

# METHODS IN MOLECULAR BIOLOGY

*Series Editor*

John M. Walker

School of Life and Medical Sciences

University of Hertfordshire

Hatfield, Hertfordshire, UK

For further volumes:  
<http://www.springer.com/series/7651>

For over 35 years, biological scientists have come to rely on the research protocols and methodologies in the critically acclaimed *Methods in Molecular Biology* series. The series was the first to introduce the step-by-step protocols approach that has become the standard in all biomedical protocol publishing. Each protocol is provided in readily-reproducible step-by-step fashion, opening with an introductory overview, a list of the materials and reagents needed to complete the experiment, and followed by a detailed procedure that is supported with a helpful notes section offering tips and tricks of the trade as well as troubleshooting advice. These hallmark features were introduced by series editor Dr. John Walker and constitute the key ingredient in each and every volume of the *Methods in Molecular Biology* series. Tested and trusted, comprehensive and reliable, all protocols from the series are indexed in PubMed.

# **Immunogenetics**

## **Methods and Protocols**

Edited by

**Anton W. Langerak**

*Department of Immunology, Erasmus MC, Rotterdam, The Netherlands*

 **Humana Press**

*Editor*

Anton W. Langerak  
Department of Immunology  
Erasmus MC  
Rotterdam, The Netherlands



ISSN 1064-3745

Methods in Molecular Biology

ISBN 978-1-0716-2114-1

<https://doi.org/10.1007/978-1-0716-2115-8>

ISSN 1940-6029 (electronic)

ISBN 978-1-0716-2115-8 (eBook)

© The Editor(s) (if applicable) and The Author(s) 2022. This book is an open access publication.

**Open Access** This book is licensed under the terms of the Creative Commons Attribution 4.0 International License (<http://creativecommons.org/licenses/by/4.0/>), which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license and indicate if changes were made.

The images or other third party material in this book are included in the book's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the book's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder.

The use of general descriptive names, registered names, trademarks, service marks, etc. in this publication does not imply, even in the absence of a specific statement, that such names are exempt from the relevant protective laws and regulations and therefore free for general use.

The publisher, the authors and the editors are safe to assume that the advice and information in this book are believed to be true and accurate at the date of publication. Neither the publisher nor the authors or the editors give a warranty, expressed or implied, with respect to the material contained herein or for any errors or omissions that may have been made. The publisher remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

This Humana imprint is published by the registered company Springer Science+Business Media, LLC, part of Springer Nature.

The registered company address is: 1 New York Plaza, New York, NY 10004, U.S.A.

---

## Preface

Adaptive immune cells (lymphocytes) are equipped with unique antigen receptors, termed immunoglobulins (IG) and T cell receptors (TR), which collectively form a highly diverse repertoire. In the lymphocytes, IG/TR diversity is actually created at the DNA level, thus giving rise to an enormous adaptive immune receptor repertoire (also known as the *immunome*) that can be studied in healthy and diseased subjects in the context of research questions and clinical applications. This field of (fundamental and translational) research is known as *immunogenetics*.

The immunogenetics domain has rapidly evolved in the last ten years or so, mainly through the introduction of high-throughput technologies. With these new technologies, unprecedented insight into the adaptive immune receptor repertoire could be obtained with much more sequencing depth and coverage of the repertoire than ever before. In this volume, many chapters are dedicated to lab protocols, bioinformatics, and immunoinformatics analysis of this high-resolution immunome analysis, exemplified by many different applications. Additionally, the newest technological variations on these protocols are discussed, including non-amplicon, single-cell, and cell-free strategies. Collectively, the chapters illustrate the impact that immunogenetics has achieved and will further expand in all fields of medicine, from infection and (auto)immunity, to vaccination, to lymphoid malignancy and tumor immunity.

As the guest editor of this volume on immunogenetics in the *Methods in Molecular Biology* book series, I am very pleased with the content and quality of this book. I am grateful to all authors who contributed to the success of this book volume with their valuable and informative chapters that collectively cover a broad spectrum of methodologies for applications in research and clinical diagnostics. I sincerely hope that readers will find the protocols and the method descriptions as useful as I did, for their own laboratory studies. Enjoy reading!

