

# Dietary Patterns Affect the Gut Microbiome – The Link to Risk of Cardiometabolic Diseases

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#### Abstract

Clusters of bacterial species within the gut microenvironment, or gut enterotype, have been correlated with cardiometabolic disease risk. The metabolic products and metabolites that bacteria produce, such as short-chain fatty acids, secondary bile acids, and trimethylamine, may also affect the microbial community and disease risk. Diet has a direct impact on the gut microenvironment by providing substrates to and promoting the colonization of resident bacteria. To date, few dietary patterns have been evaluated for their effect on the gut microbiome, but the Mediterranean diet and Vegetarian diets have shown favorable effects for both the gut microbiome and cardiometabolic disease risk. This review examines the gut microbiome as a mediator between these dietary patterns and cardiometabolic disease risk. *J Nutr* 2018;148:1402–1407.

**Keywords:** microbiome, Mediterranean diet, Vegetarian diet, cardiovascular disease, trimethylamine N-oxide, secondary bile acids, SCFAs

## Introduction

The gut microbiome (GM) plays a central role in health (1, 2). Disruption of the GM, or gut dysbiosis, is associated with an increased risk of cardiometabolic diseases, such as metabolic syndrome (MetS), cardiovascular disease (CVD), and type 2 diabetes mellitus (T2DM) (3-5). Numerous studies have reported correlations between gut bacteria and CVD risk (4, 6-11) and despite the relation between the GM and cardiometabolic disease being of great scientific interest, this area of research is in its infancy, and there are many questions that await answers. At present it is proposed that specific bacterial metabolites produced in the GM vary depending on the hosts' gut microbial environment (12), which is influenced by many factors including age, genetics, cohabitation, medication, and diet (13-15). The metabolites produced via the microbiota are also affected by diet. Therefore, the combination of foods and nutrients consumed may change the composition and functionality of the GM. Thus, the role that the GM plays in nutrient metabolism and production of primary and secondary metabolites is likely to mediate the relation between diet and cardiometabolic disease risk (12, 16–18). The objective of this review is to summarize recent evidence about how dietary patterns affect the GM and, in turn, how the GM affects cardiometabolic disease risk.

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Abbreviations used: CVD, cardiovascular disease; DCA, deoxycholic acid; GM, gut microbiome; MetS, metabolic syndrome; TMA, trimethylamine; TMAO, trimethylamine *N*-oxide; T2DM, type 2 diabetes mellitus.

There is substantial individual variation in the GM, although differences in the distinct clusters of bacterial species within the gut ecosystem (gut enterotype) exist between healthy individuals and those with chronic diseases, such as T2DM and colorectal cancer (19-21). Thus, the gut enterotype may be related to the development of disease but may also affect the response to pharmacologic and dietary interventions (7, 22–27). The production of gut-derived metabolites is reliant on the bacteria present in the GM and their function within the gut ecosystem (28–30); this is a plausible explanation for how the GM may influence disease risk. Specifically, there is evidence to support an association between disease risk and gut-derived metabolites including SCFAs, trimethylamine (TMA), and secondary bile acids, which will be a focus of this review.

Greater production of SCFAs is associated with lower risk of T2DM, chronic kidney disease, and CVD (31-36). Conversely, TMA may increase disease risk (37-39). TMA can be produced in the GM through metabolism of choline, phosphatidylcholine, and L-carnitine; it is further oxidized to trimethylamine Noxide (TMAO), a proatherogenic molecule associated with the development of CVD (40). Finally, the conversion of primary to secondary bile acids is GM-dependent (41) and the type of secondary bile acids produced depends on diet and the composition of the GM. The secondary bile acids produced may also alter the GM composition. For example, a high dietary intake of saturated fat increases bile acid secretion and bile acids in the intestine, which results in the production of hydrophobic secondary bile acids (e.g., deoxycholic acid—DCA). These hydrophobic secondary bile acids can also change the composition and structure of the GM by affecting microbial growth (42). Secondary bile acids enter the circulation and

