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Topical issue on:

**BIOAVAILABILITY AND TISSUE-TARGETING DIETARY LIPIDS:
NEW APPROACHES TO THEIR FORMULATION?
BIODISPONIBILITÉ ET CIBLAGE TISSULAIRE DES LIPIDES ALIMENTAIRES :
NOUVELLES STRATÉGIES POUR LA FORMULATION ?**

PROCEEDINGS

OPEN ACCESS

Towards infant formula biomimetic of human milk structure and digestive behaviour

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Abstract – Lipids of human milk or infant formula convey most of the energy necessary to support the newborn growth. Until recently, infant formula chemical composition had been optimized but not their structure. And yet, more and more proofs of evidence have shown that lipids structure in human milk modulates digestion kinetics and is involved in metabolic programming. Indeed there is a striking difference of structure between human milk which is an emulsion based on dispersed milk fat globules (4 μm) secreted by the mammary gland and submicronic neoformed lipid droplets (0.5 μm) found in infant formula. These droplets result from a series of operation units. This difference of structure modifies digestion kinetics and emulsion disintegration in the intestinal tract of the newborn. This difference persists along gastric phase which is mainly dominated by acid and enzyme-induced aggregation. Lipid droplets size is thus the key parameter to control gastric lipolysis and emptying and intestinal lipolysis. This parameter also controls proteolysis since adsorbed proteins are more rapidly hydrolyzed than when in solution. In animal models, these differences of lipid structure would also impact digestive and immune systems' maturation and microbiota. Lipid structure during neonatal period would also be involved in the early programming of adipose tissues and metabolism. The supplementation of infant formulas with bovine milk fractions (milk fat globule membrane extracts, triacylglycerol) or recent development of large droplets infant formula, along with new fields of innovation in neonatal nutrition, are here reviewed.

Keywords: human milk fat globules / lipid structure / infant formula / neonatal digestion / programming

Résumé – Vers des formules infantiles mimant la structure et le profil de digestion du lait maternel.

Les lipides laitiers du lait maternel ou de formules infantiles fournissent la plupart de l'énergie disponible pour la croissance du nouveau-né. Jusqu'alors, la composition chimique des formules infantiles a été optimisée mais pas leur structure. Or il est de plus en plus démontré que la structure des lipides dans le lait maternel module les cinétiques de digestion et participe à la pré-programmation métabolique. Il existe en effet une différence majeure entre les émulsions natives de globules gras laitiers (4 μm) sécrétées par la glande mammaire et la structure des lipides dans les formules infantiles qui résultent d'une succession d'opérations unitaires de transformation et sont constituées de gouttelettes submicroniques (0.5 μm) avec des interfaces néoformées. Cette différence de structure modifie les cinétiques de digestion et de déstructuration des émulsions dans le tractus digestif des nouveau-nés. De façon générale, la différence de structure initiale entre lait maternel et formules infantiles se maintient pendant toute la phase gastrique, qui est surtout dominée par des mécanismes d'agrégation acide et enzymatique. La taille des gouttelettes est le paramètre clé pour contrôler les lipolyses gastrique et intestinale et module également la vitesse de vidange gastrique néonatale. Ce paramètre contrôle également les cinétiques de protéolyse puisque les protéines adsorbées sont hydrolysées plus rapidement que les protéines solubilisées. Ces différences de structure impacteraient également chez l'animal le développement des systèmes digestif, immunitaire et la mise en place du microbiote. La structure des lipides ingérés pendant la période néonatale pourrait également

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participer à la pré-programmation des tissus adipeux et du métabolisme chez l'adulte. La réintroduction d'extraits membranaires et de triacylglycérols issus de lait bovin dans les formules, le développement de formules nouvelles basées sur des émulsions moins fines ainsi que les dernières tendances de développement de formules infantiles biomimétiques sont reportées et discutées dans cette synthèse.

Mots clés : globules gras / lait maternel / structure lipidique / formules infantiles / digestion néonatale

1 Introduction

Human milk (HM) chemical composition was established for the first time in 1838 (Cone, 1981). Since then, this chemical composition has been the basis for preparing infant formula (IF). However, proofs of evidence of the part played by HM structure in the concept of early programming have been produced over the last decade (Oosting *et al.*, 2011). These new findings have refocused a large part of innovations in targeting IF than are biomimetic of HM in terms of structure but also in terms of digestive behaviour.

Diet in the early life modulates not only short-term outcomes such as growth, but also long-term health outcomes along life (Victoria *et al.*, 2016). Factors which are potentially influenced by diet during neonatal period, include neuro-development, metabolic health, immune competency, atopic disease, establishment of the mucosal microbiome and behavioural responses to foods and eating (Maslowski and Mackay, 2011; Arrieta *et al.*, 2014; Raiten *et al.*, 2016).

Despite the great advances in the optimization of IF composition, IF cannot provide all the numerous immune protective and bioactive factors present in HM. In addition, IF lipids structure has remained up to now very different from HM, which can lead to a difference in the kinetics of digestion and assimilation of nutrients (Armand *et al.*, 1996; Bourlieu and Michalski, 2015; Bourlieu *et al.*, 2015a). Indeed lipids in HM are found under the form of dispersed droplets called Milk Fat Globules (MFG, \varnothing 0.1–20 μm , average = 4 μm). This native MFG is a unique biophysical element differing from other lipoproteic objects (lipoproteins, oleosomes, etc.) by its external trilayered membrane inherited from its secretion by the lactocyte. The MFG membrane (MFGM) is tensioactive and consists in potentially bioactive polar lipids (glycerophospholipids, sphingolipids and glycosphingolipids such as gangliosides), cholesterol, proteins, glycoproteins, enzymes and minor components such as RNA/miRNA (Lopez and Ménard, 2011; Bourlieu and Michalski, 2015; Lopez *et al.*, 2015). Due to its heterogeneous chemical composition, this membrane presents a specific physical-state, with phase coexistence: nanodomains called lipid rafts (*i.e.* sphingolipids and cholesterol that aggregate along sections of the membrane) are in the lipid-ordered phase and are characterized by a high degree of rigidity; besides, other parts of the MFGM concentrating more unsaturated molecules are in liquid-disordered state (Gallier *et al.*, 2010; Lopez and Ménard, 2011). On the contrary, the IF emulsion is based on several reassembled dairy and non-dairy fractions, resulting from successive technological operations. IF lipids are generally composed of submicronic droplets (0.3–0.8 μm) stabilized by a neoformed membrane mainly based on milk proteins and non-dairy tensioactive molecules.

Since HM is a complex and dynamic fluid very difficult to mimic, most improvements of IF have consisted in the addition of components individually identified as bioactive in HM, in

the optimized IF (Committee on the evaluation of the addition of ingredients new to infant formula, 2004; Gallier *et al.*, 2015; Zou *et al.*, 2016). Notably, components such as long chain polyunsaturated fatty acids (LCPUFA), lactoferrin, nucleotides, essential amino-acids (AA), oligosaccharides (OS), prebiotics, probiotics can be added into some IF.

The interest for HM lipid structure started in the 70s with studies showing specific triacylglycerol (TG) structure in HM and resulting improved fatty acids absorption (Tomarelli *et al.*, 1968). Understanding the higher levels of organization of human MFGM or MFG has been delayed until the development of modern biophysical tools (among which confocal laser scanning microscopy). And yet, despite these developments and the obvious nutritional importance of HM, data on HM lipid structure remains limited (Simonin *et al.*, 1984; Michalski *et al.*, 2005b; Zou *et al.*, 2012; de Oliveira *et al.*, 2015, 2016, 2017). Many structural characteristics of HM MFG have been inferred from cow's MFG which paradoxically has been more studied.

Over the last decade, going beyond HM individual lipids has been the new priority. Understanding human MFG's fate and behaviour in the gastro-intestinal tract, and the part MFG play in intestinal, immune and metabolic programming has become central question in neonatal nutrition. How milk emulsion droplets interact with each other and respond to different environmental conditions depends on the emulsion properties (*i.e.* size of MFG and quantity of interface, type and organization of proteins) and on the characteristics of the interfacial layers (*i.e.* thickness, composition, charge) (Singh *et al.*, 2009; Bourlieu and Michalski, 2015; Gallier *et al.*, 2016). These characteristics differ among milks (*e.g.* HM and bovine milk) and are modified after milk technological treatments (heat treatments, homogenization) (Innis *et al.*, 2010; Gallier *et al.*, 2012, 2013a, 2013b; Garcia *et al.*, 2014; Bourlieu *et al.*, 2015b; Le Huërou-Luron *et al.*, 2016). Designing IF that mimic HM digestive behaviour supposes prior characterization of this behaviour. Recent work proposed by (de Oliveira *et al.*, 2015, 2016, 2017) have described HM structural change during its passage through the upper gastrointestinal tract.

In this context, the objectives of the present reviews is to highlight recent findings about HM lipid structure, its evolution during digestion and putative part in nutritional programming which have opened the way to recent innovations in IF processing.

