

[®] Cores in Support of Impactful Innovation: the Healthy Oregon Project

JACKILEN SHANNON, PHD, & CHRIS HARRINGTON, PHD

WESTERN ASSN OF CORE DIRECTORS

ANNUAL MEETING 2023



Healthy Oregon Project (HOP) A Resource for Research

Presented by: Jackilen Shannon, PhD

I have no financial disclosures, real or perceived conflicts with the work I am presenting today.

What is the Healthy Oregon Project?

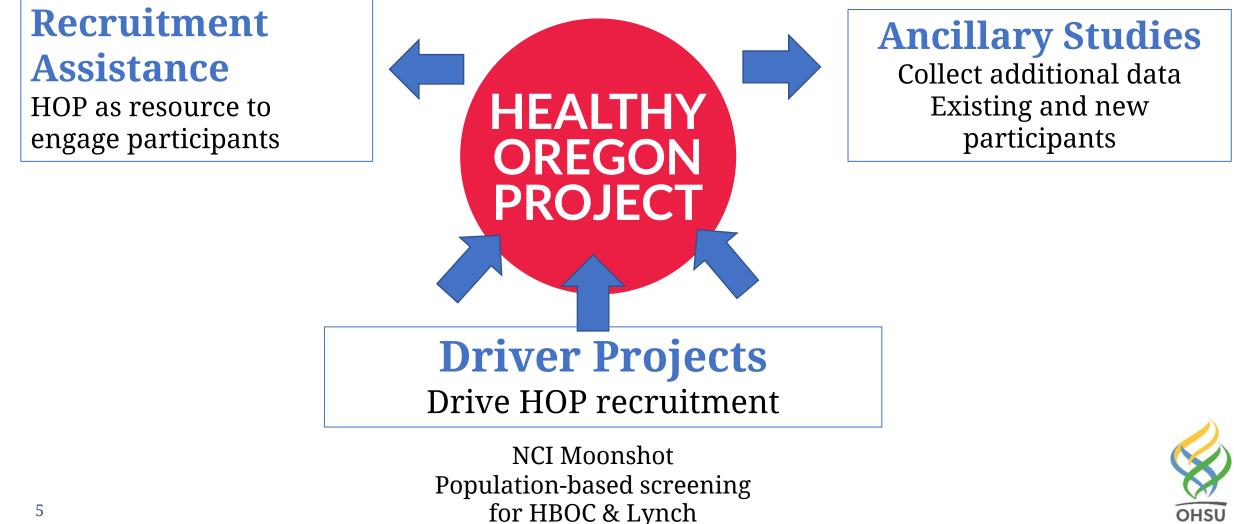
HOP is a research project that aims to involve >100,000 Oregonians in a large, sustainable cohort.

To learn about how a person's **genetics, health, behaviors** and **environment**, affect **risks** of cancer and chronic diseases.

Data saved to a **secure and privacyprotected repository** of information that can be used to answer many different questions about health.



Supporting Collaboration and Innovation by Design



HOP Driver Scientific Project #1 Funded as a Moonshot Project by the National Cancer Institute:

MPI – Paul Spellman, Jackilen Shannon

Question: Is screening the general population effective and sustainable.

- *Effective*== do screened individuals alter their health care to reduce their risk of dying from cancer
- *Sustainable* == are the costs per QALY under \$50,000 and is the health care system capable of meeting demand

Comparing HOP participants to patients receiving standard guideline based genetic screening, and existing cancer patients offered screening

- 1. Adherence to current guidelines for screening and prophylactic intervention.
- 2. Efficiency of cascade testing.
- 3. Costs and effectiveness, associated with genetic screening models.





HOP Genetic – Our First Driver Project WHY ?

