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Dietary lipids and reward processes

1	Dietary lipids as regulators of reward processes: multimodal integration matters
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Abstract

The abundance of energy-dense and palatable diets in the modern food environment tightly contributes to the obesity pandemic. The reward circuit participates to the regulation of body homeostasis by integrating energy-related signals with neural substrates encoding cognitive and motivational components of feeding behaviors. Obesity and lipids-rich diets alter dopamine (DA) transmission leading to reward dysfunctions and food overconsumption. Recent reports indicate that dietary lipids can act, directly and indirectly, as functional modulators of the DA circuit. This raises the possibility that nutritional or genetic conditions affecting "lipid sensing" mechanisms might lead to maladaptations of the DA system. Here, we discuss the most recent findings connecting dietary lipid sensing with DA signaling and its multimodal influence on circuits regulating food-reward processes.

Dietary signals: all roads lead to the brain

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Physiological mechanisms optimizing feeding behaviors have been selected during 40 41 evolution. This is exemplified by the natural tropism of mammals toward energy-dense 42 food (rich in lipids and sugar) which represents a promise of maturation and survival. 43 Nowadays, the ubiquitous availability of palatable and energy-dense diets challenges the highly conserved and sophisticated systems controlling food intake. It is clear that, 44 although inheritable genetic traits might contribute as risk factors for the development of 45 metabolic disorders, the consumption of calorie-dense food beyond metabolic needs 46 47 arises as a primary culprit in the obesity pandemic. During the last decades, the brain has emerged as the master regulator of energy 48 49 homeostasis by integrating metabolic demands with the reinforcing nutritional properties 50 of food [1]. In fact, while the hypothalamic melanocortin circuit ensures the homeostatic 51 control of ingestive behaviors and metabolism, the mesocorticolimbic dopamine (DA) 52 system integrates and translates metabolic signals into reward values [2] so that 53 associated cues are imbued with incentive salience and the power to drive choices and 54 behaviors [3]. Thus metabolic and reinforcing systems are dynamically integrated so that 55 metabolic requirement is translated into motivated behaviors to obtain nutrients [4]. The 56 ability of organisms to store energy extends this process beyond current needs, making 57 overconsumption adaptive throughout most of history. 58 Dietary signals can modulate brain circuits and functions in distinct ways. In fact, nutrients 59 sensing chemosensory mechanisms operate in compartmentalized cellular elements 60 located throughout our body, notably in the mouth, gastrointestinal (GI) tract and brain, 61 thereby playing a key role in the multimodal regulation and control of ingestive behavior(s) (Figure 1). This multimodal integration has been recently highlighted in humans where 62 63 immediate orosensory and subsequent post-ingestive signals derived from a palatable food (milkshake) led to two temporally separated waves of DA release in the ventral and 64 65 dorsal striatum, respectively [5]. Among nutrients (e.g. glucose, amino acids), dietary and circulating lipids have emerged 66 as critical modulators of brain functions. Indeed, foods rich in fat are highly reinforcing. 67 Despite orosensory perception and organoleptic components of food are known to be 68 69 intimately associated with the reinforcing values of food, novel physiological routes are

emerging as instrumental in conveying the reinforcing aspects of dietary lipids through direct and/or indirect connections with the mesolimbic DA circuit. However, dietary lipids-related functions seem to strongly depend on the nature and source of circulating lipids (pre- to post-prandial availability of circulating lipids, **Box 1**).

BOX 1: Triglycerides or free fatty acids: the physiology of timing and nutritional states.

Triglyceride (TG)-rich lipoproteins and free fatty acids (FFA) bound to albumin represent the two main sources of circulating lipids in the body [6]. While FFA are released upon adipose tissue lipolysis during periods of nutrients deficiency, lipoproteins are particles released from the liver and the gut to serve as hydrophobic carriers of TG, which are glycerol-esterified long chain fatty acids (LCFA) [7]. TG-rich lipoproteins arise from the gut as chylomicrons (CM) or from the liver as very-low-density lipoproteins (VLDL). As lipoprotein lipase (LPL) hydrolyses TG to provide lipids to metabolically active tissues, CM evolve towards intermediate (IDL) and high-density lipoproteins (HDL) (**Figure 1**). In physiological conditions plasma FFA and TG levels fluctuate in opposite directions. Indeed, plasma FFA drop in post-prandial periods due to the antilipolytic action of insulin onto adipose tissue. On the other hand, plasma TG rise due to lipid re-esterification and complexification into CM in enterocytes after a meal [7].

Both FFA and TG-particles were shown to access the brain. The transport of FFA in the brain was demonstrated using radiolabeled lipids [8] and an increased in brain FFA transport was found associated to metabolic disease [9].

