

Health consequences of early-onset compared with late-onset type 2 diabetes mellitus

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ABSTRACT

Background: Although cumulating evidence has suggested that early-onset type 2 diabetes mellitus (T2DM) conferred on patients a broader tendency for complications beyond vascular ones, a comprehensive analysis of patterns of complications across all relevant systems is currently lacking.

Method: We prospectively studied 1 777 early-onset (age at diagnosis ≤ 45 years) and 35 889 late-onset (>45 years) T2DM patients with matched unexposed individuals from the UK Biobank. Diabetes-specific and -related complications were examined using phenome-wide association analysis, with patterns identified by comorbidity network analysis. We also evaluated the effect of lifestyle modifications and glyceamic control on complication development.

Results: The median follow-up times for early-onset and late-onset T2DM patients were 17.83 and 9.39 years, respectively. Compared to late-onset T2DM patients, patients with early-onset T2DM faced a significantly higher relative risk of developing subsequent complications that primarily affected sense organs [hazard ratio (HR) 3.46 vs. 1.72], the endocrine/metabolic system (HR 3.08 vs. 2.01), and the neurological system (HR 2.70 vs. 1.81). Despite large similarities in comorbidity patterns, a more complex and well-connected network was observed for early-onset T2DM. Furthermore, while patients with early-onset T2DM got fewer benefits (12.67% reduction in pooled HR for all studied complications) through fair glyceamic control (median $\text{HbA}_{1c} \leq 53$ mmol/mol) compared to late-onset T2DM patients (18.01% reduction), they seemed to benefit more from favorable lifestyles, including weight control, healthy diet, and adequate physical activity.

Conclusions: Our analyses reveal that early-onset T2DM is an aggressive disease resulting in more complex complication networks than late-onset T2DM. Aggressive glucose-lowering intervention, complemented by lifestyle modifications, are feasible strategies for controlling early-onset T2DM-related complications.

Keywords: early-onset type 2 diabetes mellitus, phenome-wide association analysis, comorbidity network analysis, late-onset type 2 diabetes mellitus

Introduction

Early-onset type 2 diabetes mellitus (T2DM) is becoming a fundamental public health issue worldwide. The landmark SEARCH Study¹ reported that early-onset T2DM accounted for 13.7% of all people with diabetes in youth and recorded a 4.8% annual surge in incidence.² Early-onset T2DM appears to be a more aggressive disease by unclear mechanisms in terms of frequency of complications that are not commensurate merely with the age of onset.^{3,4} Previous studies have primarily focused on the microvascular and macrovascular complications of early-onset T2DM.^{5,6} However, recent data suggest that the complications are perhaps widely distributed throughout the human system,^{7–10} even in the early stages of the disease, but the evidence-based data gap is the Achilles' heel of these studies.

A way to overcome this challenge is to perform phenome-wide association analyses (PheWAS) to identify the characteristics of subsequent complications, the essential prerequisite to developing effective interventions. Taking advantage of the rich in-

formation on demographics and lifestyles as well as the complete medical record in the UK Biobank, we investigated the health consequences of early-onset T2DM (age at diagnosis ≤ 45 years) compared to late-onset T2DM (age at diagnosis > 45 years) through systemically computing the phenotypic risk and patterns of morbidities. Furthermore, we explored whether the heightened risk of complications in early-onset T2DM may be modifiable by glyceamic control or favorable lifestyles. To our knowledge, this is the first PheWAS study on T2DM, with potentially far-reaching aims that provide not only an overview of potential complications but also has implications for effective intervention strategies.

Methods

Study design and data sources

We analyzed data from the UK Biobank, a community-based prospective study described in detail elsewhere.¹¹ Briefly, the UK Biobank recruited 502 507 participants aged between