*Rotterdam, The Netherlands*

*Anton W. Langerak*

---

# Contents

<i>Preface</i> .....	v
<i>Contributors</i> .....	xi
<b>1</b> The Advent of Precision Immunology: Immunogenetics at the Center of Immune Cell Analysis in Health and Disease .....	1
<i>Anton W. Langerak</i>	
<b>2</b> Next-Generation Sequencing-Based Clonality Detection of Immunoglobulin Gene Rearrangements in B-Cell Lymphoma .....	7
<i>Diede A. G. van Bladel, Jessica L. M. van der Last-Kempkes, Blanca Scheijen, Patricia J. T. A. Groenen, and on behalf of the EuroClonality Consortium</i>	
<b>3</b> One-Step Next-Generation Sequencing of Immunoglobulin and T-Cell Receptor Gene Recombinations for MRD Marker Identification in Acute Lymphoblastic Leukemia.....	43
<i>Patrick Villarese, Chrystelle Abdo, Matthieu Bertrand, Florian Thonier, Mathieu Giraud, Mikael Salson, and Elizabeth Macintyre</i>	
<b>4</b> Immunoglobulin/T-Cell Receptor Gene Rearrangement Analysis Using RNA-Seq .....	61
<i>Vincent H. J. van der Velden, Lorenz Bastian, Monika Briüggemann, Alina M. Hartmann, and Nikos Darzentas</i>	
<b>5</b> Minimal Residual Disease Analysis by Monitoring Immunoglobulin and T-Cell Receptor Gene Rearrangements by Quantitative PCR and Droplet Digital PCR .....	79
<i>Irene Della Starza, Cornelia Eckert, Daniela Drandi, and Giovanni Cazzaniga, and on behalf of the EuroMRD Consortium</i>	
<b>6</b> Quality Control for IG/TR Marker Identification and MRD Analysis .....	91
<i>Eva Fronkova, Michael Svaton, and Jan Trka</i>	
<b>7</b> cfDNA-Based NGS IG Analysis in Lymphoma .....	101
<i>Christiane Pott, Michaela Kotrova, Nikos Darzentas, Monika Briüggemann, Mouhamad Khouja, and on behalf of the EuroClonality-NGS Working Group</i>	
<b>8</b> Targeted Locus Amplification as Marker Screening Approach to Detect Immunoglobulin (IG) Translocations in B-Cell Non-Hodgkin Lymphomas .....	119
<i>Elisa Genuardi, Beatrice Alessandria, Aurora Maria Civita, and Simone Ferrero</i>	
<b>9</b> Immunoglobulin/T Cell Receptor Capture Strategy for Comprehensive Immunogenetics .....	133
<i>James Peter Stewart, Jana Gazdova, Shambhavi Srivastava, Julia Revolta, Louise Harewood, Manisha Maurya, Nikos Darzentas, and David Gonzalez</i>	

