

The specificity of the interaction with intestinal bacterial fermentation by prebiotics determines their physiological efficacy

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The concept of prebiotic food ingredients is an important recent development in nutrition. The concept has attracted a great deal of attention, and many food ingredients (mainly dietary carbohydrates) have been claimed to be 'prebiotic'. It is emphasised that in order to be called prebiotic, a compound should be: (1) non-digestible; (2) fermentable; (3) fermentable in a selective way. These properties should be demonstrated in human volunteers in at least two independent dietary intervention trials. On the basis of published and unpublished results, it is shown in the present paper that the way in which a prebiotic influences intestinal fermentation is the key to its physiological properties. This statement is illustrated mainly by considering an established group of prebiotics, the $\beta(2-1)$ fructans. These linear molecules show a strong discontinuity in physicochemical properties as the chains become longer. The $\beta(2-1)$ fructans with a chain length of up to ten monomer units are very soluble and are particularly 'bifidogenic'. Longer chains (ten to sixty-five monomer units) are poorly soluble in water, they have less pronounced bifidogenic properties, and they are fermented more slowly. It was observed that a combination of short-chain and long-chain fructans (Synergy1) is physiologically (for example, increasing mineral absorption, suppressing carcinogenesis, modulating lipid metabolism, etc) more active than the individual fractions. A possible mechanism is described in the present review. From an in-depth overview of the literature it is confirmed that for prebiotic action, the 'selectivity principle' for intestinal fermentation is determinative for the type and for the efficiency of physiological activity. It is confirmed that prebiotics act through their influence on intestinal fermentation.

Prebiotics: Intestinal fermentation: Colonic bacteria: Inulin

Introduction

Prebiotics

Since the introduction of the term prebiotics (Gibson & Roberfroid, 1995), a vast amount of literature related to the topic has been generated. Prebiotics have been defined as non-digestible food ingredients that affect the host beneficially by selectively stimulating the growth and/or activity of one or a limited number of bacteria in the colon, thus improving host health. As research has progressed, the definition has been fine-tuned.

There are three criteria that need to be fulfilled for a food ingredient to be designated a prebiotic.

First, the food ingredient must be non-digestible by host enzymes. Food ingredients need to be resistant to attack by

digestive enzymes, such as carbohydrases from the brush border or those of pancreatic origin; as such, they remain available to the intestinal flora. The non-digestibility can be demonstrated by means of tests *in vitro*, using intestinal scrapings from various animals (pigs, rats, birds, fish). For human purposes, the best way to demonstrate non-digestibility is with ileostomised volunteers (Ellegard *et al.* 1997).

Second, the food ingredient must be fermented in the gastrointestinal (GI) tract. This can best be shown by means of fermentation tests *in vitro*. These tests show whether an ingredient is fermented, but *in vitro* models alone cannot be considered adequate for the study of the complex ecosystem of the colon (Newton *et al.* 1998).

Third, there must be selectivity in stimulation of the intestinal flora and of metabolic activity. This criterion is of

Abbreviations: ACF, aberrant crypt foci; DP, degree of polymerisation; DP_{av} , average degree of polymerisation; GI, gastrointestinal; HP-inulin, high-polymer inulin; OF, oligofructose; RS, resistant starch; SCFA, short-chain fatty acids; ST-inulin, native inulin; SYN1, Synergy1.

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utmost importance. A prebiotic ingredient, by definition, is not available to all bacteria of the intestinal ecosystem. Prebiotics must be particularly readily available to some groups of bacteria that are not related to a diseased state of the intestine (of which lactobacilli and bifidobacteria traditionally are considered indicator organisms). There are indications that this selectivity criterion originates in the genome of bacteria. Certain colonic bacteria may have specific transporters for certain types of prebiotics. This has been demonstrated with *Lactobacillus acidophilus* for oligofructose (OF; Barrangou *et al.* 2003). As a result, a prebiotic is less available to other groups of bacteria such as proteolytic bacteria, of which several are pathogenic. It is often the case that stimulating the growth of certain groups of bacteria allows them to take a more dominant position in the GI ecosystem, which they substantiate by producing metabolites, including bacteriocin-type products, or compounds related to quorum-sensing cascades.

A demonstration of the fulfilment of this criterion is much more difficult, and can be achieved only by means of repeated human dietary intervention studies. Whereas data from studies *in vitro* can give some mechanistic indications (production of metabolites such as short-chain fatty acids; SCFA), they cannot be fully representative of what happens in the human (or animal) GI environment. Data from experimental models cannot be extrapolated to man, as the GI architecture of different species of animals is dramatically different.

Prebiotic activity

The primary 'effect of prebiotics' (prebiosis) is an interaction with the GI ecosystem, which basically is an interaction with the fermentative capacity of the GI ecosystem. It is thought that all physiological effects that take place following prebiotic consumption originate in this altered fermentative functioning of the GI ecosystem. The initiation of physiological change originating in an altered fermentative capacity of the gut could happen through microbe–microbe interactions, host–microbe interactions and/or host–bacterial metabolite interactions. The latter may have the most important impact. The main metabolites are the SCFA: acetate, propionate and butyrate, which are all biologically active compounds (Cummings *et al.* 1995). Prebiotics can easily double the pool of SCFA in the GI tract, and the molar ratio between the different SCFA is altered (Djouzi & Andrieux, 1997; Femia *et al.* 2002).

