

Precision Diabetes Is Slowly Becoming a Reality

Viswanathan Mohan^a Venkatesan Radha^b

^aDepartment of Diabetology, Madras Diabetes Research Foundation and Dr. Mohan's Diabetes Specialities Centre, Chennai, India; ^bDepartment of Molecular Genetics, Madras Diabetes Research Foundation, Chennai, India

Significance of the Study

- Till recently, a “one-size-fits-all” approach was used to treat most forms of diabetes. With the availability of big data and analytic tools like genomics, epigenetics, metabolomics and proteomics, precision diabetes is now slowly becoming a reality. Precision diabetes is already being applied to monogenic forms of diabetes like maturity-onset diabetes of the young, neonatal diabetes mellitus and congenital hyperinsulinaemic hypoglycaemia. In many such cases, this can be a life changer, as insulin can be stopped completely and be replaced with oral glucose-lowering agents. In other forms of diabetes like type 2 diabetes, precision diabetes promises to be of use in personalised therapy for people affected with the disorder.

Keywords

Precision medicine · Personalised medicine · Personalised diabetes · Precision diabetes · Monogenic diabetes · Neonatal diabetes · Maturity-onset diabetes of the young

Abstract

The concept of precision medicine is becoming increasingly popular. The use of big data, genomics and other “omics” like metabolomics, proteomics and transcriptomics could make the dream of personalised medicine become a reality in the near future. As far as polygenic forms of diabetes like type 2 and type 1 diabetes are concerned, interesting leads are emerging, but precision diabetes is still in its infancy. However, with regard to monogenic forms of diabetes like maturity-onset diabetes of the young and neonatal diabetes mellitus, rapid strides have been made and precision diabetes

has already become part of the clinical tools used at advanced diabetes centres. In patients with some monogenic form of diabetes, if the appropriate gene defects are identified, insulin injections can be stopped and be replaced by oral sulphonylurea drugs. In the coming years, rapid advances can be expected in the field of precision diabetes, thereby making the control of diabetes more effective and hopefully leading to prevention of its complications and improvement of the quality of life of people afflicted with diabetes.

© 2019 The Author(s)
Published by S. Karger AG, Basel

Introduction

The National Institutes of Health (NIH), USA, define precision medicine as an emerging approach to disease treatment and prevention that takes into account

individual variability in genes, environment and lifestyle [1]. This concept is not new, and in the past, patients were prescribed optimal treatment by clinicians based on the available data and resources. In its crudest sense, conducting blood transfusion based on the specific blood group of a recipient is an example of precision medicine [2]. The crux of precision medicine is custom delivery of healthcare, with tailored treatment at the level of the individual patient. However, in spite of extensive epidemiological and physiological characterisation and elucidating pathophysiological pathways, we are still far away from predicting prognoses, or even selecting the most effective medicines for a given patient. Most therapeutic approaches are based on population averages, which unfortunately do not fit everyone [3].

Since these current methods of disease categorisation are outmoded and have not achieved their objectives, a new concept has arisen in the recent years. It is recognised that individuals who share the same diagnostic labels may have very heterogeneous presentations and clinical courses and may respond differently to therapeutic interventions. This has led to the concept of personalised medicine, which, when further refined by scientific inputs, becomes precision medicine [4].

Precision medicine refers to collecting specific information about a patient and prescribing a specific treatment. This involves defining disease subtypes and biomarkers that can identify patients who are likely to benefit from a specific treatment or those who are likely to experience a particular side effect. Application of genotypic and phenotypic information forms the basis of precision medicine [5]. Personalised medicine not only incorporates the principles of precision medicine but also takes into account patients' preferences and social context, as well as other factors such as affordability of therapies.

Big Data and "Omics" Technologies in Precision Medicine

One of the important reasons for the rapid development of precision medicine in recent years is the availability of big data and the advancement of technologies to mine these data [6, 7]. It is believed that traditional sources of medical information, such as patient history, physical examination and laboratory workup, when augmented by data mining of deep phenotypic data and genomic analyses coupled with epigenetic, metabolomic, pro-

teomic and transcriptomic data, can help deliver precision medicine to individual patients [4, 8].

A plethora of transformative and disruptive technologies such as DNA sequencing, functional investigation, the emerging field of epigenetics, genome-wide association studies and miRNA scans, as well as a range of "omic" measurements, can help in the detailed *in vivo* dissection of physiology and pathology, and this improves diagnosis and treatment. They can also provide the basis for refinement of disease subtypes and support the effective optimisation of disease management to match an individual pathology or genetic make-up [9]. The field of oncology has been one of the first to adopt precision medicine, but other fields, including endocrinology, are catching up fast.

Precision Diabetes

Diabetes mellitus broadly encompasses four categories, namely, type 1 diabetes mellitus (T1DM), type 2 diabetes mellitus (T2DM), monogenic diabetes and gestational diabetes.

T1DM is an autoimmune disease with progressive destruction of β -cells, leading to polyuria, polydipsia, weight loss and hyperglycaemia. T1DM requires lifelong administration of insulin injections several times a day as endogenous insulin secretion becomes near zero. Making a proper diagnosis of T1DM with clear evidence of β -cell destruction through the presence of islet cell autoantibodies and demonstration of absent β -cell function as measured by levels of C peptide and prescribing the appropriate doses of insulin are the basis for precision medicine in the case of T1DM, although some heterogeneity exists even here.

T2DM is a heterogeneous group of disorders predominantly due to insulin resistance and some degree of an insulin secretion defect. T2DM has a range of risk factors, aetiologies and clinical presentations.

There are also several forms of monogenic diabetes, which are caused by highly penetrant single gene defects; the three major types maturity-onset diabetes of the young (MODY), neonatal diabetes mellitus (NDM) and congenital hyperinsulinaemic hypoglycaemia (CHH) are the most common. It is with these forms of diabetes that precision diabetes has already come to be used in the clinical setting [10].