10	Immunoglobulin Gene Mutational Status Assessment by Next Generation Sequencing in Chronic Lymphocytic Leukemia . . . . .	153
	<i>Anne Langlois de Septenville, Myriam Boudjogbra, Clotilde Bravetti, Marine Armand, Mikaël Salson, Mathieu Giraud, and Frederic Davi</i>	
11	NGS-Based B-Cell Receptor Repertoire AnalysisRepertoire analyses in the Context of Inborn Errors of Immunity . . . . .	169
	<i>Pauline A. van Schouwenburg, Mirjam van der Burg, and Hanna IJspeert</i>	
12	Generic Multiplex Digital PCR for Accurate Quantification of T Cells in Copy Number Stable and Unstable DNA Samples . . . . .	191
	<i>Rogier J. Nell, Willem H. Zoutman, Mieke Versluis, and Pieter A. van der Velden</i>	
13	Gene Engineering T Cells with T-Cell Receptor for Adoptive Therapy . . . . .	209
	<i>Dian Kortleve, Mandy van Brakel, Rebecca Wijers, Reno Debets, and Dora Hammerl</i>	
14	Combined Analysis of Transcriptome and T-Cell Receptor Alpha and Beta (TRA/TRB) Repertoire in Paucicellular Samples at the Single-Cell Level . . . . .	231
	<i>Nicolle H. R. Litjens, Anton W. Langerak, Zakia Azmani, Xander den Dekker, Michiel G. H. Betjes, Rutger W. W. Brouwer, and Wilfred F. J. van IJcken</i>	
15	AIRR Community Guide to Planning and Performing AIRR-Seq Experiments . . . . .	261
	<i>Anne Eugster, Magnolia L. Bostick, Nidhi Gupta, Encarnita Mariotti-Ferrandiz, Gloria Kraus, Wenzhao Meng, Cinque Soto, Johannes Trück, Ulrik Stervbo, Eline T. Luning Prak, and on behalf of the AIRR Community</i>	
16	Adaptive Immune Receptor Repertoire (AIRR) Community Guide to TR and IG Gene Annotation . . . . .	279
	<i>Lmar Babrak, Susanna Marquez, Christian E. Busse, William D. Lees, Enkelejda Miho, Mats Ohlin, Aaron M. Rosenfeld, Ulrik Stervbo, Corey T. Watson, Chaim A. Schramm, and on behalf of the AIRR Community</i>	
17	Adaptive Immune Receptor Repertoire (AIRR) Community Guide to Repertoire Analysis . . . . .	297
	<i>Susanna Marquez, Lmar Babrak, Victor Greiff, Kenneth B. Hoehn, William D. Lees, Eline T. Luning Prak, Enkelejda Miho, Aaron M. Rosenfeld, Chaim A. Schramm, Ulrik Stervbo, and on behalf of the AIRR Community</i>	
18	Bulk gDNA Sequencing of Antibody Heavy-Chain Gene Rearrangements for Detection and Analysis of B-Cell Clone Distribution: A Method by the AIRR Community . . . . .	317
	<i>Aaron M. Rosenfeld, Wenzhao Meng, Kalisse I. Horne, Elaine C. Chen, Davide Bagnara, Ulrik Stervbo, Eline T. Luning Prak, and on behalf of the AIRR Community</i>	

19	Bulk Sequencing from mRNA with UMI for Evaluation of B-Cell Isotype and Clonal Evolution: A Method by the AIRR Community .....	345
	<i>Nidhi Gupta, Susanna Marquez, Cinque Soto, Elaine C. Chen, Magnolia L. Bostick, Ulrik Stårvbo, and Andrew Farmer</i>	
20	Single-Cell Analysis and Tracking of Antigen-Specific T Cells: Integrating Paired Chain AIRR-Seq and Transcriptome Sequencing: A Method by the AIRR Community .....	379
	<i>Nidhi Gupta, Ida Lindeman, Susanne Reinhardt, Encarnita Mariotti-Ferrandiz, Kevin Mujangi-Ebeka, Kristen Martins-Taylor, and Anne Eugster</i>	
21	Quality Control: Chain Pairing Precision and Monitoring of Cross-Sample Contamination: A Method by the AIRR Community.....	423
	<i>Cheng-Yu Chung, Matías Gutiérrez-González, Sheila N. López Acevedo, Ahmed S. Fahad, Brandon J. DeKosky, and on behalf of the AIRR Community</i>	
22	Immune Repertoire Analysis on High-Performance Computing Using VDJServer V1: A Method by the AIRR Community .....	439
	<i>Scott Christley, Ulrik Stårvbo, Lindsay G. Cowell, and on behalf of the AIRR Community</i>	
23	Data Sharing and Reuse: A Method by the AIRR Community.....	447
	<i>Brian D. Corrie, Scott Christley, Christian E. Busse, Lindsay G. Cowell, Kira C. M. Neller, Florian Rubelt, Nicholas Schwab, and on behalf of the AIRR Community</i>	
24	IMGT® Immunoinformatics Tools for Standardized V-DOMAIN Analysis .....	477
	<i>Véronique Giudicelli, Patrice Duroux, Maël Rollin, Safa Aouinti, Géraldine Folch, Joumana Jabado-Michaloud, Marie-Paule Lefranc, and Sofia Kossida</i>	
25	IMGT/3Dstructure-DB: T-Cell Receptor TR Paratope and Peptide/Major Histocompatibility pMH Contact Sites and Epitope .....	533
	<i>Marie-Paule Lefranc and Gérard Lefranc</i>	
26	ARResT/Interrogate Immunoprofiling Platform: Concepts, Workflows, and Insights .....	571
	<i>Nikos Darzentas</i>	
27	Purpose-Built Immunoinformatics for BcR IG/TR Repertoire Data Analysis.....	585
	<i>Chryssi Galigalidou, Laura Zaragoza-Infante, Anastasia Chatzidimitriou, Kostas Stamatopoulos, Fotis Psomopoulos, and Andreas Agathangelidis</i>	
	<i>Index .....</i>	<i>605</i>