This process takes place in the intestine, having a surface of about 300 m² in man, compared with 2 m² for the skin or 100 m² for the lungs. The lining of the GI tract is continuous with the external part of the body. As such, the surface of the intestine is the largest interface between our body and the environment. Via this large interface, we are in intimate contact with a very dense microbial ecosystem composed of up to 10¹² bacteria/ml, which is, volumetrically, about the highest possible microbial population one can be in contact with. An improvement of the GI ecosystem thus represents an important improvement of the external environment in which we live. This improvement consists in a reduction of the numbers of pathogens and in relatively saccharolytic activity. Increased saccharolytic activity (endproducts are

SCFA, lactate and the gases CO₂ and H₂, which are vented via the breath and increased flatus) of the gut flora is concomitant with reduced proteolytic activity (production of putrefactive compounds and cytotoxic degradation products) because it is energetically more favourable to produce ATP from carbohydrates than from proteins. As long as carbohydrates are present, the bacteria need not establish a proteolytic catabolic enzyme apparatus, and hence less proteolysis takes place. It is thought that improving the composition of the intestinal ecosystem has a positive impact on the physiological functioning of an organism.

Nutritional properties of prebiotics

A direct consequence of prebiosis is an improved bowel habit. As prebiotic carbohydrates become available to the intestinal bacteria, they grow on it, increase in numbers and eventually in volume; they induce a faecal bulking effect. The SCFA products of fermentation stimulate intestinal peristalsis (Cherbut *et al.* 1991).

As a consequence of prebiotic consumption, there is a relief from constipation in chronically constipated individuals (Den Hond *et al.* 1997). As prebiotic carbohydrates arrive in the colon, they modify the composition of the intestinal flora, and they influence its metabolic activity. These processes increase the 'colonisation resistance'. This is the resistance of a certain established colonic bacterial population against other incoming bacteria (orally consumed) from finding a niche in the ecosystem (Newton *et al.* 1998). In a study with individuals travelling to 'high-risk' countries, it was observed that prebiotic consumption reduced the incidence of diarrhoea by almost 50 % (Cummings *et al.* 2001).

It is thought that an increased production of propionate is at the basis of altered lipid metabolism. Propionate, which is produced from the fermentation of prebiotics, migrates via the portal vein to the liver, where it interacts with the regulation of the expression of genes that code for digestive hormones (incretins) such as glucagon-like peptide 1 and gastric inhibitory polypeptide, and of insulin (Kok *et al.* 1998).

The human studies that have been performed fail to show very uniform experimental outcomes. This is ascribed to the complexity of the biochemistry of lipid metabolism, which certainly complicates experimental protocols and outcomes (Van Loo *et al.* 1999; Pereira & Gibson, 2002). Several studies in experimental models and in human dietary interventions, however, have shown that the parameters of lipid metabolism are not unaffected.

A consistent anticarcinogenic effect of dietary prebiotics has been observed in various experimental models.

The effect was shown in chemoprevention models in the short term (Reddy *et al.* 1997), where colonic aberrant crypt foci (ACF) are the biomarkers, as well as in long-term chemoprevention models (Verghese *et al.* 2002a,b) where the incidence of tumours is the biomarker.

In genetically predetermined Min-mice (*apc*-) models, there was a significant reduction of colonic tumours, and of tumours in the small intestine, indicating that systemic effects are involved (Pierre *et al.* 1997; M Lipkin, personal communication).

The systemic efficacy was confirmed in models where tumour cells were implanted in muscle tissue (with growth rate of the tumour as a marker) or in peritoneum (with increase in life-span as a marker) (Taper *et al.* 1997, 1998).

At present, the anticancer effect of prebiotics in combination with probiotics is being investigated in a human volunteer study (Van Loo & Jonkers, 2001).

The intake of prebiotics acidifies the intestinal contents, which solubilises minerals (mainly Ca or Mg salts) (Coudray *et al.* 2003b). The increased presence of butyrate, which is a selective source of energy for the intestinal epithelial cells, improves the absorptive capacity of the mucosa. Both these phenomena are thought to be at the basis of the repeatedly observed increase in mineral absorption from food in human volunteers (Coudray *et al.* 1997; van den Heuvel *et al.* 1998; Griffin *et al.* 2002).

Goal of the present paper

The present paper is intended to demonstrate that the particular way in which a prebiotic food ingredient interferes with the fermentation patterns of the GI ecosystem is central for improving the physiological functioning of the host. It will be shown that subtle changes in the chemical composition of the prebiotic food ingredient can have a major impact on the fermentation processes in the GI tract, which is reflected in an altered impact on certain physiological processes.

The available evidence

At present, there are only a few substrates for which a prebiotic action has been demonstrated adequately, i.e. that comply with all three prebiotic criteria as demonstrated by means of repeated human dietary intervention studies. These are the dietary oligosaccharides $\beta(2-1)$ fructans (inulins).

For the galacto-oligosaccharides there is strong evidence of selective stimulation of the intestinal flora; for isomalto-oligosaccharides and xylo-oligosaccharides, there is promising evidence. For the therapeutic disaccharide lactulose, there is good evidence.

As the inulins are dietary ingredients for which both prebiotic properties and effects on physiological function (mineral absorption, cancer experiments, immunological and lipid metabolism trials) have been demonstrated repeatedly in both experimental models and in human volunteers, they will be taken as the basis for further discussion.