This review will first describe the role of precision medicine in T2DM and then deal with the monogenic forms of diabetes.

Precision Diabetes in T2DM

Defining subgroups using molecular testing is much more difficult in T2DM, as it is a polygenic disorder and the clinical phenotype reflects environmental as well as genetic influences [11]. One approach for precision medicine in T2DM is to categorise patients based on differential responses to drugs, and to then investigate the molecular underpinnings of each subgroup using next-generation sequencing platforms and gene arrays. In clinical practice this approach is not easy to implement, although pharmacogenomic developments continue to be made [12].

In contrast to monogenic diabetes, where the gene mutation is causal and has a high phenotypic correlation, T2DM involves multiple, common, low-impact risk variants. The emerging availability of genomic and electronic health data from large populations, the so-called big data, is a powerful tool for bringing precision medicine to T2DM [13, 14]. For example, recently, at least five clusters of T2DM with differing clinical features have been identified [15].

A common nonsense mutation in the *TBC1D4* gene that substantially increases the risk of T2DM was recently identified in a Greenlandic Inuit population [16]. This variant is shown to exclusively increase postprandial glucose levels. The frequency and effect of the *TBC1D4* mutation on glucose metabolism and T2DM were determined in two related populations: Canadian Inuit and Alaskan Inuit [17]. The mutation in the *TBC1D4* gene is common among North American Inuit, and results in elevated postprandial glucose levels and an underdiagnosis of T2DM unless an oral glucose tolerance test is performed. This is one example of how precision medicine may be applied at the population level.

Some pharmacogenomic advances have been made in relation to antidiabetic drugs. Variations in the cytochrome P450 (CYP) enzymes contribute to oral antidiabetic drug metabolism in the liver and affect drug disposition and efficacy. Genetic variants in *CYP2C9*2* (I359L) and *CYP2C9*3* (R114C) have been shown to be associated with reduced blood sulphonylurea (SU) clearance [18, 19]. Furthermore, variants in *CYP2C8* were found to influence the efficacy of repaglinide and rosiglitazone [20].

Various studies have shown that rs12208357, rs34130495, rs34059508 and rs72552763 are related to the *SLC22A1* gene, which encodes organic cation transporter 1, and are genetic markers for the efficacy and excretion of metformin [21–24].

The SNP rs11212617, which is located near the ATM locus, was found to be associated with HbA_{1c} levels in response to metformin in a large-scale genome-wide association study conducted in European T2DM populations [25]. One study has demonstrated that the *PAX4* variants rs6367136 and rs10229583 and the *PSMD6* variant rs831571 were correlated with the therapeutic efficacy of repaglinide and rosiglitazone in Chinese T2DM patients [26, 27].

Dujic et al. [28] demonstrated the impact of genetic factors on gastrointestinal tolerance to metformin. The Genetics of Diabetes Audit and Research in Tayside Scotland (GoDARTS) study [29] had also shown that reduced-function alleles of the *OCT1* gene are associated with increased intolerance to metformin. Recently, it has been suggested that serotonin reuptake transporter might also be involved in intestinal metformin absorption. The number of low-expressing serotonin reuptake transporter alleles significantly increased the odds of metformin tolerance. These results suggest that gastrointestinal side effects of metformin could be related to the reduced uptake of intestinal serotonin [23].

Another pharmacogenetic approach to treatment response in T2DM was demonstrated with the use of thiazolidinediones (TZDs), compounds that are transported into the liver by OATP1B1 (encoded by the *SLCO1B1* gene) and metabolised by the CYP450 2C8 enzyme (encoded by the *CYP2C8* gene). Although variants in the *CYP2C8* gene (the *CYP2C8*3* allele) have been shown to alter TZD pharmacokinetics, the *CYP2C8*3* allele has not been shown to alter its efficacy. In an elegant study [30], 833 patients with T2DM treated with pioglitazone/rosiglitazone were genotyped for *CYP2C8* and *SLCO1B1* functional variants. The *CYP2C8*3* variant was significantly associated with reduced glycaemic response to rosiglitazone and less weight gain, whereas the *SLCO1B1* 521T>C variant was associated with enhanced glycaemic response to rosiglitazone. Patients with both genotypes (super-responders) had a significantly greater HbA_{1c} reduction. Interestingly, neither of the variants had a significant impact on pioglitazone response. This highlights the importance of studying transporter and metabolising genes as a predictor of treatment response by identifying those individuals who can benefit from the therapeutic advantages of TZDs.

There are also reports showing that genetic polymorphisms, such as *TCF7L2* polymorphisms, may influence the response to dipeptidyl peptidase-4 (DPP-4) inhibitors [31–33]. There are also some studies suggesting that gene polymorphisms may mediate the response to GLP-1 (glu-

