

MM09

Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods

This guideline, in conjunction with instructional worksheets and educational examples, provides step-by-step recommendations for design, development, validation, results reporting, and continual quality management of clinical tests based on next-generation sequencing and Sanger sequencing.

A guideline for global application developed through the Clinical and Laboratory Standards Institute consensus process.

Clinical and Laboratory Standards Institute

Setting the standard for quality in medical laboratory testing around the world.

The Clinical and Laboratory Standards Institute (CLSI) is a not-for-profit membership organization that brings together the varied perspectives and expertise of the worldwide laboratory community for the advancement of a common cause: to foster excellence in laboratory medicine by developing and implementing medical laboratory standards and guidelines that help laboratories fulfill their responsibilities with efficiency, effectiveness, and global applicability.

Consensus Process

Consensus—the substantial agreement by materially affected, competent, and interested parties—is core to the development of all CLSI documents. It does not always connote unanimous agreement but does mean that the participants in the development of a consensus document have considered and resolved all relevant objections and accept the resulting agreement.

Commenting on Documents

CLSI documents undergo periodic evaluation and modification to keep pace with advances in technologies, procedures, methods, and protocols affecting the laboratory or health care.

CLSI's consensus process depends on experts who volunteer to serve as contributing authors and/or as participants in the reviewing and commenting process. At the end of each comment period, the committee that developed the document is obligated to review all comments, respond in writing to all substantive comments, and revise the draft document as appropriate.

Comments on published CLSI documents are equally essential and may be submitted by anyone, at any time, on any document. All comments are managed according to the consensus process by a committee of experts.

Appeal Process

When it is believed that an objection has not been adequately considered and responded to, the process for appeal, documented in the CLSI *Standards Development Policies and Processes*, is followed.

All comments and responses submitted on draft and published documents are retained on file at CLSI and are available upon request.

Get Involved—Volunteer!

Do you use CLSI documents in your workplace? Do you see room for improvement? Would you like to get involved in the revision process? Or maybe you see a need to develop a new document for an emerging technology? CLSI wants to hear from you. We are always looking for volunteers. By donating your time and talents to improve the standards that affect your own work, you will play an active role in improving public health across the globe.

For additional information on committee participation or to submit comments, contact CLSI.

Clinical and Laboratory Standards Institute

P: +1.610.688.0100

F: +1.610.688.0700

www.clsi.org

standard@clsi.org

Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods

Birgit Funke, PhD, FACMG
John D. Pfeifer, MD, PhD
Sami S. Amr, PhD, FACMG
Arthur Martin Baca, MD, PhD
Mark J. Bowser, MS, MPH, MLS(ASCP)^{CM}
Jillian Buchan, MS, PhD, FACMB
Maria Laura Cremona, PhD, FACMG, CGMB
Dora Dias-Santagata, PhD, FACMG
Jianli Dong, MD, PhD, FACMG
Bert Gold, PhD, FACMG, CGMB
Elizabeth Duffy Hynes, BS
Jianling Ji, MD, MS
Sabah Kadri, PhD
Robert F. Klees, PhD
Niklas Krumm, MD, PhD

Annette Leon, PhD, MS, FACMG, CGMBS, CCS
You Li, PhD
Meredith Halks Miller, MD
Dimitri Monos, PhD
Kara Norman, PhD
Honey V. Reddi, PhD, FACMG
Avni Santani, PhD, FACMG
Timothy R. Schwartz, PhD
Junaid Shabbeer, PhD, FACMG
Birgitte Simen, PhD
Cynthe Sims, PhD, HCLD(ABB)
Zivana Tezak, PhD
Francisca Reyes Turcu, PhD
Patrik Vitazka, MD, PhD, FACMG
Richard Y. Zhao, PhD

Clinical and Laboratory Standards Institute (CLSI). *Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods*. 3rd ed. CLSI guideline MM09 (ISBN 978-1-68440-174-1 [Print]; ISBN 978-1-68440-175-8 [Electronic]). Clinical and Laboratory Standards Institute, USA, 2023.

The Clinical and Laboratory Standards Institute consensus process, which is the mechanism for moving a document through two or more levels of review by the health care community, is an ongoing process. Users should expect revised editions of any given document. Because rapid changes in technology may affect the procedures, methods, and protocols in a standard or guideline, users should replace outdated editions with the current editions of CLSI documents. Current editions are listed in the CLSI catalog and posted on our website at www.clsi.org.

