

Interactions between nutrition and the intestinal microflora

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The human gut is well colonized by a wide variety of bacteria. In the industrial nations, the stomach and small intestine have a sparse flora protected by gastric acid and a highly propulsive motility. The flora of the small intestine increases along its length and particularly in the terminal ileum (Drasar & Hill, 1974). The large intestine has a very rich, mainly anaerobic, flora capable of fermenting carbohydrate and protein, and metabolizing a wide and diverse range of endogenous and exogenous molecules such as bile acids, fats and drugs. In the less-developed countries the small intestine has heavier populations of bacteria and these may be a factor in the higher occurrence of diarrhoea and malnutrition (Gorbach *et al.* 1970; Gracey, 1979).

The interactions between nutrition and the intestinal microflora are complex. Bacteria in the gut may affect digestion and absorption, the products of bacterial fermentation may provide nutrients or affect the well-being of the host, but diet may also affect the survival and metabolism of the bacteria. The products of bacterial metabolism may be beneficial, for example butyrate which may have anti-neoplastic properties (Augeron & Laboisse, 1984), or they may be potentially harmful, for example bile acid metabolites which may be co-carcinogenic (Owen *et al.* 1987). The interaction between the intestinal microflora and nutrition begins at birth and changes with age and disease.

NEONATAL MICROFLORA AND NUTRITION

At birth a baby encounters bacteria for the first time and quickly develops a commensal microflora on its skin and along its gastrointestinal tract. The factors which determine the initial colonization are not fully understood but must relate to environment, including the birth canal, and to diet. The faecal microflora of babies fed exclusively on mother's breast milk differs in many ways from that of babies fed on formula milk derived from cow's milk. The faecal flora of a breast-fed baby is more likely to have bifidobacteria and lactobacilli as the predominant organisms with lower amounts of Enterobacteriaceae and few bacteroides species (Bullen *et al.* 1977; Balmer & Wharton, 1989a). The bottle-fed baby has a faecal flora which resembles the adult more closely, with more Enterobacteriaceae and streptococci and more bacteroides species. This difference is not always seen, however. Simhon *et al.* (1982) could find no difference in the microflora of breast-fed and bottle-fed babies in a study in London and suggested that this may be due to a difference in the obstetric practice resulting in different environmental factors which influence initial colonization. The difference between microflora of breast-fed and formula-fed babies is reflected in the profile of bacterial fermentation products in faeces, with a lower faecal pH (Bullen *et al.* 1977) and a more predominant acetic/lactic acid-type fermentation in breast-fed compared with formula-fed babies (Bullen *et al.* 1977; C. A. Edwards, S. F. Balmer, A. P. Parrett & B. A. Wharton, unpublished results).