The influence of inulin on gastrointestinal fermentation

Native inulin (ST-inulin) is obtained industrially from chicory roots by extraction with hot water, followed by refining and spray-drying. It is a polydisperse mixture of linear molecules, all with the same basic chemical structure, which is symbolised as $G-F_n$ (G, glycosyl moiety; F, fructosyl moiety; n , number of fructose units linked together through $\beta(2-1)$ bonds). The degree of polymerisation (DP) varies between three and sixty-five, and the average DP (DP_{av}) is ten. OF, which is obtained by the partial enzymic hydrolysis of ST-inulin, is a mixture of molecules with chain lengths ranging between two and eight with a

DP_{av} of four. High-polymer inulin (HP-inulin) is obtained by physically removing the lower-DP fraction from ST-inulin. The DP of HP-inulin ranges from ten to sixty-five, with a DP_{av} of twenty-five. Synergy1 (SYN1) (Raftilose®Synergy1 manufactured and commercialised for human consumption by ORAFTI NV, Tienen, Belgium) is a 1:1 mixture of OF and HP-inulin (see Fig. 1).

The bifidogenicity of each of those fractions of inulin has been demonstrated repeatedly in human dietary intervention studies (Roberfroid *et al.* 1998; Van Loo *et al.* 1999). From these reviews, it appears that the short-chain inulin (OF) increases the bifidobacteria by almost an order of magnitude more (order of magnitude 1 log colony-forming unit) than the HP-inulins (order of magnitude 0.1 log colony-forming unit), but selectivity has been demonstrated in both cases (Tuohy *et al.* 2001; Harmsen *et al.* 2002). The bifidobacteria function as indicator organisms of a well-balanced GI flora in terms of increased saccharolytic activity relative to proteolytic activity, which is monitored by quantifying the SCFA:branched SCFA ratio (Swanson *et al.* 2002). It is obvious that the OF has more impact on improving the composition of the microflora (relative amounts of different groups of GI bacteria) than the other fractions.

There is a remarkable physicochemical property of the inulin chains that is central in the explanation of differences in the way different fractions influence GI fermentation.

Whereas a DP range of three to sixty-five is relatively small, there is a big difference in the solubility of inulin chains with different lengths. Those with a DP from three to ten are very soluble (up to 85 % (w/v) in water at room temperature), while molecules with a chain length of ten to sixty-five are rather poorly soluble (up to 2 % (w/v) in water at room temperature). It is thought that this is related to the thermodynamic behaviour of inulin in solution, of which various metastable forms exist (Marchessault, 1980).

It has been shown that the *in vitro* faecal slurry fermentation of the short-chain inulins or OF occurs very quickly; 1 % (w/v) at pH 6.7 and 37°C with 2 % (w/v) inoculum is completely fermented in 5 h. However, the fermentation of the long chains or HP-inulin takes about three to five times as long (Roberfroid *et al.* 1998; Baeten, 1999).

It has been shown (J Van Loo, unpublished data) that long-chain inulin is fermented completely in the GI tract of human subjects consuming 10 g/d or more (none is recovered in faeces). Because the long chains are fermented slowly, they reach more distal regions in the colon, where they can maintain a selective metabolic activity (Tuohy *et al.* 2001; Harmsen *et al.* 2002). Whereas HP-inulin modifies the composition of the intestinal flora significantly but to a limited extent as compared with OF, it is transformed into biologically active metabolites for a longer period of time and in more distal regions of the intestine.

The hypothesis behind the formulation of SYN1 was to provide a short-chain fraction that is able to modify the composition of the colonic flora. This necessarily occurs in the proximal part of the colon, given its rapid fermentation. The remaining fraction, HP-inulin, functions subsequently as a slow but selective substrate for the modified GI ecosystem. As a direct consequence, the improved GI ecosystem can exert its beneficial effects for a longer period of time;

