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# **Genetics in Diabetes Type 2 Diabetes and Related Traits**

**Volume Editors** 

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#### **Frontiers in Diabetes**

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### **Contents**

#### VII Preface

Gloyn, A.L.; McCarthy, M.I. (Oxford)

### Gene Discovery Efforts for Type 2 Diabetes

- 1 Genome-Wide Association Studies in Type 2 Diabetes Beer, N.L.; McCarthy, M.I. (Oxford)
- **14 Fine Mapping Type 2 Diabetes Susceptibility Loci** Morris, A.P. (Oxford/Liverpool)
- 29 Whole Genome and Exome Sequencing of Type 2 Diabetes
  Gaulton, K. (Oxford); Flannick, J. (Cambridge, Mass.); Fuchsberger, C. (Ann Arbor, Mich.)

### Gene Discovery Efforts for Glycaemic and Metabolic Traits

- **42 Genome-Wide Association Studies of Glycaemic Traits: A MAGICal Journey** Florez, J.C. (Cambridge, Mass./Boston, Mass.); Barroso, I. (Hinxton/Cambridge)
- **58 Genome-Wide Association Studies of Obesity and Related Traits** Mohlke, K.L. (Chapel Hill, N.C.); Lindgren, C.M. (Oxford)

### Gene Discovery Efforts for Monogenic Disorders of $\beta\text{-Cell}$ Dysfunction and Insulin Resistance

71 Next-Generation Sequencing for the Diagnosis of Monogenic Diabetes and Discovery of Novel Aetiologies

Ellard, S.; De Franco, E. (Exeter)

**87 Whole-Exome Sequencing of Patients with Severe Disorders of Insulin Action** Semple, R. (Cambridge); Barroso, I. (Hinxton/Cambridge)

### 'Omics' of Type 2 Diabetes and Related Traits

- **102 Epigenetic Modifications and Type 2 Diabetes in Humans** Ling, C. (Malmö)
- 111 Insights into β-Cell Biology and Type 2 Diabetes Pathogenesis from Studies of the Islet Transcriptome

van de Bunt, M. (Oxford); Morán, I.; Ferrer, J. (London/Barcelona); McCarthy, M.I. (Oxford)

122 Genomics of Adipose Tissue

Pinnick, K.E.; Karpe, F. (Oxford)

### Insights into Molecular Mechanisms and Pathophysiology from Genetics

## 133 Translating Genetic Association Signals for Diabetes and Metabolic Traits into Molecular Mechanisms for Disease

Rees, M.G. (Oxford/Bethesda, Md.); Gloyn, A.L. (Oxford)

### 146 Understanding Molecular Mechanisms for Diabetes and Obesity through Mouse Models

Merkestein, M. (Oxford); Cox, R. (Harwell); Ashcroft, F. (Oxford)

#### **Clinical Translation**

### 158 Genetics of Drug Response in Diabetes

Pearson, E.R. (Dundee); Florez, J.C. (Boston, Mass./Cambridge, Mass.)

### 173 Translating Advances in Our Understanding of the Genetics of Diabetes into the Clinic

Gardner, D.S. (Oxford/Singapore); Owen, K.R.; Gloyn, A.L. (Oxford)

- 187 Author Index
- 188 Subject Index

VI Contents

### **Preface**

In the 1960s, the American Geneticist J.V. Neel referred to diabetes as the 'geneticists nightmare' owing to the high probability that the phenotype was heterogeneous, not clearly defined with a variable age of onset and a strong environmental influence. In the 1970s, the distinction between autoimmune (type 1 diabetes) and non-autoimmune (type 2 diabetes) was made clarifying a major cause of the disease heterogeneity. Through the 1990s, the discovery of the genes involved in mendelian forms of diabetes demonstrated the enormous power of human genetics to uncover fundamental insights into glucose homeostasis and to inform on treatment and prognosis for patients with particular genetic subtypes of diabetes. The genetic basis of type 2 diabetes, however, remained elusive.

In recent years, the field of human genetics discovery has been revolutionised by publically funded initiatives such as the Human Genome Project, HapMap and 1000 Genomes projects. These, in tandem with technological advances such as array genotyping and next-generation sequencing, have enabled genome-wide studies of genetic variation in previously unimaginable sample numbers. This has in turn led to an explosion in the number of genetic loci robustly implicated in type 2 diabetes risk.

In writing this book, we have called upon a number of our colleagues who, over the years, have been part of highly collaborative international efforts to advance our understanding of the genetic basis of type 2 diabetes and related traits. We are indebted to them for agreeing to help us with capturing this journey. They have described the huge progress that has been made, whilst at the same time outlining the substantial challenges that lie ahead if we are to fully capitalise on the these discoveries and translate our improved understanding of the genetic basis of type 2 diabetes into advances in clinical care.

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