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Importance of digestive mucus and mucins for designing new functional food ingredients 4 Mounira Kebouchi^(a, b), Zeeshan Hafeez^(a), Yves Le Roux^(b), Annie Dary-Mourot^(a, c), 5 Magali Genay $(a, c)^*$ (a) Université de Lorraine, CALBINOTOX, F-54000, Nancy, France (b) Université de Lorraine, INRA, URAFPA, F-54000, Nancy, France $8 \t\t(c)$: Co-senior authors (*) Corresponding author: Magali Genay. Composés Alimentaires : Biofonctionnalités et Risques Neurotoxiques (CALBINOTOX), Faculté des Sciences et Technologies, B.P. 70239, F-54506 Vandœuvre-lès-Nancy, France*. Phone: +33 (0) 3 72 74 51 90. E-mail:*

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Abstract

The mucus, mainly composed of the glycoproteins mucins, is a rheological substance that covers the intestinal epithelium and acts as a protective barrier against a variety of harmful molecules, microbial infection and varying lumen environment conditions. Alterations in the composition or structure of the mucus could lead to various diseases such as inflammatory bowel disease or colorectal cancer. Recent studies revealed that an exogenous intake of probiotic bacteria or other dietary components (such as bioactive peptides and probiotics) derived from food influence mucus layer properties as well as modulate gene 22 expression and secretion of mucins. Therefore, the use of such components for designing new functional ingredients and then foods, could constitute a novel approach to preserve the properties of mucus. After presenting some aspects of the mucus and mucins in the gastrointestinal tract as well as mucus role in the gut health, this review will address role of dietary ingredients in improving mucus / mucin production and provides new suggestions for further investigations of how dietary ingredients / probiotics based functional foods can be developed to maintain or improve the gut health.

Keywords: intestinal mucosa; mucins; digestive tract; probiotic bacteria; food ingredients; functional foods

1. Introduction

The human digestive tract consists of a stable alliance among the epithelial barrier, the immune system and the resident microbiota. Besides its implication in absorption and assimilation of nutrients, exclusion of undigested food, protective role against harmful molecules and pathogenic microorganisms, the contribution of the digestive tract to sensory functions including the innate and adaptive immune responses have also been well documented. The epithelium of the gastrointestinal tract (GIT) consists of enterocytes, goblet cells, enteroendocrine cells, Paneth cells, microfold cells (M cells), cup and tuft cells, and is coated with a gel layer, called mucus (Ahluwalia, Magnusson, & Lohman, 2017; McGuckin, 41 Thornton, & Whitsett, 2015). Furthermore, the human GIT also harbors more than 10^{13} of resident bacteria per gram of feces colonizing the mucus layer (Sender, Fuchs, & Milo, 2016). The number of microbial cells as well as the number of species varies throughout the GIT (Li et al., 2015) with few resident bacteria in the stomach and small intestine (Bik et al., 2006), and a dramatic increase in the distal ileum and the colon (Ermund, Schütte, Johansson, Gustafsson, & Hansson, 2013; Li et al., 2015). The association of bacteria with the host have 47 led to co-evolution with mutual benefits and impacts (Bäckhed, Ley, Sonnenburg, Peterson, & Gordon, 2005; Liang, Leung, Guan, & Au, 2018). Hence, the resident bacteria could influence the composition and structuration of the mucus layer and finally the mucus properties (Jakobsson et al., 2015).

The digestive mucus, mostly consisting of mucins, plays a major role in the good functioning of the digestive tract and thereby the human health. Whatever the reasons, alterations in composition or structure of the mucus result in a wide variety of diseases including inflammatory bowel deseases or colorectal cancer (Johansson, Sjövall, & Hansson, 2013; Velcich et al., 2002). Hence, mucins, such as MUC2 ("MUC" refers to human mucins

whereas "Muc" corresponds to mucins in mice and other animals), constitute a target for the design of new therapeutic molecules because of their involvement in tumorigenesis (Aithal et al., 2018; Li et al., 2018).