2 Proofs of evidence of perinatal lipid nutrition and HM's structure being involved in programming and consequences for neonatal nutrition

At birth, the gastrointestinal tract is immature and its development continues during the first year of life. Although genetically programmed, the infant's intestinal and immune

development is modulated by the diet (Le Huërou-Luron *et al.*, 2010). The critical part played by essentials FA (linoleic acid LA and alpha-linolenic acid ALA) and LCPUFA (especially docosahexaenoic acid DHA and arachidonic acid ARA) during neonatal period in brain growth, cognitive skills development, motor and retinal functions were established early (Yehuda *et al.*, 2005). Infant brain growth and associated DHA accumulation is significant during the last trimester of gestation and continues during early infancy (Lapillonne, 2007). And yet, the capacity of bioconversion of LA and ALA to LCPUFA is limited during neonatal period inducing a dependence on the dietary intakes from HM or IF lipids. The impact of lipids on the early programming of intestinal functions and metabolism was questioned later than for carbohydrates or proteins. Early works of Ailhaud and collaborators established that among LCPUFA, ARA was a precursor of prostacyclin in preadipocytes and was very adipogenic *in vitro* (Gaillard *et al.*, 1989). Contribution of ARA to adipose tissue development during perinatal period was further investigated in rodents. This investigation allowed demonstrating that mother's diet with variable LA/LNA ratios (LA being ARA precursor) impacted pup's body composition (fat mass, adipocytes size, etc.) (Massiera *et al.*, 2003). Since gestation and first year of life are the most important windows of adipose tissue proliferation, they set the hypothesis that this difference in body composition persisted later in life. These observations showed that ARA *via* metabolism of LA was a key determinant in fat mass and globally underlined that LA/LNA ratios during perinatal period could have important programming effect on body composition (Ailhaud *et al.*, 2006). Further proofs of evidence of lipids being involved in programming other functions were obtained in 2010 in animal models (rodents) by two research groups (Innis *et al.*, 2010; Oosting *et al.*, 2010). These results were obtained on non-dairy lipids but paved the way for further investigations on 'dairy lipid structure and programming' conducted just after by Oosting *et al.* (2011, 2012, 2014).

2.1 The *n*-6/*n*-3 ratio in key factor of perinatal nutrition and programming

Innis *et al.* (2010) proposed that the increasing incidence of inflammatory bowel disease observed in Western countries over the last decades could be linked to dietary factors including perinatal lipid nutrition. They set the hypothesis that perinatal lipid nutrition affects, directly or indirectly through microbiota modulation, the epithelial barrier function and development, immune system and inflammatory response. Thus perinatal lipid nutrition could alter early intestinal development and programs future response to inflammatory diseases in adulthood in animal model (rats). In their study, perinatal supply of *n*-6 and *n*-3 polyunsaturated FA was modified through the mother's diet. Diets was administrated throughout gestation and lactation and contained as % energy either 20% safflower oil (SO), 20% canola oil (CO), or 8% fish oil (FO) plus 2% SO (10% FO), or 18% FO plus 2% SO (20% FO). At 15 days of age, pups in the 20% and 10% FO groups had lower ARA (20:4 *n*-6) and higher eicosapentaenoic acid (20:5 *n*-3) and DHA (22:6*n*-3) in colon phospholipids ($P < 0.01$), shorter intestinal crypts ($P < 0.05$), and higher

paracellular permeability than in SO or CO groups. Later on, after weaning to standard diet (21 days) and at a stage representing adulthood (*i.e.* 3 months of age) male offspring in the FO groups showed homogeneous colon phospholipid FA composition but lasting reduction of crypt depth and an enhanced inflammatory response when challenged with dinitrobenzene sulfonic acid (chemically-induced colitis). The eicosanoid metabolite prostaglandin E2 (PGE2) synthesized *via* cyclooxygenase COX-1 and COX-2 from ARA and influencing intestinal cell crypt survival, proliferation and permeability was proposed as putative mechanisms for this programming effect.

At the same time, Oosting *et al.* (2010) investigating further Ailhaud's group hypothesis (Ailhaud *et al.*, 2006), established still in a rodent model, that *n*-3 LCPUFA-rich neonatal diet programmed adult body composition (reduction of 30% fat accumulation during western style diet challenge) and metabolic homeostasis. Healthier plasma lipid profile, plasma glucose parameters and less adipocytes hypertrophy were observed in the group of mice that had received the *n*-3 enriched diet during neonatal period. Oosting *et al.* (2010) set the hypothesis that unbalanced *n*-6/*n*-3 neonatal intake favoured LA conversion to ARA and its eicosanoids metabolites. These metabolites may directly stimulate adipogenesis through activation of peroxisome proliferator-activated receptor delta and gamma.

Breastfeeding is associated with specific growth pattern compared to classic IF feeding, this pattern is marked by moderate early growth and lower fat mass (Nommsen-Rivers and Dewey, 2009). Several meta-analyses have shown that breast-feeding has a small protective effect against obesity later in life (Agostoni *et al.*, 2009). Relying on these positive outcomes and considering the strong difference of lipid structure between IF and HM, Oosting *et al.* (2014) set the hypothesis that HM lipid structure was involved in programming.

2.2 Difference in lipid structure between human milk and classic infant formula and its potential influence on programming

Human milk is an emulsion of lipid in water and lipids are organized under the form of MFG stabilized by a trilayered membrane (Fig. 1). This membrane is mainly composed of polar lipids (phospholipids and glycosphingolipids) but also of apolar lipid such as cholesterol. MFG structure has been presented as a key element for HM lipid paradoxical metabolic fate: MFG is a rapid conveyor of energy through its triacylglycerol (TG) core but contains some low-digestible bioactive complex lipids and proteins which influence lipid metabolism and contribute to intestinal and systemic health. On the contrary IF formula are generally based on a blend of vegetable lipids and found under the form of submicronic droplets stabilized by dairy proteins or non-dairy tensioactive (vegetable lecithins, functionalized lecithins or esters of partial glycerides mainly) (Fig. 2). Taken individually, milk polar lipids present in MFGM have proven bioactivities, with reported impact on cell metabolism, on brain development and cognitive functions, and on immunity and gut health (Hirabayashi and Furuya, 2008; Kullenberg *et al.*, 2012; Lonnerdal, 2014). Some benefits of the MFGM have been

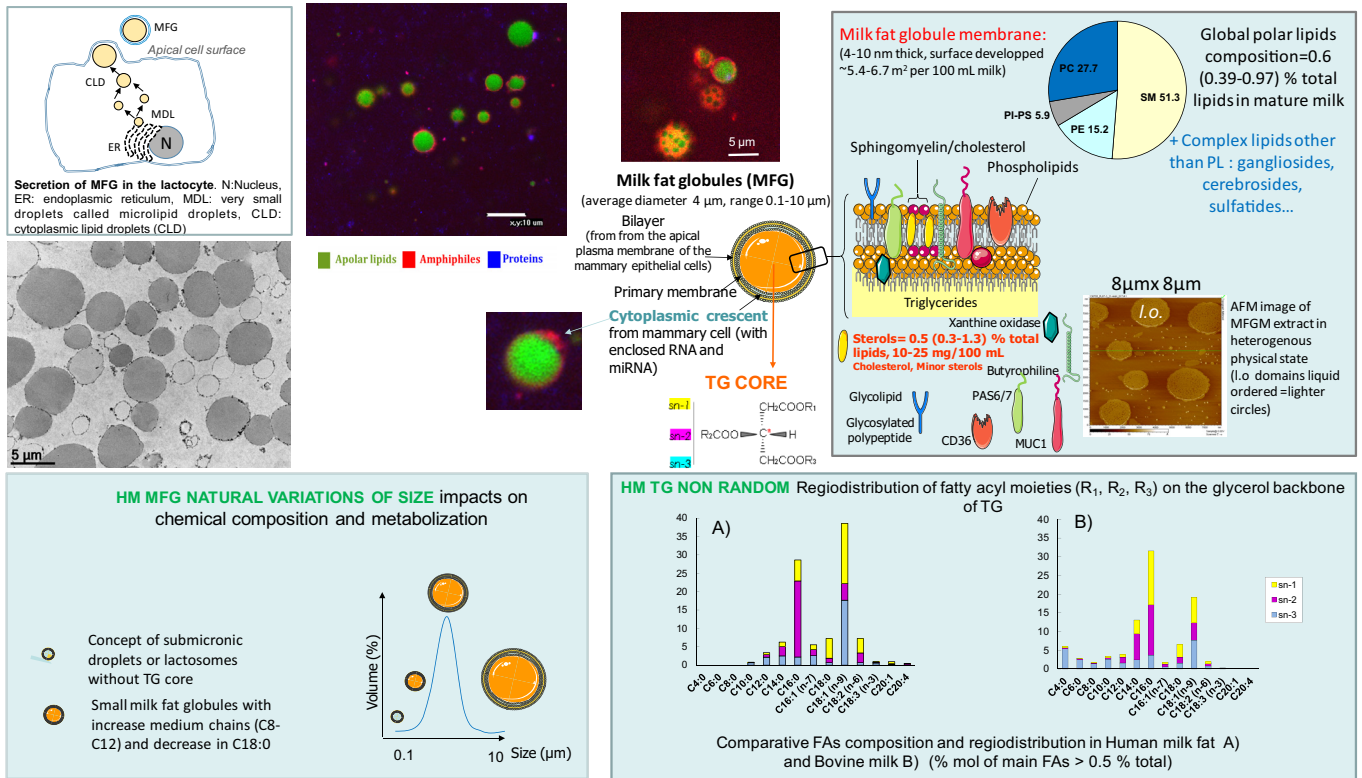


Fig. 1. Origin and structure of lipids in human milk. Adapted with modifications from Bourlieu and Michalski (2015) and Bourlieu *et al.* (2015a).

