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Brain-specific microRNA-mediated regulation of metabolic homeostasis in *Drosophila melanogaster*

To maintain optimal metabolic balance, an organism must coordinate its food intake, fat storage, and appropriate energy mobilization to meet physiological demands and support overall well-being. These metabolic processes require precise regulation to ensure timely responses to environmental and internal cues. Small non-coding microRNAs (miRNAs) are well suited for this role, as they can regulate physiological and adaptive processes that require rapid or localized control of gene expression.

We have identified a novel neural circuit in *Drosophila* that plays an important role in metabolic homeostasis, mediated by the conserved Central Nervous System (CNS)-specific miRNA miR-1000. We previously identified miR-1000 to be a neuroprotective miRNA by regulating Glutamatergic signaling in the brain (Verma et al., Nature Neuroscience, 2015). In addition to its role in neuroprotection, we recently discovered that miR-1000 also regulates key aspects of metabolism, including body weight, fat storage, survival during nutrient deprivation, and longevity (Verma et al, Genes and Development, 2026). miR-1000 achieves this by repressing the Neuropeptide Like Peptide 1 (Nplp1) gene in the CNS. Loss of miR-1000 and consequent increase in Nplp1 expression result in flies with higher fat storage and extended survival under nutrient-deprived conditions. While the biological function of Nplp1 neuropeptides has been largely unknown, this study highlights their critical role in maintaining metabolic homeostasis in *Drosophila*. Additionally, we observed that miR-1000 and Nplp1 levels change dynamically during refeeding after starvation, suggesting their involvement in adaptive metabolic responses. miR-1000 regulates triacylglycerol (TAG) synthesis and storage in lipid droplets, supports fat mobilization during starvation, and facilitates its storage upon refeeding. These findings underscore the importance of miR-1000 and Nplp1 in metabolic regulation and provide new insights into the mechanisms of energy balance.