Scientific community and public are now more interested in natural products to maintain consumer health. This is reflected by the ever-growing market of functional food products containing health-promoting ingredients and / or bacteria (Douillard, & de Vos, 2019; Vicentini, Liberatore, & Mastrocola, 2016). Therefore, it would be interesting to explore the possibility of identifying functional ingredients able to modulate mucus production as the drug treatments are usually associated with adverse effects. After a brief description of the mucus, its beneficial role in maintaining human health and the control of this layer, this review will address the effects of food ingredients on mucus or mucins production and the possible ways to design / improve functional ingredients, a first step in developing functional foods, summarizing thereby the recent insights in this domain.

2. Overview of digestive mucus and its beneficial role

Mucus, secreted by intestinal goblets cells, is characterized by its viscosity and 72 rheological properties. It consists in water $(\approx 95\%)$, electrolytes, enzymes, salts, DNA, lipids, growth factors, antibacterial peptides and proteins such as defensins, immunoglobulins, lysozymes and trefoil factors (McGuckin, Thornton, & Whitsett, 2015), and especially glycoproteins called mucins. It not only constitutes a lubricant facilitating the transport of the chyme along the GIT (Corfield, 2015; Lai, Wang, Wirtz, & Hanes, 2009) but also serves as a barrier that protects the epithelial mucosa against mechanical injuries as well as from harmful enteric pathogenic bacteria, viruses and parasites (Birchenough, Johansson, Gustafsson, Bergström, & Hansson, 2015; Ermund, Schütte, Johansson, Gustafsson, & Hansson, 2013; Liévin-Le Moal & Servin, 2006; Pelaseyed et al., 2014). Morover, it also plays a key role in 81 maintaining intestinal homeostasis (Faderl, Noti, Corazza, & Mueller, 2015).

Studies conducted in rodent models and humans have shown that the mucus organization and composition differ along the GIT (Figure 1). In the small intestine, there is a single, discontinuous and disorganized mucus layer which is continuously secreted from the crypt openings (Atuma, Strugala, Allen, & Holm, 2001; Ermund, Schütte, Johansson, Gustafsson, & Hansson, 2013; Pelaseyed et al., 2014). In the stomach and the colon, the mucus is thicker and composed of two layers. The inner dense layer is strongly attached to the epithelium, devoid of bacteria and is constantly renewed every 1-2 hours by surface goblet cells of the colon (Ermund, Schütte, Johansson, Gustafsson, & Hansson, 2013; Gustafsson et al., 2012; Johansson, 2012). The outer loose layer, facing the lumen, is disorganized, more hydrated and thicker than the inner layer and is colonized by the commensal bacteria in the colon (Atuma, Strugala, Allen, & Holm, 2001; Ermund, Schütte, Johansson, Gustafsson, & Hansson, 2013; Hansson & Johansson, 2010; Johansson et al., 2008). It results from the inner layer probably through the proteolytic cleavage occurring within the mucins (Johansson et al., 2008).

As major constituents of mucus, mucins represent large and highly glycosylated proteins expressed in different organs. In the adult human digestive tract, a wide variety of mucins are secreted at the epithelium surfaces (Figure 1). They are clustered into three distinct subfamilies: i) secreted gel-forming mucins, ii) secreted non-gel forming mucins, iii) membrane-bound mucins, and their physiological roles are described in Table 1 (Moniaux, Escande, Porchet, Aubert, & Batra, 2001). Being highly glycosylated molecules, a high proportion of mucin's molecular weight corresponds to carbohydrates (oligosaccharides or glycan chains). Their structure described in Figure 2 is important for both viscoelasticity (Demouveaux, Gouyer, Gottrand, Narita, & Desseyn, 2018) and bioactivity of mucus (Petrou, & Crouzier, 2018). Biosynthesis and secretion of all types of mucins present some similarities such as N-glycosylation in the endoplasmic reticulum and O-glycosylation at PTS sequence (proline, threonine and serine rich region) in the Golgi apparatus. Afterwards, membrane bound mucins are inserted into the membrane *via* C-terminal transmembrane domain whereas secreted mucins are oligomerized and packed into Goblet cell granules for secretion (Figure 3).