associated specifically to the combination of its components and their structural organization (Bourlieu and Michalski, 2015). MFG membrane configuration is reported to protect some bioactive molecules and retard their digestive breakdown, allowing them to exert their function until the distal intestinal part. For instance, the limited susceptibility of sphingomyelin to digestion favours the persistence of sphingomyelin-cholesterol complexes in the proximal part of the intestine. In addition, sphingomyelin and its metabolites (ceramide, sphingosine, ceramide-1-P and sphingosine-1-P) influence TG hydrolysis, cholesterol absorption, lipoprotein formation and mucosal growth in the gut (Nilsson and Duan, 2006). It was also hypothesized that milk sphingolipids could play a protective effect against bacterial toxins and bacterial development *via* a competition for bacteria binding sites as many bacteria binds to epithelial cell *via* glycosphingolipids (El-Loly, 2011). Milk sphingolipids transiting in the gut may decrease pathogens adherence to the intestinal mucosa facilitating pathogens elimination (Sprong *et al.*, 2001) and thus influence the gut microbiota composition. The bactericidal activity of digestion products of sphingolipids and more specifically of lysosphingomyelin was evidenced *in vitro* by Sprong *et al.* (2001). Along sphingolipids, glycosylated proteins and lipids that also escape upper intestinal digestion are also crucial factors contributing to MFGM bioactivity. Additionally, the protective structure of the MFG membrane contributes to the high oxidative stability of expressed HM, despite its high content in LCPUFAs (Michalski *et al.*, 2008). Considering the striking difference in size and interfacial composition between classic IF and HM, (Oosting *et al.*, 2012)

investigated, still in an animal model (mice), whether the administration of a concept ‘more breastmilk like’ formula compared to a standard formula could program adult body composition and metabolism. The concept IF Nuturis[®] (WO2013135739A1) is based on large vegetable fat droplets (mode, *i.e.* most frequent size in the distribution in volume, of 6.25 μm) covered by milk phospholipids (2 g/L). Thus the particle size distribution, emulsion interfacial composition and organization is very close, although not totally similar, to HM (Gallier *et al.*, 2015). The administration during the neonatal period (from day 16 to 42) of this Nuturis[®] formula in mice, followed by a western diet challenge, resulted during adulthood to lower fat accumulation, lower fasting plasma leptin, resistin, glucose and lipid (TG and total cholesterol) than in the group fed conventional formula (Oosting *et al.*, 2012). The mechanisms underlying this nutritional programming effects of Nuturis[®] remained at this stage unclear and very likely plurifactorial. However, authors hypothesized that emulsion droplets size and interface could modify digestion kinetics. At this stage, limited and quite conflicting results had been published on milk emulsion digestion (Armand *et al.*, 1996, 1999) suggesting that adult would digest faster small droplets based emulsion which would not be the case for infants. Since then, these results have been fully clarified and will be summarized in Section 3. At this stage, however, the fact that kinetics of lipid digestion could further modulate plasma TG appearance (Michalski *et al.*, 2006) and β-oxidation rate (Michalski *et al.*, 2005a) was already established in animal models and later on confirmed in human (Vors *et al.*, 2013). In a further study (Oosting *et al.*, 2014), still using a very close design study in mice, it was shown

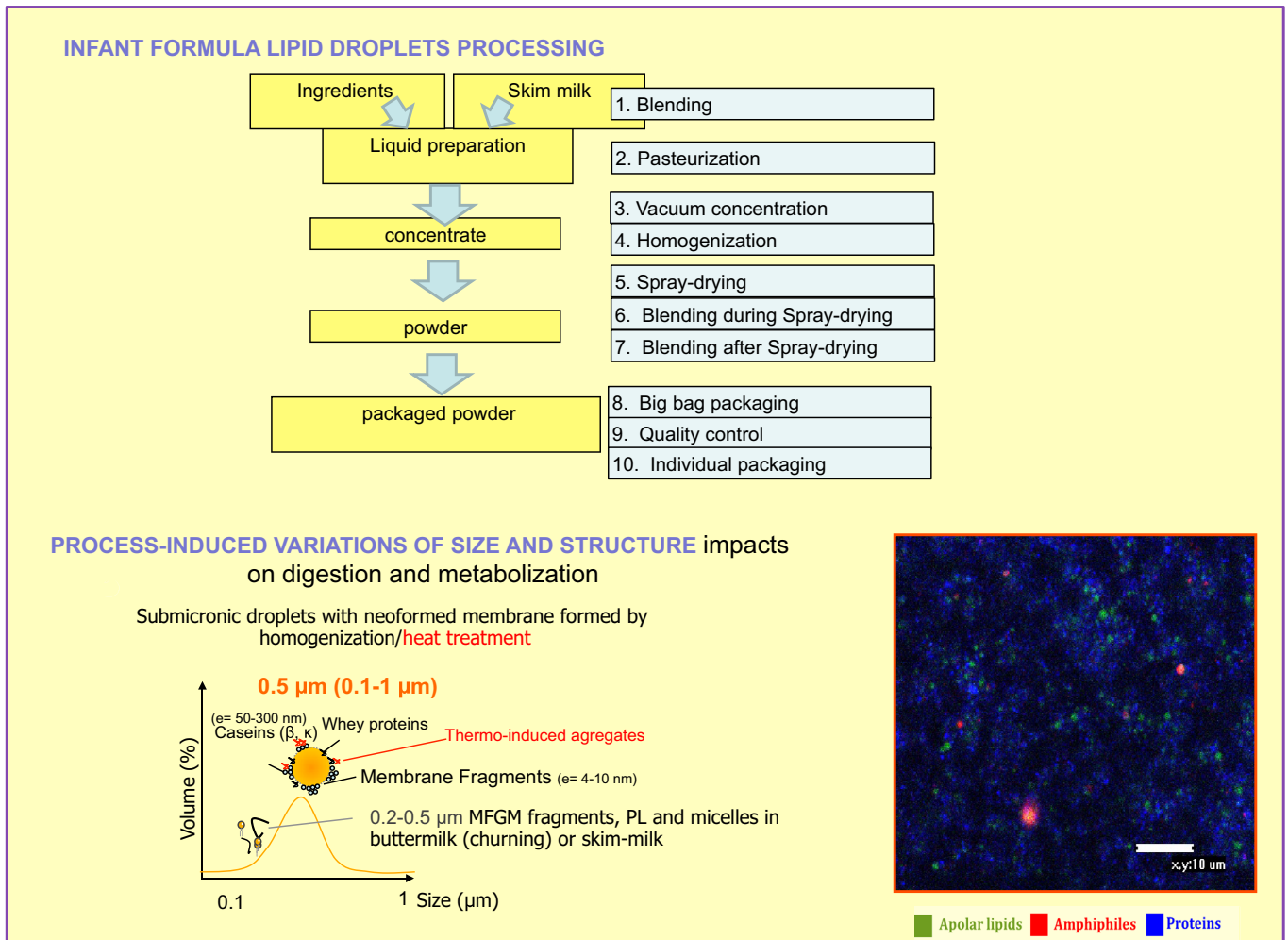


Fig. 2. Origin and structure of lipids in infant formulas. Adapted with modifications from Bourlieu and Michalski (2015) and Bourlieu *et al.* (2015a).

that neonatal administration of Nuturis[®]IF induced in adulthood, reduced adipocyte size without affecting their number in epididymal white adipose tissue. This was accompanied by the modulation of several transcription factors involved in metabolic regulation such as peroxisome proliferator-activated receptor gamma, CCAAT/enhancer-binding protein and retinoid X receptor.

The last recent study on the physical structure of lipids and its programming effect, aimed at evaluating whether the protective effect against obesity in adulthood was arising from the droplet size and/or from the droplet membrane (Baars *et al.*, 2016). Only the administration of large droplets with phospholipid coating, *i.e.* the combination of the two factors under the Nuturis[®] formula form, during the neonatal period reduced fat accumulation in adulthood. A direct effect of MFGM contained in the concept formula, was also weakened by a recent study comparing the growth of infants fed with a standard IF *vs.* a formula enriched with MFGM fragments (Timby *et al.*, 2014a). The study did not reveal any difference in infant growth, weight gain and body fat at 12 months. Conversely, HM lipids affected preadipocyte differentiation *in vitro* in the absence of the standard adipogenic compounds, which was not observed with IF lipids (Fujisawa *et al.*, 2013).

Thus, among the structural aspects of HM lipids, both lipid droplet size as well as the MFGM coating, may contribute to its reported protective effect against obesity (Baars *et al.*, 2016). Nuturis[®]IF is currently being testing in an interventional clinical trial in healthy term infants (NCT01609634) for its tolerance, safety and impact on growth and body composition up to 24 months.

3 Approaches to develop biomimetic IF

3.1 Mimicking milk fat globule coverage

Since the stability of large droplet-based emulsion like Nuturis[®]IF is a real technological challenge, another more straightforward strategy to produce more biomimetic IF has consisted in supplementing classic IF with bovine MFGM extract, without considering the lipid droplet size.

On a technological point of view, MFGM fractions are isolated from by-products of the butter industry, such as buttermilks and butterserums (El-Loly, 2011; Conway *et al.*, 2014). These fractions can be concentrated thanks to the development of micro and ultrafiltration technologies (Morin *et al.*, 2006, 2007a, 2007b; Gassi *et al.*, 2016). These compounds

appear as an attractive alternative to non-dairy tensioactive or milk proteins to better approach HM ultrastructure. These compounds can be added in the IF premix. An important aspect of such addition is not only their protein/phospholipids ratios but also the order of introduction of these two amphiphiles in the emulsion which can impact on the emulsion interfacial composition and structure (Bourlieu *et al.*, 2015a; Gallier *et al.*, 2015). The precise interfacial composition of such biomimetic IF remains to be characterized as well as its susceptibility to digestion.

In terms of clinical impact, six double-blinded randomized controlled Clinical trials in infants and children have investigated the safety and efficacy of such IF supplementation with bovine MFGM (Hernell *et al.*, 2016). Results are mostly positive and demonstrate benefits in terms of cognition, behaviour, gut health and immunity (Zavaleta *et al.*, 2011; Gurnida *et al.*, 2012; Veereman-Wauters *et al.*, 2012; Billeaud *et al.*, 2014; Poppitt *et al.*, 2014; Timby *et al.*, 2014a, 2014b, 2015). Although the intervention strategy and MFGM concentrates added are still too heterogeneous, these results provide evidence of the beneficial effects of individual components of MFGM.