~1 M People Have HBOC, Lynch or another Significant Inherited Cancer Syndrome

- Only about 20-30% are aware of their syndrome
- HBOC >50% lifetime risk of breast cancer
- HBOC > 30% lifetime risk of ovarian cancer. >20% die of ovarian cancer
- Lynch > 50% lifetime risk of colon cancer
- Knowing you are Lynch + can result in 14 QALY saved
- Currently only high-risk families are tested because of real and perceived costs and significant limitations in clinical service because of limited # of counselors
- Clinical trial to test efficacy of screening all adults for HBOC and Lynch



NCT04494945: Approaches to Identify and Care for Individuals With Inherited Cancer Syndromes

Aim: Evaluate the effectiveness and sustainability of heritable cancer syndrome testing in the two novel testing populations as compared to current practice.

- •Enrollment via Healthy Oregon Project App
- Population 1: Everyone
- Population 2: Those with cancer
- •Test: 90% sensitive. Very low-cost.
- •NOT for those who meet current genetic testing guidelines.

Success Will Change Standard of Care

WHAT Is HOP Screening For?

Inherited Cancer Syndromes (5-10% of all cancers)

Population-based **genetic testing** for known cancer-related genetic mutations.

- **32 genes** associated with increased risk of cancer
- Medically actionable per NCCN guidelines





HOP Gene Panel

Genes chosen based off of Color's/Myriad's Hereditary Cancer Panels and by HOP team

ATM	MEN1	BAP1	*Only 2 probes covering a specific site:
MITF*	RAD15C	BARD1	CDK4: 12:g.58145429-58145431 (codon 24)
MLH1	RAD15D	BMPR1A	MITF: chr3:g.70014091 POLD1:chr19:g.5090713 POLE:chr12:g.133250250 **Only probes covering areas known to contain large deletionsGREM1: 2 probes every 2.5KB from GREM1 5' UTR to SCG5 and coding exons of SCG5 and GREM1. See Myriad Approach Extra Region: Chr15:g.32988815-33021692 ***Exons 12-15 not analyzed in PMS2 due to high homology with pseudogene
MSH2	RB1	BRCA1	
MSH6	RET	BRCA2	
MUTYH	SMAD4	BRIP1	
NBN	STK11	CDK4*	
PALB2	TERF2IP	CDKN2A	
PMS2***	TERT	CHEK2	
POLD1*	TP53	POT1	Hg19 coordinates. Refseq Annotation
APC	POLE*	PTEN	
TSC1	TSC2		OREGON

PROJE

Return of Genetic Screening Results

NEGATIVE: provided via a **notification letter** that is provided **in the Healthy Oregon Project App**.

POSITIVE: any participant with a positive result will
be contacted by an OHSU genetic counselor.
Results will be explained in detail, including
recommendations during a phone consultation.



Receiving Positive Results

OHSU Genetic Counselor

HOP Participant Navigator Ryan Lutz







Amelia Mulford



Consult for Results

Resource Connector

How to participate in **HOP**



STEP 1: Download secure app on iOS/Android

STEP 2: Review information about **HOP** and **sign** an electronic **consent** in the app to participate

STEP 3: Create a secure account

STEP 4: Explore your options for participating (DNA screening & Surveys)

STEP 5: If you choose to donate your DNA for genetic screening, **request a HOP Kit and receive it in the mail.**



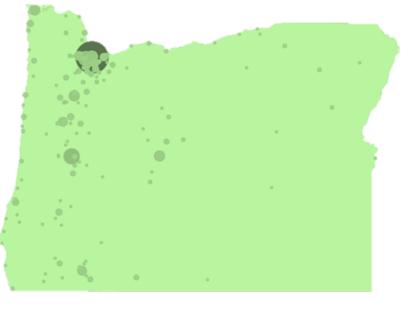
HOP app is HIPAA compliant!



How are we Doing? Enrollment as of 05/01/23= 38,140

- Pre-COVID 3,827 (~12months of recruitment)
 - Events and vending machines
- Post-COVID (since October 2020 ~30 months)
 - 34,313 participants enrolled
 - Social media, word-of-mouth, attendance at community events
 - 31,609 (92%) Consented to genetics screening and 29,880 (87%) requested a mailed kit
 - 23,415 Samples received in the lab (78% return rate)
 - 20,177 Tests results returned to date
 - 835 (4.1% -though anticipate closer to 5%) received genetic counseling for a positive result.

