

Assessing the Clinical Validity of Genes Implicated in Hereditary Breast and Ovarian Cancer Susceptibility Using the ClinGen Framework

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Clinical Validity: What do we do with the information?

ON CANCER

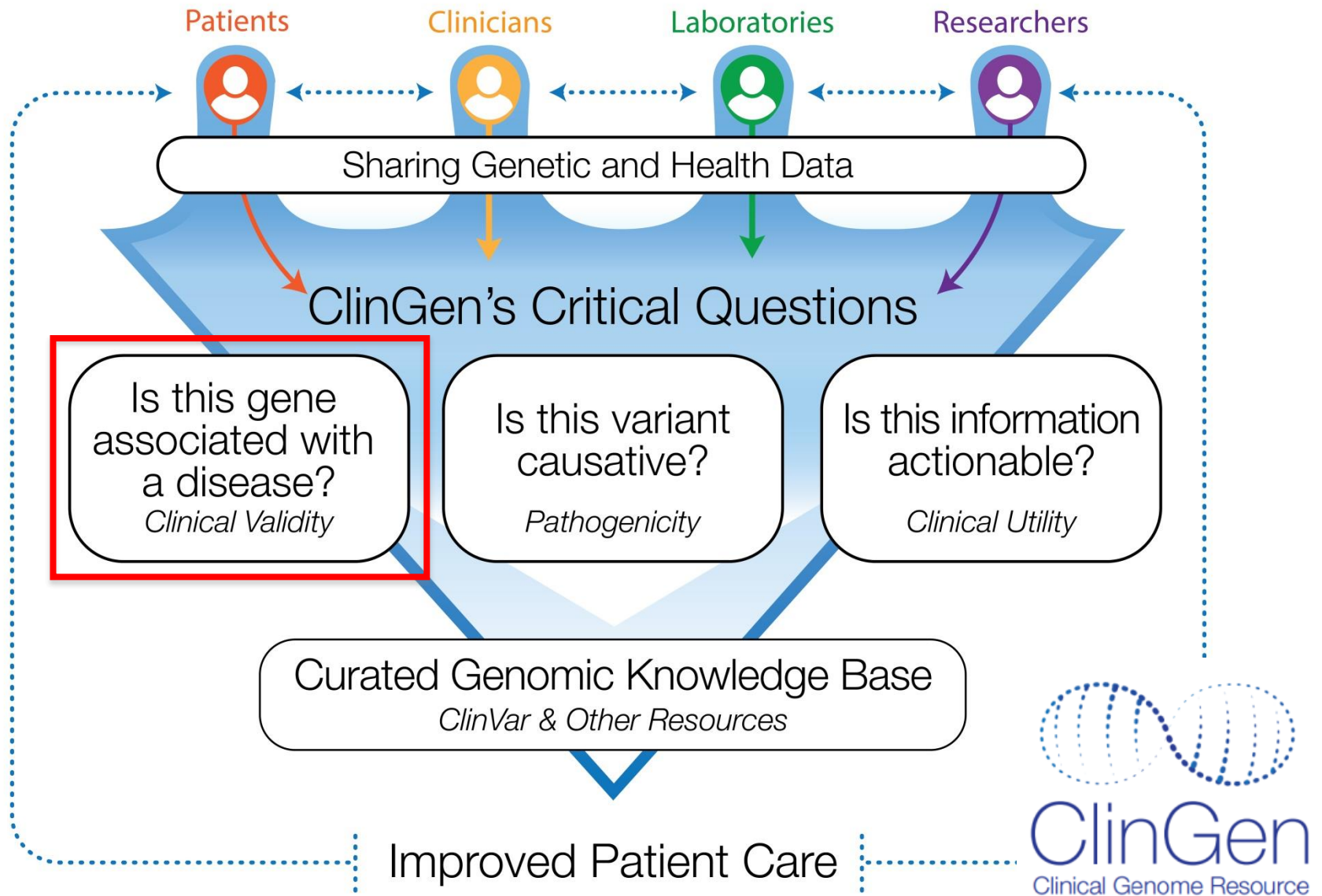
When to Consider Multigene Panel Testing

By [Andrea Peirce](#) on Thursday, April 23, 2015



Multigene panel testing enables people to learn about not just one but many inherited mutations at once.