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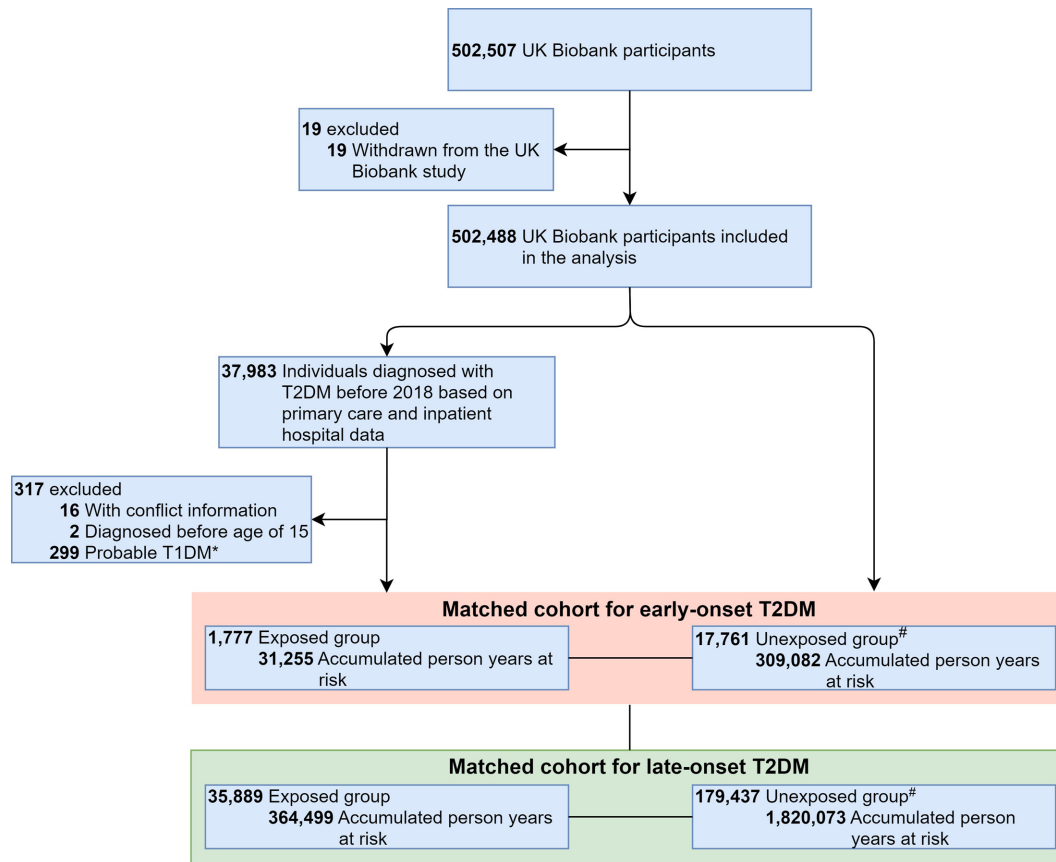


Figure 1. Flowchart for population selection. *The algorithm for identifying probable T1DM patients is shown in Supplementary Fig. 1. #For each T2DM patient, we randomly selected up to 5 (for late-onset T2DM patients) or 10 (for early-onset T2DM patients) individuals without T2DM diagnosis at the diagnosis date of the index patient (i.e. the index date) from the study population, individually matched by date of birth (± 1 year), sex, and decile of Townsend deprivation index.

40–69 years between 2006 and 2010, with sociodemographic information, lifestyle factors and medical conditions collected at baseline. The National Health Service (NHS) Digital and the NHS Central Register data were combined to supplement mortality information. Inpatient hospital information from 1997 onwards was obtained from nationwide databases, including the Hospital Episode Statistics, the Patient Episode Database for Wales, and the Scottish Morbidity Record.¹¹ The primary care records as of 2017 were integrated from multiple general practice (GP) databases, covering 45% of the UK Biobank participants.¹² The UK Biobank has full ethical approval from the NHS National Research Ethics Service (reference number: 16/NW/0274), and the biomedical research ethics committee of West China Hospital approved the current study (reference number: 2019–1171).

Disease categories

Initially, 37 983 individuals were diagnosed with T2DM based on primary care and inpatient records (diagnosis codes are given in Supplementary Table 1, see online supplementary material). We excluded 299 suspected T1DM patients to reduce misdiagnosed or misclassification as T2DM,¹³ according to a validated algorithm for T1DM identification in UK Biobank data (Supplementary Fig. 1, see online supplementary material).¹⁴ We further excluded 18 cases with conflicting information or diagnosed before the age of 15, leaving 1 777 early-onset and 35 889 late-onset T2DM patients. We then randomly selected 5 or 10 age, sex, and Townsend deprivation index-matched unexposed individuals for each T2DM

patient (Fig. 1). With satisfiable accuracy for T2DM diagnosis in the linked databases (82%–90%),^{15–17} the UK Biobank database has been widely used for diabetes-related studies.^{18, 19}