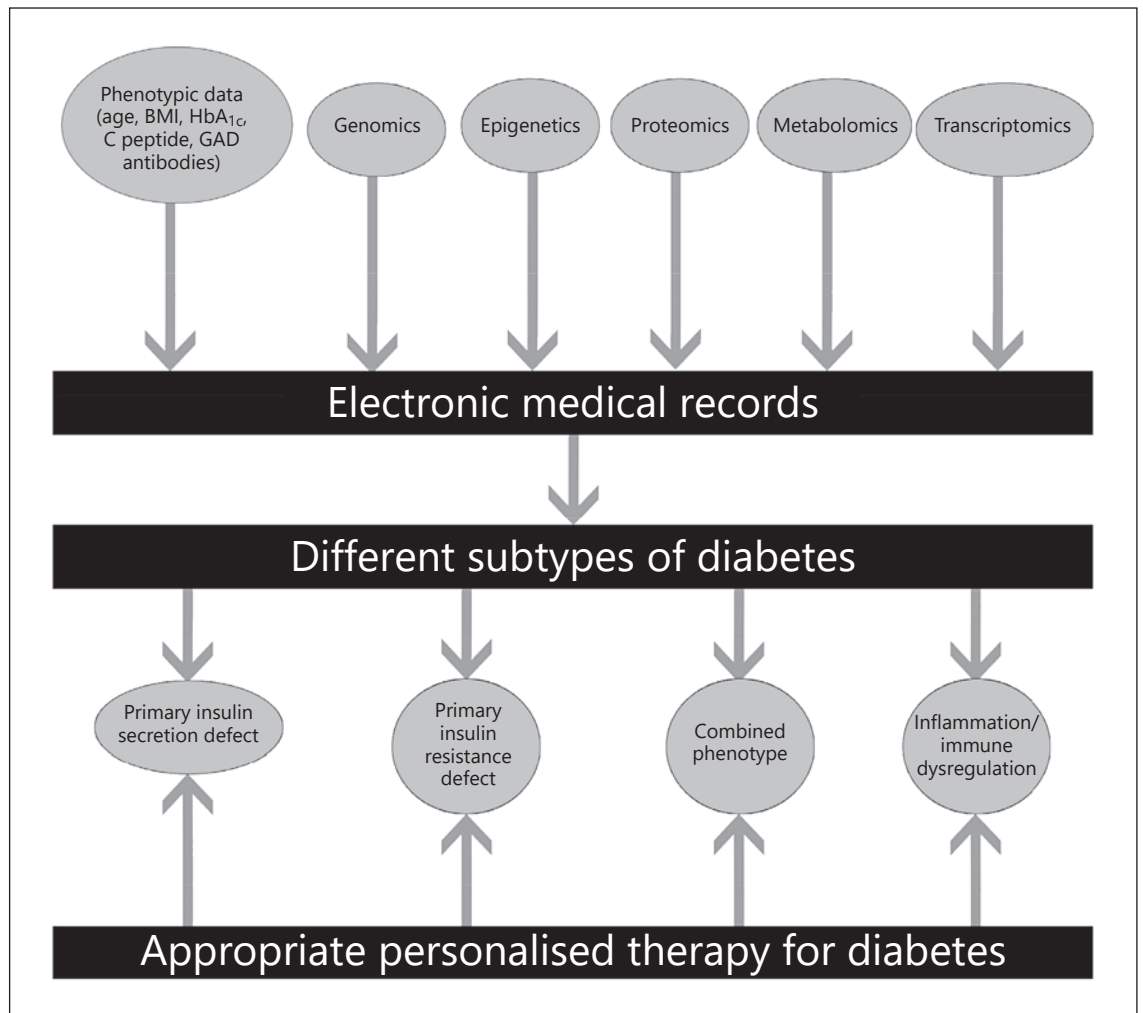


Fig. 1. Precision diabetes: a futuristic model. Adapted from Schwartz et al. [44].

cagon-like peptide-1) receptor agonists [34] and DPP-4 inhibitors [35].

A multitude of newer antidiabetic agents are now available. These include DPP-4 inhibitors, sodium/glucose cotransporter 2 (SGLT2) inhibitors and GLP-1 receptor agonists. It is now known that there are at least 8 pathophysiological defects in T2DM. These drugs help address several of the defects. It is also known that there could be ethnic differences in the response to these drugs; for example, the DPP-4 drugs seem to work better in Asians than in Europeans [36–40]. This could be because β -cell dysfunction is more severe in Asians, particularly in Indians [41]. Indeed, it has been shown that Asian Indians and Pima Indians represent two extreme phenotypes of T2DM, one leaner with more insulin deficiency

and the other obese with severe insulin resistance [42]. This suggests that an insulin secretagogue would be preferable for the former and insulin sensitisers for the latter. This is one sample application of personalised or precision medicine at the population level.

In a brilliant review, Schwartz et al. [43] make out a case for a new classification system of diabetes and discuss the rationale for and implications of the β -cell-centred classification schema for T2DM. They also emphasise the concept of epigenetics and inflammation, as well as the role of inflammation and immune dysregulation, in the pathophysiology of T2DM [44]. They suggest how data from “multiomics” could be integrated with electronic medical records to provide “dynamic decision support” for recommendations regarding the treatment

of T2DM. They envisage that in the not-too-distant future, data on genomic, epigenomic, proteomic and metabolomic markers could be available on a chip. This will help to classify individuals with diabetes more accurately, and thus make diabetes treatment more patient centred and precision medicine in diabetes a reality.

Figure 1 summarises the concept of a futuristic model of precision diabetes which could be applied at a diabetes clinic to offer personalised diabetes care [44].

Monogenic Diabetes: The Field where Precision Diabetes Has Already Become a Reality

In contrast to the polygenic forms of diabetes like T2DM and T1DM, where precision diabetes is still an evolving field, there are many forms of monogenic diabetes where precision diabetes has already found a place in clinical practice. We will deal with three of these conditions: (1) NDM, (2) CHH, and (3) MODY.

Neonatal Diabetes Mellitus

NDM is defined as diabetes which develops in neonates under 6 months of age. About 50% of the patients with NDM have mutations in the K_{ATP} channel genes, namely, *KCNJ11* and *ABCC8* [45]. Patients with mutations in these two genes respond to high-dose SUs and show excellent glucose control without any increase in hypoglycaemia and glucose variability [46]. Improvements in neurological function are also seen in these patients after treatment with SU [47, 48]. In such patients, treatment with insulin reduces the glucose level but does not elicit an optimal response – and, more specifically, the neurological functions do not show any improvement [49]. SUs close the SU channel in the cells so that exocytosis of insulin takes place, thereby correcting the defect. Since K_{ATP} channels are present in β -cells, the brain, the liver, etc., all cells are corrected simultaneously, thus improving their functions as well. This is an excellent illustration of precision diabetes and one of the most exciting clinical situations in diabetology. Switching a child aged just a few weeks or months to oral tablets, from what would otherwise have meant lifelong insulin injections, is nothing short of a miracle to the child and the family, and we have been able to help many such children [50].

