. **For example, the immune system has been implicated as an important determinant of several traits including extraversion, neuroticism, conscientiousness, and emotional response (e.g., anxiety, depression, disgust), motivation, novelty-seeking, self-regulation, and aggression [25,52–54]. Some of these associations may reflect action of inflammatory molecules to inhibit the synthesis of serotonin, leading to a bias towards negative affectivity (e.g., neuroticism). That is, the presence of chronically raised inflammation, as implicated in several non-communicable diseases (physical and psychological), has been shown to alter tryptophan catabolism, reducing serotonergic synthesis and activating kynurenine metabolic pathways. This alters brain chemistry to affect development, increase vulnerability to**

³ Alpha diversity is the ecological diversity of a single sample, taking into account the number of different taxa and their relative abundances.

negative affectivity, damage tissue, and predispose to developmental, psychological and neurological disorders [52,55,56].

On-the-other-hand, dopamine agonism, which is implicated in reward processing and Extraversion, is shown to have anti-inflammatory effects [57]. Thus, brain mechanisms underpinning such traits, may indeed act to protect against chronically raised inflammation, to a certain extent. However, some evidence implies an inverse “U” curve between certain cytokines and personality [25,52–54] as well as cognitive function [58] in the general population, which may be moderated by sex, ethnicity [59], and co-occurring personality traits [60]. Thus, both very low and very high reward sensitivity is associated with elevated inflammation. Important distinctions may be between the ‘pleasure’ obtained from reward and the ‘drive’ to obtain reward [61], which are differentiated in more recent developments of reinforcement sensitivity theory (RST; [62]). BAS drive, for example, interacts with ruminative response style to increase inflammatory molecules (e.g., IL-6) in response to social stress [61]. Thus, complex bidirectional relationships exist between the many different inflammatory cytokines, tryptophan catabolites (e.g., Kynurenine, Quinolinic acid) other molecules (e.g., endocannabinoids), distinct subdomains of personality and psychopathology, that some authors have courageously attempted to unravel [52]. Mayer [63] for example, present elegant models of the bidirectional communication between the gut and the brain. Whilst progress has been made with Genome and Connectome projects, we need similar maps of the enteric and immune systems, in relation to temperament, personality and psychopathology, in order to fully understand who we are.

Systemic inflammation is in part determined by the permeability of the gut wall, and thus, microbiome composition, which regulates peripheral intrusion of endotoxins and concomitant immunological responses [64]. However, afferents from the vagal nerve also directly regulate central cytokine synthesis affecting neuroinflammation [25], which may occur in the absence of peripheral response. Thus, whilst supplementation with *Bifidobacterium longum* reduced amygdala hyperactivity in adults with inflammatory bowel syndrome, it did not affect peripheral inflammatory markers [42], suggesting the microbiome may affect limbic function irrespective of systemic inflammation [8].

Furthermore, various psychological routes to illness might operate, that are associated with distinct physiological pathways. For example, Heym et al. [65] show that harsh self-judgement (poor self-compassion), a vulnerability trait for depression, is strongly inversely related to faecal numbers of *Lactobacillus*. This may reflect an observed effect of *Lactobacillus* (along with *Bifidobacterium*) in supporting positive emotional bias [66] and reducing ruminative thoughts [67], but could also be due to the impact of raised psychological stress (e.g., through self-blame) on the ecology of the gut microbiome [68]. Brain networks implicated in self-judgment, self-awareness and self-referential processes (e.g., Insula, ACC, PFC, Amygdala) show strong bidirectional connections with the enteric nervous system (for review see [63]).

Whilst self-Judgement appeared unrelated to a general peripheral marker of inflammation known as C-reactive protein (CRP), CRP (but not *Lactobacillus* or *Bifidobacterium*) was associated with poor cognitive empathy [65], another psychometric vulnerability marker for depression (as well as autism [69], dark personality traits and personality disorders [70,71]). In

Heym et al. [65], cognitive empathy may have acted as a proxy for cognitive flexibility [72], which would hold ties to plasticity (extraversion, openness [73,74]). **In line with this**, Sumich et al. [75] report associations between plasticity and certain fatty acids with established anti-inflammatory action, whilst those with pro-inflammatory action were associated with Instability (neuroticism, low agreeableness, low conscientiousness). Intake of long chain polyunsaturated omega-3 fatty acids (associated with anti-inflammatory effects) is inversely related to reactive aggression [76] and associated traits [77].

Endocrine Function

Complex relationships exist between the endocrine, enteric and immune systems that influence psychological and physical health [78,79]. Hormones likely act as bidirectional conduits of communication between the microbiome and brain, impacting temperament, personality and vulnerability to psychopathology [79]. For example, sex steroid hormone levels in humans are correlated with gut microbial composition and diversity [80], and animal studies show transference of gut microbiota from male to female rats alters the microbiome and elevates testosterone [81]. Conversely ovarian hormones (e.g., estradiol supplementation) alter gut microbiota, as can stress hormones, such as cortisol, norepinephrine and epinephrine [21]. Thus, following puberty, sex differences in the composition of the gut microbiome become more prominent, reflecting the maturation of microbial, neurological, endocrinological and immunological systems [79,80] and may contribute to subsequent sex differences in psychopathologies [82]. **Oxytocin (released during social affiliation) regulates the immune and threat response systems, adaptively improving amygdaloid accuracy in responding to real threats, whilst attenuating response to 'non-threats'** [83]. Disentangling causal relationships between these multidirectional systems is in its infancy [23], but urgently required to fully understand mechanisms underpinning temperament, personality and psychopathology.

Pre- and probiotics

With regard to clinical trials in humans, meta-analyses suggest that whilst probiotics [84] have a significant effect on mood in clinical populations [85], their effects in non-clinical populations remain unclear [85]. Nevertheless, some positive findings exist. For example, in Petrochemical workers, probiotic yogurt (*Lactobacillus* and *Bifidobacterium* species) or supplement with several species led to improvement in depression, anxiety and stress [86]. A prebiotic [87] intervention to increase *Lactobacillus* and *Bifidobacterium* species in healthy individuals shifted attention towards more positive stimuli (dot-probe task) and lowered cortisol levels [66]. Marotta et al. [88] recently reported a trial of several *Lactobacillus* and *Bifidobacterium* species in a small group (n=38) of Italian students, in which mood (depression, anger), trait depression, temperament and personality (reinforcement sensitivity) were included as outcomes, along with several other measures (sleep, coping style etc). Clear improvements in profile of mood state (depression, anger and fatigue), but not trait depression, were seen in a group taking probiotics compared to placebo. Moreover, self-reported temperament and personality did not significantly change over the course of the study (9 weeks); unsurprising over such a short period. **Such small studies with multiple comparisons are prone to both type I and II errors. On-the-other-hand, whilst studies with**

larger sample sizes offer greater power, very large studies are at risk for spurious findings that may reach statistical significance with negligible effect sizes. There is therefore a need for further adequately powered, top-down, hypothesis-driven replication studies.

Summary and Conclusion

Whilst several associations have been identified between the gut microbiome, temperament, personality and psychopathology across the lifespan, the direction of these relationships are currently unknown. At least in part, however, they likely reflect a role for temperament and personality on diversity and composition of the microbiome by influencing social and environmental exploration during development. This may, in turn, influence vulnerability to physical and psychological clinical conditions through alteration in inflammatory environments and regulation of frontal and limbic function. However, further work manipulating the microbiome (through diet or supplementation) over longer periods of time is needed before any conclusions can be drawn about direct effects of the microbiome on temperament and personality.

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