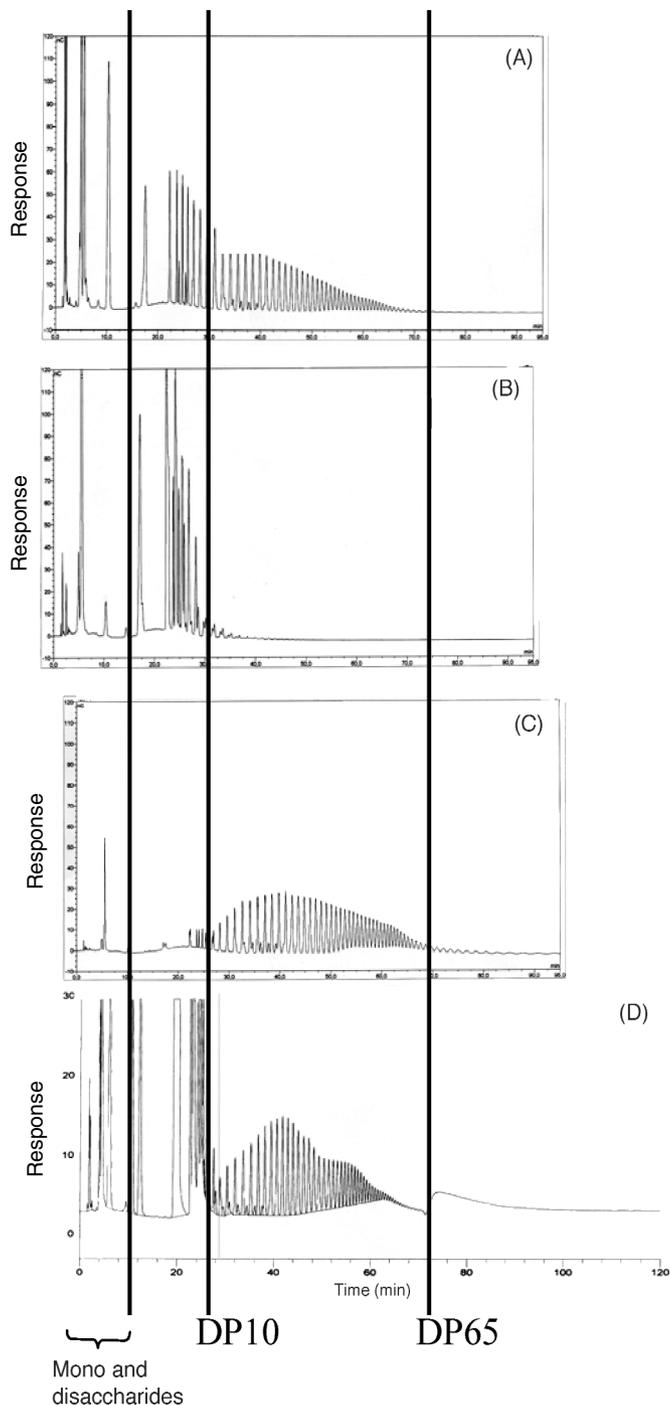


Fig. 1. High-performance, anion-exchange chromatograms (obtained with a pulsed amperometric detector; Van Loo *et al.* 1995) of the different chicory inulin fractions that were tested in this work. The technique is not quantitative, but shows accurately the distribution of various chain lengths. The monosaccharides are fructose and glucose; the disaccharide is sucrose. ST-inulin (A) is the inulin as extracted from the chicory root. The degree of polymerisation (DP) varies from three to sixty-five; the average DP (DP_{av}) is ten. Oligofructose (B) is partially hydrolysed ST-inulin (DP between two and eight; DP_{av} 4). HP-inulin (C) is ST-inulin devoid of the oligofructose fraction (DP between ten and 65; DP_{av} 25). Synergy1 (D) is a 1:1 mixture of oligofructose and HP-inulin (ORAFTI NV, Tienen, Belgium).

an implicit consequence is that the improved GI ecosystem can extend its metabolic activity into more distal areas of the GI tract. Until now it has been impossible to demonstrate these phenomena *in situ* and *in vivo* but, as will be seen later, there are indirect indications of the validity of the hypothesis.

So far, the observation and interpretation of the prebiotic properties of inulin reveals that subtle differences in the molecular structure (chain length) of carbohydrates can have a major impact on fermentation processes.

The fact that non-digestible carbohydrates with different molecular composition and binding properties have dramatically different influences on fermentation has been demonstrated (Wang & Gibson, 1993; Gibson *et al.* 1995; Hopkins *et al.* 1998). For inulin there is at present a vast set of experimental evidence from both fermentation and dietary intervention studies. Therefore the rest of the present paper will focus on inulin. It is emphasised that this does not imply that other prebiotics could not contribute to the discussion. There however is no evidence available to date. The present discussion on inulin allows us to show the impact of prebiotics that only differ in one aspect (chain length). All other chemical parameters are similar (composing monosaccharide, type of binding), which makes the discussion from this point of view scientifically more relevant.

Impact of the type of fermentation (inulin fraction) on physiological processes

Anticarcinogenesis studies

The anticarcinogenic properties of inulin have been demonstrated by means of various experimental models (chemoprevention, genetically predetermined mouse model and tumour implantation models; Van Loo & Jonkers, 2001). In the chemoprevention studies, the effect of inulin fractions with different chain length has been studied (Verghese *et al.* 2003). The idea was to replace some of the starch (10 % with no effect on the growth rate of the animals) in a standard rat diet with OF, ST-inulin, HP-inulin and SYN1.

It was observed that the greater the proportion of chains with DP of ten to sixty-five (series OF (0 % DP > 10); ST-inulin (50 % DP > 10); HP-inulin (100 % DP > 10)), the smaller the number of total aberrant crypts observed in the colon. From Fig. 2, it appears that at the same time the ACF with high multiplicity (three, four, and five and more ACF/focus) decreases markedly.

The observation with SYN1 (50 % DP > 10) is not in accord with these observations; whereas the relative amount of long chains is similar to ST-inulin, its effect on the reduction of ACF is decreased markedly. It may be observed from Fig. 2 that the reduction of foci with high multiplicity is much more pronounced for SYN1 than for the other inulin fractions.

Fig. 3, which is derived from the same experiment, shows that the higher the fraction with DP > 10, the more important becomes the effect of inulin in the distal part of the colon (series OF (0 % DP > 10); ST-inulin (50 % DP > 10); HP-inulin (100 % DP > 10)). Again, there is a disparity