Not all mutations in the *KCNJ11* and *ABCC8* genes, however, lead to NDM. Only some mutations are clinically actionable. This needs to be investigated by functional methods and through segregation analysis. The knowledge of causal mutations has helped in the success-

ful transfer of children from insulin to SUs, as shown in studies by Hattersley and Ashcroft [51] and recently by our group [52]. Patients with transient neonatal diabetes due to 6q24 methylation abnormalities can be treated with low-dose SU when they relapse, while patients with other subtypes of neonatal diabetes require insulin treatment [53]. The ability to explain additional clinical abnormalities that are associated with the underlying genetic cause is one of the major benefits of precision medicine. One such example is cardiac defects in patients with *GATA6* mutations and remission of transient diabetes in patients with 6q24 methylation abnormalities [54].

Patients with transient neonatal diabetes due to 6q24methylation abnormalities can be treated with low-dose SUs [55] when they relapse, whereas patients with NDM due to K_{ATP} channel mutations require high-dose SU and those with non- K_{ATP} channel mutations require insulin treatment [56].

Congenital Hyperinsulinaemic Hypoglycaemia

Another important area where precision diabetes is playing a role is CHH. Here, hypoglycaemia (low blood glucose) manifests just after birth or in the first few months of life. This hypoglycaemia can be severe, debilitating and persistent. In many of these neonates with hypoglycaemia, definite mutations can be identified that are responsible for the condition. We reported on a series of CHH cases from our centre [57]. Some of these neonates respond to diazoxide, while others require subtotal pancreatectomy. Before such a drastic procedure, it is useful to confirm the diagnosis by genetic testing, and this is where precision diabetes plays a role [57].

In the case of conditions such as thiamine responsive megaloblastic anaemia syndrome, an early genetic diagnosis and treatment with thiamine help in the control of diabetes [58]. For IPEX and some severe monogenic autoimmune syndromes, a genetic diagnosis helps in early detection and allows consideration of early curative bone marrow transplantation before patients become severely affected [59].

Maturity-Onset Diabetes of the Young

By its classic definition, MODY presents in non-obese individuals before the age of 25 years, is inherited in an autosomal dominant manner, is primarily due to β -cell dysfunction and can be treated without insulin for at least 5 years [60]. Since the presentation of MODY has characteristics of both type 1 (early onset, lean) and type 2 (family history of diabetes, with β -cell function), it is often misdiagnosed. After the discovery of the MODY genes, a

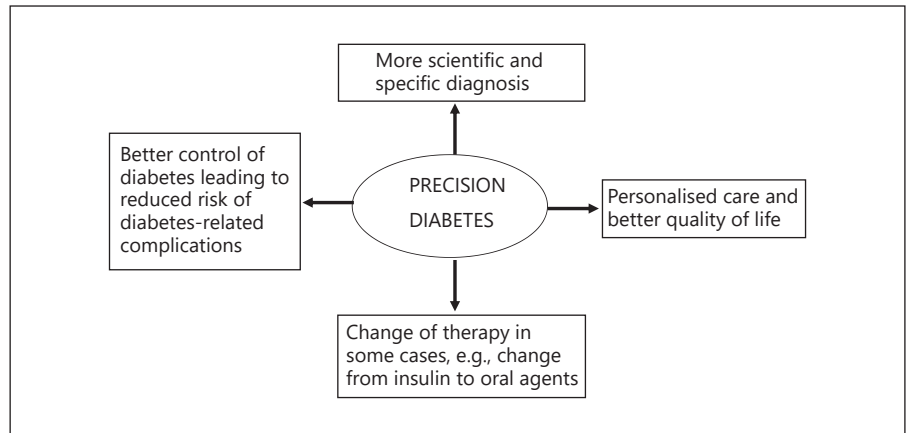


Fig. 2. Advantages of precision diabetes applications.

Table 1. Usefulness of precision diabetes in monogenic diabetes

I. Neonatal diabetes mellitus

1. Helps to differentiate transient neonatal diabetes from permanent neonatal diabetes
2. Helps to stop insulin administration and switch to high-dose sulphonylurea if the children have *KCNJ11* or *ABCC8* mutations
3. Diabetes caused by *SLC19A2* gene mutations can be successfully treated with thiamine
4. Early detection and treatment of Wolcott-Rallison syndrome ensures earlier detection and management of liver abnormalities
5. Developmental disorders of the exocrine pancreas require replacement of pancreatic enzyme in addition to insulin
6. Helps detect various genetic syndromes associated with neonatal diabetes mellitus

II. MODY

1. Patients with *HNF1A* and *4A* MODY can be taken off insulin and treated with low-dose sulphonylurea
2. GCK MODY patients require only a diet regimen and all pharmacological treatment can be stopped, except if they become pregnant

MODY, maturity-onset diabetes of the young.

genetic workup helps in classifying MODY accurately. This is very important, since some of the MODY subtypes entail specific therapeutic recommendations based on the genetic aetiology. The transcription factor genes *HNF1A* (MODY 3) and *HNF4A* (MODY 1) promote transcription of multiple genes related to glucose metabolism,

insulin secretion and insulin production. Diagnosis of MODY 1 and 3 is important for clinical therapy, as patients with those types are very sensitive to SUs [61–67]. This is due to decreased expression of the target genes *HNF1A* and *4A* in the liver, leading to decreased uptake of SUs, in turn resulting in increased circulating levels of SUs [68]. Therefore, *HNF1A* and *4A* MODY patients need lower doses of SUs, which are the first-line treatment for MODY 1 and 3 [69]. It has been demonstrated that switching of treatment from insulin to SU improved HbA_{1c} levels in MODY 3 and MODY 1 patients. In a randomised trial, Pearson et al. [70] showed that SUs lead to a 4-fold greater reduction of fasting plasma glucose levels in *HNF1A* MODY patients compared to T2DM patients matched for age, BMI and blood glucose levels.