Mucins play a major role in the protection and functioning of different parts of the GIT. At the oral sphere level, mucins of saliva, e.g. MUC5B and MUC7 are involved in maintenance of its rheological properties and protect the floor of the mouth against mechanical forces during eating and speaking. They also protect from different diseases by reducing bacterial attachement and / or their biofilm formation capacity (Table 1 and Figure 1) (Amerongen, Bolscher, & Veerman, 1995; Dawes et al., 2015; Plummer & Douglas, 2006; Stinson et al., 1982; van't Hof, Veerman, Nieuw Amerongen, & Ligtenberg, 2014). Other mucins of oral sphere (MUC1, MUC16 and MUC4) that line the duct of parotid, submandibular and minor salivary glands are involved in cell signal transduction and binding of secreted mucins MUC5B and MUC7 (Frenkel & Ribbeck, 2015b; Liu et al., 2002; O'Brien, Beard, Underwood, & Shigemasa, 2002; Sengupta et al., 2001). In the rest of the GIT, the mucins safeguard the cell surface epithelium from the endogenous toxic molecules and pathogenic bacteria (Table 1 and Figure 1) (Allen & Flemström, 2005; Bevins, 2006). Consequently, they prevent from different diseases such as ulcerative colitis and colon cancer (Johansson et al., 2008; Sluis et al., 2006; Velcich et al., 2002). In addition, the alteration of mucus glycosylation pattern impacts both the composition of intestinal microbiota and the gut architecture in mouse models (Arike & Hansson, 2016; Sommer et al., 2014; Staubach et al., 2012). For example, it has been demonstrated that the loss of mucin glycosylation especially the Core 1–derived *O*-glycans (the predominant form of mucin *O*-glycans in mice) led to the development of spontaneous colitis in a mouse model (Fu et al., 2011).

3. Control of mucus layer, mucin synthesis and associated deseases

Under normal physiological conditions, mucus is regularly renewed and this phenomenon results from a balance between mucus degradation and synthesis. Intestinal bacteria take important part in regenerating the mucus layer. The establishment of mucus-degrading bacteria in the host intestine starts few months after birth and is completed gradually with age (Palmer, Bik, DiGiulio, Relman, & Brown, 2007). This process is not only linked with the individual age but also with the diet. Midtvedt et al. (1994) have observed that the establishment of mucin degradation bacteria is delayed at least 4 months in breast-fed children than those fed with formula milk (Midtvedt, Carlstedt-Duke, & Midtvedt, 1994). Among intestinal bacteria known to have a mucolytic activity, *Akkermansia muciniphila* is one of the bacteria having large enzymatic machinery for mucins oligosaccharides breakdown (Derrien, Vaughan, Plugge, & de Vos, 2004; Derrien, Collado, Ben-Amor, Salminen, & de Vos, 2008). In mouse, it is one of the first users of mucus (Berry et al., 2013). This bacterium hydrolyzes mucins oligosaccharides and releases sugar residues such as N-acetylglucosamine, N-acetylgalactosamine, fucose or sialic acid, which can be used by other bacteria such as *Bifidobacterium infantis* (Derrien, Belzer, & de Vos, 2017; Garrido, Kim, German, Raybould, & Mills, 2011) or *Bacteroides thetaiotaomicron* (For comprehensive review see Koropatkin, Cameron, & Martens, 2012). Some other intestinal bacteria such as *B. bifidum*, *Ba. fragilis* and *Ruminococcus gnavus* also present a mucolytic activity (Katoh et al., 2017; Macfarlane & Gibson, 1991; Png et al., 2010).

The bacteria and / or their metabolites could modulate the expression and / or the secretion of mucins (Liévin-Le Moal et al., 2002; Smirnova, Guo, Birchall, & Pearson, 2003), the development of the mucus secreting goblet cells as well as the mucin glycosylation profile. It has been shown that *Ba. thetaiotaomicron* influenced the development of goblet cells in mono-associated rats by stimulating the expression of *klf4* gene encoding a transcription factor involved in goblet cell differentiation. This study also reported that this bacterium induced the expression of some genes encoding mucins (*muc2* and *muc4*) in the gnotobiotic rats compared to germ free model. Furthermore, the presence of two metabolically complementary bacteria, *Ba. thetaiotaomicron* and *Faecalibacterium prausnitzii* in gnotobiotic rats modulates the mucin glycosylation and goblet cells development in the colonic epithelium compared to germ free rats (Wrzosek et al., 2013).