Identification of subsequent medical conditions was based on inpatient hospital data, with date of diagnosis recorded as the date of the first identified hospital episode. We restricted the original International Classification of Diseases-10th revision (ICD-10) codes to chapters 1–14 (i.e. excluding the codes related to pregnancy, perinatal conditions, symptoms or signs, or external causes of morbidity and mortality). The ICD-10 codes were mapped to ‘phecodes’, a coding system considered more relevant to medical conditions mentioned in clinical settings.²⁰ To sustain sufficient data power, we only used the first level ‘phecodes’ (3-character ‘phecodes’), resulting in 413 specific medical conditions in the analysis (Supplementary Table 2, see online supplementary material). These medical conditions were summarized as diabetes-specific complications (i.e. microvascular complications and macrovascular complications, according to a previous study²¹) and complications related to diabetes, which were further summarized as 13 broad categories based on the mainly affected systems. The diagnoses of medical conditions in inpatient data in the UK have been validated, demonstrating an overall high diagnostic accuracy.²²

Sociodemographic characteristics and lifestyle factors, including birth, sex, household income, family history, smoking status, drinking status, and fruit/vegetable consumption, were collected at baseline. The anthropometric data used for body mass index (BMI) calculations were measured during the baseline visit.

Each participant was assigned a Townsend deprivation index based on the postal codes provided at recruitment as an index of population-level deprivation.²³ For individuals who completed the physical activity questionnaire, total physical activity was calculated by summing the metabolic equivalent task (MET)-weighted time spent in vigorous, moderate, and walking activity.²⁴ We further extracted HbA_{1c} (Hemoglobin A_{1c}) test results recorded both at baseline assessment and in the primary care database (Supplementary Table 3, see online supplementary material). We defined 'fair glycemic control' as a median HbA_{1c} ≤ 53 mmol/mol,²⁵ calculated based on all available HbA_{1c} tests conducted at least 6 months after the T2DM diagnosis.

Statistical analysis

PheWAS for identifying T2DM-associated complications

Hazard ratio (HR) of diseases per types of complications (i.e. microvascular complications, macrovascular complications, or complications related to diabetes) was first assessed using Cox models, which were stratified by matching identifier (i.e. date of birth, sex, and Townsend deprivation index) and partially or fully adjusted for household income, BMI, smoking, and alcohol-drinking status. PheWAS for specific medical conditions with a prevalence ≥1% were then performed using fully adjusted Cox models. Specifically, we removed individuals with a history of all related medical conditions indicated in the previous study during the analysis of each medical condition of interest (Supplementary Table 2).²⁰ Medical conditions with *P*-value < the Bonferroni corrected threshold and HR > 1.0 were considered complications significantly associated with the prior T2DM diagnosis. To quantify the overall magnitude difference between PheWAS results of the matched cohort for early-onset T2DM and those of late-onset T2DM, we calculated the changed percentages of pooled HRs obtained using the inverse-variance weighting method (i.e. $(\frac{HR_{\text{early-onset}}^{\text{pooled}}}{HR_{\text{typical}}^{\text{pooled}}} - 1) \times 100\%$).

Comorbidity network analyses

The comorbidity network was constructed to investigate the diversity of disease clusters associated with a prior diagnosis of early-onset and late-onset T2DM.²⁶ All possible pairs were compiled using identified complications in the PheWAS step. We estimated relative risk (RR) and ϕ -correlation as the measures of strength of comorbidity associations for pairs with a prevalence ≥1.0%. The comorbidity networks were then constructed based on the pairs of complications with RR ≥ 2.0 and ϕ -correlation ≥ 0.1. Next, to clarify the comorbidity patterns, the networks were partitioned into groups with high intrinsic connectivity (i.e. modules), using the community detection algorithm Louvain.²⁷

Sub-analyses and sensitivity analyses

We performed sub-analyses to evaluate the prioritized blood glucose control strategies on subsequent complications for early-onset and late-onset T2DM patients. Likewise, we explored the potential modifying effect of lifestyle factors by performing sub-analyses for T2DM patients with adequate or inadequate fruit/vegetable consumption,²⁸ for those with BMI ≥ 29.9 or < 29.9, and for those with total physical activity ≤ 798 or > 798 MET.min/week. Furthermore, we also repeated the sub-analyses for different lifestyle factors among T2DM patients with poor glycemic control (*n* = 1 089 and 8 901 for early-onset and late-onset T2DM, respectively).