GCK MODY (MODY 2) patients do not require insulin or oral drugs, as these patients have a regulated blood glucose set at a higher threshold level [71, 72]. They respond to diet and exercise and rarely develop complications from diabetes. Only during pregnancy do GCK MODY women need insulin [73]. However, its validation in non-European populations has not been tested.

In Asian Indians, T2DM occurs earlier and often overlaps with MODY, and hence understanding and recognising MODY subtypes in this population gains more importance. Our work has revealed important insights into the molecular genetics of MODY by showing the presence of novel mutations in the *HNF1A* (MODY 3) [74] and *HNF4A* (MODY 1) [61] genes. We showed that about 9% of clinically diagnosed MODY subjects had MODY 3 [75], while 3% had MODY 1. We also identified MODY 5 in some of our patients [76]. We have recently reported on a comprehensive analysis of all 14 MODY genes and have also reported on what is likely to be a new MODY

gene, with a mutation in the *NKX6-1* gene, located on chromosome 7 [77].

Table 1 summarises the usefulness of precision diabetes in monogenic forms of diabetes. Advancements in genetic testing due to the development of next-generation sequencing have made comprehensive analyses of all known genetic bases for monogenic diabetes possible. This technology has also helped in reducing the turnaround time for genetic testing. A rapid and precise genetic diagnosis is now possible even before the development of all clinical features [49, 51]. Figure 2 summarises the advantages of precision diabetes.

Conclusions

Rapid strides are being made in the field of precision diabetes. While it is still too early to say that we have reached a stage where every patient with diabetes can be given a personalised form of treatment, we are beginning

to understand that one size does not fit all. In the case of T2DM, clustering of the disorder has now been identified as the first step in precision diabetes. In the case of monogenic forms of diabetes, particularly in MODY, NDM and CHH, genetic testing has come to stay and can be life-changing for those individuals with certain genetic variants. In the years to come, precision diabetes can be expected to play a big role in improving the quality of life of people with all forms of diabetes and help them achieve better control of their diabetes and thereby prevent complications arising from the disorder.

Disclosure Statement

Our work on Precision Diabetes is supported by the INSPIRED project, a grant from NIHR, UK. Financial support for the Monogenic Diabetes study by ICMR, India, is gratefully acknowledged. We also thank ICMR for support from the ICMR Center for Advanced Research (CAR) on Diabetes.

References

- 1 NIH: What is precision medicine? [Internet] [accessed 2018 Dec 22]. Available from: <https://ghr.nlm.nih.gov/primer/precision-medicine/definition>.
- 2 Bayne-Jones S. Dr. Karl Landsteiner Nobel Prize Laureate in Medicine, 1930. *Science*. 1931 Jun;73(1901):599–604.
- 3 Florez JC. Precision medicine in diabetes: is it time? *Diabetes Care*. 2016 Jul;39(7):1085–8.
- 4 Mirnezami R, Nicholson J, Darzi A. Preparing for precision medicine. *N Engl J Med*. 2012 Feb;366(6):489–91.
- 5 Klonoff DC. Precision medicine for managing diabetes. *J Diabetes Sci Technol*. 2015 Jan;9(1):3–7.
- 6 Khan N, Yaqoob I, Hashem IA, Inayat Z, Ali WK, Alam M, et al. Big data: survey, technologies, opportunities, and challenges. *ScientificWorldJournal*. 2014;2014:712826.
- 7 Klonoff DC. Twelve modern digital technologies that are transforming decision making for diabetes and all areas of health care. *J Diabetes Sci Technol*. 2013 Mar;7(2):291–5.
- 8 Ashley EA. Towards precision medicine. *Nat Rev Genet*. 2016 Aug;17(9):507–22.
- 9 McCarthy MI. Painting a new picture of personalised medicine for diabetes. *Diabetologia*. 2017 May;60(5):793–9.
- 10 Shepherd M, Shields B, Ellard S, Rubio-Cabezas O, Hattersley AT. A genetic diagnosis of HNF1A diabetes alters treatment and improves glycaemic control in the majority of insulin-treated patients. *Diabet Med*. 2009 Apr;26(4):437–41.
- 11 Florez JC. Pharmacogenetics in type 2 diabetes: precision medicine or discovery tool? *Diabetologia*. 2017 May;60(5):800–7.
- 12 Huang C, Florez JC. Pharmacogenetics in type 2 diabetes: potential implications for clinical practice. *Genome Med*. 2011 Nov;3(11):76.
- 13 Klonoff DC. Personalized medicine for diabetes. *J Diabetes Sci Technol*. 2008 May;2(3):335–41.
- 14 Maruthur NM, Gribble MO, Bennett WL, Bolen S, Wilson LM, Balakrishnan P, et al. The pharmacogenetics of type 2 diabetes: a systematic review. *Diabetes Care*. 2014;37(3):876–86.
- 15 Ahlqvist E, Storm P, Käräjämäki A, Martinell M, Dorkhan M, Carlsson 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 May;6(5):361–9.
- 16 Dash S, Sano H, Rochford JJ, Semple RK, Yeo G, Hyden CS, et al. A truncation mutation in TBC1D4 in a family with acanthosis nigricans and postprandial hyperinsulinemia. *Proc Natl Acad Sci USA*. 2009 Jun;106(23):9350–5.
- 17 Manousaki D, Kent JW Jr, Haack K, Zhou S, Xie P, Greenwood CM, et al. Toward precision medicine: TBC1D4 disruption is common among the inuit and leads to underdiagnosis of Type 2 diabetes. *Diabetes Care*. 2016 Nov;39(11):1889–95.
- 18 Holstein A, Plaschke A, Ptak M, Egberts EH, El-Din J, Brockmöller J, et al. Association between CYP2C9 slow metabolizer genotypes and severe hypoglycaemia on medication with sulphonylurea hypoglycaemic agents. *Br J Clin Pharmacol*. 2005 Jul;60(1):103–6.
- 19 Ragia G, Petridis I, Tavridou A, Christakidis D, Manolopoulos VG. Presence of CYP2C9*3 allele increases risk for hypoglycemia in Type 2 diabetic patients treated with sulfonylureas. *Pharmacogenomics*. 2009 Nov;10(11):1781–7.
- 20 Niemi M, Leathart JB, Neuvonen M, Backman JT, Daly AK, Neuvonen PJ. Polymorphism in CYP2C8 is associated with reduced plasma concentrations of repaglinide. *Clin Pharmacol Ther*. 2003 Oct;74(4):380–7.
- 21 Nies AT, Koepsell H, Winter S, Burk O, Klein K, Kerb R, et al. Expression of organic cation transporters OCT1 (SLC22A1) and OCT3 (SLC22A3) is affected by genetic factors and cholestasis in human liver. *Hepatology*. 2009 Oct;50(4):1227–40.
- 22 Shu Y, Sheardown SA, Brown C, Owen RP, Zhang S, Castro RA, et al. Effect of genetic variation in the organic cation transporter 1 (OCT1) on metformin action. *J Clin Invest*. 2007 May;117(5):1422–31.
- 23 Pawlyk AC, Giacomini KM, McKeon C, Shuldiner AR, Florez JC. Metformin pharmacogenomics: current status and future directions. *Diabetes*. 2014 Aug;63(8):2590–9.