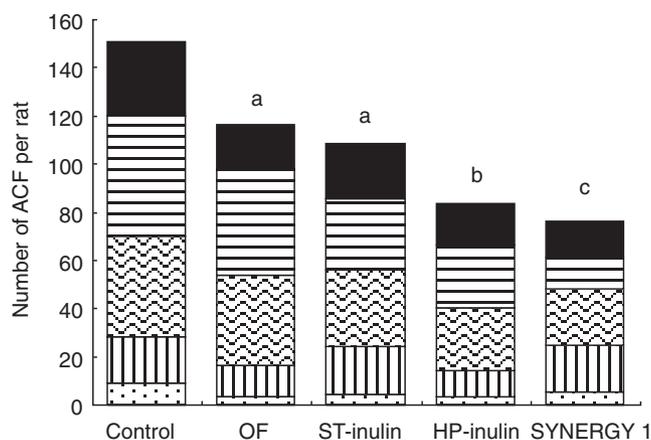


Fig. 2. A chemoprevention trial in which rats (n 12) were challenged twice (1-week interval) with azoxymethane (15 mg/kg body weight). They were given the respective diets (control; oligofructose (OF); native inulin (ST-inulin); high-polymer inulin (HP-inulin); Synergy1 (1:1 mixture of OF and HP-inulin; ORAFIT NV, Tienen, Belgium)) 2 weeks before carcinogen application, and then for 8 weeks more, after which the aberrant crypt foci (ACF) were counted, and classified as foci with one, two and up to five ACF per focus. The more ACF per focus, the higher the probability of development into a tumour. (■), Five or more ACF/focus; (▨), four ACF/focus; (▩), three ACF/focus; (▧), two ACF/focus; (▦), one ACF/focus. ^{a,b,c} Values with unlike superscript letters are significantly different; all are significantly different from the control ($P < 0.05$) (from Verghese *et al.* 2003).

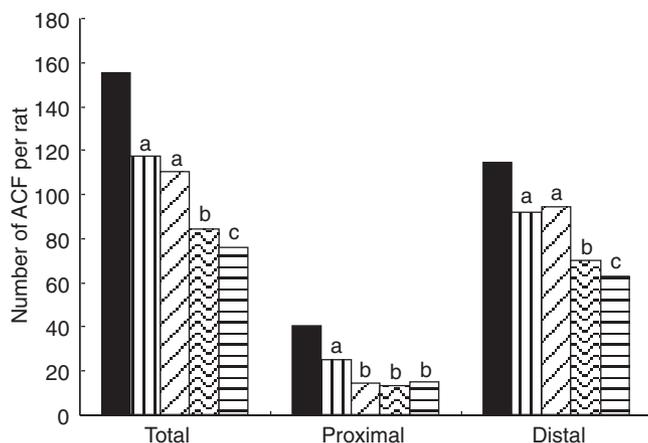


Fig. 3. Data from the same experiment as in Fig. 2, but rearranged to show the effects in the proximal and distal colon. The diets were: control (■), oligofructose (OF; ▨), native inulin (ST-inulin; ▩), high-polymer inulin (HP-inulin; ▧), and Synergy1 (1:1 mixture of OF and HP-inulin; ORAFIT NV, Tienen, Belgium) (▦). It is observed that short chains (OF) exert beneficial effects particularly in the proximal colon. The higher the proportion of long chains, the more important becomes the anticarcinogenic effect in the distal part of the colon (series ST-inulin and HP-inulin). There is a smaller proportion of long chains in Synergy1 than in HP-inulin, but the additional effect on the composition of the flora in the proximal colon induced by the short-chain fraction of Synergy1 (see Fig. 1) could explain the difference. ^{a,b,c} Values with unlike superscript letters in each group (total, proximal, distal) are significantly different; all are significantly different from the control (from Verghese *et al.* 2003).

in this observation with SYN1, where the effect in the distal colon is more pronounced than with any of the other fractions (Fig. 3).

This difference can be explained by the additional effect of SYN1; i.e. the modification (with its short-chain fraction) of the flora in the proximal colon to a greater degree than the other prebiotic food ingredients. This would allow the fermentation of its long-chain fraction ($DP > 10$) to occur by a more potent anticarcinogenic intestinal population. Also, the long-chain fraction ($DP > 10$) in SYN1 is, on average, longer (DP_{av} 25) than in ST-inulin (DP_{av} 14), but is similar to that in HP-inulin (DP_{av} 25).

Earlier, Reddy *et al.* (1997) reported a difference between OF and HP-inulin in a similar protocol, which these authors attributed to a difference in fermentation rate between OF and HP-inulin, as the carcinogen azoxymethane induces ACF in more distal parts of the colon. As long chains are fermented more slowly, the observed effect could be an indirect demonstration that they also migrate to more distal parts in the colon, where they can maintain a more saccharolytic (production of SCFA) fermentation.

Klinder *et al.* (2003) quantified cellular parameters of survival (differentiation, tumour progression, and invasive growth) in HT29 cells and Caco2 cells, which they challenged with supernatant fractions of fermentation broths originating from a three-stage colon simulation model. They compared a fermentation broth that had glucose as a C-source with a fermentation broth that had SYN1 as the C-source. It was observed that SYN1 fermentation broths reduced the markers of tumour progression significantly more than the glucose fermentation broth.

Glucose is a commonly available C-source. The microflora developed on glucose is different from that established on SYN1. The different effect of the supernatant fractions of both on the molecular processes taking place in the tested cell lines demonstrates that fermentation pattern (bacteria and metabolites) plays a central role in the influence that these exert on physiological processes.

Conclusion. The results of these experiments suggest that the modification of the gut microflora is key in causing the anticarcinogenic effect of inulin. Slowly and selectively fermented long chains, however, sustain the effect (selectively maintain the metabolic activity), which is reflected in a stronger physiological impact (ACF reduction in this case).

It is observed that the longer the chain length, the smaller the number of crypts with high multiplicity.

Whereas the fraction of long chains is smaller in SYN1, there are markedly fewer ACF with high multiplicity than are seen with HP-inulin. The additional effect of the short-chain fraction on the intestinal flora in the proximal part of the colon could explain the difference.

The type of fermentation determines how the physiological processes are influenced.