On the other hand, at a very different level, *i.e.* the molecular level, it is well known that interfacial composition of milk lipids influence digestive lipases activity. The way MFGM organization modulates digestive lipase activity is now well described. Considering first gastric lipase, (Bourlieu *et al.*, 2016) established that its adsorption was favoured in more fluid like part of the MFGM, *i.e.* in liquid-disordered phase of the membrane where products of lipolysis, free fatty acids, concentrate. In addition, the presence of residual negative charges in the MFGM could also favour lipolysis kinetics. On the contrary gastric lipase would not adsorb onto liquid-ordered domains where sphingomyelin and cholesterol concentrate. Quite similar observation in intestinal phase has been reported for human pancreatic lipase which selectively adsorbed with colipase onto liquid disordered phase (Chu *et al.*, 2010). The close packing of the hydrophilic lipid headgroups in liquid ordered domains prevented lipase adsorption by steric hindrance. These liquid ordered domains concentrating SM and cholesterol are thus supposed to be transported along the intestine, playing potential antimicrobial activities. Conversely, liquid disordered domains where free fatty acids concentrates constitutes evolving and preferential zone for pancreatic lipase, colipase and carboxylester lipase adsorption (Sugar *et al.*, 2001, 2003). Considering this high effect of MFGM composition and organization on digestive lipase activity, the non-dairy tensioactives (vegetable lecithins, functionalized lecithins or esters of partial glycerides mainly) authorized in IF can be questioned. These non-dairy tensioactive are notably used in hydrolyzed IF where tensioactive dairy protein can not play their stabilizing part. Indeed the non-digestibility of some of these constituents, citric acid esters of mono- and diglycerides, was recently evidenced by Amara *et al.* (2014).

3.2 Mimicking MFG core: the importance of TG regiodistribution

The core of MFG is composed mainly of TG. These TG have a specific structure induced by the non-random regiodistribution of the FA on the glycerol backbone. This specific regiodistribution is a fairly constant feature of HM

that does not evolve with lactation stage (Martin *et al.*, 1993). Proofs of evidence of the nutritional importance of such regiodistribution have been demonstrated as soon as the 70s (Tomarelli *et al.*, 1968). This regiodistribution differs between bovine and HM (Fig. 1), but also more importantly between IF and HM, since the source of fat in IF is in most cases a combination of blended vegetable oils which mimics the FA composition of HM (Bourlieu *et al.*, 2015a; Zou *et al.*, 2016). Vegetable fats also show non-random regiodistribution but in the opposite way to dairy fat: their saturated FA are typically positioned on the external *sn*-1,3 positions, whereas unsaturated FA are mainly located in *sn*-2 position (Straarup *et al.*, 2006; Innis, 2011). As a result, most commercial IF based on blends of vegetable fats present a TG structure that differs largely from the one of HM fat (Kurvinen *et al.*, 2002). The regiodistribution of the FA on TG affects the FA lipolysis and its subsequent uptake (Bourlieu *et al.*, 2009; Zou *et al.*, 2012), as digestive lipases are also regio- or stereo-selective. For instance, MCFA (C8:0 to C12:0) are mainly esterified in *sn*-3 position in HM, and thus selectively released by the *sn*-3 stereospecific human gastric lipase (HGL). These medium chain FA are then absorbed *via* the portal vein, transported to the liver to be β -oxidized, and thus constitute a rapid source of energy for the newborn (Bourlieu *et al.*, 2015a). Oleic acid (C18:1 *n*-9) is usually found at the *sn*-1 and *sn*-3 positions, which corresponds to the stereo-specificity of the human pancreatic lipase. Palmitic acid (C16:0) mostly present at the *sn*-2 position favours its intestinal absorption as *sn*-2 monoglyceride (Jensen *et al.*, 1978; Bernback *et al.*, 1990; Bourlieu and Michalski, 2015) and limits loss of calcium/palmitate salts in stools.

In order to get closer to the structure of the TG found in HM, fat analogs (Betapol[®] and INFAT[®]) are commercially available; these analogs are obtained after enzymatic or chemical modifications and aimed to mimic the FA regiodistribution found in HM. Notably, a special attention is paid to the positional distribution of palmitic acid, which presents substantial benefits for the infant growth when esterified at the *sn*-2 position, as in HM (Innis, 2011; Zou *et al.*, 2016). However, these analogues are still not found in commercial IF (Straarup *et al.*, 2006) probably due to their cost but also to the reject of health claim petitions for beta-palmitate (increase in calcium absorption and softening of stool) by the European food Safety authority in 2011 and 2014 (Zou *et al.*, 2016).

The reintroduction of bovine milk fat which share close TG regiodistribution with HM up to 50% total fat in IF has been advocated by various research groups (Bourlieu *et al.*, 2015a; Delplanque *et al.*, 2015; Hernell *et al.*, 2016; Lonnerdal, 2016). In addition dairy fat enriched with ALA was proven superior to plant oil blend for DHA brain restoration in animal model (Delplanque *et al.*, 2011, 2013). ALA deficient rat pups fed by a palm-oil-blend-based diet (0.4% ALA) over gestation and lactation were, at weaning, fed either a palm-oil-blend-based diet or dairy fat supplemented with sunflower and rapeseed oils. Both diets contained recommended amounts of ALA *i.e.* 1.5%, 14–16% LA and LA/ALA ratio ~9–10. The dairy fat diet induced more DHA accumulation than the vegetable oil based diet. Supplementation of the vegetable-oil-based diet with DHA (0.12) and ARA (0.4) to mimic LCFA enriched IF did not modify this trend. No modulation of desaturase or elongase activity was detected in the various diets. Authors

thus set the hypothesis that short chain FA contained in dairy fat which are good substrate for β -oxydation could prevent ALA from β -oxydation. ALA partitioning towards elongation and desaturation pathways would thus be favoured.

3.3 Mimicking milk fat globule coverage and core

Le Huërou-Luron *et al.* (2014) hypothesized and checked in animal model (piglets) that IF with milk fats stabilized by MFGM closer to HM would impact differently neonatal digestive physiology and microbiota. The presence of milk lipids in IF led to a higher resistance to digestion of dairy proteins (casein and β -lactoglobulin) as compared to formula with vegetable lipids. This effect can be due to differences in the emulsion interface composition between the two formulas, with more dairy proteins adsorbed in the vegetable fat based IF devoid of MFGM; indeed both lactoglobulin and caseins are more susceptible to proteolysis when adsorbed to an oil-water interface than in solution (Macierzanka *et al.*, 2009). The developmental profile of intestinal immune cells was also modulated by the lipid composition of formula. Pro-inflammatory cytokine secretion of mesenteric lymph node cells was enhanced in milk fat formula-fed piglets compared to vegetable fat fed ones. Overall, the addition of milk fat to IF led to the maturation of the immune system of milk fat fed piglets was closer to the one of mother-fed piglets. Finally, fat source also impacted the microbiota: Piglets fed the formula with a blend of vegetable fat and milk lipids had a higher percentage of fecal *Proteobacteria* and *Bacteroidetes* and lower *Firmicutes* than piglets fed the vegetable fat formula. The incorporation of both milk fat and MFGM fragments in IF by modifying gut digestion, its dynamic of immune maturation and faecal microbiota composition, changes formula immune functionality (Le Huërou-Luron *et al.*, 2016).

Addition of dairy fat and MFGM also induced the addition of sterol enclosed in the MFG core. Sterols range from 10 to 25 mg/100 ml. Cholesterol stands for around 90% of sterol and globally declines in % of total fat as lactation progresses (Jensen, 1999; Michalski, 2013). It was suspected very early that milk fat cholesterol ingested during neonatal period and before total maturation of hepatobiliary system contributed to cholesterol homeostasis in the adult (Reiser *et al.*, 1979). It was demonstrated later on, that the high cholesterol content in HM leads to transiently higher total serum cholesterol concentration in infancy. However, later on in adulthood, the grown-up breastfed infants have lower total serum cholesterol concentration in comparison with a formula-fed group (Owen *et al.*, 2002). More recently, Timby *et al.* (2014b) showed in a double-blinded randomized trial that increasing cholesterol intake between 2 and 6 months of age *via* bovine MFGM addition in IF induced an higher total serum cholesterol concentration similar to the one observed in breast-fed infants without modification of the LDL:HDL ratio.

4 Data collected on human milk digestive behaviour *in vitro* and *in vivo*

In the forefront of these systemic and long-term health effects of HM, characterizing its behaviour in the upper digestive part seems a priority to formulate IF having close

digestive behaviour and properties. *In vitro* studies about HM and IF have contributed to clarify some points about the influence of the quantity and quality of the dairy emulsion interface on the digestion process (Bourlieu and Michalski, 2015; Gallier *et al.*, 2016).

4.1 Static or semi-dynamic neonatal model of digestion

Though several pertinent static *in vitro* digestion models mimicking the infant digestive conditions have been proposed (Dupont *et al.*, 2010; Bouzerzour *et al.*, 2012; Bourlieu *et al.*, 2014; Levi *et al.*, in press), very few studies have investigated the behaviour of HM during static *in vitro* digestion. Most digestion models were employed to evaluate IF digestion (Lueamsaisuk *et al.*, 2014). Chatterton *et al.* (2004), compared the protein breakdown of HM, whey protein concentrate and IF during *in vitro* gastric digestion using gastric juice of neonate. Gastric pH modulated strongly gastric proteolysis. Several HM proteins, including osteopontin and mucin present in the MFGM, resisted to digestion at pH 4 and above. The resistance of some proteins of the MFGM to gastric proteolysis is a central element to explain the persistence of human MFG throughout gastric digestion. Globally protein from human or whey protein concentrate from bovine milk had a comparable behaviour during gastric digestion, conversely IF's protein which undergo several heat-treatment during IF processing, were more susceptible to proteolysis.