- 24 Mahrooz A, Parsanasab H, Hashemi-Soth MB, Kashi Z, Bahar A, Alizadeh A, et al. The role of clinical response to metformin in patients newly diagnosed with type 2 diabetes: a monotherapy study. *Clin Exp Med*. 2015 May;15(2):159–65.
- 25 Zhou K, Bellenguez C, Spencer CC, Bennett AJ, Coleman RL, Tavendale R, et al.; GoDARTS and UKPDS Diabetes Pharmacogenetics Study Group; Wellcome Trust Case Control Consortium 2; MAGIC Investigators. Common variants near ATM are associated with glycemic response to metformin in type 2 diabetes. *Nat Genet*. 2011 Feb;43(2):117–20.
- 26 Chen M, Hu C, Zhang R, Jiang F, Wang J, Peng D, et al. Association of PAX4 genetic variants with oral antidiabetic drugs efficacy in Chinese type 2 diabetes patients. *Pharmacogenomics J*. 2014 Oct;14(5):488–92.
- 27 Chen M, Hu C, Zhang R, Jiang F, Wang J, Peng D, et al. A variant of PSMD6 is associated with the therapeutic efficacy of oral antidiabetic drugs in Chinese type 2 diabetes patients. *Sci Rep*. 2015 May;5(1):10701.
- 28 Dujic T, Zhou K, Donnelly LA, Tavendale R, Palmer CN, Pearson ER. Association of organic cation transporter 1 with intolerance to metformin in type 2 diabetes: a GoDARTS Study. *Diabetes*. 2015 May;64(5):1786–93.
- 29 Zhou K, Donnelly LA, Kimber CH, Donnan PT, Doney AS, Leese G, et al. Reduced-function SLC22A1 polymorphisms encoding organic cation transporter 1 and glycemic response to metformin: a GoDARTS study. *Diabetes*. 2009 Jun;58(6):1434–9.
- 30 Dawed AY, Donnelly L, Tavendale R, Carr F, Leese G, Palmer CN, et al. CYP2C8 and SLC01B1 variants and therapeutic response to thiazolidinediones in patients with type 2 diabetes. *Diabetes Care*. 2016 Nov;39(11):1902–8.
- 31 Aghaei Meybodi HR, Hasanzad M, Larijani B. Path to personalized medicine for type 2 diabetes mellitus: reality and hope. *Acta Med Iran*. 2017 Mar;55(3):166–74.
- 32 Zimdahl H, Itrich C, Graefe-Mody U, Boehm BO, Mark M, Woerle HJ, et al. Influence of TCF7L2 gene variants on the therapeutic response to the dipeptidylpeptidase-4 inhibitor linagliptin. *Diabetologia*. 2014 Sep;57(9):1869–75.
- 33 Prasad RB, Groop L. Genetics of type 2 diabetes – pitfalls and possibilities. *Genes (Basel)*. 2015 Mar;6(1):87–123.
- 34 't Hart LM, Fritsche A, Nijpels G, van Leeuwen N, Donnelly LA, Dekker JM, et al. The CTRB1/2 locus affects diabetes susceptibility and treatment via the incretin pathway. *Diabetes*. 2013 Sep;62(9):3275–81.
- 35 Dawed AY, Zhou K, Pearson ER. Pharmacogenetics in type 2 diabetes: influence on response to oral hypoglycemic agents. *Pharmacogenomics Pers Med*. 2016 Apr;9:17–29.
- 36 Mohan V, Yang W, Son HY, Xu L, Noble L, Langdon RB, et al. Efficacy and safety of sitagliptin in the treatment of patients with type 2 diabetes in China, India, and Korea. *Diabetes Res Clin Pract*. 2009 Jan;83(1):106–16.
- 37 Sudhakaran C, Kishore U, Anjana RM, Unnikrishnan R, Mohan V. Effectiveness of sitagliptin in Asian Indian patients with type 2 diabetes – an Indian tertiary diabetes care center experience. *Diabetes Technol Ther*. 2011 Jan;13(1):27–32.
- 38 Kim YG, Hahn S, Oh TJ, Kwak SH, Park KS, Cho YM. Differences in the glucose-lowering efficacy of dipeptidyl peptidase-4 inhibitors between Asians and non-Asians: a systematic review and meta-analysis. *Diabetologia*. 2013 Apr;56(4):696–708.
- 39 Park H, Park C, Kim Y, Rascati KL. Efficacy and safety of dipeptidyl peptidase-4 inhibitors in type 2 diabetes: meta-analysis. *Ann Pharmacother*. 2012 Nov;46(11):1453–69.
- 40 Kim YG, Hahn S, Oh TJ, Park KS, Cho YM. Differences in the HbA1c-lowering efficacy of glucagon-like peptide-1 analogues between Asians and non-Asians: a systematic review and meta-analysis. *Diabetes Obes Metab*. 2014 Oct;16(10):900–9.
- 41 Staimez LR, Weber MB, Ranjani H, Ali MK, Echouffo-Tcheugui JB, Phillips LS, et al. Evidence of reduced β -cell function in Asian Indians with mild dysglycemia. *Diabetes Care*. 2013 Sep;36(9):2772–8.