Studies on increased mineral absorption

Experimental models. The increase of mineral absorption (Ca, Mg) from food by the dietary administration of inulin or

its fractions to experimental models (Delzenne *et al.* 1995; Ohta *et al.* 1998; Coudray *et al.* 2003b) and human volunteers (Coudray *et al.* 1997; van den Heuvel *et al.* 1999; Griffin *et al.* 2002; Cashman, 2003; L Holloway, S Moynihan and A Friedlander, unpublished results) is well documented.

Reconsidering these studies with an emphasis on chain length, or on the fermentation properties of the particular inulin types used, can contribute to the goal of the present paper.

In an exploratory mineral absorption experiment, Delzenne *et al.* (1995) used ST-inulin and OF as test substances. They observed significant increases in the retention of Ca, Mg and Zn and they did not observe significant differences in efficacy between OF and ST-inulin.

Recently, Coudray *et al.* (2003b) reported the results of a study where the effect of OF, HP-inulin and SYN1 on Ca and Mg absorption were compared with a control (no non-digestible oligosaccharides added). They confirmed the results reported by Delzenne *et al.* (1995) for OF for both Ca and Mg absorption. They observed additionally that HP-inulin was as efficient as OF with respect to increased Ca absorption, but that it increased Mg absorption significantly more than OF. The effect of SYN1 in this experiment was significantly more efficient in increasing Ca absorption from the diet than both OF and HP-inulin. There was no additional effect on Mg absorption, but this may be due to saturation, as over 75 % of all dietary Mg is absorbed (Fig. 4).

In the experimental models on mineral absorption, there seems to be a trend similar to that seen in the anticarcinogenesis studies. The more slowly fermented longer chains increase the length of time of the physiological observation. In the mineral absorption model, however, the effect of the 1:1 OF:HP-inulin combination product is markedly more pronounced than in the anticancer models.

Whereas the caecal soluble Ca pool was increased 20- to 30-fold in all inulin-type fed groups, the increased solubility of Ca and Mg did not correlate with the increased mineral absorption capacity of the different inulin types. There was no observable difference in caecal SCFA composition. This suggests that the difference in improving physiological properties of the different inulin fractions is taking place more distally than the caecum, presumably in the colon. Ohta *et al.* (1998) showed an increase of calbindin in the caecum and colorectum. It should be investigated whether long chains in HP-inulin or SYN1 affect calbindin presence in the colorectal area more than OF does.

The difference in the Ca increase between the SYN1 diet and the OF diet is much greater than the difference between the HP-inulin diet and the OF diet. This suggests that the combination of rapidly fermented prebiotics (OF chains in caecum) and slowly fermented prebiotics (HP-inulin chains in colon and rectum) enhances mineral absorption significantly.

Younes *et al.* (2001) published the results of a study in rats where ST-inulin, resistant starch (RS) and a combination of the two were compared. They observed that both ST-inulin (+61 %) and RS (+34 %) increased Ca absorption, and that the combination of the two increased Ca absorption significantly more (+130 %; from 23 to 53 % absorption of the dietary Ca) than their component units.

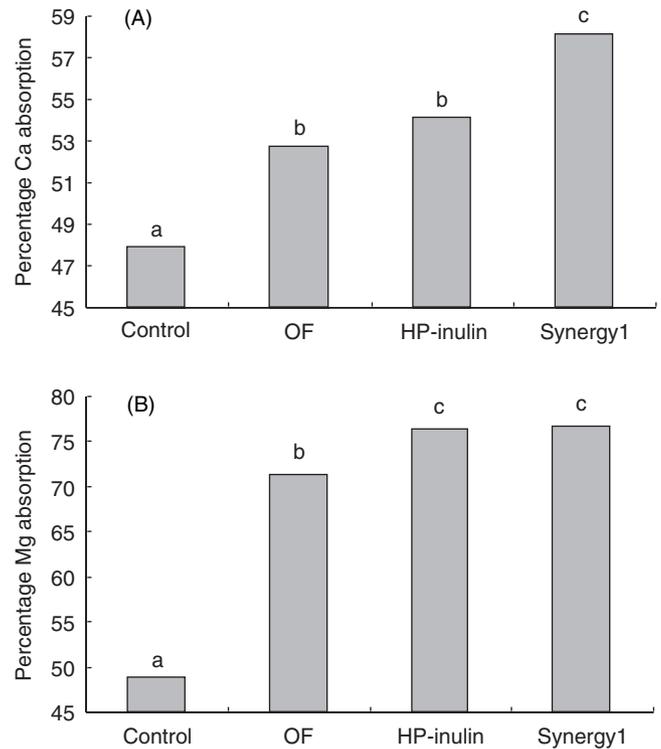


Fig. 4. Increase of percentage Ca absorption (A) and percentage Mg absorption (B) upon intake of 10 % oligofructose (OF), high-polymer inulin (HP-inulin) and Synergy1 (1:1 mixture of OF and HP-inulin; ORAFIT NV, Tienen, Belgium) by rats. The presence of long chains increases the mineral absorption, but for Synergy1 there is a markedly more important effect on Ca absorption. ^{a,b,c} Values with unlike superscript letters are significantly different ($P < 0.05$) (from Coudray *et al.* 2003b).

All three diets increased Mg absorption significantly but there was no significant additional effect due to the combination.

There was significantly more caecal SCFA with the RS diet and the combination diet than there was with the ST-inulin diet (and in all three cases significantly more than in control; about 3-fold). RS had significantly more SCFA produced than ST-inulin, but the percentage Ca absorption was (not significantly) lower with RS. This suggests that SCFA and other fermentative factors play a role (other metabolites, types of bacteria that are stimulated).