---

## Contributors

CHRYSTELLE ABDO • *Hôpital Necker Enfants-Malades, Laboratoire d’Onco-Hématologie, Assistance Publique– Hôpitaux de Paris, Paris, France*

ANDREAS AGATHANGELIDIS • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; Department of Biology, School of Science, National and Kapodistrian University of Athens, Athens, Greece*

BEATRICE ALESSANDRIA • *Hematology Division, Department of Molecular Biotechnologies and Health Sciences, University of Torino, Torino, Italy*

SAFA AOINTI • *IMGT®, the international ImMunoGenetics information system®, Laboratoire d’ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France; Clinical Research and Epidemiology Unit, CHU Montpellier, Univ Montpellier, Montpellier, France*

MARINE ARMAND • *AP-HP, Pitié-Salpêtrière Hospital, Laboratory of Hematology, Paris, France; Sorbonne Université, Paris, France*

ZAKIA AZMANI • *Center for Biomics, Erasmus MC University Medical Center, Rotterdam, The Netherlands; Department of Cell Biology, Erasmus MC University Medical Center, Rotterdam, The Netherlands*

LMAR BABRAK • *Institute of Biomedical Engineering and Medical Informatics, School of Life Sciences, FHNW University of Applied Sciences and Arts Northwestern Switzerland, Muttenz, Switzerland*

DAVIDE BAGNARA • *Department of Experimental Medicine, University of Genoa, Genoa, Italy*

LORENZ BASTIAN • *Department of Hematology, University of Schleswig-Holstein, Kiel, Germany*

MAITHEU BERTRAND • *Hôpital Necker Enfants-Malades, Laboratoire d’Onco-Hématologie, Assistance Publique– Hôpitaux de Paris, Paris, France*

MICHAEL G. H. BETJES • *Erasmus MC Transplant Institute, Division of Nephrology and Transplantation, Department of Internal Medicine, Erasmus MC University Medical Center, Rotterdam, The Netherlands*

MAGNOLIA L. BOSTICK • *Takara Bio USA, Inc., San Jose, CA, USA; PACT Pharma, Inc., South Francisco, CA, USA*

MYRIAM BOUDJOGHRA • *AP-HP, Pitié-Salpêtrière Hospital, Laboratory of Hematology, Paris, France*

CLOTILDE BRAVETTI • *AP-HP, Pitié-Salpêtrière Hospital, Laboratory of Hematology, Paris, France; Sorbonne Université, Paris, France*

RUTGER W. W. BROUWER • *Center for Biomics, Erasmus MC University Medical Center, Rotterdam, The Netherlands; Department of Cell Biology, Erasmus MC University Medical Center, Rotterdam, The Netherlands*

MONIKA BRÜGGEMANN • *Department of Hematology, University of Schleswig-Holstein, Kiel, Germany; Medical Department II, University Hospital Schleswig-Holstein, Kiel, Germany*

CHRISTIAN E. BUSSE • *Division of B Cell Immunology, German Cancer Research Center (DKFZ), Heidelberg, Germany*