How TG-rich particles access the brain is still debated. CM are likely too large to cross the blood brain barrier (BBB), but smaller HDL and remnant CM (rCM) were shown to access the brain [6]. The brain expresses high levels of receptors for VLDL, low-density lipoprotein (LDL), oxidized HDL or LDL receptor-related protein (LRP) [6]. Thus, the presence of membrane and mitochondrial machineries dedicated respectively to the transport and oxidation of lipids in neurons suggest a role for FFA as signaling molecules [10]. For example, the TG-processing enzyme LPL is expressed in very defined brain structures [6,11] suggesting that LPL-expressing neurons are selectively equipped to detect and respond to TG-rich particles. Since circulating FFA and TG can access the

brain during pre- or post-prandial periods, dietary lipids under either forms seem good candidates to regulate feeding behaviors through a continuum process embedding signals staring from taste buds, gut-brain axis and/or through direct actions on brain cells (neurons and astrocytes).

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From the mouth to the brain: a taste for fat

Taste is a sense assessing the nature, quality, texture and composition of food, thereby driving either consumption or avoidance. Although five taste modalities are commonly recognized (sour, salty, sweet, bitter, and umami), several studies have shown that dietary LCFA can trigger brain responses and affect digestive functions and feeding behavior [12.13] through the interaction with dedicated receptors located on taste bud cells (TBC) [14] (Figure 2). The ability of TG to be locally and rapidly hydrolyzed into FFA represents the major stimulus for the perception of fat-containing food in both humans and rodents [15,16]. Importantly, fat sensing requires complex inter-organized mechanisms since FFA of different chains length can elicit different behavioral responses (consumption vs avoidance/irritation) [17]. In the lingual epithelium, taste buds are highly specialized bulb-like structures composed of 50-100 TBC organized into cell types-assembled clusters functionally connected to sensory afferent fibers of cranial nerves VII and IX, also known as the chorda tympani (CT) and glossopharyngeal (GL) nerves (Figure 2). At the cytoarchitectural level, the nutrients-exposed apical region of taste buds is mainly composed of type I (glia-like cells), II (gustatory receptors-expressing cells) and III (neuron-like cells) TBC which, through a complex and dynamic inter-TBC networks of transmitters (ATP, acetylcholine, endocannabinoids, serotonin, GABA and norepinephrine), represent the main chemosensory processors [18]. Ingestion of very-low concentration of LCFA elicits DA release in the nucleus accumbens (NAc) [19] and fat consumption is rapidly accompanied by the activation of limbic, hedonic and motivational circuits in rodents and humans [12,20-22]. Therefore, it is unsurprising that a preference for high-fat foods is a widely shared and conserved trait in mammals. How do TBCs sense fat? During the prandial period, although TG are the major

components of dietary fats, TG-hydrolyzed LCFA seem to drive the oral chemosensory

stimulation and its associated behavioral outputs [23,24]. This chemosensory detection/reception of dietary lipids relies, at least in part, on TBC-expressing membrane proteins, notably: GPR40 (or FFA1), GPR120 (or FFA4), CD36 and TRPM5 [17,25-27] (Figure 2). While Gg-coupled receptors GPR40/FFA1 (rodent TBCs) and GPR120/FFA4 (human and rodent TBCs) have been reported to mediate fat-induced behavioral and cellular responses [25,28], other studies have not led to the same conclusions [29,30], thus complicating the molecular view of fat taste. More is known about the functions of lingual CD36 (also known as fatty acid translocase (FAT)), a transmembrane glycoprotein with high affinity for FFA and whose single nucleotide polymorphism (rs1761667) lowers CD36 expression and fat gustatory perception in humans [24]. At the transductional level, LCFA-activated CD36 and GPR120 both lead to increased Ca²⁺ levels, through two different but complementary intracellular signaling paths [31]. In particular, activation of CD36 promotes PLC-dependent hydrolysis of PIP2 which gives rise to IP3 and DAG. In turn, IP3, by binding to its endoplasmic reticulum receptor, promotes Ca2+ release. The rise in Ca2+ participates in activating the TRPM5 ion channel which, by depolarizing/activating TBCs, mediates the detection and transmission of gustatory signals [27,32].

- Fat signal transduction also relies on the activation of MAP kinase cascade (MEKs-ERKs)
- via a mechanism requiring Ca²⁺ signaling [33]. Importantly, down-regulation of ERK1/2
- signaling cascade significantly decreased fat preference in mice [33], thereby indicating
- that chemosensory oral detection of fat requires both Ca²⁺ and ERK dynamics.
- Once stimulated, TBCs activate sensory afferent fibers conveying fat-taste information to
- the rostral nucleus tractus solitarius (NTS) which in turn relays such information to
- 155 secondary brain structures, allowing integration and elaboration of fat signals and
- encoding of their rewarding/reinforcement values.