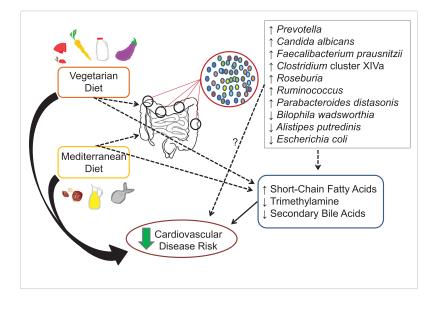


FIGURE 1 Schematic of known and proposed interactions between diet and cardiometabolic risk illustrates the known risk-reducing properties of Vegetarian and Mediterranean diets on cardiometabolic diseases and the proposed interactions between the Vegetarian and Mediterranean diets with the gut microbiome and gut-derived metabolites that can reduce cardiovascular disease risk. The thick, solid-black arrows represent a large pool of evidence to support a pathway; narrow, solid-black arrows represent an intermediate amount of evidence to support a pathway; narrow, dotted black lines represent emerging evidence to support a pathway; and narrow, dotted black lines with a question mark represent other possible mechanistic pathways. Vegetarian and Mediterranean diets can alter the presence or absence of various bacteria and, in turn, also alter the gut metabolome. The direct effect of the gut microbial environment on cardiovascular disease risk is unknown. Vegetarian and Mediterranean diets can also affect the production of gut metabolites by serving as substrate for the resident bacteria.

have been implicated in atherosclerosis, diabetes, and other cardiometabolic diseases (42-44).

There is substantial evidence showing that diet modifies the GM; however, less well characterized are the functional consequences of these modifications and the relation with cardiometabolic diseases (15). Recent clinical trials have reported the individual effects of dietary components on the GM (45–48). Fewer studies have examined the effect of dietary patterns on the GM. However, examination of dietary patterns takes into consideration the combinations and quantities of foods and nutrients consumed and their cumulative effects on health, and thus is a more robust way of understanding dietdisease relations (49, 50). The aim of this article is to summarize how dietary patterns affect the GM and, in turn, how the GM affects cardiometabolic disease risk. Figure 1 presents an overview of proposed interactions among diet, the microbiome and cardiometabolic risk.

To date, few dietary patterns have been evaluated for their effect on the GM, although several studies have examined the GM in relation to the Mediterranean diet and Vegetarian diets (22, 51, 52). Notably, these dietary patterns are also associated with lower CVD risk (53, 54). Thus, it is reasonable to hypothesize that the GM may be an important mediator between these dietary patterns and cardiometabolic disease risk. Because studies of the effect of dietary patterns on the GM and CVD risk have predominantly investigated Mediterranean and Vegetarian diets, this review will focus on these dietary patterns.

## **Vegetarian Diet**

A large body of evidence has demonstrated the cardiovascular benefits of following a Vegetarian diet (55–58), including lacto-ovo-vegetarian, lacto-vegetarian, ovo-vegetarian, and vegan diets, and it is plausible that the GM plays a role in the observed effects because the many plant-based constituents (including dietary fiber) of a Vegetarian diet provide substrates to microbes in the lower gastrointestinal tract.

Observational data show consumption of a Vegetarian diet compared with an omnivorous diet differentially affects the gut enterotype (22, 59). de Moraes and colleagues (22) examined if gut enterotype correlated with diet and cardiometabolic risk by comparing vegans, vegetarians, and omnivores. The investigators separated individuals into 3 enterotypes based on the abundance of the most prevalent bacterial strains: Bacteroides, Prevotella, and Ruminococcaceae. The authors reported that the frequency of vegans was greater in Prevotella than in the Bacteroides and Ruminococacceae enterotypes; however, the frequencies in vegetarians and omnivores did not differ (22). The authors also correlated the enterotypes with markers of CVD risk and found a favorable association between the Prevotella enterotype and LDL cholesterol; the vegans and vegetarians in the Prevotella enterotype had lower LDL cholesterol compared with omnivores. Matijašić et al. (59) reported that individuals following a Vegetarian diet had higher fecal Prevotella present and the proportion of Bacteroides + Prevotella (grouped together) relative to all other bacteria was greater, whereas the percentage of Clostridium coccoides (Clostridium cluster XIVa) was lower compared with omnivores. Prevotella bacteria have previously been associated with a carbohydrate-rich diet and produce SCFAs (22, 60) whereas C. coccoides have been found in higher concentrations in individuals with irritable bowel syndrome (61). Finally, De Filippo et al. (62) reported significant, advantageous differences in gut microbial composition in children living in rural Africa compared with children residing in Europe. The authors suggested that the African children (aged 1-6 y) had a microbial advantage due to a highfiber diet rich in cereals, legumes, and vegetables for several vears.