- GIOVANNI CAZZANIGA • *Centro Ricerca Tettamanti, Fondazione Tettamanti, Centro Maria Letizia Verga, Monza, Italy; Genetics, Department of Medicine and Surgery, University of Milan Bicocca, Monza, Italy*
- ANASTASIA CHATZIDIMITRIOU • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; Department of Molecular Medicine and Surgery, Karolinska Institute, Stockholm, Sweden*
- ELAINE C. CHEN • *Department of Pathology, Microbiology, and Immunology, Vanderbilt Vaccine Center, Vanderbilt University Medical Center, Nashville, TN, USA*
- SCOTT CHRISTLEY • *Department of Population and Data Sciences, UT Southwestern Medical Center, Dallas, TX, USA*
- CHENG-YU CHUNG • *Department of Pharmaceutical Chemistry, The University of Kansas, Lawrence, KS, USA; Department of Chemical Engineering, The University of Kansas, Lawrence, KS, USA*
- AURORA MARIA CIVITA • *Hematology Division, Department of Molecular Biotechnologies and Health Sciences, University of Torino, Torino, Italy*
- BRIAN D. CORRIE • *Biological Sciences, Simon Fraser University, Burnaby, BC, Canada*
- LINDSAY G. COWELL • *Department of Population and Data Sciences, UT Southwestern Medical Center, Dallas, TX, USA; Department of Immunology, UT Southwestern Medical Center, Dallas, TX, USA*
- NIKOS DARZENTAS • *Department of Hematology, University of Schleswig-Holstein, Kiel, Germany; Medical Department II, University Hospital Schleswig-Holstein, Kiel, Germany*
- FREDERIC DAVI • *AP-HP, Pitié-Salpêtrière Hospital, Laboratory of Hematology, Paris, France; Sorbonne Université, Paris, France*
- RENO DEBETS • *Laboratory of Tumor Immunology, Department of Medical Oncology, Erasmus MC-Cancer Institute, Rotterdam, The Netherlands*
- BRANDON J. DEKOSKY • *Department of Pharmaceutical Chemistry, The University of Kansas, Lawrence, KS, USA; Department of Chemical Engineering, The University of Kansas, Lawrence, KS, USA; Bioengineering Graduate Program, The University of Kansas, Lawrence, KS, USA*
- XANDER DEN DEKKER • *Center for Biomics, Erasmus MC University Medical Center, Rotterdam, The Netherlands; Department of Cell Biology, Erasmus MC University Medical Center, Rotterdam, The Netherlands*
- DANIELA DRANDI • *Hematology Division, Department of Molecular Biotechnology and Health Sciences, University of Torino, Torino, Italy*
- PATRICE DUROUX • *IMGT®, the international ImmunoGenetics information system®, Laboratoire d'ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France*
- CORNELIA ECKERT • *Department of Pediatric Oncology Hematology, Charité - Universitätsmedizin Berlin, Berlin, Germany; German Cancer Consortium, and German Cancer Research Center, Heidelberg, Germany*
- ANNE EUGSTER • *Center for Regenerative Therapies Dresden, Faculty of Medicine, TU Dresden, Dresden, Germany*
- AHMED S. FAHAD • *Department of Pharmaceutical Chemistry, The University of Kansas, Lawrence, KS, USA*
- ANDREW FARMER • *Takara Bio USA, Inc., San Jose, CA, USA*

SIMONE FERRERO • *Hematology Division, Department of Molecular Biotechnologies and Health Sciences, University of Torino, Torino, Italy; Hematology Division, AOU “Città della Salute e della Scienza di Torino”, Torino, Italy*

GÉRALDINE FOLCH • *IMGT®, the international ImMunoGenetics information system®, Laboratoire d’ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France*

EVA FRONKOVA • *CLIP - Childhood Leukaemia Investigation Prague, Department of Paediatric Haematology and Oncology, Second Faculty of Medicine, Charles University and University Hospital Motol, Prague, Czech Republic*

CHRYSI GALIGALIDOU • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; Department of Molecular Biology and Genetics (MBG), Democritus University of Thrace, Alexandroupolis, Greece*

JANA GAZDOVA • *Patrick G Johnston Centre for Cancer Research, Queen’s University Belfast, Belfast, UK*

