

New mediators in diabetes pathogenesis: Exosomes and metabolites

Type 2 diabetes is a systemic disorder characterized by metabolic impairment in multiple organs¹. Various factors trigger insulin resistance in muscle, liver and adipose tissue resulting in increased insulin demand and de-repression of hepatic gluconeogenesis. When pancreatic β -cells fail to compensate their mass and function to the increased insulin demand, β -cell apoptosis and dedifferentiation occurs, which leads to hyperglycemia. As the diabetes progresses, systemic complications involving microvessels (retinopathy, nephropathy and neuropathy) and macrovessels (cardiovascular, cerebrovascular and peripheral vascular diseases) develop, which leads to increased mortality. As simple it might seem, the mechanism of how diabetes develops and progresses has not been fully elucidated. Cumulative evidence suggests that inter-organ communicating factors, such as exosomes (or extracellular vesicles) or metabolites, might participate in the process of developing type 2 diabetes and its complications (Figure 1).

Exosomes are 30–100 nm sized lipid bi-layered vesicles containing proteins, lipids, carbohydrates and nucleic acids (micro ribonucleic acids, long non-coding ribonucleic acids and messenger ribonucleic acids)². Exosomes are formed through the endosomal pathway and can be secreted by most cell types. Importantly, exosomes secreted from a certain cell can be delivered to adjacent or distant cells in an autocrine, paracrine or endocrine manner to serve as an intercellular communicator. Interestingly, the circulating exosome level was significantly higher in patients with diabetes compared with euglycemic controls, suggesting the potential role of exosome in diabetes³. A well-designed rodent study showed that adipose tissue is an important source of circulating exosomes⁴. Along with adipokines or free fatty acids, exosomes are one of the important mediators of how adipose tissues contribute to the systemic insulin resistance. Adipose tissue-derived exosomes can develop insulin resistance by stimulating macrophages to express inflammatory cytokines (interleukin-6, tumor necrosis factor- α)⁵. Exosomal micro ribonucleic acid-27a from adipocytes can induce insulin resistance in muscles⁶. Adipose tissue macrophage derived exosomal micro ribonucleic acid-155 has the potential to impair insulin signaling in muscle, liver and adipose tissue by peroxisome proliferator-activated receptor γ suppression. Muscle-derived exosomes also play an important role in systemic metabolism. Muscle-derived exosomes from high palm oil-fed mice can incorporate to the pancreatic β -cells to regulate β cell mass⁷. Muscle-derived exosomes can also modulate myoblast proliferation and differentiation in a paracrine manner⁸. Research on pancreatic islets and exosomes have been heavily focused on the effect of other tissue-originated

exosomes on pancreatic islets or β -cells. However, some evidence suggests that pancreatic islets also excrete exosomes, which act in a paracrine manner to regulate glucose homeostasis. Exosomal neutral ceramidase are secreted from β -cells when treated with pro-inflammatory cytokines, which protects β -cells from palmitate- or cytokine-induced apoptosis⁹. Healthy human donor islet-derived exosomes suppress amyloid deposition in a paracrine manner, which is impaired in islets of patients with type 2 diabetes¹⁰. Exosomal Lnc-364 from β -cells can regulate insulin secretion and β -cell proliferation¹¹. As such, exosomes from various tissues can modulate biological processes in cells in an endocrine or paracrine manner, which can trigger insulin resistance and β -cell failure to develop type 2 diabetes. As many of the exosome studies were carried out *in vitro* or *ex vivo*, more carefully designed *in vivo* or translational research should be carried out.