These observational data are supported by a clinical trial showing that after short-term (5-d) consumption of an animalbased or plant-based diet, *Bilophila wadsworthia*, *Alistipes putredinis*, and a *Bacteroides sp*. were more prevalent in fecal samples after consumption of the animal-based diet (63). These bacteria are all bile-resistant, which may be the reason they are more prevalent after the higher-fat, animal-based diet. However, a 1-mo study in obese individuals reported there were no changes in the gut enterotype after following a vegan diet, despite reduced CVD risk (64). Currently, it is unclear how long it takes for the gut enterotype to change and how stable the GM is over time. There is a need for tightly controlled studies to explore functional changes that occur in response to a vegetarian or vegan diet and how long it takes for these changes to occur.

Enterotypic differences appear to also affect the gut metabolome, which could contribute to the protective effects of a Vegetarian diet (22, 52). TMA and secondary bile acid production are dependent on the GM, and following a Vegetarian or vegan diet can reduce concentrations of both compounds; however, it is unclear whether the alteration in substrate (dietary components) or changes to the gut enterotype are associated with changes in gut-derived metabolites. Previous data showed vegan and Vegetarian diets resulted in lower concentrations of fecal secondary bile acids (65, 66) compared with omnivorous diets. Recently, David et al. (63) reported elevated secondary bile acids, including DCA, and the bacterial gene expression involved in their production, including bile salt hydrolases and sulfite reductases, after 5 d of consumption of an animal-based diet compared with a plant-based diet. DCA has been shown to reduce intestinal barrier function, increase inflammation, and promote liver cancer (67). Evidence suggests a plant-based diet may create a different enterotype compared with an omnivorous diet and promote lower production of secondary bile acids.

Reduced concentrations of TMA have also been reported subsequent to following a vegan diet compared with an omnivorous diet (52). Koeth and colleagues (52) provided vegan/vegetarian and omnivorous participants with oral carnitine and reported that vegans and vegetarians had a reduced capacity to produce TMA compared with the omnivores. Vegan and Vegetarian diets are low in choline, phosphatidylcholine, and L-carnitine owing to the absence of red meat and fish; therefore, they would be expected to produce less TMA and TMAO. However, the finding that vegans/vegetarians produce a lower amount of TMA postcarnitine challenge suggests that a Vegetarian diet may favorably alter the GM and create a microenvironment that does not metabolize carnitine into TMA.

In summary, the GM may be an important mediator in the reduced cardiometabolic disease risk associated with a Vegetarian diet, but more studies are needed to understand the mechanism. A plant-based diet can favorably affect the GM through changes in bacteria, such as increases in *Prevotella* taxa, and reduction of harmful metabolites, such as TMA, but the effect on gut enterotype and SCFA production requires further study. The key question that follows is mechanistically how changes in the GM modulate risk factors for cardiometabolic disease.

## **Mediterranean Diet**

Prominent components of the Mediterranean diet include fruits, vegetables, legumes, olive oil, nuts, seafood, and wine. Evidence demonstrates that this eating pattern is associated with a lower risk of CVD and that adherence to a Mediterranean diet can affect gut microbial patterns and the fecal metabolic profile of microbial-derived phenolics (51, 68–70). Thus, the GM may partly mediate the relation observed between the Mediterranean diet and reduced risk of CVD.

Emerging research suggests that following a Mediterranean diet is associated with a more-favorable GM composition (68, 69). Mitsou and colleagues (68) examined the association between the Mediterranean diet and gut microbiota characteristics in an adult population. The authors reported that participants with higher Mediterranean diet scores had a lower fecal Escherichia coli count, a higher ratio of Bifidobacteria to E. coli, and increased concentrations and prevalence of Candida albicans. Gutiérrez-Díaz et al. (69) also investigated the association between the Mediterranean diet and the GM, but reported that better adherence to a Mediterranean diet was associated with higher Faecalibacterium prausnitzii and Clostridium cluster XIVa compared with lower adherence. The authors note that the Clostridium cluster XIVa group contains butyrate-producing species and F. prausnitzii is a known butyrate producer, therefore suggesting the Mediterranean dietassociated GM may be made up of more beneficial bacteria. A Mediterranean-style diet has been shown to lower the risk of CVD (53), which may be partly dependent on the GM. Haro et al. (70) investigated the GM differences between individuals with coronary heart disease with and without MetS and obesity from the CORDIOPREV study, following a Mediterranean diet or low-fat diet for 2 y. This prospective randomized trial showed that consumption of a Mediterranean diet or a low-fat diet may restore some bacterial populations that are reduced in MetS patients with obesity (70). In addition, obese subjects with MetS in the Mediterranean diet group had an increased abundance of Roseburia and Ruminococcus genera, Parabacteroides distasonis, and F. prausnitzii; the low-fat diet did not change these genera (70). Cross-sectional studies suggest that the Mediterranean diet may be associated with favorable changes in the GM, and these observational data are supported by a prospective trial that showed reduced gut dysbiosis in obese individuals with MetS. However, the implications of these shifts in the GM for cardiometabolic health still need to be elucidated. Thus, more evidence is needed to understand the microbial shifts that are important for a healthy GM, and how this, in turn, affects cardiometabolic disease risk and management.