ELISA GENUARDI • *Hematology Division, Department of Molecular Biotechnologies and Health Sciences, University of Torino, Torino, Italy*

MATHIEU GIRAUD • *Université de Lille, CNRS, UMR 9189—CRISTAL, Inria, Lille, France*

VÉRONIQUE GIUDICELLI • *IMGT®, the international ImMunoGenetics information system®, Laboratoire d’ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France*

DAVID GONZALEZ • *Patrick G Johnston Centre for Cancer Research, Queen’s University Belfast, Belfast, UK*

VICTOR GREIFF • *Department of Immunology, University of Oslo, Oslo University Hospital, Oslo, Norway*

PATRICIA J. T. A. GROENEN • *Department of Pathology, Radboud University Medical Center, Nijmegen, The Netherlands*

NIDHI GUPTA • *Takara Bio USA, Inc., San Jose, CA, USA*

MATÍAS GUTIÉRREZ-GONZÁLEZ • *Department of Pharmaceutical Chemistry, The University of Kansas, Lawrence, KS, USA; Department of Chemical Engineering, The University of Kansas, Lawrence, KS, USA*

DORA HAMMERL • *Laboratory of Tumor Immunology, Department of Medical Oncology, Erasmus MC-Cancer Institute, Rotterdam, The Netherlands*

LOUISE HAREWOOD • *Patrick G Johnston Centre for Cancer Research, Queen’s University Belfast, Belfast, UK*

ALINA M. HARTMANN • *Department of Hematology, University of Schleswig-Holstein, Kiel, Germany*

KENNETH B. HOEHN • *Department of Pathology, Yale School of Medicine, New Haven, CT, USA*

KALISSE I. HORNE • *Department of Pathology and Laboratory Medicine, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, USA*

HANNA IJSPEERT • *Department of Immunology, Laboratory Medical Immunology, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands; Academic Center for Rare Immunological Diseases (RIDC), Erasmus University Medical Center, Rotterdam, The Netherlands*

JOUMANA JABADO-MICHALOUD • *IMGT®, the international ImMunoGenetics information system®, Laboratoire d’ImmunoGénétique Moléculaire LIGM, Institut de Génétique*

- HUMAINE, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France
- MOUHAMAD KHOUJA • Medical Department II, University Hospital Schleswig-Holstein, Kiel, Germany
- DIAN KORTLEVE • Laboratory of Tumor Immunology, Department of Medical Oncology, Erasmus MC-Cancer Institute, Rotterdam, The Netherlands
- SOFIA KOSSIDA • IMGT®, the international ImMunoGenetics information system®, Laboratoire d'ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France
- MICHAELA KOTROVA • Medical Department II, University Hospital Schleswig-Holstein, Kiel, Germany
- GLORIA KRAUS • Center for Regenerative Therapies Dresden, Faculty of Medicine, TU Dresden, Dresden, Germany
- ANTON W. LANGERAK • Laboratory Medical Immunology, Department of Immunology, Erasmus MC, University Medical Center, Rotterdam, The Netherlands
- ANNE LANGLOIS DE SEPTENVILLE • AP-HP, Pitié-Salpêtrière Hospital, Laboratory of Hematology, Paris, France
- WILLIAM D. LEES • Institute of Structural and Molecular Biology, Birkbeck College, University of London, London, UK
- GÉRARD LEFRANC • IMGT®, the international ImMunoGenetics information system®, Laboratoire d'ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine IGH, UMR 9002, CNRS, Université de Montpellier, Montpellier cedex 5, France
- MARIE-PAULE LEFRANC • IMGT®, the international ImMunoGenetics information system®, Laboratoire d'ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine IGH, UMR 9002, Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier cedex 5, France
- IDA LINDEMAN • Department of Immunology, Oslo University Hospital and K.G. Jebsen Coeliac Disease Research Centre, University of Oslo, Oslo, Norway
- NICOLLE H. R. LITJENS • Erasmus MC Transplant Institute, Division of Nephrology and Transplantation, Department of Internal Medicine, Erasmus MC University Medical Center, Rotterdam, The Netherlands
- SHEILA N. LÓPEZ ACEVEDO • Department of Pharmaceutical Chemistry, The University of Kansas, Lawrence, KS, USA; Department of Chemical Engineering, The University of Kansas, Lawrence, KS, USA
- ELINE T. LUNING PRAK • Department of Pathology and Laboratory Medicine, Perelman School of Medicine, The University of Pennsylvania, Philadelphia, PA, USA
- ELIZABETH MACINTYRE • Hôpital Necker Enfants-Malades, Laboratoire d'Onco-Hématologie, Assistance Publique–Hôpitaux de Paris, Paris, France
- ENCARNITA MARIOTTI-FERRANDIZ • INSERM, Immunology-Immunopathology-Immunotherapy (i3), Sorbonne Université, Paris, France
- SUSANNA MARQUEZ • Department of Pathology, Yale School of Medicine, New Haven, CT, USA
- KRISTEN MARTINS-TAYLOR • 10x Genomics, Pleasanton, CA, USA
- MANISHA MAURYA • Patrick G Johnston Centre for Cancer Research, Queen's University Belfast, Belfast, UK
- WENZHDAO MENG • Department of Pathology and Laboratory Medicine, Perelman School of Medicine, The University of Pennsylvania, Philadelphia, PA, USA

