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Lipid droplet biogenesis and functions in health and disease

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Supplementary Information:

Supplementary box 1: Adipose heterogeneity: size (hyperplasia) vs number (hypertrophy)

Unified by the ability to store energy in lipid droplets, adipose tissue is a remarkably heterogeneous organ varying in colour, anatomical distribution, and function (**Supplementary table B1**). It is comprised of lipid droplet occupying adipocytes among a stromal vascular fraction of fibroblasts, nerve tissue, blood vessels and immune and endothelial cells¹. Though, adipocyte turnover stabilises to 8.3 years in adulthood², adipose tissue expansion may occur by increasing adipocyte size (hypertrophy), number (hyperplasia) or both. In the former, existing adipocytes become engorged through enhanced lipid droplet growth and accumulation, before supporting the development of the latter, formation of new adipocytes through increased adipogenesis of precursors³. As angiogenesis precedes hyperplasia⁴, and thereby enables sufficient vascularisation and adipokine and cytokine signalling upon expansion, hyperplasia is considered an adaptive response⁵. Hypertrophy, on the other hand, is linked to an onslaught of negative metabolic outcomes. It is positively correlated with cell senescence, strongly implicated in the development of inflammation, associated with mitochondrial dysfunction, elevated basal lipolysis, hypoxia, fibrosis, and increased leptin but reduced adiponectin production and secretion^{6,7}. Interestingly, hypertrophy predominates visceral adipose tissue expansion, while hyperplasia drives the growth of subcutaneous depots⁸. By regulating the vascular supply of adipose tissue, the sex hormone, estrogen may influence the mode of hypertrophic or hyperplasic expansion, in addition to modulating the typical gynoid or android adipose tissue distribution of premenopausal women or men, respectively⁹. According to the adipose tissue expandability hypothesis, which concurs with both the adipokine and inflammation hypotheses^{10,11}, it is this capacity of expansion, rather than the absolute amount of adipose tissue one possesses that may lead to such adverse metabolic complications. This is because beyond the critical point where further expansion is limited, lipotoxicity may ensue through increased circulating lipids and ectopic lipid accumulation in non-adipose insulinsensitive tissues such as the liver, heart, and skeletal muscle 12 .

Supplementary table B 1: lipid droplet diversity across mammalian cells in health and disease

Supplementary box 2: lipid droplet biogenesis vs adipogenesis: mutual factors and mechanistic differences

Broadly involved in the anabolic arm of lipid metabolism, lipid droplet biogenesis and adipogenesis are mechanistically distinct processes united by several mutual factors. Unlike lipid droplet biogenesis which constitutes the creation of neutral lipid storage organelles, adipogenesis involves the generation of new adipocytes via two main steps. In the initial commitment phase, mesenchymal precursors commit to the adipocyte lineage, becoming preadipocytes with unaltered morphology³⁰. During differentiation, preadipocytes then change conformation to enable lipid accumulation and fatty acid binding protein 4, glucose transporter 4, leptin and adiponectin expression 31 . Countless extranuclear factors regulate this process, including local insulin, triiodothyronine, glucocorticoid, brain natriuretic peptide, Wnt, transforming growth factor-β and hedgehog signalling, as well as systemic influences from circadian rhythms, reactive oxygen species and chronic inflammation^{5,32}. However, the most critical factor for white, brown, and beige adipocyte differentiation remains the transcriptional master regulator, PPAR- γ^{33-35} . Whilst PPAR- γ is both necessary and sufficient for adipogenesis³⁶, it can be functionally synergised by its coactivator, $C/EBP\alpha^{37}$, and potently inhibited by phosphatidic acid and its derivative, cyclic-phosphatidic acid^{38,39}.

