

ly suggested by I. Feinberg, who postulated aberrant peri-adolescent pruning of synapses (resulting in either too much or too little pruning) as underlying schizophrenia⁵. In a subsequent paper, we suggested that an exaggerated pruning of synapses during adolescence/young adulthood could explain the onset of the disorder at that age⁶. This view is indirectly supported by phosphorus magnetic resonance spectroscopy studies that showed greater neuropil contraction in first episode schizophrenia⁷, which was associated with a gene-dosage effect of C4A and C4B copy numbers⁸.

While these observations may help connect several previously murky “dots” in our understanding of the pathophysiology of schizophrenia, several caveats are worth considering. First, the pathophysiology of schizophrenia may not simply be related to synapse loss. Substantive evidence show that abnormalities in myelin, neurons, oligodendrocytes, astrocytes and endothelial cells may also be involved. Human post-mortem studies that demonstrated dendritic spine loss, a proxy measure of synaptic pruning, are primarily localized to the basilar dendrites in the deeper layers of cortex, but not the entire cortex. Second, complement cascade alterations may not be unique to schizophrenia, with recent observations suggesting similar pathophysiological mechanisms in Alzheimer’s disease and bipolar disorder.

Third, genetic factors underlying C4 expression may be only one among several possible mechanisms underlying alterations in synaptic pruning. Environmental factors, including intrauterine infections, may lead to complement and inflammatory alterations via maternal immune activation. Sleep deprivation may lead to synapse elimination via microglial phagocytosis. Traumatic brain injury could result in immune and complement activation with loss of synapses. Other genetic factors besides complement component genes affect synaptic pruning, such as genes that code for gamma-aminobutyric acid (GABA) and N-methyl-D-aspartate (NMDA) receptors (all of which are implicated in risk for schizophrenia). Furthermore, OTX2, which is associated with risk for bipolar disorder, impacts timing of synapse elimination via peri-neuronal nets.

Fourth, while complement alterations may be a useful starting point in understanding the schizophrenia puzzle, we are far from developing actionable biomarkers. Peripheral alterations in complement proteins are inconsistently seen, and vary across

illness phases. Further, peripheral complement proteins do not cross the intact blood-brain barrier, and are not a proxy for complement activity in the brain. However, activated complement factors may lead to blood-brain barrier dysfunction which may further affect the progression of disease. Thus, future studies also need to examine cerebrospinal fluid samples, across prodromal, early and chronic psychotic states.

Finally, innovative studies are needed to directly demonstrate increased pruning in schizophrenia. Recent observations using a unique ligand for synaptic vesicle glycoprotein-2 showed reduced binding in schizophrenia that is interpreted as reduced synapse density⁹. These findings are awaiting replication.

Thus, many paths may lead to the hypothesized excess of synaptic pruning, and complement abnormalities may be only one such path. Further, accelerated synaptic pruning may be only one of many mechanisms underlying what we call schizophrenia, may not be unique to this illness, and may not be central to this collection of disease entities. The etiopathology of schizophrenia and related disorders is best conquered piecemeal (i.e., by identifying pathophysiologically distinct transdiagnostic subtypes, given their daunting heterogeneity). While the synaptic pruning model may be a promising step in the right direction, there are miles to go before we rest in this pursuit, and many more promises to keep.

Matcheri Keshavan¹, Paulo Lizano¹, Konasale Prasad²⁻⁴

¹Beth Israel Deaconess Medical Center, Harvard Medical School, Boston, MA, USA;

²Department of Psychiatry, University of Pittsburgh School of Medicine, Pittsburgh, PA, USA;

³Department of Bioengineering, Swanson School of Engineering, Pittsburgh, PA, USA;

⁴Veterans Affairs Pittsburgh Healthcare System, Pittsburgh, PA, USA

This work was supported by grants from the US National Institute of Mental Health (KL2TR002542, R01MH112584 and R01MH115026).

