



# Impact of *Helicobacter pylori*-related Microbial Dysbiosis in the Pathogenesis of Metabolic Syndrome and Gastrointestinal Dysmotility Disorders

**TO THE EDITOR:** Singh et al<sup>1</sup> reviewed data on gut dysbiosis and/or interactions involved in the pathogenesis of gastrointestinal tract (GIT) dysmotility and metabolic syndrome (MetS) conditions including irritable bowel syndrome (IBS), functional dyspepsia (FD), and type 2 diabetes mellitus (T2DM). They concluded that MetS is greatly influenced by the gut, thereby metabolic disorders could begin there.

In this respect, *Helicobacter pylori* infection, a global high burden, induces GIT dysmotility (mainly abnormal gastric emptying).<sup>2</sup> Likewise, *H. pylori*-associated MetS conditions are related with dysmotility-stimulated gastrointestinal microbial overgrowth, potentially leading to bacteremia with systemic conditions.<sup>3</sup> As mentioned by the authors,<sup>1</sup> MetS occurs commonly in patients with small intestinal bacterial overgrowth (SIBO).<sup>3</sup> Gut microbiota has been connected with MetS parameters, and *H. pylori*-related T2DM, a parameter of MetS, provokes diabetic gastroparesis which may contribute to hepatic encephalopathy (HE) by altering GIT motility and inducing SIBO and subsequent bacterial translocation.<sup>4</sup> *H. pylori*-associated insulin resistance (IR), the main MetS key component,<sup>5</sup> may further provoke diabetic gastroparesis, and IR-associated diabetes is linked with bacteremia and sepsis triggering subsequent systemic conditions. In this respect, dysbiosis of the GIT microbiome, including oral and gastric *H. pylori* bacteria and several non-*H. pylori* microorganisms, is related with systemic inflammatory processes and conditions, including HE. In the setting of defensins, also mentioned by the authors,<sup>1</sup> the upregulation of human  $\beta$  defensin-1 (hBD-1) may present a biomarker of bacterial translocation involved in the pathogenesis of HE.<sup>6</sup> Moreover, *H. pylori* infection appears to induce hBD-1 mRNA expression,<sup>6</sup>

which also may contribute to the pathophysiology of the central nervous system pathologies, by altering both innate and adaptive immune system reactions.<sup>7</sup>

Regarding *H. pylori* and MetS-related FD accompanied by GIT dysbiosis, *H. pylori* eradication may reduce the risk of MetS criteria in FD patients.<sup>8</sup> Likewise, *H. pylori* and MetS-related IBS may be associated with dysbiosis, and *H. pylori* eradication therapy may decrease the risk of IBS. Interestingly, *H. pylori* infection and MetS-related dysbiosis may represent major risk indicators of Parkinson's disease (PD).<sup>5</sup> In this respect, *H. pylori* infection may affect the bioavailability of L-dopa, a dopamine precursor used for therapy of PD, by injuring the duodenal mucosa, the main place of L-dopa main absorption. On the other hand, *H. pylori* eradication seems to improve L-dopa onset plus duration time, and PD's GIT motor fluctuations. Besides, MetS and *H. pylori* infection are connected with systemic sclerosis (SSc).<sup>9</sup> SSc, by inducing GIT dysmotility and SIBO, may trigger the development of complications with high morbidity/mortality. Moreover, SSc is associated with multiple sclerosis, the latter of which is also connected with dysbiosis. Eradication of *H. pylori* infection may influence the pathophysiology of SSc and multiple sclerosis, at least in certain populations.<sup>10</sup> Important to note, however, is that the relationship between *H. pylori* infection and the aforementioned chronic diseases has not been established yet, and, thus, further large-scale studies are needed.

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