- ENKELEJDA MIHO • *Institute of Biomedical Engineering and Medical Informatics, School of Life Sciences, FHNW University of Applied Sciences and Arts Northwestern Switzerland, Muttenz, Switzerland; SIB Swiss Institute of Bioinformatics, Lausanne, Switzerland; aiNET GmbH, Basel, Switzerland*
- KEVIN MUJANGI-EBEKA • *INSERM, Immunology-Immunopathology-Immunotherapy (i3), Sorbonne Université, Paris, France*
- ROGIER J. NELL • *Department of Ophthalmology, Leiden University Medical Center, Leiden, The Netherlands*
- KIRA C. M. NELLER • *Health Sciences, Simon Fraser University, Burnaby, BC, Canada*
- MATS OHLIN • *Department of Immunotechnology, Lund University, Lund, Sweden*
- CHRISTIANE POTT • *Medical Department II, University Hospital Schleswig-Holstein, Kiel, Germany*
- FOTIS PSOMOPOULOS • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; Department of Molecular Medicine and Surgery, Karolinska Institute, Stockholm, Sweden*
- SUSANNE REINHARDT • *DRESDEN-concept Genome Center, DFG NGS Competence Center, c/o Center for Molecular and Cellular Bioengineering (CMCB), Technische Universität Dresden, Dresden, Germany*
- JULIA REVOLTA • *Patrick G Johnston Centre for Cancer Research, Queen's University Belfast, Belfast, UK*
- MAËL ROLLIN • *IMGT®, the international ImMunoGenetics information system®, Laboratoire d'ImmunoGénétique Moléculaire LIGM, Institut de Génétique Humaine, (IGH), Centre National de la Recherche Scientifique (CNRS), Université de Montpellier (UM), Montpellier, France*
- AARON M. ROSENFELD • *Department of Pathology and Laboratory Medicine, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, USA*
- FLORIAN RUBELT • *Roche Sequencing Solutions, Roche, Pleasanton, CA, USA*
- MIKAËL SALSON • *Université de Lille, CNRS, UMR 9189—CRIStAL, Inria, Lille, France*
- BLANCA SCHEIJEN • *Department of Pathology, Radboud University Medical Center, Nijmegen, The Netherlands; Radboud Institute for Molecular Life Sciences, Radboud University Medical Center, Nijmegen, The Netherlands*
- CHAIM A. SCHRAMM • *Vaccine Research Center, National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, MD, USA*
- NICHOLAS SCHWAB • *Department of Neurology with Institute of Translational Neurology, University of Muenster, Muenster, Germany*
- CINQUE SOTO • *The Vanderbilt Vaccine Center and Department of Pediatrics, Vanderbilt University Medical Center, Nashville, TN, USA*
- SHAMBHAVI SRIVASTAVA • *Patrick G Johnston Centre for Cancer Research, Queen's University Belfast, Belfast, UK*
- KOSTAS STAMATOPOULOS • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; Department of Molecular Medicine and Surgery, Karolinska Institute, Stockholm, Sweden*
- IRENE DELLA STARZA • *Hematology, Department of Translational and Precision Medicine, “Sapienza” University of Rome, Rome, Italy; GIMEMA Foundation, Rome, Italy*
- ULRIK STERVBO • *Center for Translational Medicine, Immunology, and Transplantation, Medical Department, and Immundiagnostik, Marien Hospital Herne, University Hospital of the Ruhr-University Bochum, Herne, Germany*