- 42 Narayan KM. Type 2 diabetes: why we are winning the battle but losing the war? 2015 Kelly West Award Lecture. *Diabetes Care*. 2016 May;39(5):653–63.
- 43 Schwartz SS, Epstein S, Corkey BE, Grant SF, Gavin JR 3rd, Aguilar RB. The time is right for a new classification system for diabetes: rationale and implications of the β -cell-centric classification schema. *Diabetes Care*. 2016 Feb;39(2):179–86.
- 44 Schwartz SS, Epstein S, Corkey BE, Grant SF, Gavin JR III, Aguilar RB, et al. A unified pathophysiologic construct of diabetes and its complications. *Trends Endocrinol Metab*. 2017 Sep;28(9):645–55.
- 45 Hattersley AT, Patel KA. Precision diabetes: learning from monogenic diabetes. *Diabetologia*. 2017 May;60(5):769–77.
- 46 Pearson ER, Flechtner I, Njølstad PR, Malecki MT, Flanagan SE, Larkin B, et al.; Neonatal Diabetes International Collaborative Group. Switching from insulin to oral sulfonylureas in patients with diabetes due to Kir6.2 mutations. *N Engl J Med*. 2006 Aug;355(5):467–77.
- 47 Sagen JV, Raeder H, Hathout E, Shehadeh N, Gudmundsson K, Baevre H, et al. Permanent neonatal diabetes due to mutations in KCNJ11 encoding Kir6.2: patient characteristics and initial response to sulfonylurea therapy. *Diabetes*. 2004 Oct;53(10):2713–8.
- 48 Beltrand J, Elie C, Busiah K, Fournier E, Boddaert N, Bahi-Buisson N, et al.; GlidKir Study Group. Sulfonylurea therapy benefits neurological and psychomotor functions in patients with neonatal diabetes owing to potassium channel mutations. *Diabetes Care*. 2015 Nov;38(11):2033–41.
- 49 Gloy AL, Diatloff-Zito C, Edghill EL, Bellan -Chantelot C, Nivot S, Coutant R, et al. KCNJ11 activating mutations are associated with developmental delay, epilepsy and neonatal diabetes syndrome and other neurological features. *Eur J Hum Genet*. 2006 Jul;14(7):824–30.
- 50 Jahnavi S, Poovazhagi V, Mohan V, Bodhini D, Raghupathy P, Amutha A, et al. Clinical and molecular characterization of neonatal diabetes and monogenic syndromic diabetes in Asian Indian children. *Clin Genet*. 2013 May;83(5):439–45.
- 51 Hattersley AT, Ashcroft FM. Activating mutations in Kir6.2 and neonatal diabetes: new clinical syndromes, new scientific insights, and new therapy. *Diabetes*. 2005 Sep;54(9):2503–13.
- 52 Radha V, Ramya B, Gopi S, Kavitha B, Preetika S, Thai K, et al. Successful transition to sulphonylurea therapy from insulin in a child with permanent neonatal diabetes due to a KCNJ11 gene mutation. *J Diabetol*. 2018;9(2):65–7.
- 53 De Franco E, Flanagan SE, Houghton JA, Lango Allen H, Mackay DJ, Temple IK, et al. The effect of early, comprehensive genomic testing on clinical care in neonatal diabetes: an international cohort study. *Lancet*. 2015 Sep;386(9997):957–63.
- 54 Villamayor L, Rodr guez-Seguel E, Araujo R, Carrasco M, Bru-Tar  E, Mellado-Gil JM, et al. GATA6 controls insulin biosynthesis and secretion in adult β -cells. *Diabetes*. 2018 Mar;67(3):448–60.
- 55 Cav  H, Polak M, Drunat S, Denamur E, Czernichow P. Refinement of the 6q chromosomal region implicated in transient neonatal diabetes. *Diabetes*. 2000 Jan;49(1):108–13.
- 56 Rubio-Cabezas O, Flanagan SE, Damhuis A, Hattersley AT, Ellard S. KATP channel mutations in infants with permanent diabetes diagnosed after 6 months of life. *Pediatr Diabetes*. 2012 Jun;13(4):322–5.
- 57 Jahnavi S, Poovazhagi V, Kanthimathi S, Balamurugan K, Bodhini D, Yadav J, et al. Novel ABCC8 (SUR1) gene mutations in Asian Indian children with congenital hyperinsulinemic hypoglycemia. *Ann Hum Genet*. 2014 Sep;78(5):311–9.
- 58 Olsen BS, Hahnemann JM, Schwartz M, Østergaard E. Thiamine-responsive megaloblastic anaemia: a cause of syndromic diabetes in childhood. *Pediatr Diabetes*. 2007 Aug;8(4):239–41.
- 59 Bacchetta R, Barzaghi F, Roncarolo MG. From IPEX syndrome to FOXP3 mutation: a lesson on immune dysregulation. *Ann N Y Acad Sci*. 2018 Apr;1417(1):5–22.