No static or semi-dynamic digestion model was applied to HM allowing characterizing MFG digestive behaviour. However, a semi-dynamic model was applied to concept IF containing either bovine MFG (mode $\sim 6 \mu\text{m}$), or submicronic droplets resulting from high pressure homogenization alone or combined with post-homogenization heat-treatment (bimodal distribution with mods $\sim 0.2\text{--}0.6 \mu\text{m}$) (Bourlieu *et al.*, 2015b). This study established that the quantity of interface developed by milk lipids, directly influence gastric lipolysis and proteolysis kinetics. For milk emulsion having the same composition, initial rate of lipolysis was strictly proportional to the quantity of interface developed. Caseins (α_{s1} – α_{s2} , β and κ) were hydrolyzed faster than whey proteins (β -lactoglobulin and α -lactalbumin) whatever the type of emulsion. However, caseins were even faster proteolyzed in homogenized emulsions compared to the emulsion based on minimally processed large bovine MFG. This enhanced susceptibility to proteolysis of casein when adsorbed at an hydrophobic interface had already been presented by (Macierzanka *et al.*, 2009). The kinetics of gastric aggregation and internal structure of the aggregates were easily visualized on confocal scanning laser images. Magnifications and decompositions of the images (with or without protein labelling) revealed the persistence of preserved MFG or submicronic neoformed droplets. These lipid particles were the core elements around which protein aggregates built up.

4.2 Dynamic models

Zhang *et al.* (2014) investigated HM protein digestion using the Dynamic Gastric Model (DGM) to mimic the gastric conditions of infants aged of 9 to 12 months. Focusing on the

MFGM proteins, this study highlighted that among the numerous proteins detected in HM, proteins such as lipoproteins or mucins RNASE linked to the MFGM were more resistant to gastric proteolysis and could exert complementary bioactivity similarly to intestinal mucin. The lower susceptibility to pepsin of the MFGM's proteins was explained by their glycolysation or the presence of disulphide bounds.

Fondaco *et al.* (2015) compared the behaviour of different IF and HM during dynamic gastro-intestinal digestion using the TIM-1 model (TNO, Zeist, The Netherlands). HM and IF were characterized by very different initial structure notably in terms of particles size distribution with main modes of 7 μm in HM compared to $\sim 0.4 \mu\text{m}$ in IFs. HM developed less viscosity than IF over the range of pH encountered during neonatal gastric phase (pH 6 to 4 mainly). Whether this limited viscosity could impact on gastric emptying and satiety was questioned by the authors. Bio-accessible FFA released in jejunum and ileum data were quantified. An initial rapid release of FFA was observed in each compartment for IF whereas a significant lag (~ 90 min) was observed for HM. In addition, a correlation between the rate of lipolysis and specific surface developed by IF which directly influence digestive lipase access. However, at comparable specific surface, HM presented higher lipolysis rate. Altogether, these result suggested that the structure of HM lipid impacts strongly its kinetics of digestion and that the rate of lipolysis is not dictated by specific surface alone but that other factor such as endogenous maternal BSSL and the unique interfacial properties of the MFGM could contribute after an initial LAG to boost lipolysis. These results are in agreement with previous result of (Armand *et al.*, 1996) showing reduced lipolysis in the gastric aspirate of preterm after administration of IF compared to HM.

de Oliveira *et al.* (2015, 2016) used an *in vitro* dynamic system (DIDGI[®]) to study the gastrointestinal digestion of pooled mature HM ($n=5$) either raw (RHM) or pasteurized (PHM). The digestive behaviour of HM was characterized using either term and preterm infant digestive conditions. Gastric digestion was marked by a strong emulsion destabilization, due to acid and enzymatic-induced aggregation. The pasteurization of HM reduced its gastric destabilization: whereas an important destabilization was observed in RHM, PHM particle size profile remained quasi-steady during all the gastric phase. Gastric aggregates formed during RHM digestion presented slightly larger modes in the preterm (mode 1 = 96.3, mode 2 = 5.4 at 30 min) than in the term (mode 1 = 76.5 μm , mode 2 = 6.5 μm at 60 min) conditions. As observed by confocal laser scanning imaging, these aggregates were based on clusters of milk fat globules around which protein aggregates built up (Gallier *et al.*, 2013a; Bourlieu *et al.*, 2015b). Presenting lower density than milk aqueous phase, these lipoproteic clusters were prone to creaming in the gastric compartment. Such gastric destabilization in the adult favors a rapid gastric emptying of the aqueous low caloric watery phase, whereas the creamy phase is emptied more slowly (Golding and Wooster, 2010). The early gastric destabilization of raw HM observed here could be a protective mechanism favoring emptying and compensating the immature gastric motility in preterm newborns. Despite inactivation of endogenous lipase BSSL by pasteurization (activity levels of 23.4 ± 3.3 U/ml in RHM and 0 U/ml in PHM as assessed by pH-stat and using tributyrin as substrate), similar gastric

hydrolysis rate and quasi similar hydrolysis degree were observed in RHM or PHM. Final gastric hydrolysis degrees ranged between $13.1 \pm 1.4\%$ in preterm model to 24.1 ± 3.2 in full-term model. In addition, MFG were observed until at the end of gastric phase and delivered as preserved structure in the duodenum, which corresponds to a very specific delivery system of lipids unique to HM. With regards to the specificity of released FA short to medium-chains FA (C8:0 to C14:0) and LA were more released from RHM in the gastric phase, and are positioned mainly in the external positions (*sn-1,3*) of the glycerol backbone (Innis, 2011; Bourlieu and Michalski, 2015). Such positions correspond to the regioselectivity of the gastric lipase (*sn-3*) and of the BSSL (mainly *sn-1* and *sn-3*), which suggest a concerted action of gastric lipase and BSSL in the RHM (Bemback *et al.*, 1990). When comparing the released FA with the acyl chains initially esterified in HM, it appeared that oleic acid (C18:1 *n-9*) and stearic acid (C18:0) were selectively released, as their relative amount were higher than that in the total esterified FA profile. In both gastric and intestinal digestion, C8:0 presented the higher percentage of release.

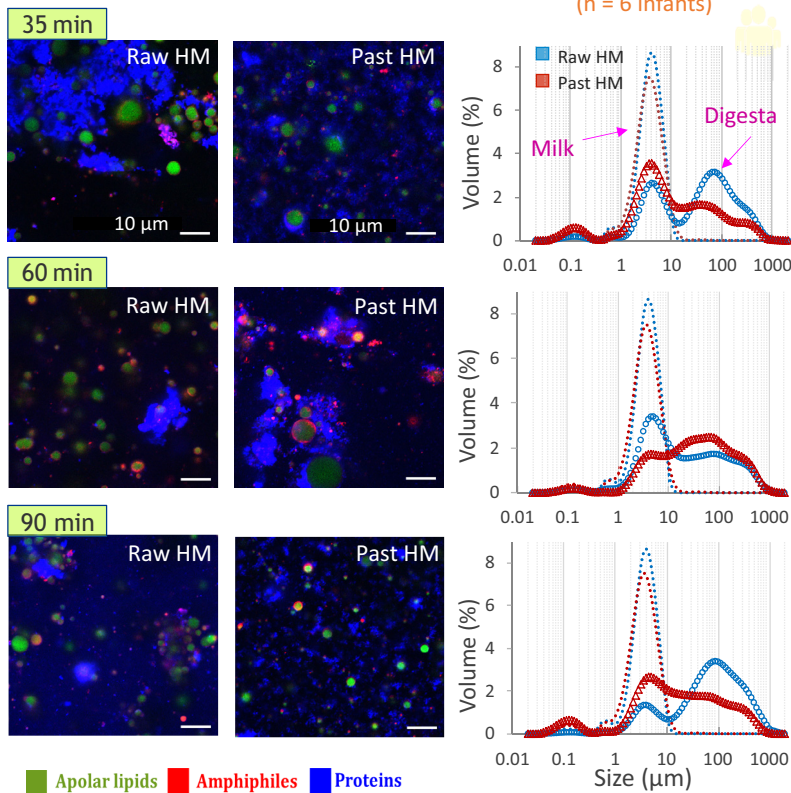
4.3 Data collected on HM digestion *in vivo* in gastric compartment

In an *in vivo* study on the preterm newborn, Armand *et al.* (1996) reported higher gastric lipolysis levels in HM compared with IF (which is homogenized), with values around $\sim 17\%$ and 9% at 30 min, respectively. These results suggested a lower accessibility of HGL to the TG core in the droplet of IF compared to HM, which was justified by the differences on the quality of the interface. Observations by transmission electron microscopy of gastric aspirates also allowed observing persistence of lipid structure in gastric phase. The quality of FFA released in gastric aspirate was not characterized.