Along with the exosomes, metabolites are circulating biomolecules that have the potential to activate biological processes in various tissues. Metabolomics studies in prospective cohorts suggest the potential role of various metabolites on the development and progression of type 2 diabetes. A large-scale metabolomics study in the Framingham Offspring cohort suggested that five branched chain amino acids (tyrosine, isoleucine, leucine, phenylalanine and valine) are highly associated with the development of diabetes¹². A metabolomics study showed the association between specific metabolites (branched chain and aromatic amino acids, triacylglycerol in very low-density lipoprotein, non-esterified cholesterol in high-density lipoprotein and linoleic n-6 fatty acid) and diabetes risk¹³. Also, serum levels of metabolites (phospholipids, adenosine monophosphate) were associated with the preventative (pharmacological and lifestyle intervention) effect on the development of type 2 diabetes¹⁴. Despite epidemiological studies strongly suggesting the role of metabolites in the pathogenesis of diabetes, the mechanism of how these metabolites affect whole-body glucose metabolism is poorly understood. Recently, radioisotope-tracing combined metabolomics study enabled researchers to understand the physiological or pathological flux of metabolites¹⁵. This technological advancement will encourage researchers to further study the mechanism how metabolites affect whole-body glucose metabolism.

In conclusion, exosomes and metabolites are the new emerging mediators of diabetes development. Recently, a novel subclassification of newly diagnosed diabetes patients was documented¹⁶. This new subclassification (severe autoimmune, severe insulin deficient, severe insulin resistant, mild obesity-related and age-related) was based on pathophysiological

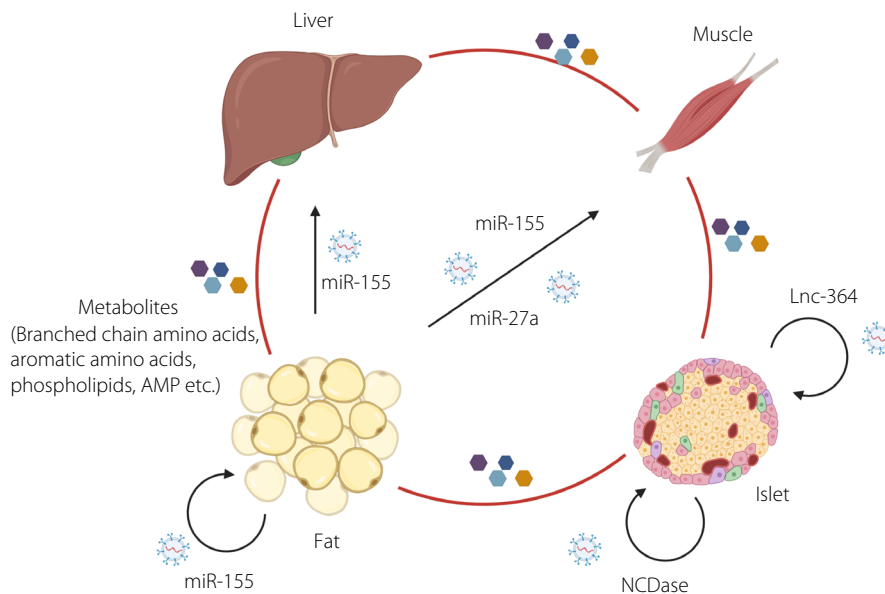


Figure 1 | Examples of the role of exosomes and metabolites in the pathogenesis of diabetes. Circulating exosomes and metabolites mediate inter-communicating signals between endocrine organs (liver, muscle, fat, islets), which are associated with the development type 2 diabetes. AMP, adenosine monophosphate; miR, micro ribonucleic acid; NCDase, neutral ceramidase.

heterogeneity, and this heterogeneity was shown to exist even before the development of diabetes^{16,17}. Future studies comparing the difference of exosomes and metabolites between different diabetes subclasses might show valuable information regarding the pathogenesis of diabetes. These novel mediators might serve their role as a biomarker to predict the development and progression of diabetes. Future well-designed mechanistic studies will encourage the development of therapeutic targets associated with these novel mediators.

ACKNOWLEDGMENTS

This work was supported by grants from the National Research Foundation of Korea (2018R1D1A1B07043223, 2021R-1F1A1061197) to SHL.

DISCLOSURE

The authors declare no conflicts of interest.
 Approval of the research protocol: N/A.
 Informed consent: N/A.
 Approval date of registry and the registration no. of the study/trial: N/A.
 Animal studies: N/A.

