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CLIA lab certification and CLIA test validation

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National Human Genome Research Institute The Forefront of Genomics[®]

Current disclosures

- NIH/NHGRI research funding (intramural)
- As outside activities: Editor-in-Chief, American Journal of Medical Genetics; textbook/publishing royalties through Wiley, Inc.



My (relevant) background

- Previously ran ~500-person CLIA/CAP-certified clinical genetics/genomics lab (which also conducted large-scale genomic research); ~100K samples/year (>200K tests/year), primarily exomes/genomes/large panels
- PI on several research studies involving WGS in healthy and affected individuals, involving an array of clinical and non-clinical analyses



Agenda

- 1. Certification and validation
- 2. Processes and examples



Certification and validation



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CLIA: see https://www.cms.gov/regulations-and-guidance/legislation/clia





Some basic categories of "official" clinical lab certification in genetics/genomics

- CLIA (CMS, but involves FDA, CDC; various types of certification)
- CAP (peer-based inspection process)
- FDA (options for IDE research; specific assays; overarching approach continues to be discussed, and some major sequencers may proceed down this route)
- Some states require additional steps or can be exempt (eg, New York, Washington)

Licensing



Laboratories within the US are certified under the Clinical Laboratory Improvement Amendments (CLIA) and accredited by the College of American Pathologists (CAP). genetic testing in the US as well as internationally.





usiness & Policy Technology Research Diagnostics

e » Tools & Technology » Sequencing

AMP, CAP Release Guidelines for Use of In Silico Approaches for **Validating NGS Data Pipelines** Oot 18, 2022 | Andrew P. Han



New York pprove

CLIA framework

- Required if **individual** results are returned (not related to whether patients are charged for results or if sequencing is done on a purely clinical or research basis)
- Focuses on analytic validity
- Test categories:

NIH See:

O Waived tests - simple, no risk of harm
O Moderate/high complexity - includes genomics
O Specific specialty areas - includes cytogenetics

NHGRI https://www.genome.gov/Pages/PolicyEthics/GeneticTesting/The_CLIA_Framework.pdf

Components

- Proficiency testing (PT) (nonspecialty areas: biannual self-testing, PT exchange)
- Quality systems

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- (FDA-related requirements)
- Personnel: Lab director, <u>technical</u> supervisor, <u>clinical</u> consultant, general supervisor
- CLIA inspections (biennial, usually w/ 2 weeks notice) – paperwork/documentation, procedure manuals, personnel interviews, real charts





Reality at a mid-size genomics lab (~500 staff, ~100K samples/year)

- Lab director (for sign-off)
- QC director
- 3-4 staff on lab QC team
- 1 person for NY state requirements
- 1 person on "launch team"
- Integrated with other teams per requirements and by necessity (wet lab, bioinformatics, IT, individual clinical groups, legal/compliance, etc.)

Processes and examples



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Case study

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- A research study plans to sequence the genomes of 10,000 people
- Consent includes possible receipt of individual, clinically relevant results from genome sequencing



1. Pure CLIA (or similar) model

- Everything (e.g., DNA extraction, sequencing, orthogonal confirmation, etc.) handled in CLIA compliant manner from the outset
- Logistic frontloading
- Easier and more flexible downstream (e.g.,may obviate orthogonal confirmation in some situations), but may be difficult to set up and maintain
- Doesn't offer a path for many
- "historic" biobanks

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2. More frequent model (research \rightarrow CLIA)

- Most common model
- Logistic backloading (doesn't require CLIAcompliant sample handling from the start)
- Recontact typically involves letting participant know there may be a finding of interest that requires technical validation, then collecting a **new sample for variantspecific, CLIA-based orthogonal confirmation**
- Requires ongoing communication





Example of hybrid approach

- Geisinger MyCode project
- Samples collected in CLIAcompliant manner (since 2015)
- Analyses done on research basis
- Data and CLIA-compliant samples shared with clinical lab for confirmation and reporting
- If samples had not been collected in CLIA-compliant manner, new samples could be obtained for confirmation/reporting



Kelly et al., AJMG part C 2021

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A few lessons learned...

- Sometimes, **partnering** with a CLIA/etc.-certified "clinical lab" with lots of experience in this area (and data) can be helpful
 - Logistics of sample confirmation and regulatory management
 - Databases of variants for comparison
- Don't forget **non-SNVs**
 - Medically important, even if more challenging
 - Often requires different assays
- Sample swaps!

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• Challenging participant-specific situations **will** arise: have an experienced advisory panel on hand!

Questions/Discussion







Performance establishment

CLIA performance specifications

- Accuracy
- Precision
- Analytical sensitivity and specificity
- Reportable range of results
- Reference intervals
- Any other required performance characteristics
- (No specific clinical validation requirements)
- (Reagent stability addressed through calibration, QC)

See: https://www.cdc.gov/labtraining/onelab/docs/6-30_EventSlideD.pdf



"Clinical" versus research

Genetics	
inMedicine	2

www.nature.com/gim			
	Check for updates		

ACMG STATEMENT

ACMG SF v3.0 list for reporting of secondary findings in clinical exome and genome sequencing: a policy statement of the American College of Medical Genetics and Genomics (ACMG)

David T. Miller^{1,20}, Kristy Lee^{2,20}, Wendy K. Chung³, Adam S. Gordon⁴, Gail E. Herman⁵, Teri E. Klein⁶, Douglas R. Stewart⁷, Laura M. Amendola⁸, Kathy Adelman⁹, Sherri J. Bale¹⁰, Michael H. Gollob¹¹, Steven M. Harrison¹², Ray E. Hershberger¹³, Kent McKelvey¹⁴, C. Sue Richards¹⁵, Christopher N. Vlangos¹⁶, Michael S. Watson¹⁷, Christa Lese Martin¹⁸ and ACMG Secondary Findings Working Group¹⁹*

Genetics in Medicine (2021) 23:1381-1390; https://doi.org/10.1038/s41436-021-01172-3

Additionally, the recommendation to return SF only applies in the clinical setting. The decision to return SFs in a research setting is left to the research team and relevant local institutional review board (IRB).