Abnormal synthesis and / or processing of mucins leads to disruption of mucus structure and imbalance of mucosal barrier function. This could favour the development of a pathogen bacterium which becomes able to circumvent the protective function of the mucus barrier (Navabi, Johansson, Raghavan, & Lindén, 2013). Likewise, the abnormal functioning of the mucosal barrier leads to the development of intestinal diseases such as Inflammatory Bowel Disease (Ulcerative colitis, Crohn's disease) and eventually colorectal cancer (Bergstrom et al., 2017; Velcich et al., 2002; Wenzel et al., 2014). Patients with ulcerative colitis have shown modifications in mucus layer properties, mucin gene expression, mucin proteins synthesis as well as mucin's glycosylation structure (O-glycosylation, sulfation, and silylalation) leading thus to a deficient mucus barrier (Bergstrom et al., 2017; YamamotoFurusho, Ascaño-Gutiérrez, Furuzawa-Carballeda, & Fonseca-Camarillo, 2015). For instance, it has been reported that patients with active ulcerative colitis showed a significant decrease in gene and protein expression and production of MUC20 in epithelial cells from mucosa as well as submucosa (Yamamoto-Furusho, Ascaño-Gutiérrez, Furuzawa-Carballeda, & Fonseca-Camarillo, 2015). In addition, in mice model lacking core 2 β1,6-N-acetylglucosaminyl-transferase enzyme which initiates core 2 type protein O-glycosylation in mucin protein, an increase in intestinal permeability and susceptibility to experimental colitis (Dextran sulfate sodium induced colitis) has been observed (Stone et al., 2009). Furthermore, in Muc2- deficient mice, the bacterial contact with the epithelium explains the inflammation and the development of the cancer in these animals (Johansson et al., 2008).

4. Impact of food components on mucus and mucins

As discussed before, the control of mucus layer plays an essential role in maintaining health and should constitute a major target for conception of preventive strategies, including specific diets. Although knowledge about the impact of food ingredients on the digestive mucus layer remain relatively scarce. However, different dietary components such as milk proteins, bacteria or their metabolites, bioactive molecules (released either by bacteria in certain fermented foods or derived directly from food after digestion), fiber and short chain fatty acids have been shown to modulate the mucin gene expression and secretion *via* interaction with intestinal mucosa (Mack, Michail, Wei, McDougall, & Hollingsworth, 1999; Martinez-Maqueda, Miralles, Cruz-Huerta, & Recio, 2013; Rokana, Singh, Mallappa, Batish, & Grover, 2016; Burger-van Paassen et al., 2009). Hence, the positive effects of different food components on intestinal mucus properties particularly mucin modulation are discussed hereafter.

