ARTICLE



Clinically meaningful and lasting HbA_{1c} improvement rarely occurs after 5 years of type 1 diabetes: an argument for early, targeted and aggressive intervention following diagnosis

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Abstract

Aims/hypothesis Our objectives were to explore whether the phenomenon of HbA_{1c} 'tracking' occurs in individuals with type 1 diabetes, how long after diagnosis does tracking take to stabilise, and whether there is an effect of sex and age at diagnosis on tracking.

Methods A total of 4525 individuals diagnosed with type 1 diabetes between 1 January 1995 and 1 May 2015 were identified from The Health Improvement Network (THIN) database. Mixed models were applied to assess the variability of HbA_{1c} levels over time with random effects on general practices (primary care units) and individuals within practices.

Results 4525 individuals diagnosed with type 1 diabetes were identified in THIN over the study period. The greatest difference in mean HbA_{1c} measurement (-7.0 [95% CI -8.0, -6.1] mmol/mol [0.6%]) was seen when comparing measurements made immediately after diagnosis (0–1 year since diagnosis) with those at 10 or more years (the reference category). The mean difference in HbA_{1c} for the successive periods compared with 10 or more years after diagnosis declined and was no longer statistically significant after 5 years. In the stratified analysis using sex and age group there was considerable heterogeneity with adult onset type 1 diabetes appearing to track earlier and at a lower mean HbA_{1c}.

Conclusions/interpretation In individuals with type 1 diabetes, glycaemic control measured by HbA_{1c} settles onto a long-term 'track' and this occurs on average by 5 years following diagnosis. Age at diagnosis modifies both the rate at which individuals settle into their track and the absolute HbA_{1c} tracking level for the next 10 years.

Keywords Glycated haemoglobin \cdot HbA_{1c} \cdot Tracking \cdot Type 1 diabetes mellitus

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Abbreviation

THIN The Health Improvement Network

Introduction

Clinical practice suggests that HbA_{1c} remains remarkably stable in individuals with type 1 diabetes. Some individuals are consistently able to achieve good glycaemic control at repeated clinical follow-up, while others struggle to do so for any meaningful period of time. In these latter individuals, life events (such as pregnancy) or planned interventions (including structured education or continuous subcutaneous insulin infusion therapy) associate with improvements in HbA_{1c} . However, this improved HbA_{1c} is often not maintained beyond a few years [1, 2]. This stability of HbA_{1c} that can



Research in context

What is already known about this subject?

Glycaemic control, measured by HbA_{1c}, may settle into a long-term 'track'

What is the key question?

• Does the phenomenon of HbA_{1c} 'tracking' occur in individuals with type 1 diabetes, how long after diagnosis does tracking take to stabilise and what is the effect of sex and age at diagnosis on tracking?

What are the new findings?

- This is the first study to demonstrate the phenomenon of glycaemic tracking in a large number of newly diagnosed individuals with type 1 diabetes across a broad age range and over a long period of follow-up
- Glycaemic tracking occurs on average at 4–5 years following diagnosis
- The age of diagnosis modifies both the rate at which individuals settle into their track and the absolute HbA_{1c} tracking level for the next 10 years

How might this impact on clinical practice in the foreseeable future?

There is a 4–5 year window during which long-term HbA_{1c} level and therefore risk of diabetes complications is
determined. Thus, urgent and appropriate targeting of therapies for this early period of type 1 diabetes should be
considered

manifest over decades, or even a lifetime, has been referred to as glycaemic 'tracking' [3].

Glycaemic tracking is distinct from HbA_{1c} variability. Year to year variability in HbA_{1c} is a characteristic described in individuals with pre-existing diabetes and associates with both micro- and macrovascular disease [4, 5]. Glycaemic tracking is also not simply the inverse of glucose variability, which is the daily change in blood glucose that includes symptomatic hypo- and hyperglycaemia.

It is vitally important to explore the phenomenon of glycaemic tracking. If tracking is an inherent part of the natural history of type 1 diabetes, this would emphasise the importance of early metabolic control after diagnosis. There may therefore be a window of opportunity where focused interventions set the scene for long-term glycaemic control and facilitate cost effective allocation of time and resources. Exploring the phenomenon of glycaemic tracking may also provide insights into underlying mechanisms, and from there, testable approaches to influencing the final level of HbA_{1c} tracking.