Roman *et al.* (2007) analyzed products of lipolysis and human gastric lipase intragastric levels in preterm infants ($n=9$, GA = 29 + 1 wk, BW = 1.5 kg, medium age = 34.3 ± 1.8) fed either HM or medium chain-enriched IF. Medium chain-enriched IF contained 20–40% (wt/wt) medium chain whereas HM contain only around 10%. These medium chain FA are supposed to be better absorbed by the newborn as they can be directly absorbed through the gastric mucosa and transported *via* the portal vein and in intestinal phase their adsorption is less limited by biliary salt concentration compared to long chain FA. They observed similar rates of lipolysis for the IF and HM with lipolysis level reaching a maximum of $18 \pm 4\%$ after 180 min. The FFA and partial glycerides present in infant stomach during the 90–180 min time period were characterized. FFA profile was dominated in IF or HM group by palmitic acid (17.0 ± 0.2 vs. $24.8 \pm 6.2\%$ (wt/wt) in IF and HM respectively) and oleic acid (28.2 ± 1.2 vs. $27.1 \pm 3.3\%$ (wt/wt) in IF and HM respectively). Released caprylic acid (C8:0) was not much higher in the enriched IF (5.5 ± 1.2 vs. $3.5 \pm 1.5\%$ (wt/wt) in IF and HM respectively) and analysis of partial glycerides confirmed this trend. This limited release can probably be explained by the limited amount of caprylic acid present in *sn-3* position in vegetable fat compared to dairy fat.

de Oliveira *et al.* (2017) compared the gastric behaviour of pasteurized or raw HM in the preterm infant ($n=12$,

A) Gastric structural evolution of HM



B) Instantaneous lipolysis and FA bioaccessibility

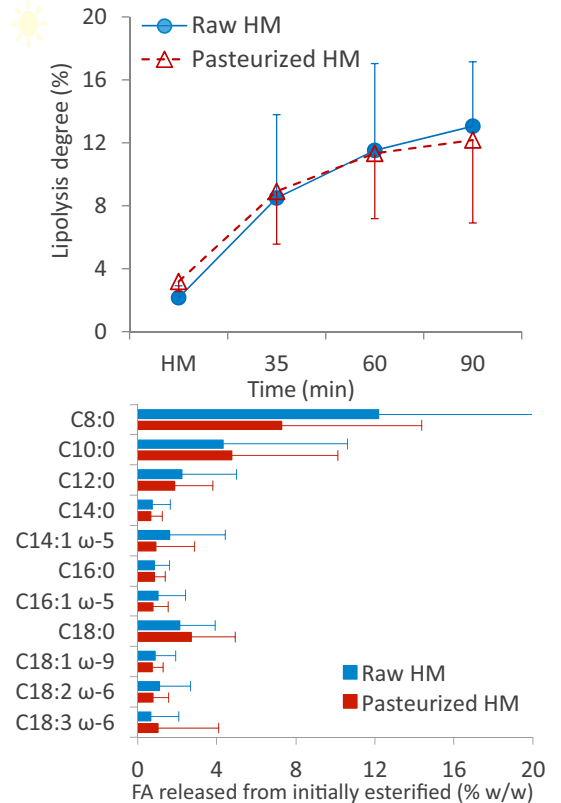


Fig. 3. Evolution of structure and lipolysis during gastric phase *in vivo* of raw or pasteurized human milk (de Oliveira *et al.*, 2017).

GA = 30.0 ± 1.1 wk, BW = 1.4 ± 0.3 kg, medium age = 27 ± 12 wk. Trial was conducted over a 6 day period and each infant was its own control. Lipolysis, FFA, proteolysis and gastric emptying rates were characterized (Fig. 3). Despite inactivation of endogenous lipase BSSL, lipolysis rate was not affected by pasteurization and averaged $12.6 \pm 4.7\%$ after 90 min digestion. BSSL was responsible for pre-lipolysis occurring prior to pasteurization and was higher for PHM than in RHM (respectively $3.2 \pm 0.6\%$ vs. $2.2 \pm 0.8\%$). Emulsion destabilization was observed from 35 min onwards and characterized by smaller aggregates and higher specific surface in PHM. The gastric emptying of HM was not modified by pasteurization with half time of ~ 30 min in range with the average values reported for HM in preterm infants (25–47 min). This rapid emptying is opposed to the longest time reported for IF (72 min on average).

5 Evolutions of IF in between scientific, technological and regulatory evolutions

IF are substitute of HM which have to totally support infant growth and thus are strictly regulated by different authorities worldwide (Codex Alimentarius Commission; US Food and Drug Administration; the European commission; the European Society for Pediatric, Gastroenterology, Hepatology and Nutrition, etc.). Composition and authorized additives allowed by these authorities were already summarized by (Zou *et al.*, 2016).

However scientific and technological advances are pushing forwards new fields of innovations in neonatal nutrition (Fig. 4). These fields are summarized on Figure 5. Details of corresponding clinical trials subcategorized in these fields of innovations are presented in Table 1. Among the emerging trends, the specific functions and metabolic fate of nanometer-sized lipid-protein particles termed ‘lactosomes’ are still unsolved (Argov-Argaman *et al.*, 2010). These particles are devoid of a TG core, have a density equivalent to plasma high-density lipoproteins, are enriched in immuno-modulatory proteins and are characterized by a distinct proteome and lipidome compared with MFG and high-density lipoproteins. Whether these lactosomes are similar to exosomes containing miRNA (small non-coding RNAs), which regulate several biological processes is still unknown. The cytoplasmic crescent of the MFGM (Fig. 1) entraps RNA and miRNA that would be involved in metabolic and immune regulation in the infant (Munch *et al.*, 2013). Their putative part in lipid metabolism preprogramming remains to be elucidated.

5.1 Fields of innovation based on recent analysis of clinical trials about infant formula (Tab. 1)

Most clinical trials (27%) focussed on protein composition of IF with particularly high number of trials addressing cow’s milk allergy and assessing the safety of extensively hydrolyzed IF or IF based on alternate source of proteins. Other important

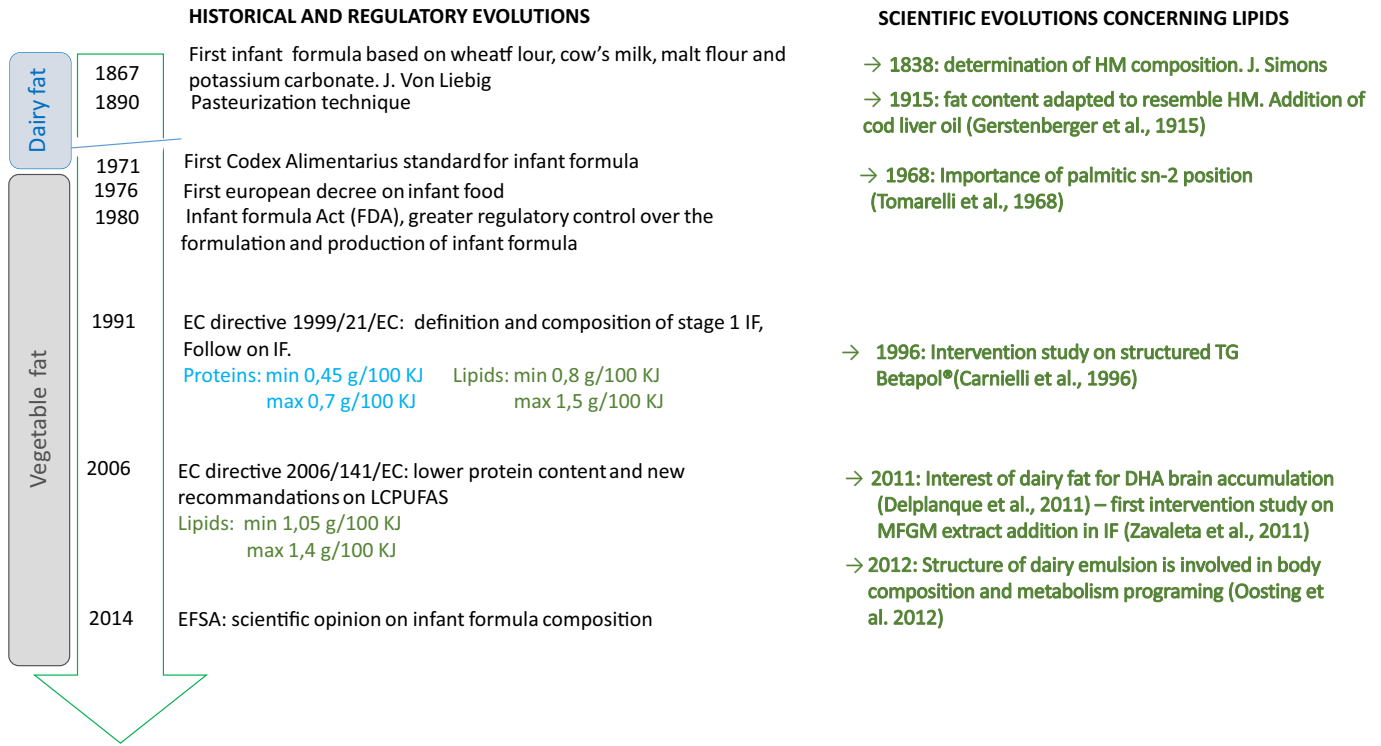


Fig. 4. Important steps in historical, regulatory evolutions and scientific research which have influenced infant formula production.

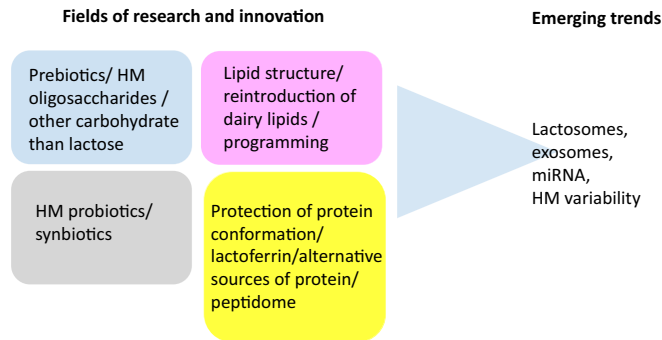


Fig. 5. Summary of fields of innovation based on recent analysis of clinical trials, WOS and patent database about infant formula.

fields of research concern prebiotics (oligosaccharides from HM notably) which promote the development of bifidogenic microbiota, probiotics possibly isolated from HM and combination of both (synbiotics) have been investigated. Clinical trials targeting lipids in the IF represent 14% of the total trials. Topics of these trials cover supplementation with dairy lipids or fraction, structured TG, beta-palmitate, LC-PUFA supplementation, structure of lipids and programming. Combining factors to make biomimetic IF is also a trend in recent clinical trials (for instance NCT00624689 ‘effects of milk fat globule membrane-enriched IF with reduced energy and protein content’). Carotenoids supplementation including lutein (NCT00913406) important for brain and visual development (retina epithelium) have also been assessed (Ramirez, 2016).