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From the gut to the brain: interoceptive and integrative fat-sensing mechanisms

Food palatability, which promotes positive reinforcement and overeating, is traditionally described in terms of sensory (smell, taste) pleasantness, thereby embracing into the definition of "reward" conscious features such as liking, wanting, predictive value and insensitive salience [3]. However, it is of paramount importance to mention that

reinforcement mechanisms can also derive, at least in part, from unconscious and interoceptive processes which rely on the detection of "reinforcers" (nutrients, energy contents, circulating signals) from specialized internal detectors/sensors usually inaccessible to conscious palatable features [2]. This can be easily summarized by the fact that food consumption, preference and/or seeking, although facilitated by sensory pleasantness, tightly depend on energy availability and contents which, by acting as secondary reinforcers, represent conditioned signals predictive of the metabolic effects of ingested nutrients [34–36].

The gastrointestinal (GI) tract, a highly heterogeneous chemosensory organ, is specialized in sensing and absorbing nutrients, and, in virtue of its complex endocrine/paracrine networks, initiates several physiological processes (**Figure 3**). Dietary lipids occupy a key place in the complex GI functions that control energy homeostasis and feeding behaviors. Once in the GI lumen and upon the action of intestinal lipases, TG-derived lipids (diacylglycerol, monoacylglycerol and FFA) stimulate specialized gut cells [37], notably enteroendocrine cells (EEC), which initiate hormonal and gut-to-brain responses [38]. Indeed, EEC types are well-equipped with dietary fat cellular sensors as they express FFA1, FFA4, GPR119 and CD36 [39–42] (**Figure 3**). In line with this evidence, post-oral delivery of dietary lipids triggers the release of GLP-1 and CCK [35,43]. Such capability of EEC to detect dietary lipids contributes to their modulatory functions.

Originally believed to be specific to distinct secretory mediators (CCK, GLP-1, GIP,

secretin, PYY), recent breakthroughs have shown that this "one mediator-one cell type" theory is outdated as EEC located throughout the GI tract express and actively co-release a combination of different hormonal mediators and neurotransmitters [44–47] and even signal using specialized entero-neuronal synapses named neuropods [48,49]. Overall, these recent evidence indicate that fat-sensing at EEC level may trigger multiple modulatory responses.

Beside EEC, dietary lipids are also sensed, absorbed and metabolized by enterocytes (**Figure 3**). In fact, after gastric TG hydrolysis, FFA and 2-monoacylglycerol (2-MAG) are transported into the enterocytes either by passive diffusion or by protein-dependent mechanisms (CD36, FATP4). Within the enterocytes, 2-MAG and the majority of LCFA

- are re-esterified as TG, and either stored as cytosolic lipid droplets or packaged into
- lipoproteins (mostly CM). Lipoproteins are therefore secreted into the circulation through
- the lymphatic system to ultimately reach all organs [50].
- 197 Aside of FFA transport and metabolism, lipids can directly signal onto specialized
- receptors. In that regards, studies using intragastric fat infusions have highlighted several
- redundant mechanisms through which dietary lipid sensing in the gut regulates feeding
- 200 (**Figure 3**):
- 201 (1) Luminal TG-hydrolyzed oleic acid is transported into enterocytes via a CD36-
- 202 dependent mechanism and converted into oleoylethanolamide (OEA). OEA acts as a
- 203 satiety signal through a mechanism that is still not fully elucidated, mostly via the
- 204 activation of the lipid-activated transcription factor peroxisome proliferator-activated
- receptors- α (PPAR- α) and the mediation of the vagus nerve [51–54]. However, some
- studies have also suggested alternative mechanisms by which LCFA and byproducts of
- TG (FFA and 2-MAG), rather than OEA, may function as immediate signals of feeding
- onto the DA system [35], while non-vagal afferent mechanisms could relay OEA-induced
- satiety effects [55].
- 210 (2) The inhibition of gastric emptying by lipids-triggered GPCRs-dependent
- 211 enteroendocrine release of GLP-1, CCK and PYY [39,42,56].
- 212 (3) The action of lipid oxidation byproducts from enterocytes or microbiota, such as ketone
- bodies/beta-hydroxy butyrate (BHB), which signal through spinal afferent fibers [57].

