Paper Summary

<!--META_START-->

Title: A semiquantitative metric for evaluating clinical actionability of incidental or secondary findings from

Authors: Jonathan S. Berg, Ann Katherine M. Foreman, Julianne M. O'Daniel, Jessica K. Booker, Lacey

DOI: 10.1038/gim.2015.104

Year: 2016

Publication Type: Journal Article

Discipline/Domain: Genomic Medicine / Medical Genetics

Subdomain/Topic: Clinical Actionability Assessment in Genomic Sequencing

Eligibility: Eligible

Overall Relevance Score: 95

Operationalization Score: 100

Contains Definition of Actionability: Yes

Contains Systematic Features/Dimensions: Yes

Contains Explainability: Yes

Contains Interpretability: Yes

Contains Framework/Model: Yes

Operationalization Present: Yes

Primary Methodology: Conceptual Framework Development and Application

Study Context: Evaluation of incidental/secondary findings in clinical genome-scale sequencing

Geographic/Institutional Context: University of North Carolina at Chapel Hill, USA

Target Users/Stakeholders: Clinical geneticists, genomic testing laboratories, healthcare providers, policy

Primary Contribution Type: Framework/method for assessing clinical actionability of gene-disease pairs

CL: Yes

CR: Yes

FE: Yes

TI: Partial

EX: Yes

GA: Partial

Reason if Not Eligible: N/A

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Title:

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A semiquantitative metric for evaluating clinical actionability of incidental or secondary findings from geno
**Authors:**
Jonathan S. Berg et al.
**DOI:**
10.1038/gim.2015.104
**Year:**
2016
**Publication Type:**
Journal Article
**Discipline/Domain:**
Genomic Medicine / Medical Genetics
**Subdomain/Topic:**
Clinical Actionability Assessment in Genomic Sequencing
**Contextual Background:**
The paper addresses the challenge of systematically evaluating the clinical actionability of genomic varia
**Geographic/Institutional Context:**
University of North Carolina at Chapel Hill, USA
**Target Users/Stakeholders:**
Clinical geneticists, genetic counselors, genomic laboratories, healthcare providers, professional organization
**Primary Methodology:**
Conceptual framework and scoring metric development, applied analysis of multiple gene lists
**Primary Contribution Type:**
Development and validation of a semiquantitative scoring framework for clinical actionability
## General Summary of the Paper
This paper presents a semiquantitative metric for evaluating the clinical actionability of incidental or secon
## Eligibility
Eligible for inclusion: **Yes**
## How Actionability is Understood
Actionability is conceptualized as a **continuum** rather than a binary state, integrating the potential seven
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- > "The LVBC developed a semiquantitative metric for determining the clinical actionability of gene-diseas
- > "The subcategories... approximate the clinical utility of revealing incidental/secondary findings in a pres

- ## What Makes Something Actionable
- High severity of the potential health outcome
- Moderate to high likelihood of disease manifestation
- Availability of effective preventive or therapeutic interventions
- Low burden or acceptable risk of intervention
- Substantial and reliable knowledge base supporting decision-making

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- ## How Actionability is Achieved / Operationalized
- **Framework/Approach Name(s):** Semiquantitative Metric for Clinical Actionability
- **Methods/Levers:** Five criteria scored 0-3 (severity, likelihood, efficacy, burden, knowledge base)
- **Operational Steps / Workflow:** Evidence review → Assign scores for each criterion → Consensus sc
- **Data & Measures:** Literature from OMIM, GeneReviews, PubMed, clinical guidelines
- **Implementation Context:** NCGENES project and ACMG incidental findings recommendations
- > "All five criteria are scored on a scale of 0–3... The outcome and intervention are defined in advance...
- > "The LVBC chose to consider genes with a score ≥11... as meeting the threshold of actionability." (p. 4

- ## Dimensions and Attributes of Actionability (Authors' Perspective)
- **CL (Clarity):** Yes Clarity through structured scoring definitions (Table 1)
- **CR (Contextual Relevance):** Yes Explicit to presymptomatic, incidental/secondary findings contex
- **FE (Feasibility):** Yes Evaluated through "burden of intervention" score
- **TI (Timeliness):** Partial Implied in presymptomatic intervention consideration
- **EX (Explainability):** Yes Transparent, evidence-based scoring with defined criteria
- **GA (Goal Alignment):** Partial Implicit alignment with clinical utility and patient benefit
- **Other Dimensions Named by Authors:** Knowledge base strength

- ## Theoretical or Conceptual Foundations
- Clinical utility concepts from genomic medicine
- Evidence-based assessment models
- Prior ACMG deliberative consensus recommendations

Indicators or Metrics for Actionability

- Total score (0-15)
- Threshold ≥11 for high actionability
- Subscores for severity, likelihood, efficacy, burden, and knowledge base

Barriers and Enablers to Actionability

- **Barriers:** Limited evidence base; subjective burden assessment; rare diseases with insufficient pene
- **Enablers:** Structured metric; multidisciplinary consensus; adaptability to different contexts

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Relation to Existing Literature

The paper builds on and critiques earlier expert consensus models like the ACMG recommendations, add

Summary

This paper delivers a rigorous, transparent framework for assessing the clinical actionability of incidental

Scores

- **Overall Relevance Score:** 95 Clear conceptualization of actionability as multidimensional, detailed
- **Operationalization Score:** 100 Fully developed metric with applied examples and scoring workflow

Supporting Quotes from the Paper

- "Actionability is a continuum, not a binary state." (p. 468)
- "The LVBC established five core characteristics of clinical actionability..." (p. 469)
- "The LVBC chose to consider genes with a score ≥11... as meeting the threshold of actionability." (p. 47)

Actionability References to Other Papers

- ACMG Recommendations for Reporting of Incidental Findings (Green et al., 2013)
- Evidence-based Genomic Applications in Practice and Prevention Working Group (Goddard et al., 2013
- NCGENES project preliminary outputs (Berg et al., 2013)