In sensitivity analyses, to test the robustness of the main results with the definition of T2DM, we repeated the PheWAS after removing T2DM patients with any documented diagnosis of other types of diabetes (codes listed in Supplementary Table 3), and those with only one documented diagnosis of T2DM. Additionally, with a concern that the observed difference in the RR of subsequent complications after early-onset and late-onset T2DM was mainly attributed to longer disease duration (or more extended surveillance period) for early-onset than late-onset T2DM patients, we further re-assessed these estimates by restricting to the first 10 years of follow-up. Finally, to relieve the concern that the observed results could be largely influenced by systematic difference of comorbidity status between early-onset and late-onset T2DM patients, we also re-assessed these estimates after additionally excluding T2DM patients with a history of hypertension, disorders of lipid metabolism, overweight, obesity and other hyperalimentation, chronic kidney disease, or cardiovascular disease at the index date, along with their matched controls.

All the statistical analyses were conducted using SciPy (version 1.4.1), Statsmodels (version 0.11.1), and Lifelines (version 0.25.2) in Python 3.8.

Results

The matched cohort for early-onset T2DM included 1 777 exposed patients and 17 761 matched unexposed individuals. In the matched cohort for late-onset T2DM, the numbers of exposed patients and matched unexposed individuals were 35 889 and 179 437, respectively (Fig. 1 and Table 1). In addition to an equal sex distribution (60.32% vs. 59.54%), better socioeconomic conditions (−1.26 vs. 0.95 for median Townsend deprivation index), and more smokers (55.12% vs. 42.26%) and drinkers (91.05% vs. 83.57%) were observed in typical, compared with early-onset T2DM patients. A higher percentage of obesity (61.34% vs. 56.86% for BMI ≥ 29.9), low physical activity (29.77% vs. 26.60%), and poor glycemic control (61.28% vs. 24.80%) were found in early-onset T2DM patients.

Results of PheWAS

The median follow-up times for early-onset and late-onset T2DM patients were 17.83 and 9.39 years, respectively. Except for diabetes-specific complications, early-onset T2DM was also significantly associated with increased risk of subsequent diabetes-related complications in multiple systems of the body (Fig. 2 and Supplementary Table 4, see online supplementary material). Compared with typical T2DM, the top three involved high-risks systems in order were sense organs (HR: 2.86 vs. 1.59), endocrine/metabolic (HR: 3.09 vs. 2.01), and respiratory system (HR: 1.73 vs. 1.49), respectively. Among 413 medical conditions widely distributed in all systems of the body, a total of 181 and 142 passed the 1% prevalence threshold. PheWAS indicated up to 101 diabetes-specific or -related complications of various systems among early-onset T2DM patients and 129 among late-onset T2DM patients (Supplementary Table 5, see online supplementary material), although most of the complications were significantly associated with both types of T2DM (Fig. 3). The overall difference in association magnitude (i.e. between matched cohorts for early-onset and late-onset T2DM) was 44.36%. Notably, we observed substantial differences for complications specific to diabetes (overall magnitude difference: 131.80% for microvascular and 125.42% for macrovascular complications; see HRs for specific complications in Supplementary Table 5).

Table 1. Basic characteristics of the individuals. The values are reported as median (lower quantile–upper quantile) for continuous variables or number (%) for categorical variables.