Glycaemic tracking remains to be well characterised. Crucially, previous reports [3–18] have largely examined individuals with pre-existing type 1 diabetes (i.e. not from the time of diagnosis). Those few studies that have followed individuals from the time of diagnosis have been of limited sample size and short duration, have examined individuals across a narrow age group (primarily paediatric) and have not adjusted for important confounders (e.g. socioeconomic status). These studies are summarised in the electronic supplementary material (ESM) Table 1 and illustrate the need to study the phenomenon of glycaemic tracking in a more robust manner.

To explore the concept of glycaemic tracking formally, we performed a large, UK population-based cohort study involving over 4000 individuals with newly diagnosed type 1 diabetes and over 50,000 longitudinally collected HbA_{1c} measurements. We wished to establish: (1) whether the phenomenon of HbA_{1c} tracking occurs in individuals with type 1 diabetes; (2) how long after diagnosis does tracking take to stabilise; and (3) the effect of sex and age of diagnosis at tracking.

Methods

The Health Improvement Network (THIN) database is a large primary care dataset derived from general practices (primary care units) across the UK [19]. More than 675 practices contribute to the dataset resulting in over 14 million patient records of which around 4 million are active participants. The database consists of individuals' demographic details, diagnosis, prescriptions and laboratory results. The database is generalisable to the UK population, and has been utilised for numerous epidemiological studies, including type 1 diabetes [20].

The use of the THIN data for research was approved by the South-East Multicenter Research Ethics Committee in 2003, without the need for informed consent. As per the requirements for ethical approval, further registration and authorisation for this project were obtained from the Scientific Review Committee of the data provider (IQVIA: 17THIN015).

Population Individuals diagnosed with type 1 diabetes between 1 January 1990 and 1 May 2015 were identified using

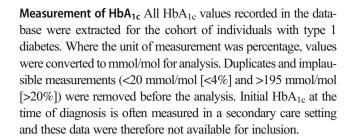


Table 1 Baseline characteristics of participants and HbA_{1c} summary statistics

Characteristics	All participants $N = 4525$	Participants contributing for 10 years and above $n = 938$
Age group (years)		
0–10	1294 ± 28.6	293 ± 31.3
10–20	1737 ± 38.4	324 ± 34.5
20–30	798 ± 17.6	138 ± 14.7
30–40	696 ± 15.4	183 ± 19.5
Sex		
Male	2743 ± 60.6	574 ± 61.2
Female	1782 ± 39.4	364 ± 38.8
Townsend index		
1	1020 ± 22.5	229 ± 24.4
2	869 ± 19.2	178 ± 19.0
3	923 ± 20.4	190 ± 20.3
4	828 ± 18.3	175 ± 18.7
5	650 ± 14.4	137 ± 14.6
Missing	235 ± 5.2	29 ± 3.0
Follow-up period (years)		
Mean (SD)	6.0 ± 4.6	13.2 ± 2.4
Median (25th–75th percentile)	5.0 (2.1–9.2)	13.2 (11.2–14.8)
HbA _{1c} (mmol/mol) ^a		
Mean (SD)	72.6 ± 20.6	73.7 ± 19.5
Median (25th-75th percentile)	69.4 (58.5–82.5)	70.5 (60.7–83.6)
HbA1c (%) ^a		
Mean (SD)	8.8 ± 4.0	8.9 ± 3.9
Median (25th–75th percentile)	8.5 (7.5–9.7)	8.6 (7.7–9.8)

Data presented as mean \pm SD or median (25th–75th percentile)

appropriate Read codes (https://digital.nhs.uk/article/1104/ Read-Codes) and based on an algorithm recently published by Sharma et al 2016 [21]. To be classified as having type 1 diabetes, participants need to have a type 1 diabetes specific Read code, be aged less than 40 years at diagnosis and have been prescribed insulin but not oral hypoglycaemic medications. Individuals were eligible to be included in the cohort if they were diagnosed at least 1 year after registration with the practice or a year after the practice was eligible to take part, whichever was the latest. The date of diagnosis used was the one recorded in the database. Participating general practices were eligible to take part following: (1) introduction of the computerised system; (2) the date practices were deemed as having acceptable mortality rates. This was important to ensure that practices were recording important information and comorbidities accurately.