Joonyub Lee¹, Seung-Hwan Lee^{1,2*}

¹Division of Endocrinology, Department of Internal Medicine, College of Medicine, Seoul St. Mary's Hospital, The Catholic

University of Korea, Seoul, Korea; ²Department of Medical Informatics College of Medicine, The Catholic University of Korea, Seoul, Korea
 *E-mail: hwanx2@catholic.ac.kr

REFERENCES

- DeFronzo RA. From the triumvirate to the ominous octet: a new paradigm for the treatment of type 2 diabetes mellitus. *Diabetes* 2009; 58: 773–795.
- Cocucci E, Meldolesi J. Ectosomes and exosomes: shedding the confusion between extracellular vesicles. *Trends Cell Biol* 2015; 25: 364–372.
- Freeman DW, Noren Hooten N, Eitan E, et al. Altered extracellular vesicle concentration, cargo, and function in diabetes. *Diabetes* 2018; 67: 2377–2388.
- Thomou T, Mori MA, Dreyfuss JM, et al. Adipose-derived circulating miRNAs regulate gene expression in other tissues. *Nature* 2017; 542: 450–455.
- Deng Z-B, Poliakov A, Hardy RW, et al. Adipose tissue exosome-like vesicles mediate activation of macrophage-induced insulin resistance. *Diabetes* 2009; 58: 2498–2505.
- Yu Y, Du H, Wei S, et al. Adipocyte-derived exosomal MiR-27a induces insulin resistance in skeletal muscle through repression of PPARγ. *Theranostics* 2018; 8: 2171–2188.
- Jalabert A, Vial G, Guay C, et al. Exosome-like vesicles released from lipid-induced insulin-resistant muscles modulate gene expression and proliferation of beta recipient cells in mice. *Diabetologia* 2016; 59: 1049–1058.

8. Aswad H, Forterre A, Wiklander OPB, *et al.* Exosomes participate in the alteration of muscle homeostasis during lipid-induced insulin resistance in mice. *Diabetologia* 2014; 57: 2155–2164.
9. Tang S, Luo F, Feng YM, *et al.* Neutral ceramidase secreted via exosome protects against palmitate-induced apoptosis in INS-1 cells. *Exp Clin Endocrinol Diabetes* 2017; 125: 130–135.
10. Ribeiro D, Horvath I, Heath N, *et al.* Extracellular vesicles from human pancreatic islets suppress human islet amyloid polypeptide amyloid formation. *Proc Natl Acad Sci* 2017; 114: 11127–11132.
11. Ruan Y, Lin N, Ma Q, *et al.* Circulating lncRNAs analysis in patients with type 2 diabetes reveals novel genes influencing glucose metabolism and islet β -cell function. *Cell Physiol Biochem* 2018; 46: 335–350.
12. Wang TJ, Larson MG, Vasan RS, *et al.* Metabolite profiles and the risk of developing diabetes. *Nat Med* 2011; 17: 448–453.
13. Ahola-Olli AV, Mustelin L, Kalimeri M, *et al.* Circulating metabolites and the risk of type 2 diabetes: a prospective study of 11,896 young adults from four Finnish cohorts. *Diabetologia* 2019; 62: 2298–2309.
14. Chen Z-Z, Liu J, Morningstar J, *et al.* Metabolite profiles of incident diabetes and heterogeneity of treatment effect in the diabetes prevention program. *Diabetes* 2019; 68: 2337.
15. Jang C, Chen L, Rabinowitz JD. Metabolomics and isotope tracing. *Cell* 2018; 173: 822–837.
16. Ahlqvist E, Storm P, Käräjämäki A, *et al.* Novel subgroups of adult-onset diabetes and their association with outcomes: a data-driven cluster analysis of six variables. *Lancet Diabetes Endocrinol* 2018; 6: 361–369.
17. Wagner R, Heni M, Tabák AG, *et al.* Pathophysiology-based subphenotyping of individuals at elevated risk for type 2 diabetes. *Nat Med* 2021; 27: 49–57.

Doi: 10.1111/jdi.13654