1. Sekar A, Bialas AR, de Rivera H et al. *Nature* 2016;530:177-83.
2. Sarma JV, Ward PA. *Cell Tissue Res* 2011;343:227-35.
3. Severance EG, Gressitt KL, Buka SL et al. *Schizophr Res* 2014;159:14-9.
4. Sellgren CM, Gracias J, Watmuff B et al. *Nat Neurosci* 2019;22:374-85.
5. Feinberg I. *J Psychiatr Res* 1982;17:319-34.
6. Keshavan MS, Anderson S, Pettegrew JW. *J Psychiatr Res* 1994;28:239-65.
7. Pettegrew JW, Keshavan MS, Panchalingam K et al. *Arch Gen Psychiatry* 1991;48:563-8.
8. Prasad KM, Chowdari KV, D’Aiuto LA et al. *Transl Psychiatry* 2018;8:134.
9. Radhakrishnan R, Skosnik P, Finnema S et al. *Biol Psychiatry* 2017;81:S389.

DOI:10.1002/wps.20725

Gut microbiota: a missing link in psychiatry

The gut microbiota consists of the collection of microbes within the intestine, previously considered of little influence from a mental health perspective, but now regarded as a “virtual organ” weighing up to 1.5 kg in the adult intestine and producing molecules of primary importance for brain function and psychological well-being¹.

There are more bacteria in the human intestine than there are human cells in the body, and we feed these bacteria, while in turn they play a fundamental role in maintaining our overall health. The large intestine functions like a fermenter producing a

variety of molecules, including most common neurotransmitters such as gamma-aminobutyric acid (GABA) and serotonin, the serotonin precursor tryptophan, and the short chain fatty acids butyrate, propionate and acetate².

There are a variety of mechanisms enabling the gut microbes to communicate with the brain. These include the vagus nerve, short chain fatty acids, tryptophan and cytokines³. Certain microbes can only act centrally when the vagus nerve is intact, and can no longer do so following vagotomy. Previously, tryptophan was viewed as entirely of dietary origin, while now it has been es-

established that it is also synthesized by Bifidobacteria and enters the bloodstream, becoming available for brain entry and subsequent serotonin synthesis.

The gut microbiota has been implicated in a wide variety of neurological and psychiatric disorders, including Parkinson's disease, multiple sclerosis, depression, anxiety disorders and autism⁴. Much of what we know regarding the importance of gut microbes for brain function has been derived from studying germ-free animals, which do not have a gut microbiota. Such animals have an altered central serotonergic system, decreased dendritic spines in various brain regions, lower levels of trophic factors, along with abnormal neuron formation from progenitor cells in the hippocampus, altered myelination patterns in prefrontal cortex, and a defective blood-brain barrier.

Until relatively recently, the importance of the gut-brain-microbiota axis as a fundamental component of the stress response has largely been ignored. O'Mahony et al⁵ studied the gut microbiota in a maternal separation model of depression in rats. They reported an elevation in corticosterone in such animals, together with an increase in pro-inflammatory cytokines and a decrease in the diversity of gut microbes.

The fecal microbiota was then sequenced in a depression study⁶. Forty-six patients with depression and 30 healthy controls were recruited. High-throughput pyrosequencing showed increased faecal bacterial diversity in those currently depressed, but not in a group who had responded to treatment. This suggests that increased diversity is a state rather than trait marker for depression. Despite the extensive inter-individual variability, levels of several predominant genera differed between depressed patients and controls. The former had increased levels of Enterobacteriaceae and Alistipes, but reduced levels of Faecalibacterium.

In a study conducted at APC Microbiome Ireland, depressed patients had elevated cortisol output together with decreased faecal microbial richness. When rats were given a humanized microbiota from depressed patients, as opposed to healthy controls, they developed a depressive phenotype from both a behavioral and immune perspective⁷.

Thus, there is increasing evidence that some psychiatric disorders such as depression may be associated with a gut dysbiosis, a microbial imbalance.