Statistical analysis The analysis aimed to answer the three aforementioned questions: does HbA_{1c} tracking occur among individuals with type 1 diabetes; if tracking occurs, how soon after diagnosis does this manifest; if tracking occurs, what is the impact of sex and age at diagnosis on the natural history of tracking?

Sociodemographic characteristics, HbA_{1c} levels and time since diagnosis were summarised using descriptive statistics. Exploratory plots were produced for the mean levels of HbA_{1c} vs time since diagnosis of type 1 diabetes. Similar plots were also produced stratifying by 10 year age bands and sex. The time since diagnosis with type 1 diabetes represents the duration between the date when a particular HbA_{1c} measurement was taken and the initial date of diagnosis. This was divided into 11 categories: 0–1 years, 1–2 years, 2–3 years,..., 9–10 years and \geq 10 years to facilitate modelling because our exploratory analyses indicated that the association between HbA_{1c} level and time was likely to be non-linear.

Glycaemic tracking was defined as a period in which there was no statistically significant (p < 0.05) difference in HbA_{1c} across adjacent years in comparison with the HbA_{1c} measurement at ≥ 10 years from diagnosis.

We used linear mixed effects models to assess the variability of HbA_{1c} levels over time with random effects on practices and individuals within practices. These models, also known as multilevel or hierarchical linear models, constitute both fixed effects and random effects. The fixed part is similar to standard linear regression but the addition of random effects allows the model to account for the potential effect of variability at different grouping or clustering levels. In other words, the random effects are variance components associated with each level. In our study, there are three variance components, namely: (1) within individual variability (the residual error associated with repeated measurements from an individual), (2) between individuals within practice variation (as several individuals share the same practice), and (3) between practice variation. Individuals within the same practice may be correlated because of a shared random intercept, through a shared random slope on a covariate or both. Thus, both adjusted and unadjusted two-level random intercepts and slopes (for individuals within practices) models were fitted using the ≥10 years duration group as reference. The adjusted model included age at diagnosis (10 year bands), sex and the Townsend index.



^a Number of measurements in study period: 41,583 for all participants group, 16,989 for participants contributing for 10 years and above

The Townsend index, ranging from 1 to 5, is a measure of material deprivation, calculated using social indices such as income, education and employment specific to each participant's neighbourhood [22, 23]. Socioeconomic status has been convincingly related to glycaemic control [24] and the Townsend index has been used extensively as a covariate in diabetes studies using THIN database [25–27]. In the adjusted model, the group with the longest duration of type 1 diabetes (≥10 years) was preferred as the reference category; this is because if individuals were to track, i.e., if the proposed hypothesis was true, then we would expect to see no significant difference between this group and groups with a lower duration of type 1 diabetes. The duration after which there is no significant difference in the final HbA_{1c} will be the time point at which tracking manifests. In addition to this, a stratified analysis will be conducted using sex and age group as stratification factors.

In a sensitivity analysis, the random intercepts and slopes model were fitted including data only from individuals with 10 or more complete years of follow-up. All analyses were performed using Stata 14 (StataCorp, College Station, TX, USA).

Results

There were 4525 individuals diagnosed with type 1 diabetes in the THIN database over the study period 1995–2015 from a total of 617 practices. The majority were male (60.6%) and the highest proportion (38.4%) were diagnosed between the ages of 10 and 20 years. The mean \pm SD HbA $_{1c}$ level was 72.6 \pm 20.6 mmol/mol (8.8 \pm 4%) and the median follow-up after diagnosis was 5 years (Table 1).

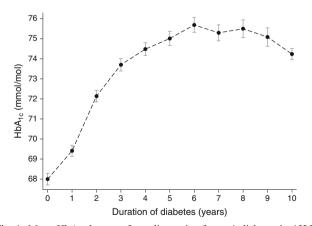


Fig. 1 Mean HbA $_{1c}$ by year from diagnosis of type 1 diabetes in 4525 patients with newly diagnosed diabetes. For example, duration time 0 represents the HbA $_{1c}$ values captured from date of diagnosis to year 1; time 1 represents the HbA $_{1c}$ measurements from year 1 to 2, and time 9 represents HbA1c measurements from year 9 to 10

HbA_{1c} increased with time from diagnosis in individuals with type 1 diabetes and stabilised by 5 years to an average of 75.0 mmol/mol (9.0%) following this period (Fig. 1). However, the timing of stabilisation was dependent on age at diagnosis and sex (ESM Fig. 1).