Characteristics	Matched cohort for early-onset T2DM		Matched cohort for late-onset T2DM	
	Diabetes patients (n = 1 777)	Matched individuals ^b (n = 17 761)	Diabetes patients (n = 35 889)	Matched individuals ^b (n = 179 437)
Age at index date, years	42.76 (40.51–44.41)	42.82 (40.50–44.49)	62.02 (56.12–67.52)	62.01 (56.08–67.50)
Townsend deprivation index ^a	0.95 (–1.96–3.62)	0.80 (–2.06–3.25)	–1.26 (–3.17–2.01)	–1.26 (–3.18–1.86)
Follow-up time, years	17.83 (13.89–22.06)	17.74 (13.55–22.29)	9.39 (5.90–13.76)	9.27 (5.92–13.66)
Sex				
Female	719 (40.46%)	7 190 (40.48%)	14 241 (39.68%)	71 203 (39.68%)
Male	1 058 (59.54%)	10 571 (59.52%)	21 648 (60.32%)	108 234 (60.32%)
Body mass index (kg/m ²)				
<24.1	119 (6.70%)	4 460 (25.11%)	2 105 (5.87%)	38 954 (21.71%)
24.1–29.9	516 (29.04%)	8 597 (48.40%)	12 913 (35.98%)	94 368 (52.59%)
≥29.9	1 090 (61.34%)	4 539 (25.56%)	20 408 (56.86%)	44 927 (25.04%)
Unknown	52 (2.93%)	165 (0.93%)	463 (1.29%)	1188 (0.66%)
Smoking status				
Ever	751 (42.26%)	7 976 (44.91%)	19 781 (55.12%)	89 724 (50.00%)
Never	998 (56.16%)	9 672 (54.46%)	15 714 (43.79%)	88 492 (49.32%)
Unknown	28 (1.58%)	113 (0.64%)	394 (1.10%)	1 221 (0.68%)
Drinking status				
Ever	1 485 (83.57%)	16 688 (93.96%)	32 677 (91.05%)	171 408 (95.53%)
Never	272 (15.31%)	968 (5.45%)	2 973 (8.28%)	7 354 (4.10%)
Unknown	20 (1.13%)	105 (0.59%)	239 (0.67%)	675 (0.38%)
Total physical activity ^c				
Low	529 (29.77%)	3 562 (20.06%)	9 547 (26.60%)	34 692 (19.33%)
Normal or high	820 (46.15%)	10 873 (61.22%)	17 604 (49.05%)	108 265 (60.34%)
Unknown	428 (24.09%)	3 326 (18.73%)	8 738 (24.35%)	36 480 (20.33%)
Fruit/vegetable consumption ^d				
Inadequate	1 206 (67.87%)	13 030 (73.36%)	24 071 (67.07%)	121 620 (67.78%)
Adequate	546 (30.73%)	4 619 (26.01%)	11 574 (32.25%)	57 132 (31.84%)
Unknown	25 (1.41%)	112 (0.63%)	244 (0.68%)	685 (0.38%)
Household income (£)				
<18 000	539 (30.33%)	3 255 (18.33%)	11 199 (31.20%)	43 494 (24.24%)
18 000–52 000	666 (37.48%)	7 644 (43.04%)	13 945 (38.86%)	77 070 (42.95%)
>52 000	234 (13.17%)	4 612 (25.97%)	3 779 (10.53%)	29 796 (16.61%)
Unknown	338 (19.02%)	2 250 (12.67%)	6 966 (19.41%)	29 077 (16.20%)
Glycemic control ^e				
Poor	1 089 (61.28%)	NA ^f	8 901 (24.80%)	NA
Good	545 (30.67%)		11 622 (32.38%)	
Unknown	143 (8.05%)		15 366 (42.82%)	

^aTownsend deprivation index was assigned to each individual based on their postcode location; a greater index score implies a greater degree of deprivation.

^bUp to 5 (for early-onset T2DM) or 10 (for late-onset T2DM) individuals free of T2DM diagnosis at the index date were randomly selected and individually matched to each patient by date of birth (within 1 year), sex, and decile of Townsend deprivation index.

^cTotal physical activity amount was calculated by summing the MET weighted time spent in vigorous, moderate, and walking activity, while low physical activity was defined as total physical activity amount ≤ 798 MET.min/week.

^dInadequate fruit/vegetable consumption was defined as eating <5 portions of fruit and vegetables per day.

^eHbA_{1c} levels were retrieved from UK Biobank baseline assessment and primary care data (using codes listed in Supplementary Table 3), and only HbA_{1c} measurements conducted at least 6 months after T2DM diagnosis were used to calculate each T2DM patient's median HbA_{1c} level; T2DM patients were further divided into good glycemic control (≤ 53 mmol/mol) group and poor glycemic control (>53 mmol/mol) group according to their median HbA_{1c} levels.