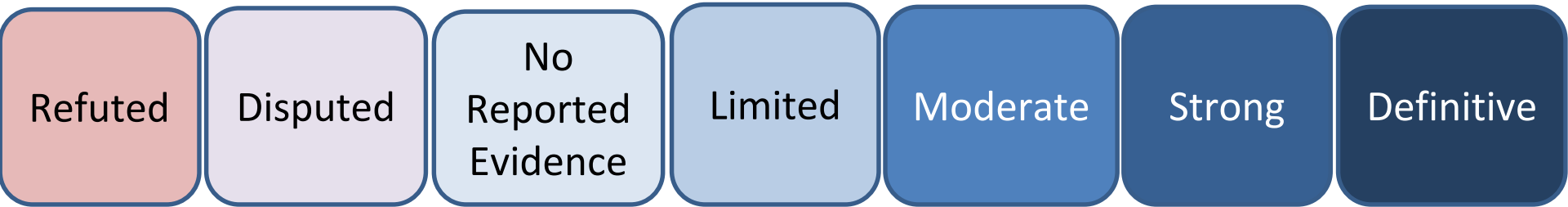
The Clinical Genome Resource



ClinGen Hereditary Cancer Working Group

- Paraganglioma/Pheochromocytoma
 - 21 genes
- Pancreatic Cancer
 - 20 genes
- Breast/Ovarian Cancer
 - 34 genes
- Polyposis/Colorectal Cancer
 - 37 genes

Clinical Validity Framework





Five Key Evidence Types

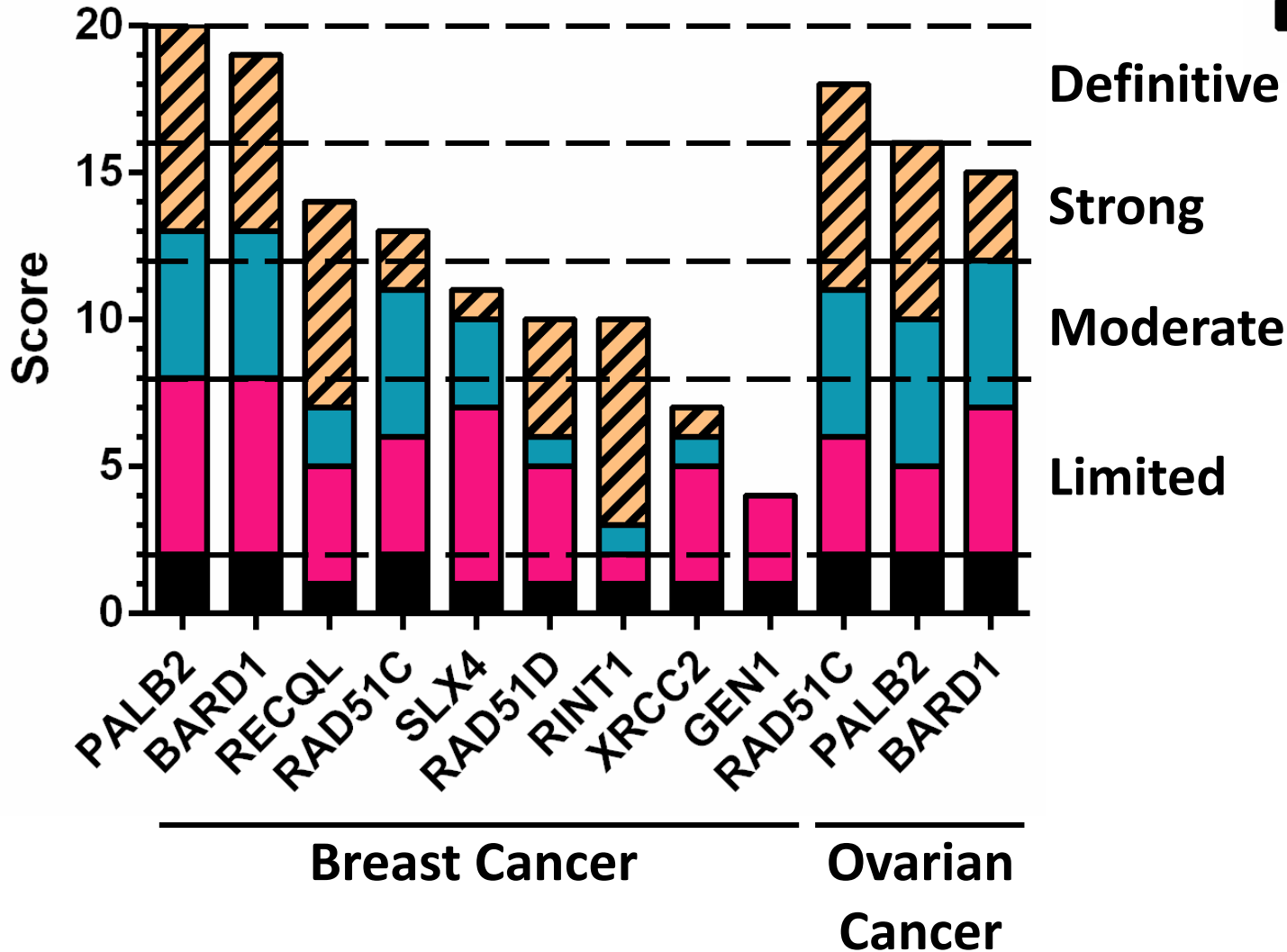
1. Genetic Evidence (proband)
2. Experimental Evidence (functional studies)
3. Publications
4. Time (first clinical publication)
5. Strength of conflicting evidence (contradictory evidence)