The dietary factors which may influence the different colonization of the gut of

breast-fed and formula-fed infants are numerous. They include the buffering capacity, the casein/whey and phosphate content, and the concentrations of oligosaccharides, bifidus factor, human proteins such as secretory IgA and lactoferrin, and other micronutrients such as Fe and nucleotides. The exact role of each of these factors is difficult to study in isolation *in vivo* but modern formulas now mimic human milk in many ways with very little real change in the faecal flora. Balmer & Wharton (1989*a,b*) and Balmer *et al.* (1989) have carried out several studies looking at the effect of dietary factors in formula milk on the faecal flora and found that whey-protein-predominant formula produced a higher prevalence of bifidobacteria and less bacteroides species than casein-predominant formula. The role of the Fe content of milk and the role of lactoferrin was less clear. Human milk contains less Fe than formula milks but it is more efficiently absorbed. Fe which remains in the gut is available for the growth of potential pathogens. Much of the free Fe in the gut of the breast-fed baby is probably bound to lactoferrin and is unavailable for bacteria. Bifidobacteria and lactobacilli do not need significant amounts of Fe for growth (Archibald, 1983). However, although whey formula without Fe reduced levels of enterococci and clostridia (Balmer & Wharton, 1991), the faeces of babies fed on low-Fe formula was more dissimilar to those of breast-fed babies than those of babies fed on a whey formula with Fe (Mevisen-Verhage *et al.* 1985; Balmer & Wharton, 1991). The nucleotides in human milk may affect the gut microflora either indirectly by increasing Fe absorption in the small intestine (Faelli & Esposito, 1970; McMillan *et al.* 1977) or by stimulating the growth of bifidobacteria. Addition of a selection of nucleotides to *in vitro* cultures increased the growth of bifidobacteria and when added to infant formula have produced a faecal flora more similar to that of a group of breast-fed babies (Gil *et al.* 1986). Other dietary factors such as the concentrations of IgA or oligosaccharides have not yet been investigated in much detail. The differences in faecal flora must relate to a whole range of dietary differences between human and formula milk and not to a single element. However, they do indicate the importance of diet in the establishment of the gut microflora and are related to a difference in the gastrointestinal infection rate in babies. Howie *et al.* (1990), showed that breast-fed babies had a lower incidence of gastrointestinal infections than formula-fed babies even after compounding factors such as social class had been accounted for.

In addition, the colonic microflora may also be important in salvaging unabsorbed carbohydrate and in cycling N in the neonate. Premature babies may have insufficient lactase (*EC* 3.2.1.108) in the small intestine and, thus, significant amounts of unabsorbed sugar enter the colon. This sugar is fermented by the colonic microflora to short-chain fatty acids (SCFA) and H₂, which are absorbed, preventing osmotic diarrhoea and conserving energy (MacLean & Fink, 1980; Mobassaleh *et al.* 1985; Kien *et al.* 1987). Protein digestion by the colonic bacteria and absorption of N may also occur in the colon of neonates (Heine *et al.* 1987). This dependence on the colonic microflora for nutrient absorption indicates a need to consider the effects of antibiotics on nutrition in small babies (Bhatia *et al.* 1986).

Little is known of the factors which affect the changes in the microflora during weaning. The flora develops towards the adult flora, the bacteria becoming more numerous and the ecosystem more complex with a greater predominance of *Escherichia coli*, streptococci, clostridia and bacteroides species (Bullen *et al.* 1976). Studies in rats have suggested that the exposure of the microflora to a particular substrate at weaning may determine the response to that substrate in adult life (Armstrong *et al.* 1992). This

study in rats looked at complex carbohydrates but the same could be true for other dietary components and endogenous secretions.

ADULT MICROFLORA AND NUTRITION

The normal adult microflora and its fermentation pathways and patterns have been described elsewhere in this symposium. In the present paper I have concentrated on the consequences of the bacterial activity and how this relates to nutrition.

Effect of diet on bacteria

Bacteria in the human colon vary substantially from person to person both in the species present and in their fermentation capacity and product profile. The capacity of a person's microflora to ferment different carbohydrates depends on past diet and the species of bacteria present. The faecal flora of adults is remarkably stable, with many studies looking at the effect of dietary change on the bacterial populations showing no real effect (Bornside, 1978; Hill, 1981). As some dietary fibres increase stool output and colonic content turnover, they increase bacterial turnover and populations. Indeed, for some fibres this increased bulk of bacterial cells is the major component of the increase in stool weight (Stephen & Cummings, 1980). The colonic fermentation capacity, rate of fermentation and range of SCFA and gaseous products are also altered by feeding fermentable carbohydrates. *In vitro* cultures of human faeces supplemented with starch, wheat bran and oat fibre produced substantial amounts of butyrate whereas arabinoglucan, ispaghula, guar gum and starch produced large proportions of propionate (Englyst *et al.* 1987; McBurney & Thompson, 1987; Adiotomre *et al.* 1990; Edwards *et al.* 1992a,b; Weaver *et al.* 1992). These patterns were also demonstrated in the faeces of rats and other animals fed on similar fibres (Mallett *et al.* 1988; Topping *et al.* 1988; Goodlad & Mathers, 1990; Edwards & Eastwood, 1992).