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- Lipid sensing in the gut-brain axis can shape how hypothalamic hunger-activated neurons
- respond to food cues. A recent study using cell type-specific *in vivo* Ca²⁺ dynamics in
- 217 Agouti-related protein (AgRP)-expressing neurons provides evidence that glucose- and
- 218 lipids-dependent signals initiated in the gut are routed through redundant and segregated
- 219 nervous afferent routes to impinge on hypothalamic responses to food cues [58]. This
- selectivity to nutrients was ascribed to distinct nervous routes of the gut-brain axis where
- vagal afferents transmitted intestinal lipid detection and spinal gut-brain inputs from the
- 222 intestinal and hepatic portal vein signaled glucose detection. Both paths serve as
- modulators of AgRP-neurons activity, thereby promoting fullness and satiation [58]. Of
- 224 note, obesity and exposure to high-fat have been shown to selectively impair lipids, but

not carbohydrates, signaling from gut→AgRP-neurons [59,60]. These results suggest that 225 226 gut-brain fat sensing represents a privileged substrate for obesity-induced 227 maladaptations. 228 Interestingly, despite promoting satiation, intragastric infusion of lipids elicits dopamine 229 release [35,61], suggesting that reward/reinforcement encoding and satiation are not 230 functional antagonists. 231 Beside their endocrine actions, gut hormones released upon intestinal lipid sensing can also act on sensory vagal afferents which highly express receptors for GLP-1, CCK and 232 233 PYY [62]. Interestingly, in addition to detecting hormonal and mechanical signals [63], 234 vagal sensory afferents can be directly modulated by dietary signals. In fact, vagal 235 neurons express the molecular machinery required to sense circulating lipids, notably 236 long- and medium-chain fatty acids [64,65], and through a polysynaptic gut-brain circuit 237 they may convey signals that are translated into reinforcement by DA-neurons [66]. 238 One would therefore argue that the blunting action of OEA over food intake (satietogenic 239 signal) may also reduce the hedonic drive to food consumption. However, recent 240 breakthroughs have elegantly shown that OEA promotes DA release in the striatum, 241 through a PPARα-mediated vagus-dependent action [61]. This DA release depends on 242 metabolic states (low-fat vs high-fat fed mice), and rerouting of the GI tract by Roux-en-243 Y Gastric Bypass (RYGB) mobilized small intestine OEA, an affect associated with a 244 vagus-dependent and diet-mediated enhancement in striatal dopamine release [53]. 245 Beside the peripheral OEA-mediating actions onto central DA release, intra-gastric or intraperitoneal administration of fat emulsions also trigger DA activity [34,36,61,67]. This 246 247 indicates that dietary and diet-derived lipids engage vagus-dependent periphery-to-brain 248 circuits which regulate dopamine release and reward events. 249 Although the detailed mechanisms underlying how gut sensing of dietary lipids may 250 dampen food intake despite promoting reinforcement-like events are not fully elucidated, 251 it is worth to mention that both vagal and spinal sensory afferents may collaborate in detecting food constituents through heterogeneous neuronal networks [68-70]. 252 253 Therefore, according to their intrinsic cell type-specificity, regionalized arborization into 254 NTS territories and specific synaptic contacts onto distinct NTS cell types, they may 255 convey different integrative information to the brainstem. In turn, NTS cell types, by

projecting to second order brain regions [71–73], may signal to different downstream structures involved in the regulation of homeostatic (hypothalamus) and hedonic feedings (mesolimbic and nigrostriatal dopamine circuits). Finally, the precise timing by which lipids (and glucose) pass through and signal into digestive segments, might directly account for the different phases by which each nutrient triggers specific brain responses.

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Central action of dietary lipids: between reward and satiety

Dietary and diet-derived signals can also directly act on distinct brain circuits and discrete cell types which regulate hunger/satiety and reinforcement. In fact, although the mechanisms underlying their entry into the brain remain largely unknown, pioneering studies have indicated that circulating lipids can reach the brain and be uptaken by neurons by crossing the BBB [8,74] (Figure 4). Moreover, although not specifically demonstrated for lipids, new evidence highlight the key role of the lymphatic system in allowing brain access to large and small molecules [75]. Neurons are fully equipped with the molecular components capable of recognizing and metabolizing dietary lipids [76]. Indeed, among central brain regions, the hypothalamus has been long recognized as a key sensor of circulating lipids [10]. Notably, hypothalamic cells (neurons and/or astrocytes) express a variety of membrane proteins capable of detecting and handling extracellular FFAs [77,78] and also respond to circulating TG [79]. Importantly, hypothalamic cells also contain the lipoprotein lipase LPL which, by locally hydrolyzing TG into 2-MAG and FFAs, contributes to the regulation of food intake and energy balance [80,81]. In fact, downregulation of LPL in hypothalamic astrocytes and/or neurons leads to increased body weight and glucose intolerance [81,82]. Beyond LPL, the hormonesensitive lipase (HSL), hydrolyzing tri- and diglycerides within the hypothalamus, may also play a fundamental role in regulating food intake and body homeostasis [83]. In fact, HSL deficiency triggers sustained food intake in high-fat diet (HFD)-fed mice, resulting in increased body weight gain [83]. Overall, these results strongly indicate that lipidssensing functions of hypothalamic cells are pivotal in regulating energy homeostasis as their silencing exacerbates pro-obesity features. Beside the hypothalamus, recent studies have clearly pointed to the midbrain dopamine