If you or your organization is not a member and would like to become one, or to request a copy of the catalog, contact us at:

P: +1.610.688.0100 **F:** +1.610.688.0700 **E:** customerservice@clsi.org **W:** www.clsi.org

Abstract

Sequencing-based clinical tests have evolved from single-gene tests to whole-genome tests. Next-generation sequencing (NGS) technologies have largely replaced Sanger sequencing and are firmly established in the medical management of hereditary disorders, as well as in tumor testing. Newer clinical NGS applications include human leukocyte antigen typing, noninvasive prenatal testing, sequencing of circulating tumor DNA in peripheral blood, and RNA sequencing. Although NGS applications have undergone major technical simplifications, clinical implementation continues to be complex. Clinical and Laboratory Standards Institute guideline MM09—*Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods* provides recommendations for design, development, validation, results reporting, and continual quality management of NGS-based tests, as well as Sanger sequencing-based tests. In conjunction with instructional worksheets and educational examples, MM09 provides step-by-step guidance to help medical laboratories translate regulatory requirements into clinical practice.

Copyright ©2023 Clinical and Laboratory Standards Institute. Except as stated below, any reproduction of content from a CLSI copyrighted standard, guideline, derivative product, or other material requires express written consent from CLSI. All rights reserved. Interested parties may send permission requests to permissions@clsi.org.

CLSI hereby grants permission to each individual member or purchaser to make a single reproduction of this publication for use in its laboratory procedures manual at a single site. To request permission to use this publication in any other manner, e-mail permissions@clsi.org.

Suggested Citation

CLSI. *Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods*. 3rd ed. CLSI guideline MM09. Clinical and Laboratory Standards Institute; 2023.

Previous Editions:

December 2004, February 2014

MM09-Ed3

ISBN 978-1-68440-174-1 (Print)

ISBN 978-1-68440-175-8 (Electronic)

ISSN 1558-6502 (Print)

ISSN 2162-2914 (Electronic)

Volume 43, Number 6

Committee Membership

Consensus Council

The Consensus Council sets priorities for CLSI standards development and votes on Final Draft documents to confirm that process requirements have been met. Consensus Council members are listed on the CLSI website: <https://clsi.org/standards-development/consensus-council/>

Document Development Committee on Nucleic Acid Sequencing

Birgit Funke, PhD, FACMG
Chairholder
Sema4
USA

John D. Pfeifer, MD, PhD
Vice-Chairholder
Washington University of Medicine
USA

Mary Pierce-Burlingame, MB(ASCP)
Committee Secretary
Oxford Gene Technologies
USA

Sami S. Amr, PhD, FACMG
 Partners HealthCare Personalized
 Medicine
 USA

Maria Laura Cremona, PhD, FACMG,
 CGMB
 Illumina, Inc.
 USA

Bert Gold, PhD, FACMG, CGMB
 USA

Jianling Ji, MD, MS
 Children's Hospital Los Angeles,
 University of Southern California
 USA

Robert F. Klees, PhD
 New York State Department of Health
 USA

Felicitas L. Lacbawan, MD, FCAP,
 FACMG
 Quest Diagnostics Nichols Institute
 USA

Annette Leon, PhD, MS, FACMG,
 CGMBS, CCS
 Color Genomics, Inc.
 USA

Honey V. Reddi, PhD, FACMG
 American College of Medical Genetics
 and Genomics
 USA

Avni Santani, PhD, FACMG
 Association for Molecular Pathology
 USA

Francisca Reyes Turcu, PhD
 FDA Center for Devices and
 Radiological Health
 USA

Karl V. Voelkerding, MD
 College of American Pathologists
 USA

Expert Panel on Molecular Diagnostics

Expert panel volunteers support the development of CLSI documents by providing technical expertise in specialty areas. Expert panel members are listed by area of expertise on the CLSI website: <https://clsi.org/standards-development/expert-panels/>

Staff

Clinical and Laboratory Standards
 Institute
 USA

Katharine I. Castagna, MS,
 MLS(ASCP)CT, MB
Program Manager

Laura Martin
Editorial Manager

Catherine E.M. Jenkins, ELS
Editor

Kristy L. Leirer, MS
Editor

Lisa M.W. Walker, MS, ELS
Editor

Acknowledgment

CLSI, the Consensus Council, and the Document Development Committee on Nucleic Acid Sequencing gratefully acknowledge the following volunteers for their important contributions to the revision of this guideline:

