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Novel Diabetes Subgroups and Risk of Diabetes-Associated Complications

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Diabetes mellitus is a heterogeneous disease with respect to clinical presentation and progression. Recent studies challenged the current classification of the disease, and this short review aims to provide a concise update on novel findings related to diabetes subgroups and potential differences in their risk to develop diabetes-associated complications.

Heterogeneity of diabetes based on pathophysiological processes

The issue of heterogeneity of diabetes has been investigated with different data-driven methods in cohorts with incident and prevalent diabetes. Two interesting studies used latent class trajectory analysis to characterise patterns of changes in diabetes risk factors in individuals who developed diabetes in order to classify these individuals based on pathophysiological processes contributing to the onset of diabetes [Hulman 2018, Vistisen 2014]. In the Whitehall II study three patterns of changes in body mass index (BMI) preceding the diagnosis could be identified. Individuals in these subgroups

differed in their trajectories of insulin resistance, beta-cell function and other cardiometabolic risk factors including interleukin-1 receptor antagonist (IL-1Ra), a biomarker of subclinical inflammation [Vistisen 2014]. In the European Group for the Study of Insulin Resistance: Relationship between

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Insulin Sensitivity and Cardiovascular Disease (EGIR-RISC) study, data from a five-point oral glucose tolerance test were used to identify distinct glucose response patterns. Individuals in these four glucose response patterns also differed with respect to insulin resistance, beta-cell function and obesity. Several biomarkers showed different distributions between the four groups including C-peptide, lipids, and C-reactive protein and adiponectin, the latter both associated with subclinical inflammation [Hulman 2018]. Thus, data-driven analyses may be useful for cardiometabolic risk stratification based on pathophysiological processes.

Five diabetes subgroups in a Swedish cohort

In 2018 a Swedish study aimed to refine diabetes classification in patients with newly diagnosed diabetes by conducting a cluster analysis based on glutamate decarboxylase (GAD) antibodies, age at diagnosis, BMI, HbA_{1c}, and homeostasis model assessment estimates of beta-cell function and insulin resistance (HOMA-B and HOMA-IR) [Ahlqvist 2018]. This analysis resulted in five subgroups which were labelled as

- severe autoimmune diabetes (SAID),
- severe insulin-deficient diabetes (SIDD),
- severe insulin-resistant diabetes (SIRD),
- mild obesity-related diabetes (MOD) and
- mild age-related diabetes (MARD).

The SAID subgroup was characterised by the presence of GAD antibodies, early age of diagnosis, low BMI, insulin deficiency and poor glycaemic control. The SIDD subgroup was similar to SAID, but without GAD antibodies. Individuals in the SIRD and MOD subgroups had the highest BMI with the strongest insulin resistance found in the SIRD patients. The MARD subgroup had the highest age at diagnosis, a lower BMI comparable to SAID and SIDD and relatively good glycaemic control [Ahlqvist 2018]. With respect to diabetes complications, the SIRD subgroup showed the highest risk of diabetic kidney disease. In contrast, SIDD had the highest risk of retinopathy. Clusters also differed with respect

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to coronary events and stroke, but this was abolished by adjusting for age and sex [Ahlqvist 2018].

Subgroup replication in the German Diabetes Study

The Swedish data were replicated in several other international cohorts [Dennis 2019, Zaharia 2019, Zou 2019]. In particular, the analysis in the German Diabetes Study (GDS) corroborated and extended the data pointing towards differences in the risk of diabetes-associated diseases between the aforementioned clusters [Zaharia 2019]. The GDS enrolls adults within 12 months of their diagnosis of diabetes [Szendroedi 2016]. In this cohort, individuals in the SIRD subgroup not only had the highest prevalence of diabetic nephropathy, but also the highest hepatocellular lipid content at study baseline and the highest prevalence of hepatic fibrosis after five years of follow-up [Zaharia 2019]. Individuals in the SIDD subgroup were characterised by the highest prevalence of confirmed diabetic sensorimotor polyneuropathy (DSPN). Both SIRD and MOD had the highest levels of C-reactive protein, i.e. the biomarker of inflammation most frequently assessed in the context of diabetes-associated complications. Importantly, this study also evaluated the stability of subgroup allocation over a 5-year follow-up period. Changes of metabolic variables, especially glycaemia, lipid levels and fatty liver index, were related to switches between clusters which were observed for 23 % of all patients for whom data from both the baseline and the 5-year follow-up examinations were available [Zaharia 2019].

Diabetes subgroups and inflammation

Future studies need to investigate in more detail which other biochemical and clinical variables distinguish these novel diabetes subgroups and to what extent these differences may explain the differences in the risk of complications in patients with diabetes. Subclinical inflammation represents one can-

didate, because inflammatory processes and biomarkers are related not only to the development of cardiovascular disease [Herder 2015], diabetic neuropathies [Herder 2019b] and nephropathy [Niewczas 2019], but also to comorbidities such as depression [Herder 2019a]. In particular, the cascade of adipose tissue dysfunction, inflammation, insulin resistance and onset of complications merits more detailed studies [Rodén 2019]. Further issues to be addressed comprise the stability of the cluster allocation over time, the reproducibility in different ethnic groups and most importantly the clinical relevance of the novel subgroups for prevention and therapy that requires testing in randomised controlled trials.

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