Human dietary intervention studies. Griffin *et al.* (2002) recently published data that contribute to the present discussion. They used a double-blind, placebo-controlled, cross-over study design with two cohorts of thirty volunteers (girls 11–14 years), who were given a high intake of Ca (1500 mg/d) in order to be above the saturation level (1200 mg Ca/d). This was done to avoid the possibility that varying sub-optimal Ca intakes would interfere with the results. One group of volunteers was given SYN1 at a dose of 8 g/d and another group received OF at 8 g/d. The percentage Ca absorption from the food was measured with a dual-stable-isotope technique, each group against its own control.

The results are presented in Fig. 5, which shows clearly that under the experimental conditions of very high Ca intake, SYN1 but not OF increases Ca absorption significantly.

In previously published studies with OF at 15 g/d for 9 d (van den Heuvel *et al.* 1999) or with ST-inulin at up to 40 g/d during 1 month (Coudray *et al.* 1997) it was shown that both have the ability to increase Ca absorption, albeit at higher levels of intake.

Tahiri *et al.* (2001) have observed an increased Mg absorption of 12 % but no effect on Ca absorption (Tahiri *et al.* 2003) with short-chain fructo-oligosaccharides (a synthetic equivalent of OF) (n 11; 10 g/d for a period of 5 weeks in a cross-over study design) in post-menopausal women. Similar results with other non-digestible oligosaccharides (of which prebiotic activity has not been proven to date) such as maltitol and hydrogenated maltitol also increased Mg absorption, but did not affect Ca absorption (Coudray *et al.* 2003a).

In a similar population of post-menopausal women, (L Holloway, S Moynihan and A Friedlander, unpublished results) have observed a significant increase in both Mg absorption of 22 % and a significant increase in Ca absorption of 39 % with SYN1 (n 15; SYN1 at 10 g/d for a period of 6 weeks in a double-blind, placebo-controlled cross-over study design).

Conclusion. Both experimental models and human dietary intervention studies show clearly that there is a link

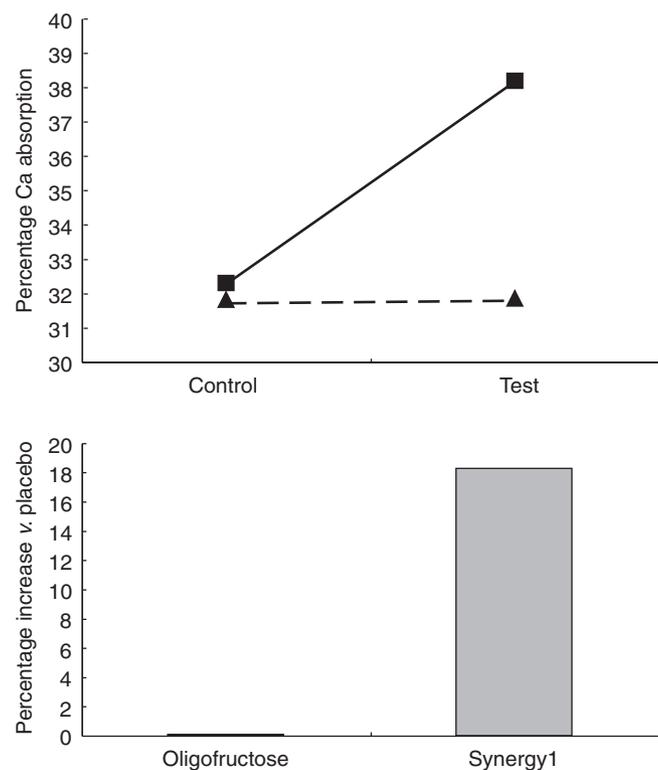


Fig. 5. Under conditions of high intake levels of Ca (1500 mg/d), Synergy1 (1:1 mixture of oligofructose and high-polymer inulin; ORAFIT NV, Tienen, Belgium) (■) but not oligofructose (▲) is able to increase significantly ($P < 0.05$) the dietary percentage Ca absorption as measured by dual-stable-isotope techniques (from Griffin *et al.* 2002).

between the observed increase in mineral absorption and the type of prebiotic used. As a consequence, there is a link between the GI fermentation pattern of a prebiotic carbohydrate and the effect on mineral absorption.

From the experiments, it follows that the effect increases with increasing chain length, which, in the case of inulin, means slower selective fermentation.

In mineral absorption studies too, the effect of the combination of a rapidly and selectively fermented substrate (OF) with a slowly and selectively fermented substrate (HP-inulin) is significantly more effective in increasing Ca absorption than the inulin fractions it is composed of.

This indicates that there is a positive interaction between the effects on mineral absorption of a rapid fermentation, which results in highly significant changes in the composition of intestinal flora, and the effects of slow fermentation, which may keep the modified flora active for a longer period of time and probably into more distal parts of the intestine.

Studies on immunology

Buddington *et al.* (2002) fed mice a control diet, a diet with 10 % HP-inulin and a diet containing 10 % OF. After 2 weeks adaptation, a group of animals were injected intraperitoneally with *Listeria monocytogenes* and another group with *Salmonella typhimurium*, in such a dose as to cause about a 50 % mortality after a certain period of time. For the *Listeria* challenge, the authors observed a significant decrease in mortality in the OF-fed animals, but for the HP-inulin the effect was significantly more pronounced, to even 0 % mortality. For *Salmonella*, there was no significant numerical decrease in mortality for the OF-fed animals, but the effect of HP-inulin remained significant (Fig. 6). In the *Listeria*-challenged mice, there was a significantly increased natural killer cell activity of the splenocytes as well as a significantly higher phagocytic activity of peritoneal macrophages (Kelly-Quagliana *et al.* 2003). Whereas the difference between OF and HP-inulin was not significant, both values were numerically more important in HP-inulin than in OF.