^fNA: Not available

Comorbidity network diversity

Among 9 870 possible comorbidity pairs, 1 025 and 479 were involved in the comorbidity network for early-onset and late-onset T2DM, respectively, after filtering based on prevalence, RR, and ϕ -correlation (Fig. 4). With 28 additional nodes and 605 additional links exclusive to early-onset T2DM (Supplementary Fig. 2, see online supplementary material), the comorbidity network for early-onset T2DM was more complex and tightly connected than that for late-onset T2DM. The modularity analysis indicated that the two networks shared similar comorbidity patterns, consisting of modules predominated by macrovascular complications (Fig. 4, dark yellow nodes), musculoskeletal (pink nodes), and digestive (brown nodes). However, early-onset T2DM was associated with increased interlinks and complications in each module and the

presence of converged modules [modules predominated by genitourinary (light brown nodes), and microvascular complications (light pink nodes)].

Sub-analyses and sensitivity analyses

Analysis by glycemic control condition showed significant reductions in risk of subsequent complications that were associated with fair glycemic control in both matched cohorts. As such the reduction was more pronounced for late-onset (pooled HRs 1.98–1.62, 18.01% reduction) than early-onset (pooled HRs 2.67–2.33, 12.67% reduction) T2DM, and a bigger magnitude difference was seen in the fair glycemic control group than in the poor glycemic control group (43.47% vs. 34.71%, Supplementary Fig. 3, see online supplementary material). Furthermore, the sub-analyses