Following a Mediterranean-style diet is associated with increased concentrations of SCFAs (51). Two studies reported that individuals with a higher Mediterranean diet score had higher concentrations of fecal SCFAs (51, 68). One study reported a greater ratio of acetate to other SCFAs with greater adherence to a Mediterranean diet (68) and the other study reported that individuals with higher Mediterranean diet adherence scores had increased SCFAs (51). Although some observational data show greater SCFA concentrations are associated with obesity, this is believed to be the result of an oversupply of energy, thus providing gut microbes with more substrate for SCFA production (71). In conditions of positive energy balance, excessive energy is consumed and, therefore, the GM is also provided with more energy than needed, and works to metabolize as much of the substrate as possible. This suggests SCFAs are not causally associated with obesity. Clinical trials have shown that intestinal SCFAs are correlated with improvements in blood lipid profile, glucose homeostasis, and inflammation (72-74), although it may take weeks to months for these changes to occur (73, 75, 76). Therefore, based on observational evidence, consumption of a Mediterranean diet could be a promising intervention to increase SCFA production and reduce cardiometabolic disease risk.

The Mediterranean diet may also reduce circulating choline metabolites, such as TMAO, and therefore reduce CVD risk.

De Filippis et al. (51) reported that higher adherence to a Mediterranean diet was associated with lower TMAO concentrations. This association was also observed in a subpopulation of the PREDIMED study. Guasch-Ferré et al. (77) examined the relation between plasma concentrations of 5 metabolites in the choline pathway (TMAO, betaine, choline, phosphocholine, and  $\alpha$ -glycerophosphocholine) and major CVD endpoints in individuals at risk for CVD. The authors reported that participants classified in the highest metabolite quartile and assigned to a low-fat control diet had a higher risk (HR: 2.37; 95% CI: 1.34–4.18) of CVD than participants in the lowest metabolite quartile and assigned to a Mediterranean diet. The large proportion of plant-based foods in the Mediterranean diet may alter the GM to produce less TMA and favorably change the metabolic profile.

The dietary pattern consumed affects the composition of the GM through the provision of substrate for the gut microbes. Various bacteria prefer different energy sources and the type of foods consumed promotes the residency of bacteria that favor the energy substrate provided. This results in gut-derived metabolic products that can alter the structure and function of the GM and affect disease risk. Moreover, changes in GM that have emerged with adherence to a Mediterranean diet suggest it may provide important dietary components that can improve GM functionality with consequent benefits on the host. Future research is needed to further investigate how the Mediterranean diet affects gut enterotype. The present evidence to date suggests that the Mediterranean diet increases SCFA production, metabolites that have been shown to reduce the risk of cardiometabolic diseases.

# Conclusions

Vegetarian and Mediterranean diets both reduce cardiometabolic risk and favorably change the GM. Current knowledge suggests that consumption of a Vegetarian or Mediterranean diet generates favorable shifts in the microbes in the lower gastrointestinal tract and beneficial gut-derived metabolites may be a product of these alterations. The temporality of this relation is not well understood and it may be bidirectional. The composition of the GM affects the gut-derived metabolic products and these metabolites may also affect the microbial community and gut enterotype. Examination of the gut enterotype, microbial functionality, and gut-derived metabolites could provide a better understanding of the changes that occur with specific dietary patterns. The entire gut environment, including present microbes, relative abundance of microbes, microbial gene expression, and microbial metabolites, is important to consider relative to the link between the GM and disease risk. However, the majority of evidence to support the correlations between these diets and changes in the GM composition and metabolites is primarily derived from epidemiologic research. Further research is needed to increase our understanding of how dietary patterns alter the GM and, importantly, how alterations in the GM mechanistically modulate risk of cardiometabolic diseases.

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