- JAMES PETER STEWART • *Patrick G Johnston Centre for Cancer Research, Queen's University Belfast, Belfast, UK*
- MICHAEL SVATON • *CLIP - Childhood Leukaemia Investigation Prague, Department of Paediatric Haematology and Oncology, Second Faculty of Medicine, Charles University and University Hospital Motol, Prague, Czech Republic*
- FLORIAN THONIER • *Inria, Rennes, France*
- JAN TRKA • *CLIP - Childhood Leukaemia Investigation Prague, Department of Paediatric Haematology and Oncology, Second Faculty of Medicine, Charles University and University Hospital Motol, Prague, Czech Republic*
- JOHANNES TRÜCK • *Division of Immunology and Children's Research Center, University Children's Hospital Zurich, University of Zurich (UZH), Zurich, Switzerland*
- DIEDE A. G. VAN BLADEL • *Department of Pathology, Radboud University Medical Center, Nijmegen, The Netherlands; Radboud Institute for Molecular Life Sciences, Radboud University Medical Center, Nijmegen, The Netherlands*
- MANDY VAN BRAKEL • *Laboratory of Tumor Immunology, Department of Medical Oncology, Erasmus MC-Cancer Institute, Rotterdam, The Netherlands*
- MIRJAM VAN DER BURG • *Department of Pediatrics, Laboratory for Pediatric Immunology, Willem-Alexander Children's Hospital, Leiden University Medical Center, Leiden, The Netherlands*
- JESSICA L. M. VAN DER LAST-KEMPES • *Department of Pathology, Radboud University Medical Center, Nijmegen, The Netherlands*
- PIETER A. VAN DER VELDEN • *Department of Ophthalmology, Leiden University Medical Center, Leiden, The Netherlands*
- VINCENT H. J. VAN DER VELDEN • *Department of Immunology, Laboratory Medical Immunology, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands*
- WILFRED F. J. VAN IJCKEN • *Center for Biomics, Erasmus MC University Medical Center, Rotterdam, The Netherlands; Department of Cell Biology, Erasmus MC University Medical Center, Rotterdam, The Netherlands*
- PAULINE A. VAN SCHOUWENBURG • *Department of Pediatrics, Laboratory for Pediatric Immunology, Willem-Alexander Children's Hospital, Leiden University Medical Center, Leiden, The Netherlands*
- MIEKE VERSLUIS • *Department of Ophthalmology, Leiden University Medical Center, Leiden, The Netherlands*
- PATRICK VILLARESE • *Hôpital Necker Enfants-Malades, Laboratoire d'Onco-Hématologie, Assistance Publique-Hôpitaux de Paris, Paris, France*
- COREY T. WATSON • *Department of Biochemistry and Molecular Genetics, University of Louisville School of Medicine, Louisville, KY, USA*
- REBECCA WIJERS • *Laboratory of Tumor Immunology, Department of Medical Oncology, Erasmus MC-Cancer Institute, Rotterdam, The Netherlands*
- LAURA ZARAGOZA-INFANTE • *Institute of Applied Biosciences, Centre for Research and Technology Hellas, Thessaloniki, Greece; First Department of Medicine, Aristotle University of Thessaloniki, Thessaloniki, Greece*
- WILLEM H. ZOUTMAN • *Department of Dermatology, Leiden University Medical Center, Leiden, The Netherlands*