- 60 Tattersall RB, Fajans SS, Arbor A. A difference between the inheritance of classical juvenile-onset and maturity-onset type diabetes of young people. *Diabetes*. 1975 Jan;24(1):44–53.
- 61 Anuradha S, Radha V, Mohan V. Association of novel variants in the hepatocyte nuclear factor 4A gene with maturity onset diabetes of the young and early onset type 2 diabetes. *Clin Genet*. 2011 Dec;80(6):541–9.
- 62 Anuradha S, Radha V, Deepa R, Hansen T, Carstensen B, Pedersen O, et al. A prevalent amino acid polymorphism at codon 98 (Ala-98Val) of the hepatocyte nuclear factor-1alpha is associated with maturity-onset diabetes of the young and younger age at onset of type 2 diabetes in Asian Indians. *Diabetes Care*. 2005 Oct;28(10):2430–5.
- 63 Murphy R, Ellard S, Hattersley AT. Clinical implications of a molecular genetic classification of monogenic beta-cell diabetes. *Nat Clin Pract Endocrinol Metab*. 2008 Apr;4(4):200–13.
- 64 Jesić MD, Sajić S, Jesić MM, Maringa M, Micić D, Necić S. A case of new mutation in maturity-onset diabetes of the young type 3 (MODY 3) responsive to a low dose of sulphonylurea. *Diabetes Res Clin Pract*. 2008 Jul;81(1):e1–3.
- 65 Pearson ER, Badman MK, Lockwood CR, Clark PM, Ellard S, Bingham C, et al. Contrasting diabetes phenotypes associated with hepatocyte nuclear factor-1alpha and -1beta mutations. *Diabetes Care*. 2004 May;27(5):1102–7.
- 66 Jahnavi S, Poovazhagi V, Mohan V, Radha V. Neonatal diabetes and hyperinsulinemia: the Indian experience. *J Neonatol*. 2013;27:15–23.
- 67 Varadarajan P, Ananthanarayanan K, Mirna K, Suresh J, Venkatesan R, Mohan V. Clinical profile and outcome of persistent hyperinsulinemic hypoglycemia of infancy. *Pediatr Oncall*. 2013;50:759–63.
- 68 Pearson ER, Starkey BJ, Powell RJ, Gribble FM, Clark PM, Hattersley AT. Genetic cause of hyperglycaemia and response to treatment in diabetes. *Lancet*. 2003 Oct;362(9392):1275–81.
- 69 Pearson ER, Liddell WG, Shepherd M, Corrall RJ, Hattersley AT. Sensitivity to sulphonylureas in patients with hepatocyte nuclear factor-1alpha gene mutations: evidence for pharmacogenetics in diabetes. *Diabet Med*. 2000 Jul;17(7):543–5.
- 70 Pearson ER, Pruhova S, Tack CJ, Johansen A, Castleden HA, Lumb PJ, et al. Molecular genetics and phenotypic characteristics of MODY caused by hepatocyte nuclear factor 4alpha mutations in a large European collection. *Diabetologia*. 2005 May;48(5):878–85.
- 71 Chakera AJ, Steele AM, Gloyn AL, Shepherd MH, Shields B, Ellard S, et al. Recognition and management of individuals with hyperglycemia because of a heterozygous glucokinase mutation. *Diabetes Care*. 2015 Jul;38(7):1383–92.
- 72 Kanthimathi S, Jahnavi S, Balamurugan K, Ranjani H, Sonya J, Goswami S, et al. Glucokinase gene mutations (MODY 2) in Asian Indians. *Diabetes Technol Ther*. 2014 Mar;16(3):180–5.
- 73 Spyer G, Macleod KM, Shepherd M, Ellard S, Hattersley AT. Pregnancy outcome in patients with raised blood glucose due to a heterozygous glucokinase gene mutation. *Diabet Med*. 2009 Jan;26(1):14–8.
- 74 Radha V, Ek J, Anuradha S, Hansen T, Pedersen O, Mohan V. Identification of novel variants in the hepatocyte nuclear factor-1alpha gene in South Indian patients with maturity onset diabetes of young. *J Clin Endocrinol Metab*. 2009 Jun;94(6):1959–65.
- 75 Balamurugan K, Bjørkhaug L, Mahajan S, Kanthimathi S, Njølstaad PR, Srinivasan N, et al. Structure-function studies of HNF1A (MODY3) gene mutations in South Indian patients with monogenic diabetes. *Clin Genet*. 2016 Dec;90(6):486–95.
- 76 Kanthimathi S, Balamurugan K, Mohan V, Shanthirani CS, Gayathri V, Radha V. Identification and molecular characterization of HNF1B gene mutations in Indian diabetic patients with renal abnormalities. *Ann Hum Genet*. 2015 Jan;79(1):10–9.
- 77 Mohan V, Radha V, Nguyen TT, Stawiski EW, Pahuja KB, Goldstein LD, et al. Comprehensive genomic analysis identifies pathogenic variants in maturity-onset diabetes of the young (MODY) patients in South India. *BMC Med Genet*. 2018 Feb;19(1):22.