5.2 Fields of innovation based on recent analysis of patents about infant formula

More than 2000 patents about IF are registered in open database (espacenet patent). Among the most recent patents, several patents target IF supplementation with bioactive components such as oligosaccharides (US2016278421 (A1), US2016278414 (A1)), bovine colostrum component (MX2015017115 (A), modified sweet whey US2016310557, structured lipid 1,3-Dioleoyl-2-Palmitoyl glycerol WO2016 176987, etc.). Six patents registered in 2016 (US2016219911, NZ631411, US2016278413, NZ627997, US2016295895, US2016302611) corresponds to technological processes to produce new IF among which formula based on large globules. Patent WO 2010/027258 details technological ways of formulating large metastable emulsion and provide powdered formulas based on these large lipid globules emulsion. This patent addresses a true challenge as such emulsion is very metastable and that particles separation (creaming/sedimentation) during transport and storage, may result in a nutrient imbalance over the time of use. The use of micronized carbohydrates (US2016295895 (A1)) allows reducing stickiness/clumping and increasing flowability of the large droplet spray-dried IF.

5.3 Fields of innovations corresponding to staging infant formula to mimic HM evolution during lactation

Human milk is a variable fluid. Hence milk lipids vary with lactation stage but also diurnally and even during a single feed

Table 1. Scientific fields of innovations reported in clinical trials dealing with IF.

Themes	Number of trials	Corresponding trial and topic
Prebiotics, GOS, other carbohydrates than lactose	22	NCT01197365, NCT01497314, NCT01515644 (inulin), NCT00486148 (GOS), NCT00836771, NCT01497314 (low lactose), NCT01934257 (lactose free soy or milk-based IF), NCT02363582 (prebiotic and lactoferrin containing IF on stool consistency), NCT01606683 (IF with GOS, beta-palmitate, acidified milk), NCT02796872 (GOS supplementation), NCT02586558 (prebiotic on colic and crying), NCT01715246 (2 Human Milk Oligosaccharides – HMOs), NCT02757924 (prebiotics and behaviour), NCT02746016 (IF and oligosaccharides), NCT02118935 (prebiotics and early behaviour), NCT02441296 (maltodextrin or lactose added IF and metabolism), NCT00808756 (fermentable carbohydrate), NCT02703987 (fermented <i>vs.</i> non fermented IF and lactose tolerance), NCT01119170 (d-lactate producing probiotics), NCT00948051 (short-chain FOS enriched IF and immunity), NCT02317406 (probiotics on digestibility and immunity), NCT01788761 (probiotics on extremely low birthweight infants)
HM probiotics, synbiotics	27	NCT01897922, NCT01625273 (synbiotics), NCT00836771, NCT02096302 (Probiotic CECT7210 on Gastrointestinal Health), NCT01017991 (probiotics for crying infants), NCT00858026 (fermented IF for weaning babies), NCT00929292 (alpha-L and probiotics' enriched for infants with colic), NCT01079208 (starter IF with synbiotics), NCT01346644 (<i>Lactobacillus fermentum</i> for newborn infants), NCT01755481 (added probiotics and whey protein concentrate), NCT01956682 (hypoallergenic with starch and probiotics), NCT01476397 (probiotics), NCT01886898 (synbiotics), starter with pro and prebiotics, NCT01081067 (probiotic), NCT01010113 (synbiotics), NCT02221687 (combiotic study), NCT01036243 (slightly hydrolyzed and probiotics), NCT01813175 (synbiotics), NCT01983072 (synbiotics and colonization), NCT00318695 (probiotics and atopy/immunological responses), NCT02031887 (synbiotics on stool microbiota), NCT00365469 (probiotics on atopy, immunological responses and gut microflora), NCT00711633 (new fermented milk for preterm infants), NCT00792090 (fermented formula for allergy prevention), NCT00810160 (GOS and <i>Bifidobacterium infantis</i>), NCT02430831 (<i>L. reuteri</i> supplementation and crying), NCT01279265 (<i>L. rhamnosus</i> GG)
Supplementation with dairy lipids or fraction, structured TG, beta-palmitate, LC-PUFA supplementation, programming	30	NCT00970398, NCT02598817 (high <i>sn</i> -2 IF), NCT02111837 (specific lipid fraction enrichment), NCT00707837 (preterm IF with added soluble lipids), NCT02092857 (Arachidonic Acid Supplementation on the Immune Response), NCT02031003 (new fat blend or new fat blend plus fiber), NCT01603719 (milkfat and prebiotics), NCT00480948 (InFat™ on stool), NCT02144402 (B-DHA IF), NCT01058187 (long chain PUFA), NCT01140243 (long chain PUFA for preterm), NCT01373541 (InFat™ on stool), NCT01116115 (InFat™), NCT00874068 (InFat™ based infant formula on bone strength), NCT02503020 (AA in very preterm infant), NCT01611649 (mix dairy lipids/plant oil on w3 in red blood cell), NCT00941564 (calcium retention and different fat blends), NCT01300130 (improving DHA and ARA in preterm), NCT00753818 (DHA and ARA and development), NCT02339727 (w6/w3 and neural development), NCT01184378 (dairy lipid and soluble milk proteins), NCT01617889 (alternate fat blend), NCT00666120 (IF with DHA/ARA and +/- iron), NCT02587702 (B-palmitate and mineral metabolism), NCT02069522 (carotenoid supplementation), NCT02332967 (structured lipid with B-palmitate), NCT01908907 (DHA supplementation in preterm), NCT00379171 (fish oil supplementation), NCT00913406 (lutein fortification), NCT00872664 (IF supplemented with carotenoids on skin and serum carotenoids), NCT02033005 (breastmilk <i>vs.</i> IF and adipose tissue volume, development and intra-hepatocellular lipids), NCT00624689 (effects of milk fat globule membrane-enriched IF with reduced energy and protein content)
Protein content, alternative source of prot, extensively hydrolyzed whey, reduced protein content, enriched protein fraction, LF-addition	60	NCT01380886, NCT01210391, NCT01155414 (hydrolysate), NCT02469402 (reduced protein content, improved quality), NCT00938483 (extensively hydrolyzed), NCT01278446, NCT01278446 (extensively hydrolyzed), NCT02433600 (enriched protein fractions), NCT01987154 (extensively hydrolyzed casein IF for preterm), NCT01573871 (hydrolyzed IF), NCT00977964 (experimental milk protein IF), NCT02006992 (hydrolyzate IF), NCT00548106 (partially hydrolyzed whey), NCT02414243 (new aa-based IF), NCT02103205 (Bovine Lactoferrin and Low Iron Concentration IF), NCT02410057 (protein-reduced, alpha-lactalbumin enriched IF), NCT00716105 (different levels of proteins), NCT01143233 (reduced proteins, hydrolyzed with pre and probiotics), NCT02456831 (soy IF), NCT01208493 (high protein content for VLBW preterm infant), NCT00465764 (alternate protein IF), NCT01354366 (hypoallergenic IF), NCT01909661 (extensively hydrolyzed rice protein or casein), NCT02274883 (enriched protein fraction), NCT01205659 (allergy/asthma), NCT02500563 (aa-based or extensively hydrolyzed on microbiota), NCT00936637 (extensively-hydrolyzed IF),

Table1. (continued).

Themes	Number of trials	Corresponding trial and topic
		NCT00338689 (early programming and low protein IF), NCT00997971 (partially hydrolyzed rice protein), NCT02431377 (alpha-lactalbumin-enriched formula), NCT02010749 (protein concentration and growth up to 2 years), NCT00798382 (soy IF), NCT02719405 (resolution of cow's milk allergy : aa-based IF, extensively hydrolyzed casein IF or Lactobacillus GG containing IF), NCT02028156 (feeding intolerance), NCT00916565 (functional proteins), NCT01507935 (hydrolysed cow's milk and microbiota), NCT01569776 (aa IF), NCT01684319 (formula free of cow's milk proteins), NCT01809951 (soy-based infant formula), NCT01583673 (aa-based formula and growth), NCT01489098 (level of dietary protein on body composition), NCT00340652 (infant diets and estrogen activity), NCT02149134 (hypoallergenic IF), NCT01727115 (cow's milk intolerance), NCT02621814 (low glycation and aggregation protein), NCT02536482 (aa-based formula), NCT00664768 (hypoallergenic new IF), NCT02397876 (partially hydrolyzed whey IF), NCT02646969 (staged protein concentration over first year of life), NCT01470768 (aa-based IF), NCT01109966 (aa-based formula), NCT01156493 (hydrolyzed protein IF), NCT01637688 (aa-based IF), NCT02626143 (nutrient-rich whey protein IF), NCT02785679 (cow's milk allergy), NCT02711163 (extensively hydrolyzed IF), NCT01998074 (extensively hydrolyzed rice protein IF), NCT01584245 (aa-based IF), NCT02405923 (rice protein hydrolysate formula), NCT02351531 (thickened extensively hydrolyzed IF)
Other (1-5-methyltetrahydrofolate, alternate source of DHA, milk hypersensitivity, iron-fortification of IF)	26	NCT02437721, NCT02132663 (alternate source of DHA), NCT00938483, NCT01216709 (iron fortified IF), NCT00658905 (rHBSSL in IF for preterm), NCT02239588 (pure canterbury stage 1 milk powder), NCT01094080 (Modified Content of Protein and Improved Fatty Acids), NCT00000873 (high calory IF for HIV-infected infants), NCT00292812 (nucleotides supplemented IF), NCT00518414 (DHA, ARA and prebiotic IF), NCT02045784 (iodine supplements), NCT01177917 (mineral absorption), NCT00366873 (calcium absorption), NCT01413581 (rHBSSL), NCT01820494 (infant with chronic diarrhea), NCT01812629 (infant with chronic diarrhea), NCT00984230 (IF + LF + probiotics + OS), NCT01025557 (isovolumetric and isocaloric preloads of milk on food intake), NCT01042561 (vitamin D status), NCT02054091 (bovine colostrum to preterm infant), NCT00554814 (iron deficiency milk supplemented with biofer), (partially hydrolyzed and with <i>L. reuteri</i>), NCT01585142 (Babynex system), NCT02679183 (honey supplemented If for preterm), NCT01627015 (novel low glycemic index formula), NCT01148667 (Casein hydrolysate added with <i>L. rhamnosus</i> GG), NCT01166451 (iron supplemented IF)
Unknown	54	NCT01137877, NCT01609634, NCT01681355, NCT02860026, NCT02322138, NCT02481531 (protein fraction), NCT02401217, NCT02860026, NCT02094547, NCT02490852, NCT00957892, NCT01808105, NCT02670863, NCT02073071, NCT01721850, NCT01162798, NCT01558440, NCT00655720, NCT00937014, NCT00705562, NCT02178189, NCT00820833, NCT01861600, NCT02658500, NCT02715895, NCT02405572, NCT01766011, NCT01370967, NCT00543673, NCT02456805, NCT00665938, NCT01300000, NCT02594683, NCT01721512, NCT00340665 (infant diet on estrogen activity), NCT01735123, NCT00712608, NCT00920166, NCT01466400, NCT02340143, NCT01762631, NCT01700205, NCT02710955 (anti-regurgitation formula), NCT00994747 (early flavour learning), NCT02066610 (selenium supplementation), NCT00503789, NCT00952328, NCT00740974, NCT00134771 (supplementation small for gestational age), NCT01177930, NCT00342303 (activity of essential fatty acid elongation/desaturation during early life), NCT02425423 (anti-regurgitation), NCT01759134, NCT00506584, NCT02759809