4.1. Effect of lactic acid and probiotic bacteria on mucin gene expression and secretion

Lactic Acid Bacteria (LAB) or probiotic bacteria present in fermented milk products can modify mucus layer by regulating mucins expression. Such modulation has been shown to 201 occur either by direct contact of bacteria with mucosal surfaces or by bacterial products (Liévin-Le Moal et al., 2002; Raja, Murali, Devaraj, & Devaraj, 2012; Smirnova, Guo, Birchall, & Pearson, 2003). It has been demonstrated that *Lactobacillus. plantarum* 299v and *Lb*. *rhamnosus* GG significantly increased the mucosal layer and mucosal integrity through upregulated expression of MUC3 which inhibited adherence of the enterohaemorrhagic *Escherichia coli* O157:H7 to intestinal epithelial cells HT29 (Mack, Ahrne, Hyde, Wei, & Hollingsworth, 2003). In the same perspective, Matter and coworkers have also reported that the expression of *MUC2* gene was induced at transcriptional and translational levels in Caco-2 cell cultures by *Lb. rhamnosus* GG (Mattar et al., 2002). Similarly, intestinal integrity was improved when mice were treated with *Lb. rhamnosus* CNCM I-3690. This was mainly due to enhanced production of goblet cells and consequently increased expression of mucin MUC2 (Martín et al., 2019). The treatment of Caco-2 cells with a cow milk, initially fermented with *Lb. paracasei* CBAL74 and heated to kill bacteria, led to increase of *MUC2* gene expression 214 and mucus layer thickness compared to non fermented milk (Paparo et al., 2018). Thus, this study highlighted that the increased mucus production is not restricted to living probiotic bacteria but their components could also preserve the mucus layer integrity. Similarly, it has been shown that administration of milk fermented with *Lb. plantarum* MTCC 5690 enhanced the intestinal barrier function by increasing expression of MUC2 in salmonella-infected mice 219 compared to those fed with unfermented probiotic milk (Rokana, Singh, Mallappa, Batish, $\&$ Grover, 2016). Besides the use of individual probiotic strains or probiotic-fermented milk, the oral administration of VSL#3 probiotic mixture (containing differents species of *Lactobacillus*, *Bifidobacterium*, and one species of *Streptococcus*) for seven days enhanced by 60 % basal luminal mucin content in rats. In addition, VSL#3 directly inoculated into 224 colonic loops significantly enhanced Muc2 mRNA expression compared to Muc1 and Muc3 for which only a slight increase was observed (Caballero-Franco, Keller, De Simone, & Chadee, 2007). Surprisingly, these results obtained *in vivo* were not reproducible *in vitro* when authors incubated VSL#3 bacterial cultures with cultured LS 174T intestinal epithelial cells suggesting that the cell culture conditions were not optimal for proper growth of probiotic bacteria. On the other hand, the authors explored the fact that bacterial metabolites could induce the mucin secretion in LS 174T cells. They reported that the bacterial secreted products from VSL#3 contained in the conditioned media remarkably stimulated mucin secretion and the stimulatory effect was similar to that obtained from the *in vivo* experiments (Caballero-Franco, Keller, De Simone, & Chadee, 2007). More recently, this probiotic mixture has been shown to reduce colonic inflammation and improve intestinal barrier function in Muc2 mucin-deficient mice indicating that VSL#3 did not require an intact mucin barrier to exert its beneficial effects (Kumar, Kissoon-Singh, Coria, Moreau, & Chadee, 2016). Finally, it has been shown that despite poor mucus-adhesion, *S. thermophilus* promotes mucus pathway in digestive tract of gnotobiotic rats by producing lactate which in turn increased the number of goblet cells and amount of mucus inducer KLF4 transcription factor. A significant increase in KLF4 protein due to lactate production by this bacterium was also reported in HT29-MTX cells (Fernandez et al., 2018).

4. 2. Impact of bioactive peptides on mucin expression and secretion

In addition to bacteria, hydrolysates from milk proteins have been shown to affect mucin expression and secretion in the intestinal mucosa (Table 2). Indeed, these proteins harbor bioactive sequences released either after digestion (*in vivo* or *in vitro*) or under the effect of proteolytic system of bacteria such as LAB during processing of fermented milks. These proteins provide peptides with different positive health effects on the host, including blood pressure lowering, immunomodulatory, antimicrobial, cholesterol-lowering and antioxidant effects. Several studies performed either *in vitro* or *in vivo* have also reported their impacts on the modulation of mucin secretion (Claustre et al., 2002; Han, Deglaire, Sengupta, & Moughan, 2008 ; Plaisancié et al., 2013; Plaisancié et al., 2015; Trompette et al., 2003). It has been shown that the luminal administration of casein and lactalbumin hydrolysates separately in the isolated perfused rat jejunum induced release of mucins. Conversely, the administration of native caseins did not significantly stimulate mucin secretion (Claustre et al., 2002). Likewise, when rats were fed for 8 days with test diets enriched either with hydrolyzed casein or a synthetic amino acid mixture as well as a protein-free diet, hydrolyzed casein enriched diet significantly up-regulated the expression of Muc3 and Muc4 mRNA in the small intestinal tissue and the colon, respectively, compared to other diets (Han, Deglaire, Sengupta, & Moughan, 2008). Fernández-Tomé and colleagues also reported significantly higher fecal mucin (*via* O-linked glycoproteins dosage in feces) content after 2- and 8-weeks administration of casein hydrolysate in rats. Indeed, long term intake of casein hydrolysate resulted in an increased expression of Muc2 and Muc3 mRNA in the ileum and the colon of rats (Fernández-Tomé et al., 2017). Furthermore, mucin secretion was escalated when β-casomorphin-7, an opioid peptide from bovine β-casein, was administered in the lumen of the