Sivakumuran Theru Arumugam, PhD,
FACMG
Phoenix Children's Hospital
USA

Rebecca Hutchins, MS
Booz Allen Hamilton
USA

Kara Norman, PhD
Thermo Fisher Scientific
USA

Arthur Martin Baca, MD, PhD
Guardant Health, Inc.
USA

Elizabeth Duffy Hynes, BS
Partners Healthcare Personalized
Medicine
USA

Jillian Murrell, PhD
University of Pennsylvania
USA

Pinar Bayrak-Toydemir, MD, PhD
University of Utah
USA

Dan Jones, MD, PhD
Association for Molecular Pathology
USA

Jennifer J. Schiller, PhD, D(ABHI)
Blood Center of Wisconsin
USA

Mark J. Bowser, MS, MPH, MLS(ASCP)^{CM}
Partners HealthCare Personalized
Medicine
USA

Sabah Kadri, PhD
Ann & Robert H. Lurie Children's
Hospital of Chicago
USA

Timothy R. Schwartz, PhD
Roche Diagnostics
USA

Jillian Buchan, MS, PhD, FACMB
University of Washington
USA

Bryan Krock, PhD, FACMG
The Children's Hospital of Philadelphia
USA

Junaid Shabbeer, PhD, FACMG
Roche Sequencing Solutions
USA

Dora Dias-Santagata, PhD, FACMG
Massachusetts General Hospital,
Harvard School of Medicine
USA

Niklas Krumm, MD, PhD
University of Washington
USA

Birgitte Simen, PhD
Ginkgo Bioworks
USA

Jianli Dong, MD, PhD, FACMG
The University of Texas Medical
Branch
USA

Marco Leung, PhD, FACMG
Nationwide Children's Hospital
USA

Cynthe Sims, PhD, HCLD(ABB)
Torreyana Corporation
USA

James Duke, PhD
The Children's Hospital of Philadelphia
USA

You Li, PhD
Thrive, An Exact Sciences Company
USA

Zivana Tezak, PhD
FDA Center for Devices and
Radiological Health
USA

Deborah Ferriola, BS
The Children's Hospital of Philadelphia
USA

Meredith Halks Miller, MD
GRAIL, Inc.
USA

Patrik Vitazka, MD, PhD, FACMG
Roche
USA

Tina Hambuch-Hawk, PhD, FACMG,
CGMB, NYCOQ
Invitae
USA

Dimitri Monos, PhD
University of Pennsylvania; The
Children's Hospital of Philadelphia
USA

Richard Y. Zhao, PhD
University of Maryland School of
Medicine
USA

Contents

Abstract	ii
Committee Membership	iii
Foreword	vii
Chapter 1: Introduction	1
1.1 Scope	2
1.2 Background	2
1.3 Standard Precautions	3
1.4 Terminology	3
Chapter 2: Next-Generation Sequencing Test Development Lifecycle	15
Chapter 3: Application-Agnostic Considerations	19
3.1 Test Familiarization	20
3.2 Test Design, Development, and Optimization	21
3.3 General Paradigms for Analytical Test Validation, Revalidation, and Verification	22
3.4 Results Confirmation	26
3.5 Quality Management System	27
3.6 Bioinformatics Infrastructure Considerations	29
Chapter 4: Clinical Applications of Sequencing	41
4.1 Sanger Sequencing	42
4.2 Next-Generation Sequencing for Hereditary Disorders and Tumors	43
4.3 Human Leukocyte Antigen Sequencing	70
4.4 Circulating DNA Applications: Noninvasive Prenatal Testing and Liquid Biopsy	81
4.5 RNA Sequencing	92
Chapter 5: Conclusion	101
Chapter 6: Supplemental Information	103
References	104
Additional Resources	119
Appendix A. Next-Generation Sequencing General Recommendations and Guidelines	120
Appendix B. Test Design and Optimization Additional Details	124
Appendix C. Somatic Applications Examples	132
Appendix D. Interpretation and Reporting Additional Details	134
The Quality Management System Approach	148

Foreword

Sequencing-based clinical tests have existed for three decades, evolving from the single-gene tests used in the late 1980s to the whole-genome tests currently in use. The introduction of next-generation sequencing (NGS) catalyzed this evolution. Increasingly, NGS is replacing Sanger sequencing, particularly when examining a large number of genes is critical for maximum clinical utility. Today, NGS is firmly established in the medical management of hereditary disorders, especially those with clinical and genetic heterogeneity, as well as in tumor testing (ie, somatic NGS). Laboratory and medical practices for these clinical applications are relatively mature, and guidance from several professional societies and other expert groups is available (see Appendix A).