(foremilk, hindmilk). Three stages are differentiated: colostrum (1–5 days post-delivery), transitional milk (6–15 days post-delivery) and mature milk (>15 days post-delivery). The difference in the composition of main macronutrients between these stages is well documented (Jenness, 1979; Jensen, 1999; Hester *et al.*, 2012; Guesnet *et al.*, 2013; Michalski, 2013). Less data is available about the variation of composition and structure of lipids in mature milk as lactation time progress. This variability in HM composition suggests adaptation to the

precise infant need at a given physiological stage and could be an important tool to diversify and improve IF (Harzer *et al.*, 1983; Nommsen *et al.*, 1991; Santoro *et al.*, 2010; Bourlieu *et al.*, 2015a; Lonnerdal and Hernell, 2016). Lonnerdal and Hernell (2016) even suggested recently a re-evaluation of the staging concept (Fig. 6B) to propose specific formula for 0–3 months (stage 1), 3–6 months (stage 2) and 6–12 months (stage 3) which could improve nutrition of formula-fed infant and make their outcomes and performance closer to HM.

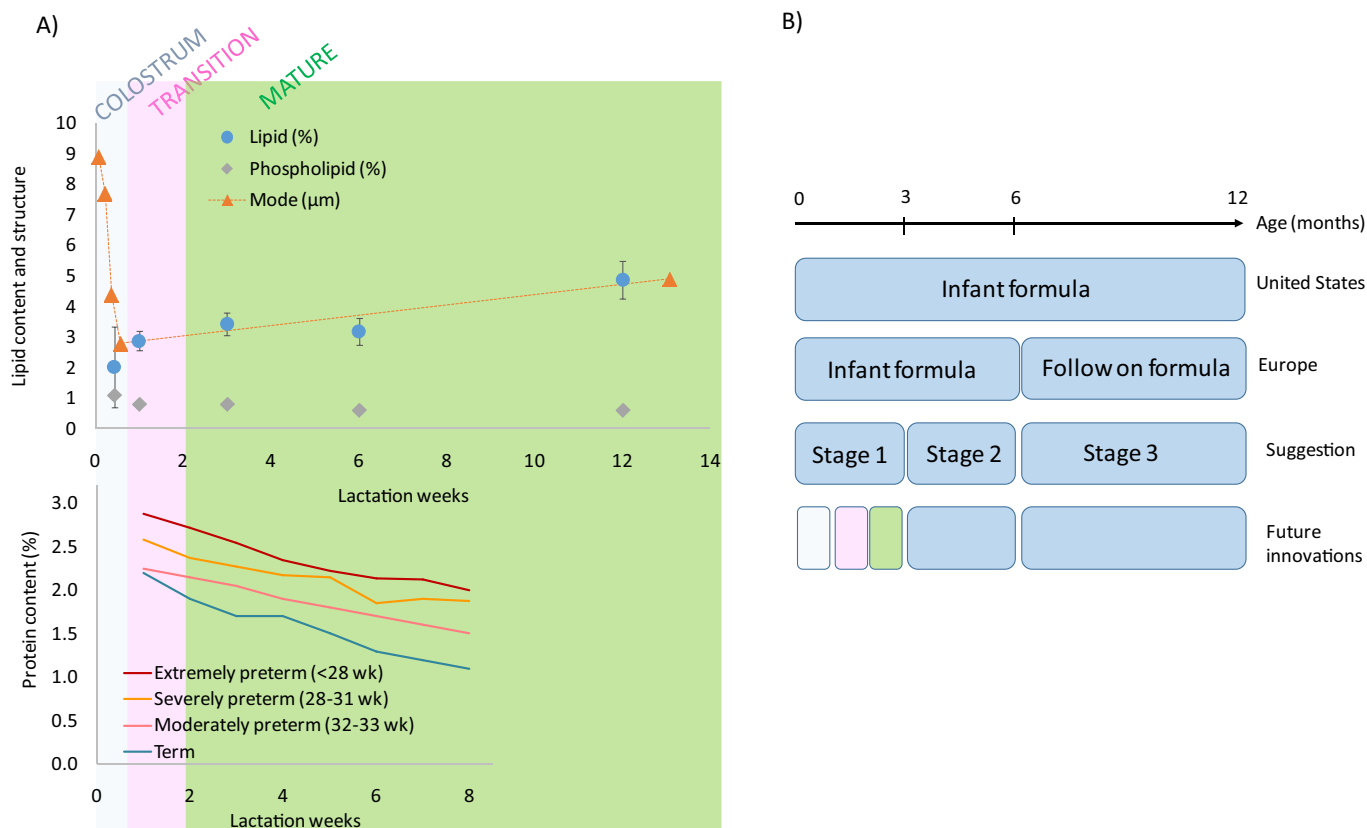


Fig. 6. Illustration of HM variability during lactation and associated concept of biomimetic staged IF. (A) Variation in lipid content and structure and in protein content of HM during lactation. (B) Concept of IF staging adapted from Lonnerdal and Hernell (2016).

Indeed when considering the total lipid content of the milk, it increases with lactation duration. This increase induces an increase in the size of the globule as the interfacial material become limiting (Fig. 6A). Typical variations reported in a study involving 120 breastfeeding Spanish mothers reported total fatty acids values of 2.02 g/100 ml in colostrum, 2.59 g/100 ml in transitional milk and 3.28 g/100 ml in mature milk (Jensen, 1999). In terms of quality of FA, saturated FA C8-C14 were reported to increase between the colostrum milk and the two other lactation periods (Lopez-Lopez *et al.*, 2002). Monounsaturated FA generally decreases between colostrum and transitional or mature milks. The total polyunsaturated content of milk do not vary with the stage of lactation. However, the ratio between the two essential FA, *i.e.* LA/ALA evolves during lactation and can overtake the acceptable range of 5:1–15:1. The diet and physiological state of the mother also impact her milk nutritional composition. The strong impact of the diet on human milk FA profile was detailed by (ESPGAN Committee on Nutrition, 1991) and puts forward as a basis to explain the regional and seasonal fatty acid profile variation in HM. DHA in milk is also highly dependent on the DHA status of the mother but would decrease with duration of lactation. Mature HM is more influenced by maternal factors (body composition, diet and parity, etc.) than milk of early lactation (Jensen, 1999).

Regarding the structure of HM lipids emulsion and its evolution with delivery term (Simonin *et al.*, 1984) or lactation stages, very limited data is available (Michalski *et al.*, 2005b; Zou *et al.*, 2012). Michalski *et al.* (2005b) reported an higher

amount of large fat globules in early colostrum with a mode of 8.9 μm at 12 h post-partum and which decreased rapidly to 2.8 μm at 96 h post-partum (Fig. 6A). The MFG size then increased regularly in mature milk, from values around 5 μm at 3 months lactation up to 6.5 μm at 18 months.

6 Conclusions

The better understanding of the composition and structure of human MFG, of their behaviour in the upper digestive tract has led to development of new IF more biomimetic of HM. These formulas include bovine milk fat fractions (MFGM, TG) or are structured under the form of large droplets having diameter similar to human MFG. Proofs of evidence of the fact that the structure of lipid ingested during neonatal period programs body composition and metabolism in adulthood has been produced in animal but must be confirmed in clinical trials.

Data on the structure of HM throughout lactation and on the digestive behaviour of specific fractions of HM lipids are still needed to build a sound basis for the optimization of IF. A better understanding of fractions that escape enzymatic digestion in the upper tract and serve as substrates for microbiota will help to control microbiota shifts that exist between breast-fed and formula-fed infants. Combinations of factors (synbiotics and lipid structure) and more staging seems important levers to improve IF but their efficacy and safety still have to be assessed in clinical trials.

Abbreviations

ALA	alpha-linolenic acid
LA	linoleic acid
ARA	arachidonic acid
DHA	docosahexaenoic acid
FA	fatty acid
FFA	free fatty acid
HM	human milk
IF	infant formula
LCPUFA	long chain polyunsaturated fatty acids
MFG	milk fat globules
MFGM	milk fat globule membrane
MPL	milk polar lipids
TG	triacylglycerides
sn	stereo-numbering

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