Clinical Validity Summary Matrix

| Assertion Criteria | Description | Score |
|--|---|------------|
| Genetic Evidence | Probands from case reports and case-control studies | 0-7 points |
| Experimental Evidence | Gene-level experimental evidence | 0-6 points |
| Publications | Quantity of clinical literature | 0-5 points |
| Time | Years since first clinical publication | 0-2 points |
| Is there valid contradictory evidence? | | |

Counting all genetic evidence from case-control studies and case reports

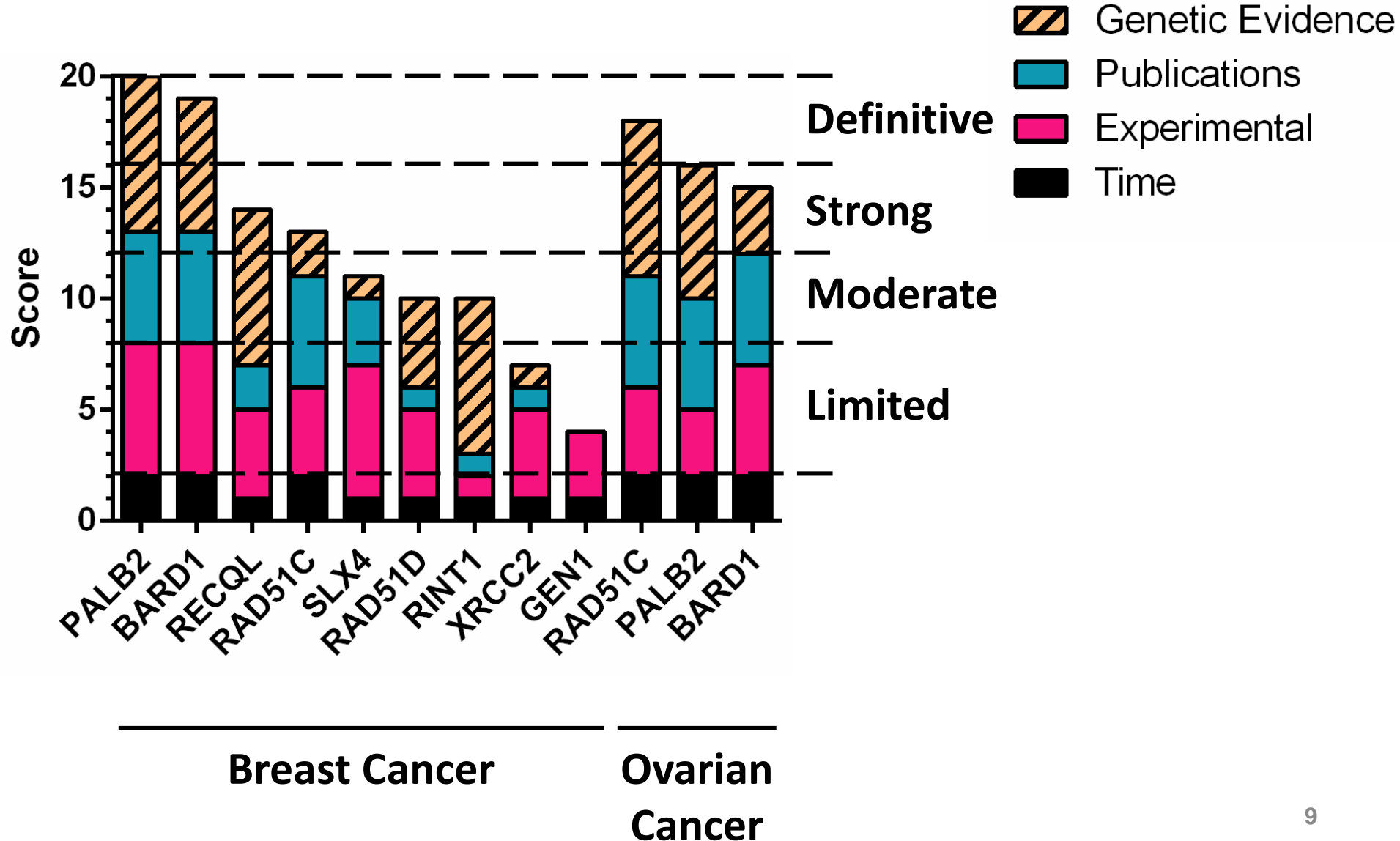
-  Genetic Evidence
-  Publications
-  Experimental
-  Time



Challenges in evaluating Assertion Criteria

- Scoring individual probands may inflate the scores if stringent criteria for pathogenicity (e.g. segregation) are not applied.
- Using only case-control studies establishes a more “conservative” threshold for genetic evidence.

Correcting the Genetic Evidence score for case-control studies reduces the evidence of a gene-disease association



Conclusions

- Assessing the Mendelian contributions to common diseases requires a hybrid approach to evaluate evidence from probands/kindreds and case-control studies.
- Many genes that are included on hereditary breast cancer panels may have limited clinical validity.
- A systematic method is needed to evaluate the strength of gene-disease associations and provide transparent assessments of clinical validity.



Acknowledgments

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