Several studies have investigated the microbiome composition in patients with bipolar disorder⁸. The first published study involved 115 patients and reported decreased levels of Faecalibacterium. This finding was replicated in an Austrian study of 32 patients. However, a Danish study of 113 patients with newly-diagnosed bipolar disorder compared to unaffected first-degree relatives and healthy individuals found no differences in Faecalibacterium, while Flavonifractor, a bacterial genus that may induce oxidative stress and inflammation, was associated with the disorder.

Interestingly, two recent clinical trials have demonstrated a beneficial effect of adjunctive psychobiotics in patients with bipolar disorder. One was an uncontrolled pilot study which reported cognitive improvements in 20 remitted individuals fol-

lowing three months consumption of nine different strains of Lactobacillus or Bifidobacterium. The second was a randomized controlled trial involving 66 patients who had recently been hospitalized for mania. After discharge, these patients were randomly assigned to receive 24 weeks of an adjunctive Lactobacillus/Bifidobacterium combination or placebo. Re-hospitalization rates were significantly lower in those individuals who were taking the psychobiotic. Thus, preliminary data support the view that probiotics of the Lactobacillus and Bifidobacterium genera hold therapeutic potential in bipolar disorder.

Unlike genes in human cells, we can readily change genes in our microbiota by altering diet. There is increasing evidence that a poor quality diet may bring about the altered microbiota observed in mood disorders. Narrowing of dietary diversity with reduced intake of essential nutrients can reduce the availability of substrates for specific microbial growth and this may contribute to the intestinal dysbiosis of depression and other psychiatric disorders.

Over recent decades, dietary patterns in the West and elsewhere have undergone major compositional changes, with increased intakes of red meat, high fat foods, and refined sugars. This "Westernization" of diets results in dysbiosis, which may at least partially contribute to the increasing incidence of chronic inflammatory disorders, such as depression. The Mediterranean diet is associated with lower rates of depression and impacts optimally on the gut microbiota. Preliminary evidence indicates that such a diet may have antidepressant effects.

Individuals with depression or vulnerability to depression should be encouraged to enhance a plant-based diet with a high content of grains and fibres⁹. A decreased consumption of red meat, especially of processed meat, and a regular intake of fish and fermented foods, is optimal from a mental health perspective. The intake of refined sugars should be restricted.

Incorporating the gut microbiota in our studies of stress-related psychiatric illnesses expands the range of therapeutic targets, not only for pharmacological interventions, but also for nutritional ones. This may be one of the missing links that have restricted therapeutic advances in psychiatry during the past decades.

Timothy G. Dinan^{1,2}, John F. Cryan^{1,3}

¹APC Microbiome Ireland, University College Cork, Cork, Ireland; ²Department of Psychiatry, University College Cork, Cork, Ireland; ³Department of Anatomy and Neuroscience, University College Cork, Cork, Ireland

Both authors are funded by Science Foundation Ireland (grant no. SFI/12/RC/2273).

1. Dinan TG, Cryan JF. Psychosom Med 2017;79:920-6.
2. Dinan TG, Cryan JF. Neuropsychopharmacology 2017;42:178-92.
3. Cryan JF, Dinan TG. Nat Rev Neurosci 2012;13:701-12.
4. Rogers GB, Keating DJ, Young RL et al. Mol Psychiatry 2016;21:738-48.
5. O'Mahony SM, Marchesi JR, Scully P et al. Biol Psychiatry 2009;65:263-7.
6. Jiang H, Ling Z, Zhang Y et al. Brain Behav Immun 2015;48:186-94.
7. Kelly JR, Borre Y, O'Brien C et al. J Psychiatr Res 2016;82:109-18.
8. Gondalia S, Parkinson L, Stough C et al. Psychopharmacology 2019;236:1433-43.
9. Dinan TG, Stanton C, Long-Smith C et al. Clin Nutr 2019;38:1995-2001.

DOI:10.1002/wps.20726