The polysaccharidase enzymes required for the breakdown of some carbohydrates are subject to dietary regulation; their activity is induced by exposure to the substrate and expression may be repressed by products of the fermentation reaction (Salysers & Leedle, 1983). Florent *et al.* (1985) intubated the caecum of human volunteers and fed them on lactulose for 1 week. They found that the pH of the colon was reduced and the pattern of fermentation was changed at the end of the week, with a more rapid degradation of lactulose, a faster accumulation and clearance of intermediates and decreased H₂ production. Other researchers report increased capacity to ferment fibres after ingestion for 1 week (Read & Eastwood, 1992) and studies in rats indicate at least 4 weeks of feeding is necessary before full fermentation capacity is achieved (Walter *et al.* 1986).

Nutritional consequences of colonic fermentation

The major consequences of carbohydrate fermentation in the colon are the loss of water-holding capacity (WHC) of certain fermentable dietary fibres, production of SCFA and gases, decreased pH, release of bound molecules from dietary fibre, production of bacterial cells and, hence, changes in the expression of other bacterial enzymes that have important physiological implications.

The most obvious effect of colonic bacteria on a food is the fermentation of dietary

fibre and resistant starch. The impact of bacterial fermentation on the physiological effects of these can be seen when fermentable substrates are eaten by people taking antibiotics. Kurpad & Shetty (1986) showed that fibre had a much greater effect on stool output when subjects were also taking antimicrobial agents. The capacity of individuals to ingest lactulose without an increase in stool output was also decreased by ampicillin (Rao *et al.* 1988). The WHC of a dietary fibre is an important factor in its effect on stool output. However, many dietary fibres with high WHC have very little effect on stool output. This is because of bacterial degradation of the fibre in the colon. The post-fermentation WHC (measured *in vitro*) is the best indicator of action on stool output (McBurney *et al.* 1985).

An increase in stool output has several nutritional implications. Increased faecal mass includes an increased loss of N, energy, water and electrolytes. There may be an increased loss of bile acid metabolites and other potential toxins which may decrease the risk of mucosal damage and alter the bile acid pool.

The SCFA produced during carbohydrate fermentation also have several possible nutritional actions. They are rapidly absorbed by the colonic mucosa promoting water and Na absorption (Ruppin *et al.* 1980) and preventing osmotic diarrhoea in situations of small intestinal malabsorption of sugars. The colonic mucosa may use the SCFA as a preferential energy source, especially butyrate (Roediger, 1982). Very little butyrate is seen in the blood draining from the large intestine and propionate is metabolized in the liver, leaving acetate as the only SCFA to be seen in the systemic blood under normal conditions (Cummings *et al.* 1987). The SCFA may also have several other effects in the colonic lumen. SCFA in the colonic lumen may increase cellular proliferation of the colonic epithelia (Sakata, 1987) and may increase proliferation of the mucosa of the small intestine by a blood-borne factor (Sakata, 1989). Butyrate has been shown to stimulate differentiation of cancer cell lines (Augeron & Laboisse, 1984). SCFA in the lumen may also have effects on colonic motility but their exact role on propulsive or segmenting contractile activity is unclear. In isolated rat colon muscle strips propionic and butyric acid stimulated contraction (Yajima, 1985), whereas in an isolated rat caecal/colon model a mixture of SCFA decreased overall colonic motor activity (Squires *et al.* 1992). SCFA have also been shown to dilate arterial capillaries and may, thus, affect blood flow and absorption rate from the colon (Mortensen *et al.* 1990).