The largest difference between mean HbA_{1c} measurements was between those taken in the first year following diagnosis (time: 0 years in Fig. 2) and those taken after 10 or more years (the reference category) (-7.0 [95% CI -8.0, -6.1] mmol/mol [0.6%]; Fig. 2). The mean HbA_{1c} difference for the successive periods after diagnosis (1–2 years, 2–3 years, etc.) compared with after 10 or more years declined considerably and was no longer statistically significant 5 years following diagnosis (a duration time of 4 years in figures illustrates HbA_{1c} s measured between 4 and 5 years after diagnosis); the mean HbA_{1c} difference for this duration after diagnosis was -0.8 (95% CI -1.8, 0.2) mmol/mol (0.7%). The findings remained similar when analysis was restricted only to participants contributing HbA_{1c} measurements for 10 years and above (ESM Fig. 2).

There was considerable heterogeneity in the timing of tracking in the stratified analysis using sex and age group. Adult onset type 1 diabetes seem to track earlier (Fig. 3) and at a lower mean HbA_{1c} (ESM Fig. 1). Children diagnosed between 0 and 10 years tracked from 6–7 years and 8–9 years after diagnosis for boys and girls, respectively (Fig. 3a, e). For the 10–20 years diagnosis age group, tracking occurred at 9–10 years in males and 7–8 years in females (Fig. 3b, f). For the 20–30 years group, tracking occurred at 2–3 years for both sexes (Fig. 3c, g). Among the 30–40 years age group, tracking was at 4–5 years in men (Fig. 3d) and 5–6 years in women (Fig. 3h).

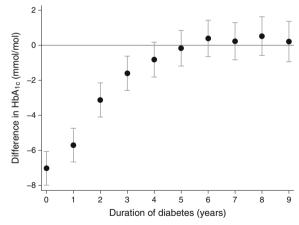
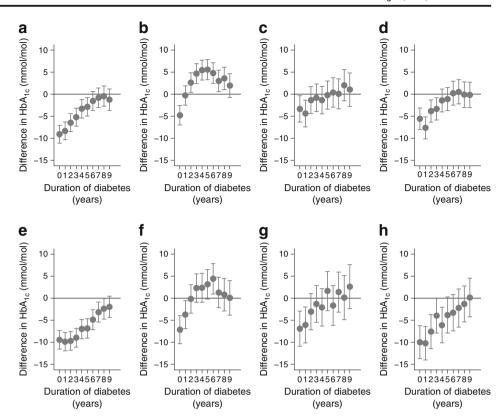


Fig. 2 Mean (95% CI) difference in HbA_{1c} between the mean for the reference group (≥10 years post diagnosis) and the mean for each year after diagnosis. Duration time 0 represents the HbA_{1c} values captured from date of diagnosis to year 1; time 1 represents the HbA_{1c} measurements from year 1 to 2, etc. Models were constructed using a random intercept and slopes model adjusting for age, sex and Townsend index. The time point at which the 95% CI for the yearly difference crosses the null value (zero) is considered the starting point of tracking



Fig. 3 Difference in HbA_{1c} between the reference group (≥10 years) and each year after the time of diagnosis of type 1 diabetes stratified by age at diagnosis and sex. Duration time 0 represents the HbA_{1c} values captured from date of diagnosis to year 1; time 1 represents the HbA_{1c} measurements from year 1 to 2, etc. (a, e) Diagnosis between ages 0 and 10 years; (b, f) between ages 10 and 20 years; (c, g) between ages 20 and 30 years; (d, h) between ages 30 and 40 years; male group (a-d) and female group (e-h). Models were constructed using a random intercept and slopes model adjusting for age, sex and Townsend index. The difference is given for each year with a point estimate and its 95% CI. The time point where the above difference crosses the null value (zero) is considered the starting point of tracking



Discussion

We show that in individuals with type 1 diabetes, glycaemic control measured by HbA_{1c} settles onto a long-term 'track' and that this occurs on average by 5 years following diagnosis. We also show that the age at diagnosis modifies both the rate at which individuals settle into their track and the absolute HbA_{1c} tracking level for the next 10 years. Our analytical approach utilising a random intercept and slope model ensures that our overall results are not confounded by trajectories at an individual level.