SYN1 was not available at the time these experiments were carried out.

Conclusion. Even in systemic physiological processes such as the immune system, the effect of different fermentation patterns of the prebiotic exerts its influence. The more slowly fermented long-chain HP-inulin seems to be more efficient in promoting effects than the short chains of OF.

Studies on factors related to lipid metabolism

Here, the data are less consistent. Some published human dietary intervention studies show effects on cholesterol, others on serum triacylglycerol levels, others on both and still other studies show no effect (Van Loo *et al.* 1999; Pereira & Gibson, 2002). From the several studies published, three were performed with OF (Alles *et al.* 1997; van Vliet, 1997; van Dokkum *et al.* 1999) and had no effect on the levels of cholesterol or serum triacylglycerols. Four were carried out with ST-inulin, of which three had an

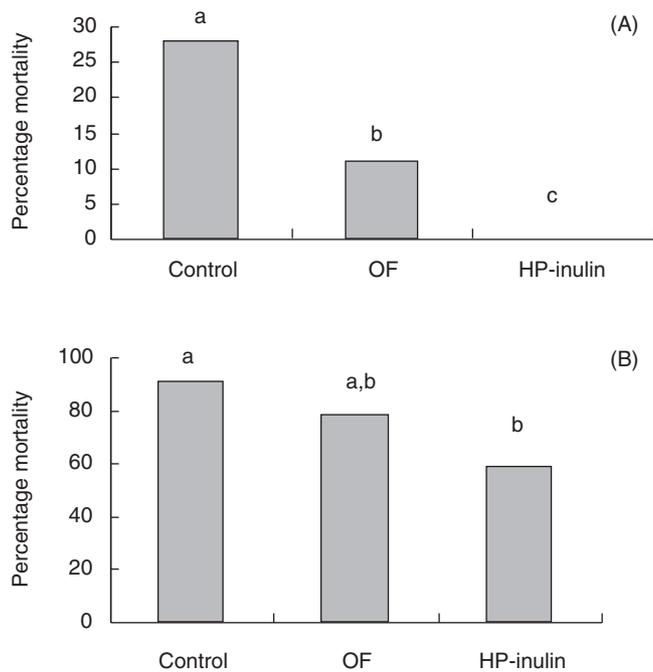


Fig. 6. In challenge experiments (*Listeria monocytogenes* (A) and *Salmonella typhimurium* (B)) where the model was fed either an oligofructose (OF)- or a high-polymer-inulin (HP-inulin)-containing diet, the percentage mortality was significantly different from the control. In the *Listeria* challenge, HP-inulin was significantly more active than OF, and this was the case numerically in the *Salmonella* challenge. ^{a,b,c} Values with unlike superscript letters are significantly different ($P < 0.05$) (from Buddington *et al.* 2002).

effect on at least one of the measured parameters of lipid metabolism (Pedersen *et al.* 1997; Brighenti *et al.* 1999; Davidson & Maki, 1999) and one had no effect (van Dokkum *et al.* 1999). Two studies were performed with HP-inulin and both were effective in affecting parameters of lipid metabolism. One showed significant effects on serum triacylglycerols and on insulin (Jackson *et al.* 1999), and one showed clear effects on hepatic lipogenesis (Letexier *et al.* 2003).

Conclusion. In none of these studies was the effect of two types of inulin compared in the same protocol. It therefore cannot be claimed on a solid basis that long chains are more efficient than short chains in affecting lipid metabolism. The observation that studies performed with long chains (ST-inulin and HP-inulin) gave more positive results than the studies performed with OF seem to confirm the trend described earlier.

General discussion and conclusion

Prebiotics selectively modify the composition and/or the metabolic activity of the GI ecosystem, which is the external environment with which the human body is in closest contact. The way in which a prebiotic interacts with that environment depends very much on its molecular structure.

There were significant effects on mineral absorption, cancer risk reduction, and immunology with prebiotics that

typically modify the flora but do not maintain the fermentation (rapidly fermented prebiotics such as OF). However, there seems to be a positive correlation between the effect on all physiological fields of interest mentioned, including lipid metabolism, and the presence of long chains.

There is a clear link between molecular structure, GI fermentation and physiological efficacy. This relationship is more complex than a simple dose–effect relationship, as the size of impact of the latter is very much dependent on the initial status (bacteriological and physiological) of a volunteer. This issue has been discussed by Roberfroid *et al.* (1998).

From these observations, it can be concluded that it is the type of GI fermentation that finally affects the physiological observations.

This hypothesis is confirmed by experiments with combination products composed of a rapidly and selectively fermented substrate (flora composition effect, physiological effects) and a slowly fermented substrate (sustained maintenance of GI flora metabolism). The physiological impact of such a combination product is more important than that of its components. This illustrates the adjuvant effect of ‘flora composition’ and ‘sustained metabolic maintenance’, and hence that impact on GI fermentation causes the physiological observations.

In other words it can be claimed that prebiotics, through their impact on GI fermentation, influence physiological processes, most of which are considered biomarkers for health status.

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