isolated vascularly perfused rat jejunum (Trompette et al., 2003). Further studies conducted with β-casomorphin-7 in intestinal mucin-producing cells of rat (DHE) and human (HT29- MTX) revealed an increased expression of different mucins in a time and a dose dependent manner. In DHE cells, Muc2 and Muc3 expression was enhanced whereas in case of HT29- MTX, an increase of transcription as well as secretion of MUC5AC was reported. The authors concluded that this peptide contributed significantly to mucin secretion *via* a direct effect on 272 intestinal goblet cells and the activation of μ -opioid receptors (Zoghbi et al., 2006). Other studies, performed with the human HT29-MTX cells, have also reported the modulation of mucin gene expression and stimulation by the total peptide pool derived from a fermented milk (yogurt) (Plaisancié et al., 2013; Plaisancié et al., 2015). Indeed, these peptides stimulated *MUC2* and *MUC4* gene expression as well as mucin secretion in HT29-MTX cells (Plaisancié et al., 2013). Interestingly, this study has identified peptide sequences carrying biological activity for further tests in rat models. Thereby, the oral administration of the 279 peptide fraction (f94-123) of β -casein to pups enhanced the number of goblet cells and Paneth cells along the small intestine which were associated with higher expression of intestinal mucins (Muc2 and Muc4) and antibacterial factors (Plaisancié et al., 2013). Furthermore, from the β-casein peptide sequence (f94-123), shorter peptides having different biological activities were also identified. These peptides up-regulated MUC4 expression in HT29-MTX cells whereas no effect was observed on the levels of secreted mucins MUC2 and MUC5AC. In contrast, other identified shorter peptide sequences increased the gene expression of *MUC2* and *MUC5AC* mucins (Plaisancié et al., 2015). In another study conducted in HT29-MTX cells, the stimulatory effect of casein hydrolysate and isolated peptides on mucin production 288 was also investigated. Casein hydrolysate and casein derived peptides mainly α_{s1} -casein 289 (f143-149) (AYFYPEL) and α_{S1} -casein (f144-149) (YFYPEL) significantly enhanced the expression of mucin gene encoding MUC5AC (over 1.7-fold basal level) after 4 h of exposure. The authors reported that casein hydrolysate containing both above cited peptides displayed a higher effect on mucin secretion as determined from mucin-like glycoproteins quantification than individual peptides (Martinez-Maqueda, Miralles, Cruz-Huerta, & Recio, 294 2013). Moreover, whey-derived peptides, α -lactorphin and β-lactorphin with described or probable opioid activity were also shown to induce secretion of MUC5AC mucin in HT29- MTX cells. Concerning the gene expression level of this mucin, it was up-regulated by α -lactorphin in these cells (Martinez-Maqueda et al., 2012). All these findings indicate that peptides derived from yogurt may play an important role in protecting intestinal lumen from damaging agents by modulating Paneth and mucus cells, which are known to play a crucial role in intestinal protection (Plaisancié et al., 2013). Hence, the presence of bioactive peptides including those probably released from LAB in dairy products from milk proteins could thereby improve gut health in the host.