Propionic acid is reported to have an influence, in the liver, on gluconeogenesis and cholesterol synthesis (Anderson & Bridges, 1984; Chen *et al.* 1984) but the concentrations necessary for these actions may not be achieved *in vivo* under normal conditions (Illman *et al.* 1988). The metabolism of SCFA by the tissues may represent a significant component of the daily energy intake up to 7% (Cummings, 1981) and the energy value of fermentable dietary fibre may be from 0 to 17 kJ/g (0 to 4 kcal/g) depending on the calculation method (Wisker & Feldheim, 1992). The energy gained by utilization of SCFA may, in some part, be offset by the increased faecal losses that often occur with the intake of these dietary fibre sources (Wisker & Feldheim, 1990).

The fermentation of dietary fibre sources by bacteria may also release sequestered molecules such as Ca and bile acids and these may then be available for absorption, in some cases after further bacterial metabolism. Bacteria deconjugate and dehydroxylate bile acids altering the composition and properties of the bile acid pool and also producing possible colonic cancer promoters. Bile acid metabolites and bacterially hydroxylated fatty acids may cause secretion (Mekhjian *et al.* 1971) and stimulate colonic motility

(Kirwan *et al.* 1975; Spiller *et al.* 1986). SCFA are potent inhibitors of pathogenic bacteria (Fay & Faires, 1975) along with deconjugated bile acids (Floch *et al.* 1972) and so the commensal bacteria help to prevent gastrointestinal infection.

It is generally believed that on a mixed diet most fermentation takes place in the proximal colon of man but experiments in rats suggest that some non-starch polysaccharides, such as xanthan and karaya, which are slowly fermented may spread the production of SCFA into the more distal colon where they may have less effect on colonic cellular proliferation (Edwards *et al.* 1992*a,b*) but may promote an increase in faecal water excretion (Edwards *et al.* 1990). Some fibres such as wheat bran, which were shown to produce significant amounts of butyrate *in vitro*, may be fermented mainly in the proximal colon so that butyrate in the distal colon, where cancer and colitic disease is more common, is unaffected (Edwards & Eastwood, 1992). Other fibres such as ispaghula, which are more often associated with propionic acid production, may in fact increase distal colonic butyrate to a greater extent (Edwards & Eastwood, 1992). It has recently been shown that rectal infusions of SCFA into the colons of patients with active colitis improved symptoms and disease activity (Breuer *et al.* 1991).

Rapid fermentation in the colon, as with lactulose or some highly fermentable fibres, may produce a substantial decrease in colonic pH which may have several effects. Bile acids may be precipitated, reducing their absorption and actions on the colonic mucosa and the 7- α -dehydroxylase enzyme will also be inhibited. Bacterial production and absorption of NH₃ may be inhibited (Swales *et al.* 1970; Vince *et al.* 1978) and colonic cellular proliferation may be stimulated (Lupton *et al.* 1988).

Increased bacterial populations may result in increased activity of certain enzymes such as β -glucuronidase, β -glucosidase (EC 3.2.1.21) and azoreductase (EC 1.6.6.7) which are involved in the metabolism of exogenous molecules such as drugs and possible carcinogens (Rowland & Mallett, 1990). β -Glucuronidase and β -glucosidase may reactivate toxins previously deactivated in the liver (Mallett & Rowland, 1990). Fermentable carbohydrate sources often increase the activity of these enzymes (Rowland & Mallett, 1990) but the effects are not always consistent and no definite role for the enzymes in carcinogenesis has been proved. However, in rats fed on carcinogens as well as fermentable fibre tumour yield was increased in the colon (Jacobs, 1990). This may be due in part to the stimulation of cellular proliferation by SCFA as well as an increase in the active carcinogen.

CONCLUSION

The intestinal microflora and the nutrition of the host have several complicated but important interactions which start at birth and develop as the populations and number of species increase, the ecosystem becomes more stable and new bacterial substrates are ingested. Some of these interactions are beneficial to the host but some may be detrimental. However, the activity of the bacteria in the human colon is complex and very little understood at present.

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