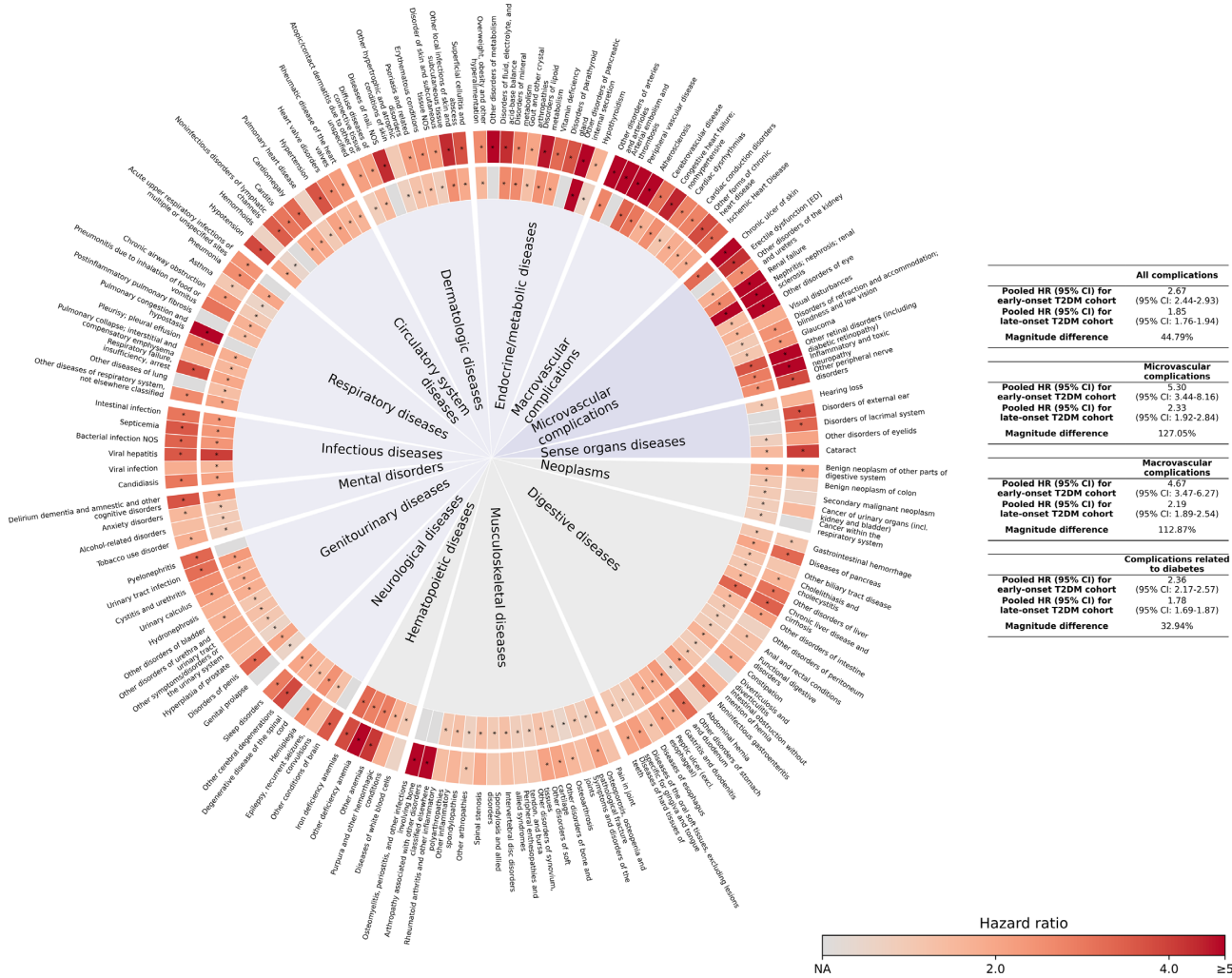


Figure 3. Risks of complications after T2DM diagnosis. The outer and inner rings show the point estimates of HRs of the corresponding complications for early-onset and late-onset T2DM respectively. Cox models were stratified by matching identifier (i.e. date of birth, sex, and Townsend deprivation index) and adjusted for household income, BMI, smoking and drinking status. Complications with P-value < Bonferroni corrected threshold are marked with an asterisk. Detailed results are shown in Supplementary Table 5.

adequate fruit/vegetable intake were significantly associated with a decreased risk of subsequent complications among early-onset T2DM patients, and were at comparable levels to those for late-onset T2DM. Therefore, we initially advocated a more aggressive glycemic control strategy for early-onset T2DM. Our data also validated that remission of early-onset T2DM is possible through weight management. Notably, due to the high prevalence of obesity in adolescents and young adults, the incidence of early-onset T2DM is expected to increase dramatically in the coming decades. Therefore, clinical guidelines focusing specifically on the treatment of early-onset T2DM and prevention of its complications are urgently needed.

The strengths of the current study include the use of a large community-based cohort to construct matched cohorts for early-onset and late-onset T2DM separately and long-term follow-up for all the subsequent complications through complete linkage to inpatient data for the UK Biobank participants since 1997. In addition, the availability of enriched data on sociodemographic information, lifestyle factors, and GP visits enabled consideration of many vital confounders in the model and further sub-analyses by modifiable lifestyles. Nevertheless, caution is needed in interpreting our results. The primary concern of the present study was that

the observed difference in the RR of subsequent complications after early-onset and late-onset T2DM could be mainly attributed to poor glycemic control status or longer follow-up time in early-onset T2DM patients. To relieve such a concern, we conducted the sub-analysis among T2DM patients with the same glycemic control status and sensitivity analysis, restricting the follow-up period to the first 10 years, and the results were generally comparable with the main results. Another concern was that the individuals included in the exposed groups could be patients with other types of DM that were misclassified as T2DM, as the ascertainment of T2DM was based solely on register data. To address this concern, we used a validated algorithm to exclude probable T1DM patients. In the sensitivity analysis, we further excluded T2DM patients with any documented diagnosis of other DM types or with only one documented diagnosis of T2DM, and such alteration did not change the main results substantially. Although many important confounders, such as sociodemographic factors, lifestyle factors, and comorbidity have been considered in the analyses, residual confounding due to undiagnosed comorbidities or other unmeasured confounders could have biased the study's results to some extent. Meanwhile, as the primary care data only covered half of the UK Biobank participants, and inpatient data were only