4.3. Impact of other food ingredients on mucin expression and secretion

Besides milk proteins, rye bran, oat bran and soybean hull (Table 2) were also shown to increase goblet cell volume density in the proximal and distal small intestine of hamsters (Lundin et al., 1993). Food fibers can also positively impact the mucus layer through the production of SCFAs, such as acetate, propionate and butyrate products, which are the metabolic products of the intestinal microbiota. Among SCFAs, butyrate appears to be the most effective in stimulating mucus secretion in rat colon (Shimotoyodome, Meguro, Hase, Tokimitsu, & Sakata, 2000). Moreover, the effect of propionate and butyrate on mucin gene expression in human goblet cell-like LS 174T cells was also demonstrated. Butyrate at 1 mM, as well as propionate induced an increase in MUC2 mRNA levels. This study has also investigated the mechanisms that regulate butyrate-mediated effect on MUC2 synthesis. The authors have revealed an active butyrate-responsive region within the MUC2 promoter and have demonstrated that the regulation of MUC2 by butyrate is associated with increased acetylation of histone H3 and H4 and methylation of H3 at the MUC2 promoter (Burger-van Paassen et al., 2009). Similar results, i.e., stimulation of MUC2 mucin and increase in histone 3 acetylation were observed after treatment of LS 174T cells with 1-2 mM sodium butyrate. The secreted mucin was detected by histological PAS staining of carbohydrate chains of mucin and expression at the protein and mRNA levels was confirmed by immunoblotting with anti-MUC2 antibody and real-time RT-PCR, respectively (Hatayama, Iwashita, Kuwajima, & Abe, 2007). More recently, the variation of four mucin gene expression 323 (*MUC2, MUC3, MUC13, and MUC17*) in coculture of Caco-2/HT29-MTX cells was analyed 324 by real-time RT-PCR in the presence of antioxidant compounds like chlorogenic acid, epicatechin gallate, and quercetin (apple, tea, and coffee polyphenols, respectively). In 326 addition, the secretion level of MUC2 was also determined by enzyme-linked immunoassay 327 (ELISA) analysis. After exposing the cells to sub-lethal concentrations (10 μ M) of each of the different polyphenolic compounds over a period of 48 h, it is reported that each phenolic compound induces differential expression patterns of mucin genes and only a significant increase in *MUC17* gene expresion was observed especially with epicatechin gallate and quercetin. However, results from ELISA analyses showed no significant differences in secretion level of mucin MUC2 (Volstatova et al., 2019). In contrast, quercetin induced the secretion of MUC2 and MUC5AC mucins in quercetin-treated LS 174T cells. Both mucins were also detected at mRNA level. The secretion of these mucins in this cell line was attributed to enhanced extracellular signal regulated kinase (ERK)1-2 activity as its inhibition by the MEK inhibitor PD98059 reverted the stimulatory effects of quercetin. Besides LS 174T cells, quercetin treatment also enhanced the mRNA levels of MUC2 and MUC5AC *via* ERK1-2 pathway in the human intestinal epithelial Caco-2 cells (Damiano et al., 2018).

5. Research approaches to improve/design new functional ingredients/foods

The increasing costs of health as well as the desire of numerous consumers to support a preventive rather than a curative approach have created favourable conditions to develop functional foods, which are claimed to display health specific properties because of the 343 presence of some specific functional and / or absence of some harmful components (Gupta $\&$ Abu-Ghannam, 2012). Although the informations presented above remain scarce, the evidence provided through the action of probiotics, bioactive peptides or some polyphenolic compounds suggest that they may be valuable ingredients / tools for strengthening the digestive mucus layer or mucus barrier that is central to consumer health. However, it must be underlined that most of the work presented in this review focused on effects of tested ingredients on mucins transcription (Table 2). Nevertheless, other parameters can also be explored such as effects of food ingredients on i) the microbiota composition and equilibrium, 351 ii) the mucus producting cells and thus on the intestinal epithelium and / or iii) glycosylation profile of mucins. The search of such ingredients constitutes a first step in designing functional foods, i.e., before including them in a food matrix, study their bioavailability in this matrix during digestion followed by their interactions with other constituents of the diet.

Two types of approaches can be used to identify functional ingredients directed towards mucus and mucins. The first is to screen high throughput probiotic strains, bioactive peptides or other dietary compounds for their adhesion capacity (for probiotics) and / or their ability to induce the production of mucins (induction of transcription/translation/secretion) using cell models. Indeed, several cell lines are known to exhibit the functional characteristics of intestinal cells found in small and large intestines and express different mucins (Table 3). Such screening can concern new probiotic strains or food ingredients, but also ingredients already described for other properties such as the opiod bioactive peptides (see above). The second approach consists of using animal models to explore the capacity of some ingredients to increase the goblet cells and then the mucus production or modulate the microbiota and the glycosylation profile of mucins, a path that seems insufficiently explored so farThis implies to be able to characterize the microbiota rapidly and on cohorts, to determine which type of flore is favored, how does it is maintained and what are the determinants. In addition; it could be interesting to explore the relationship between food ingredients and the glycosylation state of mucins.

To conclude, given the impact of diseases related to dysfunction of the digestive tract and especially the amount of mucus and mucins produced, it is necessary to intensify research in the field of functional ingredients with the prospect of exploiting them in the maintenance of gut health.

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Table 1 Mucins in the digestive tract, their localization, cell expression and physiological functions

797 **Table 2** Food ingredients and their impact on mucin expression / secretion

799 **Table 3** Summary of most cell lines *in vitro* models used in bacterial adhesion studies

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