This is the first study to demonstrate the phenomenon of glycaemic tracking on a large number of unselected, newly diagnosed individuals with type 1 diabetes across a broad age group and with a long period of follow-up. Crucially, this study is unique in that the association has been adjusted for important potential confounders such as local clinical practice and Townsend score. The study is weakened by the fact that adequate follow-up data are not available for greater than 10 years and that the diagnosis of type 1 diabetes was taken prima facie from the national THIN general practice database. For this reason, we limited the analysis to those diagnosed with type 1 diabetes under the age of 40 years, on insulin alone and with no history of oral hypoglycaemic prescriptions. Finally, the HbA_{1c} assay has evolved over the course of the time span during which this study extends. Although appropriate conversions were meticulously undertaken, inherent differences in methodologies should be taken into account

when interpreting differences in HbA_{1c} before and after 2009 in the UK. Similarly, several important changes in diabetes management have been introduced over the study period such as the introduction of novel insulin formulations, widespread adoption of intensive insulin treatment schemes and revision of education systems. These may have contributed towards an improvement of glycaemic control at a national level and although data is missing, they should be considered in the context of this study.

It is notable that the overall HbA_{1c} levels in the UK are suboptimal. This has been seen before with regard to other countries [28]. Furthermore there is a discrepancy between the sexes and this may relate to the higher insulin resistance [29] and behavioural issues [30] that may be more prevalent in younger women and girls with type 1 diabetes.

Several hypotheses, focusing on endogenous and/or exogenous factors, have been suggested to underlie the phenomenon of glycaemic tracking. Residual beta cell function, measured by stimulated C-peptide, decays with time following type 1 diabetes diagnosis [31] and low C-peptide is associated with higher HbA_{1c} [32, 33]. Therefore, the loss of C-peptide following diagnosis may explain the rising HbA_{1c} in the initial 5 years. Alternatively, or in addition, individual patient-related factors may contribute. Habituation of the day to day approach to managing chronic disease can make any long-lasting change difficult [34]. Studies involving paediatric and adolescent individuals with type 1 diabetes indicate that age, sex, body mass index, socioeconomic factors, physical activity levels,



frequency of glucose monitoring and personality traits can also influence temporal HbA_{1c} trends [16, 35–37]. It may be that a combination of these endogenous and exogenous factors contributes to the phenomenon of tracking and that both need to be addressed for effective long-term glycaemic control.

There are two clear implications of our results. First, there is a 5 year window during which longer-term HbA_{1c} and therefore risk of diabetes complications is determined. Thus, urgent and appropriate targeting of therapies to this period of type 1 diabetes natural history should be considered. This may include, for example, the development of a newly diagnosed type 1 diabetes pathway with structured education and aggressive glucose control. Second, the benefits of addressing HbA_{1c} after the first 5 years should be explored. It has been suggested that efforts would be better directed at risk factors other than HbA_{1c} in individuals with established type 1 diabetes [13]. However, such an approach may result in loss of the tracking phenomenon and deterioration in HbA_{1c} so this needs to be carefully explored. Unfortunately, data are missing with regards to both effectiveness and timing of a focused clinical intervention targeted at changing the track and this clinical need becomes even more evident in light of the study findings. Furthermore, it is worth noting that HbA_{1c} provides only one indication of glycaemic control. Day to day glucose variability is of greater concern to individuals and is poorly reflected in the HbA_{1c} measure. Indeed, 'real world' experience of new interventions such as structured education and insulin pump therapy suggest they have had a greater benefit on glucose variability than meaningful long-term changes to HbA_{1c} [38, 39].

This is the first comprehensive study of the phenomenon of HbA_{1c} tracking and supports the need to optimise glycaemic control early in the natural history of type 1 diabetes. Studies are now needed to explore the mechanisms underlying this phenomenon and how best to optimise tracking in newly diagnosed individuals.

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Data availability THIN data governance does not allow us to share individual patient data and therefore where possible metadata is presented. Researchers may apply for individual patient data access at https://www.iqvia.com/contact.

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1070 Diabetologia (2018) 61:1064–1070

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