system as a central target where circulating dietary TG may act by influencing molecular,

cellular and behavioral reinforcement features [11,84–86]. This dopamine (reward) circuit, consists of midbrain nuclei, notably the ventral tegmental area (VTA) and the substantia nigra *pars compacta* (SNc), densely projecting to dopaminoceptive structures such as the nucleus accumbens (NAc) and the dorsal striatum (DS). This brain circuit is strongly impaired by both short- and long-term exposure to HFD [87–89] as well as by perinatal dietary exposure to maternal HFD feeding [90,91].

Although the consequences of HFD exposure on the dopaminergic system are wellstudied, whether and how dietary lipids as such can modulate the activity of the dopamine circuit is still a matter of intense investigations. Interestingly, VTA DA-neurons contain FFA-associated proteins, such as the fatty acid transport proteins (FATP1 and FATP4), fatty acid-binding proteins (FABP3) [86], as well as LPL [11], thereby highlighting the capacity of DA-neurons to detect and metabolize circulating lipids. In fact, LCFA, such as the oleic and palmitic acids (OA, PA), and/or long-term central delivery of TG, modulate the activity of VTA DA-neurons by reducing their neuronal firing rate and therefore by modulating reward-associated events [11,85,86]. Hence, the presence of LPL in dopamine and dopaminoceptive neurons might represent an additional arsenal to selectively respond to postprandially released TG. It is interesting to note that while sensory information routed by the vagus nerve was shown to modulate DA release from SNc DA-neurons [66], Lpl mRNA was found to be among the 5 most robust markers differentially expressed in the VTA compared to the SNc [92], thereby suggesting that VTA could directly detect postprandial circulating TG while SNc DA-neurons would rather respond to gut-borne stimuli routed by vagal afferents.

At the postsynaptic level and throughout the rostro-caudal axis, DS and NAc dopaminoceptive neurons, also known as GABAergic spiny projection neurons (SPNs), are anatomically and functionally segregated into two major families according to the expression of either DA D1 or D2 receptors (D1R- and D2R-SPNs) [93–95]. Recently, striatal and accumbal SPNs, in both humans and rodents, have been shown to express the LPL and respond to circulating TG [11]. Notably, central delivery of dietary TG resulted in a blunted activity of DA-dependent intracellular signaling pathways (MAPK, PKA and mTOR) and concomitant increase of β -arrestin-dependent signaling (GSK3- β) within the

DS/NAc territories. In addition, TG were shown to reduce the excitability of D2R-SPNs, 318 319 sparing D1R-SPNs [11] (Figure 4). Importantly, cell type-specific deletion of LPL in D2R-320 SPNs increased neuronal excitability [11], thereby indicating that local metabolism of TG 321 may be critical in gating the neuronal activity of dopaminoceptive neurons. Of interest, 322 central delivery of dietary TG, although associated to dampened food intake and reduced 323 responses to psychostimulants [11,84], was accompanied by enhanced reward sensitivity 324 and positive reinforcement [11], indicating that circulating TG may possess reward-like 325 properties. This evidence suggests that dietary TG, and eventually their TG-derived FFA, 326 may inhibit D2R-SPNs, therefore bypassing the inhibitory action of endogenous DA onto 327 D2R of D2R-SPNs and disinhibiting D2R-SPNs-downstream targets. Such dietary 328 inhibition of D2R-SPNs may then participate in promoting reinforcement as reported by 329 cell type-specific manipulation of D2R-SPNs [96-98]. Of note, D1R-SPNs also express 330 LPL but they do not seem to respond to TG concentrations capable of modulating DRD2-SPNs excitability [11]. In line with that view, cognitive impairments caused by PUFAs 331 332 deficiency could be corrected by restoring PUFAs balance in D2R-SPNs but not D1R-SPNs [99]. 333 334 It is worth to mention that D2R is also expressed in DS/NAc cholinergic interneurons (ChINs [100]) but that this cell type seems to lack Lpl mRNA [11]. Nevertheless, future 335 336 investigations aimed at unbiasedly studying the impact of dietary lipids on the cellular 337 physiology of distinct striatal cell types are definitely needed. 338 Overall, these recent studies suggest an evolutionary acquired ability of D2R-SPNs to 339 directly respond to circulating dietary lipids. This hypothesis nicely fits with other studies indicating that D2R-neurons are highly susceptible substrates for central maladaptations 340 341 induced by obesogenic environments and altered cognitive functions. In fact, the D2R is 342 commonly pointed as a molecular culprit potentially involved in the development and/or maintenance of food-related reward deficits and associated metabolic alterations [101-343 106]. 344 345 D2R abundance predicts compulsive susceptibility in rats [103] and exposure to lipid-rich 346 diet triggers impulsive feeding independently from body weight changes [101]. Moreover, 347 knock down of striatal D2R expression leads to compulsive eating in rats exposed to high-348 fat food [103].