Originally identified for its role in adipogenesis⁴⁰, and subsequently linked to lipid droplet formation $41,42$, seipin has emerged as the central factor unifying the mutual influencers of these processes. Not only does it interact with phosphatidic acid among other phospholipids⁴³, its interaction with key enzymes involved in lipogenesis has provided mechanistic insights into its dual roles. Through binding to and putatively inhibiting the rate-limiting enzyme in glycerolipid synthesis, GPAT3/4⁴⁴, Seipin may reduce phosphatidic acid levels, preventing PPAR-γ inhibition and enabling adipogenesis. Moreover, its simultaneous and direct interaction with the PA producing, AGPAT2 and the phosphatidic acid phosphatase, lipin $1⁴⁵$ might provide a mechanism for both phosphatidic acid clearance and lipogenesis regulation by seipin. Although AGPAT2 functions to synthesise phosphatidic acid, its deficiency causes a nearly three-fold increase in the level of phosphatidic acid. The recently identified direct interaction between AGPAT2 and $CDS1/2^{46}$ explains this phosphatidic acid increase under AGPAT2 deficiency and also supports previous findings linking CDS proteins with the regulation of both cellular (lipid droplet formation) and systemic (adipogenesis) lipid storage⁴⁷.

Supplementary box 3: Lipolysis, lipophagy and chaperone-mediated autophagy

Activated during energy deprivation or stress, lipolysis, lipophagy and chaperone-mediated autophagy (CMA) are three distinct yet cooperative processes essential for regulating the degradation, and in turn size, of cLDs. In cytosolic lipolysis, triacylglycerols are sequentially hydrolysed by the neutral lipases, patatin-like phospholipase domain-containing protein 2 (PNPLA2), hormone sensitive lipase (HSL) and monoglyceride lipase⁴⁸. In both lipophagy and CMA, lysosomal degradation catalysed by lysosomal acid lipase (LAL), or the chaperone, heat shock cognate protein 70 occurs, respectively⁴⁹. Initially considered parallel processes, recent findings have revealed several functional and synergistic links between them. In particular, CMA has been shown to promote neutral lipolysis through the direct binding of heat shock cognate protein 70 to the KFERQ pentapeptide motif on perilipin 2 and $3⁵⁰$, allowing PNPLA2 unrestricted lipolytic access to the surface of cLDs. Moreover, whilst lipophagy has been found to precede lipolysis during starvation^{51,52}, lipolysis of large cLDs have been shown to supply small cytoplasmic lipid droplets for subsequent lipophagic digestion²¹. Furthermore, mammalian target of rapamycin complex 1 not only acts as the prominent negative regulator of lipophagy through its Ser142 and S211 phosphorylation and cytoplasmic sequestration of transcription factor $EB^{53,54}$, it is also implicated as an inhibitor of PNPLA2-stimulated lipolysis via its upregulation of early growth response protein 1⁵⁵.

Interestingly, both activation and inhibition of these interconnected processes is strongly and amply connected with disease. Unrestrained basal lipolysis, for example, is well established in the development of insulin resistance sequelae and cancer cachexia^{12,56}. In turn, pharmacological inhibition^{57,58} or genetic depletion^{59–61} of PNPLA2, HSL and monoglyceride lipase, have been shown to improve insulin sensitivity. Despite this, humans deficient in PNPLA2 or its coactivator, abhydrolase domain containing 5 experience systemic triacylglycerol accumulation in the form of neutral lipid storage disease with myopathy $(NLSDM)$ or ichthyosis (NLSDI), respectively^{62,63}. Likewise, LAL deficient mice demonstrate enhanced glucose tolerance⁶⁴, and humans with mutations in its encoding gene exhibit Wolman or cholesterol ester storage disease, depending on the extent of lost LAL activity⁶⁵. Finally, whereas downregulation of CMA is linked to increased lipogenesis, lipid uptake and synthesis, leading to fatty liver disease and atherosclerosis^{66,67}, upregulation of CMA may also contribute to the pathogenesis of the autoimmune disorder, lupus⁶⁸.

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