More recently, NGS has been used in additional areas of clinical practice, including human leukocyte antigen typing and noninvasive prenatal testing of fetal DNA in maternal blood to detect the presence or absence of select pathogenic variants in the fetus. Furthermore, new approaches provide additional opportunities for use in clinical areas in which NGS-based testing is already established. Innovative applications include RNA-based NGS (ie, RNA sequencing) to detect gene fusions and liquid biopsy (ie, DNA-based NGS) to detect genetic alterations in circulating tumor DNA in peripheral blood.

Although NGS applications have undergone major technical simplifications, clinical implementation continues to be challenging. Additional guidance is needed to ensure the technical and clinical validity of NGS tests. Guidance is increasingly important because genomic testing is being commoditized, and test developers vary in their interpretation and implementation of existing regulatory frameworks.

Overview of Changes

This guideline replaces the previous edition of the approved guideline, MM09-A2, published in 2014. MM09-A2 introduced NGS as a new technology. This edition has been updated beyond an introduction of NGS technology to provide practical use case and implementation guidance, as well as instructions that cover each step of the clinical NGS test development lifecycle. Several changes were made in this edition, including:

- Providing step-by-step recommendations on designing, developing, validating, and implementing a clinical NGS test
- Adding clear and specific instructions on performing steps in the clinical NGS test lifecycle
- Presenting an application-driven approach
- Providing educational use case examples, supplemented by instructional worksheets
- Updating appendixes with additional details on steps in the test development process and specific applications

NOTE: The content of this guideline is supported by the CLSI consensus process and does not necessarily reflect the views of any single individual or organization.

KEY WORDS

bioinformatics

design

development

germline

human leukocyte antigen

implementation

liquid biopsy

next-generation sequencing

noninvasive prenatal testing

optimization

quality management

RNA sequencing

somatic

validation

verification

This page is intentionally left blank.

Chapter ①

Introduction

Human Genetic and Genomic Testing Using Traditional and High-Throughput Nucleic Acid Sequencing Methods

1 Introduction

1.1 Scope

This guideline covers nucleic acid sequencing applications currently in clinical use: medical management of hereditary disorders, solid tumor and hematological malignancy testing, human leukocyte antigen (HLA) typing, noninvasive prenatal testing (NIPT), liquid biopsy, and RNA sequencing (RNAseq) applications. Most of the content in this guideline focuses on next-generation sequencing (NGS), which is the predominant platform in current use. Sanger sequencing continues to be used for certain clinical applications, so guidance on Sanger sequencing is also included. This guideline also provides introductory information on the management of computational and/or bioinformatics aspects of NGS, because these concepts are fundamental yet somewhat novel for the clinical testing community. Detailed guidance on bioinformatics will be provided in a forthcoming CLSI document.

MM09 does not cover microbial or infectious diseases applications. Detailed guidance on NGS-based infectious diseases testing is provided in CLSI document MM24.¹ This guideline also does not cover validation of confirmatory testing or mitochondrial DNA testing for inherited disorders.

This guideline is intended for developers of sequencing-based clinical tests (both Sanger sequencing and NGS), including manufacturers of commercially distributed *in vitro* diagnostic (IVD) devices and developers of laboratory-developed tests (LDTs). IVD device manufacturers might be subject to additional quality system requirements. For example, design controls are not included in this guideline, but they are well described in existing literature.^{2,3}

1.2 Background

MM09 provides step-by-step guidance on development of clinical sequencing tests. Topics include test familiarization, design, development, and optimization, as well as analytical validation and quality management. This guideline specifically focuses on explaining **how to implement** sequencing technologies in a clinical setting (ie, how to develop and analytically validate sequencing-based clinical tests) rather than providing in-depth education on **how they work**, because a large body of literature covers the latter. **NOTE:** This guideline refers to US Food and Drug Administration (FDA) requirements. FDA requirements do not apply to test developers outside the United States.

MM09 contains:

- Traditional, text-based chapters that outline the clinical test development process and provide a high-level introduction, background information, and necessary context for the test developer
- A link to instructional worksheets (shared resources with the College of American Pathologists) that provide additional information and concrete guidance, including forms adaptable by the user and educational examples (see Additional Resources)
- Appendixes with additional resources and detailed information, including descriptions of technology platforms