This observation is also supported by studies in humans showning that TG metabolism correlates with brain responses to food cues in a direction that is influenced by genetic traits affecting D2R signaling [11,107].

Whether dietary lipids as such may directly and causally be involved in promoting the dysfunctions of D2R-SPNs and D2R-coupled signaling remains to be fully established. It is interesting to note that central increase of circulating TG counteracts D2R-dependent signaling pathways but does not impact D1R-coupled signaling events [11]. Although molecular mechanistic insights are still under investigation, it is worth to mention that a role for the fatty acid binding protein FABP3 has been highlighted in modulating D2R striatal functions [108]. Indeed, how striatal SPNs compute lipids-carried information at the cellular and circuit level represent the next step in the understanding of neuronal lipid sensing mechanisms.

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Concluding remarks and future perspectives

In this review, we have highlighted and discussed the most recent discoveries related to whether and how dietary lipids control ingestive behavior. Overall, dietary lipids, notably TG and their derived FFA, can govern appetite and reinforcement through the modulation of DA release/signaling by multimodal mechanisms of lipid sensing. These mechanisms involve nervous inputs arising from the oral cavity and from various compartments of digestive tract (mouth-to-brain and gut-to-brain signals) as well as direct neuromodulatory actions of dietary TG/FFA on cell type-specific neurons of the mesolimbic circuit (central detection). Traditionally believed to solely serve as energy substrates, dietary lipids are nowadays also functionally involved in regulating brain systems. Dietary lipid-dependent actions encompass a variety of intracellular mechanisms ranging from GPCR signaling, direct and indirect allosteric regulation of receptors and channels, release of byproduct messengers from mitochondrial lipid oxidation and transcriptional regulation via lipid sensing transcription factors. This is ensured by the presence of lipids-associated molecular detectors in key cellular elements located at different strategic levels: mouth, gut, brain. Such complexity of integrative mechanisms reveals a highly sophisticated regulation of pre-prandial, prandial and postprandial fat signals which collectively contribute to the reinforcing aspects of food, and

most probably also to cognitive functions associated with the encoding of food cues. It is interesting to highlight that at the physiological level dietary lipid sensing mechanisms may follow a time-dependent process: fast (mouth-to-brain, taste), intermediate (gut-tobrain, interoception) and slow (neuronal sensing) actions. Moreover, this may strongly depend on nutritional/metabolic states and endogenous/exogenous sources of lipids as during postprandial and fed states circulating TG represent the main source of lipids, whilst in fasting and energy-deprived conditions circulating FFAs are increased. However, in conditions of abnormal circulating lipids including (but not restricted to) obesity, neural substrates involved in reward and cognitive processing are altered. It is therefore tempting to hypothesize that dietary lipid sensing mechanisms within neural circuits supporting appetite and reinforcing behaviors may undergo functional maladaptations during obesity, as already observed in the tongue [109], gut [110] and hypothalamic arcuate nucleus [59,60]. In fact, whether long-term maladaptive lipidsinduced molecular and synaptic responses may occur within the reward system during obesity remain to be established. In line with this hypothesis we have shown that shortterm binge consumption of palatable diet was able to downregulate the expression of LPL mRNA in the nucleus accumbens [11]. Future studies are now warranted to unveil indepth mechanisms of cellular lipid sensing in physiological and pathological contexts. Moreover, the emerging field of lipid sensing could also pave the way to explore the etiopathology of other reward-related psychiatric and metabolic disorders such as anorexia nervosa, bulimia and binge eating [111]. Indeed, recent discoveries and current hypotheses also raise new interesting challenges that need to be tackled to fully depict a novel conceptual framework that sees dietary lipids as key modulators of integrative functions (see **Outstanding Questions**).

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415	The authors declare no conflicts of interest.

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Figure legends

Figure 1. The physiology of fat sensing throughout the body.