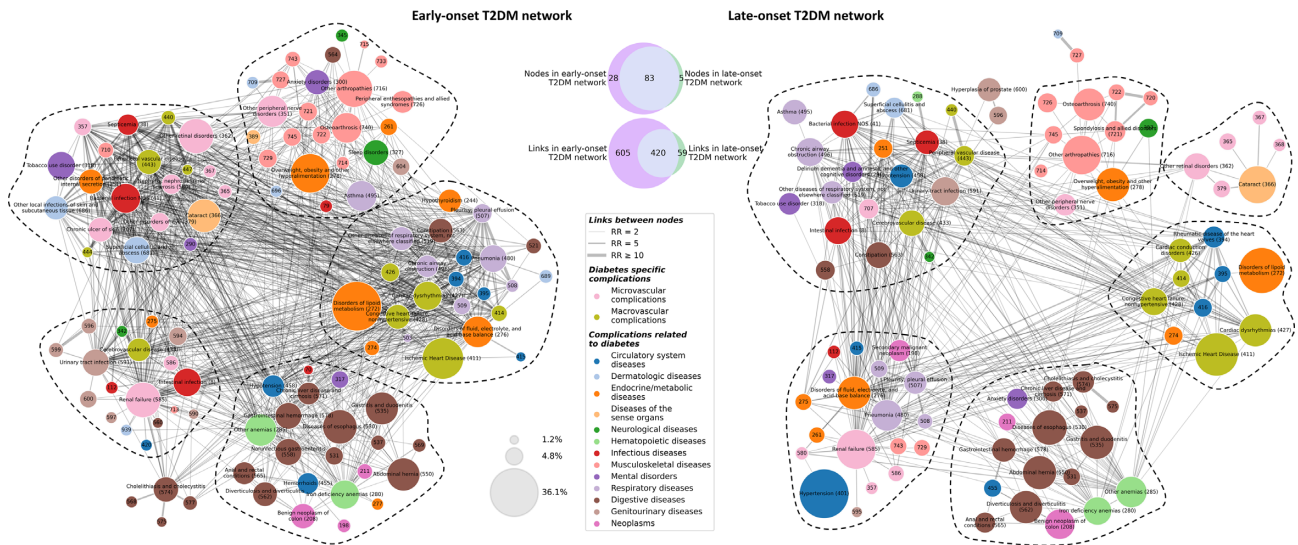


Figure 4. Comorbidity networks for early-onset and late-onset T2DM. The number of nodes and links in each network and their logical relation are shown in the Venn diagrams in the upper center of the figure. Each node represents a diabetes-associated complication, while the size and color of the node indicate the prevalence and the category of the corresponding complication (see legend). Reference nodes with percentages representing the prevalence scales are shown in the lower center of the figure. The width of the link represents the strength of comorbidity association, measured by RR. Reference links with RR are shown in the legend. Complications with prevalence $\geq 5\%$ are labeled with its phenotype name along with the 'phecode', while others are labeled only with the 'phecode' (see Supplementary Table 2 for a description of each 'phecode'). The two networks were partitioned into several modules using the Louvain algorithm, and nodes belonging to the same module are grouped together and separated from other nodes using dashed lines.

available from 1997 onwards, the identified time of first T2DM diagnosis could be later than the actual time of onset, resulting in some patients with early-onset T2DM being misclassified as late-onset T2DM patients. Other limitations of the current study include that some important variables used for matching or model adjustment (e.g. Townsend deprivation index, smoking, and drinking status) were measured at the UK Biobank study baseline and may not reflect the individuals' status at the same time as the time of T2DM diagnosis. Additionally, the small number of early-onset T2DM patients leads to a limited statistical power to detect a significantly elevated HR for some of the studied complications in the early-onset cohort, especially in the further sub-analyses. Finally, the generalizability of the study results to a broader population was limited by the inherent flaws of the UK Biobank study, including low response rate and oversampling of the White population.³⁵

Taken as a whole, we systemically reviewed the health consequences of early-onset T2DM and compared the results with those of late-onset T2DM, based on matched cohorts in the UK Biobank. Our data reveal that early-onset T2DM is more challenging in terms of glycemic control, with a higher risk of severe complications. Therefore, we advocate that guidelines for the intensive management of complications and glycemic control criteria are urgently needed.

Supplementary data

Supplementary materials are available at *PCMEDJ Journal* online.

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Author contributions

H.S. and D.L. are responsible for the study's concept and design. W.C., Y.Z., and Y.H. carried out data and project management. C.H., H.Y., and Y.Q. performed data cleaning and analysis. C.H., H.Y., Y.Q., H.S., and D.L. interpreted the data. C.H., K.M.V.N., H.S., and D.L. drafted the manuscript. All the authors approved the final manuscript as submitted and agree to be accountable for all aspects of the work.

Conflict of interest

None declared.

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