Cost-Effectiveness of Population Genomic Screening

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Reimbursement for healthcare technologies

- 1. Increasing push for value in healthcare
- 2. Difficult to quantify, but established methods
- 3. Approaches are evolving to capture broader aspects of value
- 4. In the US, formal cost-effectiveness analyses do not directly influence reimbursement decisions, but provide context and inform discussions

Cost-Effectiveness



(-) \triangle QALYs \rightarrow (+)

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Annals of Internal Medicine

Original Research

Population Genomic Screening for Three Common Hereditary Conditions

A Cost-Effectiveness Analysis

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CDC Tier 1 Conditions

Tier 1 Condition	Increased Risk For:	Risk-Reduction
Hereditary Breast and Ovarian Cancer	Breast cancer, Ovarian cancer, Other cancers	Mammography <u>+ MRI</u> , Mastectomy, Salpingo-Oophorectomy
Lynch Syndrome	Colorectal cancer, Endometrial cancer, Other cancers	Increased colonoscopy surveillance
Familial hypercholesterolemia	Myocardial infarction, Stroke	Moderate to high-intensity statin therapy



https://www.cdc.gov/genomics/implementation/toolkit/tier1.htm

Tier 1 Model Features





Inputs: Costs

Parameter	Value
Targeted Next Generation Sequencing (NGS)	\$250
Sanger confirmation Genetic Counseling	\$250

Inputs: uptake of recommended interventions

Risk-reducing intervention uptake

HBOC

	200	
	Relative mortality reduction: early- vs. late- stage breast cancer	0.94
	Cumulative mastectomy by age 30 y, %	15
	Cumulative mastectomy by age 40 y, %	30
	Cumulative mastectomy by age 50 y, %	36
	Cumulative mastectomy by age 60 y, %	36
	Cumulative salpingo-oophorectomy by age 30 y, %	8
	Cumulative salpingo-oophorectomy by age 40 y, %	48
	Cumulative salpingo-oophorectomy by age 50 y, %	68
LS	Cumulative salpingo-oophorectomy by age 60 y, %	74
	Increased colonoscopy surveillance, ages 20-75 y, %	80
FI	н	
	Proportion of tested persons who take statins, %	60

Cascade testing	
Proportion of persons who inform their family	0.70
members	
Proportion of family members who get tested	0.20
Proportion of family members with variants	0.50

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Individual model results

Potentially cost-effective not cost-effective

Model	30 years old	50 years old
HBOC*	\$87,700/QALY 🗸	\$482,100/QALY 🗙
LS	\$132,200/QALY 🗙	\$140,400/QALY 🗙
FH	\$206,700/QALY 🗙	\$463,500/QALY 🗙

*females

Combined results: Incremental QALYs per 100,000 screened



Guzauskas et al, Annals Int Med, May 2023

Cost effectiveness



Guzauskas et al, Annals Int Med, May 2023

But what if...

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Scenarios for	Scenario Inputs					Results per 100 000 30-Year-Old Persons			
Screened 30-Year-Old Persons	Assay Cost, \$	Follow-up Multiplier	Cascade Testing Uptake, %*	Prior Knowledge of Variant, %†	Total Variant Proportion, %‡	Incremental Cost (95% UI), \$ (millions)	Incremental QALYs (95% UI)	ICER (95% UI), \$/QALY	Cost-Effectiveness Probability, %§
Main (base-case) analysis	250	1	14	9	1.5	33.9 (27.0-41.1)	495 (401-757)	68 600 (41 800-88 900)	99.4
Societal perspective	250	1	14	9	1.5	25.6 (16.4-40.3)	495 (401-757)	51 700 (24 200-106 200)	99.9
Lower genetic assay cost	100	1	14	9	1.5	19.6 (15.1-24.4)	495 (401-757)	39 700 (23 500-51 800)	100
Higher genetic assay cost	500	1	14	9	1.5	57.8 (45.3-70.7)	495 (401-757)	116 800 (71 200-154 000)	44
Lower adherence to follow-up	250	0.5	14	9	1.5	31.2 (24.9-37.7)	292 (228-436)	106 800 (66 700-141 700)	57
Higher adherence to follow-up	250	1.2	14	9	1.5	35.0 (28.0-42.1)	570 (461-883)	61 400 (37 000-77 90 0)	100
Without cascade testing	250	1	0	9	1.5	32.0 (25.2-39.0)	436 (347-692)	73 300 (42 000-96 100)	98
Higher uptake of cascade testing	250	1	35	9	1.5	36.9 (29.3–44.7)	582 (478-865)	63 400 (41 100-79 700)	100
Low prior knowledge	250	1	14	7	1.5	34.5 (27.1-41.7)	512 (413-780)	67 400 (40 700-88 000)	99.4
High prior knowledge	250	1	14	11	1.5	33.4 (26.1-40.4)	477 (386-739)	69 900 (41 300-93 000)	98.9
Low variant prevalence	250	1	14	9	1.1	31.4 (24.6-37.9)	371 (303–576)	84600 (50 800-108 100)	93
High variant prevalence	250	1	14	9	2.0	36.5 (29.1-44.2)	618 (501-945)	59 000 (35 900-75 400)	100

Table 3. Base-Case and Scenario Analysis Results

False reassurance

Potential Harm Related to False Reassurance

Under the assumption that 10% of 30-year-olds without a variant subsequently avoid routine disease screening because of receipt of a negative genomic screening result, a loss of 0.05 QALY in this population would lead to genomic screening having no incremental health benefit.

Polygenic risk scores – economic value of population screening?

- Prevalence of 'high-risk' is greater than monogenic conditions
- Lifetime risk lower
- Multiple conditions

PRS vs. Tier-1

- Prevalence ~10-20x higher
- Effect size ~20-30x lower(!)
- PRS: Prevalence ~20%, Benefit ~0.03 QALYs
- Cost effectiveness likely above threshold of \$100K/QALY (not cost effective)

Tier-1 cost-effectiveness 'landscape'



PRS cost-effectiveness 'landscape'



Newborn screening

- Large number of rare conditions
- Actionability variable
- Different policy context

Implication #1

Prevalence drives economic value

- Include the most prevalent conditions
- Combine conditions

Implication #2

Clinical action is required for 'traditional' economic value

 Focus on clinical actionability for building value story and driving reimbursement

Clinical actions – eMERGE consortium

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Prospective, multi-site study of healthcare utilized monogenic findings from clinical sequencing	ation after actionable	
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Published: October 25, 2023 • DOI: https://doi.org/10.1016/j.ajhg.2023.10.0	06 • (E) Check for updates	

Summary

Keywords

References

Article info

Summary

As large-scale genomic screening becomes increasingly prevalent, understanding the influence of actionable results on healthcare utilization is key to estimating the potential long-term clinical impact. The eMERGE network sequenced individuals for

Implication #3

Screening should be efficient and relatively inexpensive

- Public or private sector reimbursement?
- Delivery and education

Summary

- Population screening for CDC Tier-1 conditions provides an excellent model for population genomic screening
- CDC Tier-1 screening likely has beneficial risk-benefit profile and provides good economic value, <u>but</u>:
 - Need further clarity on behavior of those with and <u>without</u> a variant
 - Evidence on all aspects in <u>underserved populations</u>, diverse ancestries
 - Implementation outcomes
- <u>Combining conditions</u> is essential for economic value, but <u>restricting</u> to those with good clinical or patient-centered value is critical
- Genomic population screening applications <u>will vary dramatically</u> in their economic value and evidence requirements

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