Oral perception of dietary fat requires the action of lingual lipases which hydrolyze TG into free fatty acids (FFA) whose presence is detected by taste bud cells. Encoded information is then rapidly sent toward the brain via the chorda tympani (VII) and glossopharyngeal (IX) nerves. After ingestion/digestion, dietary lipids can promote gut-to-brain communication and/or enter into the gut-liver transformative cycles. In the latter, FFA transported and re-esterified by enterocytes are packaged into TG-rich chylomicrons (CM) and released into the lymphatic and circulatory systems. The breakdown of lipoproteins through the action of lipoprotein lipase (LPL) allows the extraction of TG from large CM/VLDL leading to intermediate or high-density particles (IDL, HDL) and chylomicron remnants (rCM) recaptured by the liver. FFA and TG directly or indirectly signal to the brain. This figure was created using BioRender.

Figure 2. Fat sensing by taste bud cells.

Type 3 taste bud cells (TBC) express gustatory receptors for sweet, umami and bitter tastes. However, they also express receptors that can be activated by FFA released from dietary TG by the action of lingual lipases. FFA can activate lipid sensor CD36 and G-coupled receptor 40 (GPR40/FFAR1) and 120 (GPR120/FFAR1). The activation of Src kinase as a consequence of CD36 activation leads the recruitment of phospholipase C (PLC), the increase of intracellular Ca²⁺ and the activation of ERK1/2 through the calmodulin activated kinase (CaMKII). PLC promotes the release of intracellular Ca²⁺ through inositol 3 phosphate (IP3). Delayed Rectifying K+(DRK, Kv1.5) channels can directly or indirectly respond to FFA. Once activated by the detection of dietary lipids, TBC will release neurotransmitters and messengers allowing the transfer of lipids-associated information to the nucleus tactus solitarius (NTS) through the cranial nerves VII et IX. This figure was created using BioRender.

Figure 3. Gut lipid sensing: mechanisms and circuits.

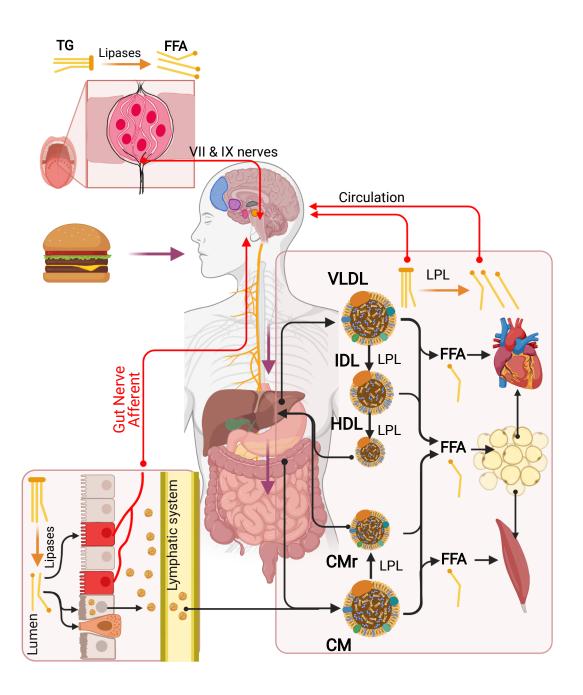
During intestinal transit, dietary TG are hydrolyzed by luminal lipases. Released 2-monoacylglycerol (2-MAG) and FFA will act on lipid sensors expressed onto enteroendocrine cells (EEC) (1, yellow circles). While TG-derived FFA can initiate signaling events by binding to CD36, G-coupled receptor 40 (GPR40/FFAR1) and 120 (GPR120/FFAR4), 2-MAG can signal by GPR119 (2, red circles). Upon activation of lipid receptors located on EEC membranes, gut hormones and neurotransmitters can be released from EEC (3, blue circles). In turn, gut hormones can enter the general circulation or directly stimulate vagal afferents.

