Obesity is a fast-emerging epidemic in the Asia-Pacific region, with numbers paralleling the rising global prevalence within the past 30 years. The landscape of gut diseases in Asia has been drastically changed by obesity. In addition to more non-specific abdominal symptoms, obesity is the cause of gastro-oesophageal reflux disease, various gastrointestinal cancers (colorectal cancer, hepatocellular carcinoma, oesophageal adenocarcinoma, gastric cardia adenocarcinoma, pancreatic cancer and gallbladder cancer) and non-alcoholic fatty liver disease. Abnormal cross-talk between the gut microbiome and the obese host seems to play a central role in the pathogenesis, but more studies are needed.

Keywords: obesity, gastrointestinal tract, Asia-Pacific, microbiota, disease
Emerging evidence has also implicated obesity as a cause of various GI cancers, among which are colorectal carcinoma, oesophageal adenocarcinoma, gastric cardia adenocarcinoma, hepatocellular carcinoma, pancreatic cancer, and gallbladder cancer (13). What drives carcinogenesis in obesity is still unclear, but a number of mechanisms may be important, including insulin resistance, which increases insulin-like growth factor levels and chronic inflammation. The latter is associated with an increase in the secretion of pro-inflammatory adipokines (leptin and adiponectin) and various other inflammatory cytokines (13).

Non-alcoholic fatty liver disease (NAFLD) is also closely linked with obesity and other components of metabolic syndrome. In a population-based study using magnetic resonance spectroscopy in Hong Kong, the prevalence of NAFLD was 27.3%, and the risk increased in a dose-dependent manner. In subjects having all five components of metabolic syndrome, the prevalence of NAFLD was above 80%, compared with only 4.5% in those without any component (14). Although the majority of patients with NAFLD have simple steatosis, NAFLD can progress to a more severe form, or non-alcoholic steatohepatitis (NASH), in one third of patients, which increases the risk of cirrhosis and liver cancer (15). What drives the progression from simple steatosis to NASH is still incompletely understood, and better diagnostic and prognostic biomarkers are clearly needed.

**Alterations in the Gut Microbiome Play a Central Role in Pathogenesis**

A major breakthrough in recent years has been a better understanding of the role of the gut microbiome in health and disease. Beyond being highly complex and diverse, the microbiome has a density that greatly outnumbers that of human cells by a factor of ten. While there are inter-individual variations, the composition of the gut microbiome is relatively stable and resilient to changes and may vary with the mode of delivery, age, and geographical area (16). Only limited data are available for the normal composition of the gut microbiome among Asian populations, but the available evidence indicates a difference at the phylum and genus levels (17). Obesity reduces the diversity of the gut microbiome, with a greater proportion of Firmicutes than Bacteroidetes (18), but more recent data may not support this observation (19).

The gut flora plays an important role in energy harvest from indigestible dietary nutrients, and nutrient overload can induce rapid changes in the gut microbial composition (increased Firmicutes/Bacteroidetes ratio). Dysbiosis in obesity is driven by the consumption of a high-fat diet (20), which in turn increases the systemic levels of lipopolysaccharide (LPS), promoting inflammation (termed as metabolic endotoxaemia), and increasing intestinal permeability. Other mechanisms of dysbiosis include low-grade inflammation through increases in toll-like receptor activity and the production of more short-chain fatty acids, and especially propionate, rather than butyrate and bile acid dysmetabolism (21).

A number of studies have suggested that intestinal dysbiosis is an important factor in the pathogenesis of inflammatory bowel disorders (Crohn’s disease and ulcerative colitis) and irritable bowel syndrome (21). Intestinal dysbiosis and altered gut interactions with the host through defective inflammasome sensing were recently found to play a central role in the progression of NAFLD (22). Significant microbial dysbiosis is also a signature feature of colon cancer (21).

**Conclusions**

There is clearly an emerging epidemic of obesity within the Asia-Pacific, and there is strong evidence for obesity being a major factor in diseases of the gut that have been relatively uncommon in this part of the world in the past. Alterations in the gut environment, and especially in the gut microbiome, seem to play a central role in the pathogenesis of obesity in gut diseases. Further studies are needed, especially in Asia, to better understand the cross-talk between the gut microbiome and the obese host.

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