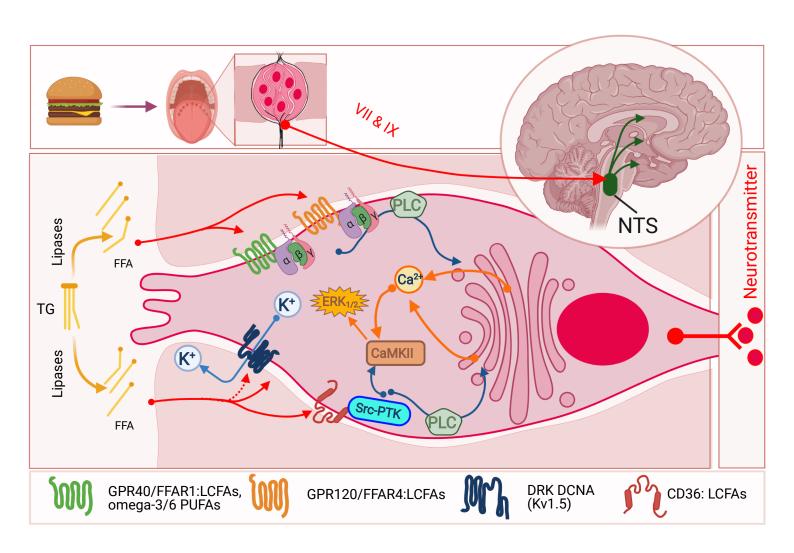
Within the enterocytes, FFA are re-esterified and packaged into chylomicrons (CM) to ultimately reach the lymphatic system (4, green circles). A portion of FFA entering the enterocytes can serve either as substrate for the synthesis of N-acylethanolamines and produce the bioactive N-oleoylethanolamine (OEA) (5, grey circles), or enter lipid oxidation processes whom beta-hydroxy butyrate (BHB) byproduct (6, brown circles) can also signal through the circulation and/or afferent fibers. Vagal afferent fibers, ascending through the nodose ganglia (NG), respond to OEA while spinal afferent fibers routed through the dorsal root ganglia (DRG) respond to BHB produced by enterocytes and microbiota. OEA, BHB, gut hormones and nutrients also signal through portal vein metabolic sensors and liver-brain afferent fibers. This figure was created using BioRender.

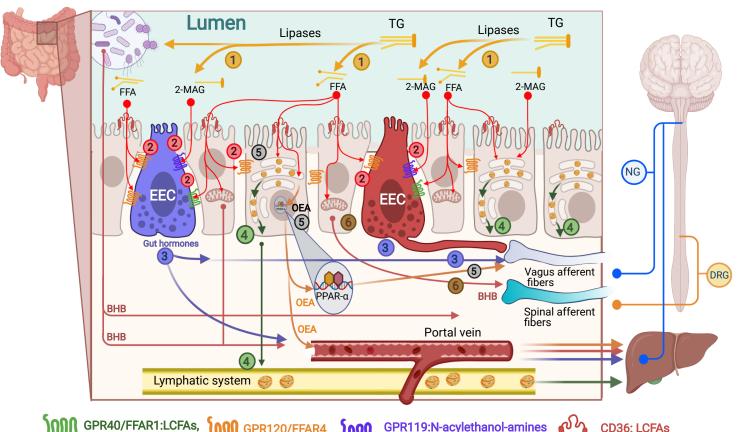
Figure 4. Direct and indirect physiological routes of lipid sensing in the reward system.

(A) Dietary lipid sensing modulates dopamine (DA) release and signaling through segregated direct and indirect physiological routes. Intestinal detection of lipids is conveyed to the striatum via a vagus→nucleus tractus solitarius (NTS)→parabrachial nucleus (PBN)→ substantia nigra (SNc) path (blue). Circulating lipids can also directly modulate DA-neurons in the ventral tegmental area (VTA) and dopaminoceptive neurons in the dorsal striatum (DS) and nucleus accumbens (NAc). (B) The lipoprotein lipase (LPL) in the brain microvasculature releases free fatty acids (FFA) from triglyceride (TG)-rich circulating particles such as chylomicrons (CM). Smaller in size, high-density lipoprotein (HDL) can cross the capillaries through transcytosis and via astrocyte endfeet.

(C) FFA can enter the neuron through membrane fatty acids transporter (FATP_{1/4}) or via the hydrolysis of TG from TG-particle docked on lipoprotein receptor (LDL-R). In the VTA, TG/FFA-mediated signaling was shown to decrease firing activity. In the striatum, dopamine D2 receptor (D2R)-expressing spiny projection neurons (SPNs) respond to TG/FFA through a mechanism that involves the LPL. TG/FFA modulate *bona fide* D2R-coupled intracellular signaling events and reduce neuronal excitability. Altogether, lipids-regulated circuits and neuronal responses concur in scaling reward-driven behaviors. Abbreviations: protein kinase A (PKA), extracellular signal-regulated kinase (ERK), ribosomal protein S6 (rpS6), ribosomal protein S6 kinase (S6K), mechanistic target of rapamycin (mTOR), dopamine- and cAMP-regulated neuronal phosphoprotein (DARPP-32), glycogen synthase kinase 3 beta (GSK-3b). This figure was created using BioRender.







GPR40/FFAR1:LCFAs, omega-3/6 PUFAs







CD36: LCFAs