With over 40,000 participants enrolled in HOP since 2018 that brings us to over 1% of the adult population in Oregon.





University Shared Resources

Integrated Genomics Laboratory & the Healthy Oregon Project

OREGON

PROJECT

PRESENTED BY: CHRIS HARRINGTON, PHD

ASSOC DIRECTOR - INTEGRATED GENOMICS LAB/DIRECTOR - GENE PROFILING SHARED RESOURCE



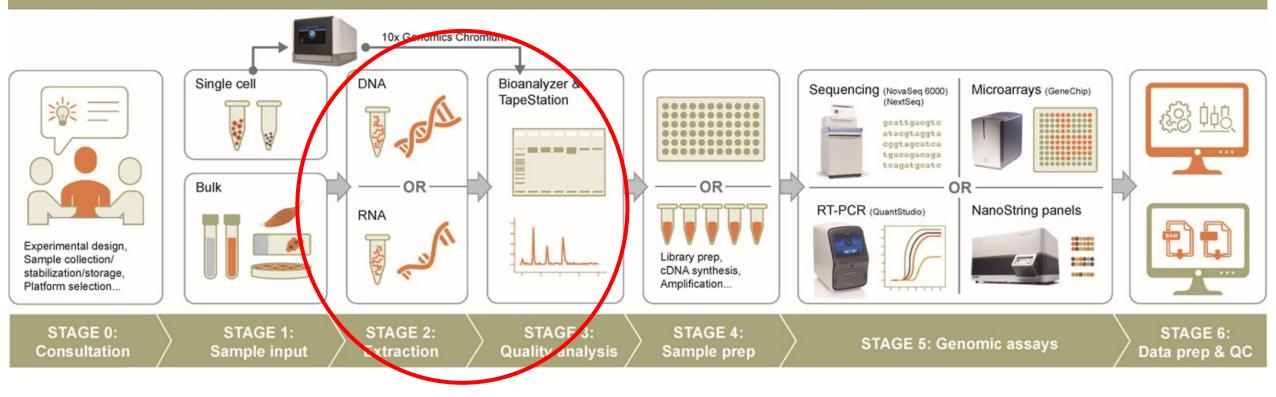
OCT 13, 2023

I have no financial disclosures, real or perceived conflicts with the work I am presenting today.

Integrated Genomics Laboratory

https://www.ohsu.edu/integrated-genomics-laboratory

INTEGRATED GENOMICS (Investigators can enter this workflow at any stage)



Coordinated pipeline of services which can be accessed at whatever point researcher requires

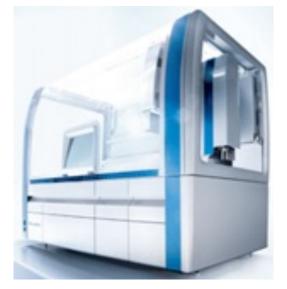
Integrated Genomics Laboratory (IGL) & the Healthy Oregon Project (HOP)

What HOP needed –

Inexpensive, reliable, high-throughput DNA extraction service for long term genetic screening project

Why reach out to IGL?

- Recent acquisition of high-throughput sample processing robot
- Expertise and experience of core staff
- Flexibility of core (compared to diagnostics lab)
- History of research support and collaboration between the IGL cores and Knight Cancer Institute researchers



QIAGEN QIAsymphony Isolation Robot Automated RNA/DNA extraction protocols



IGL/HOP Collaboration | The Plan

Use IGL resources to develop standardized DNA extraction workflow for HOP samples

Knowledgeable, experienced staff

BSL2 Hood

High-throughput equipment





Liquid handling robot



Fluorescent plate reader

Challenges:

Protocol development

- Match to saliva collection kit
- Cheap & fast

Clinical testing approved laboratory for the extractions

Dedicated staff for routine throughput

DNA extraction robot

IGL/HOP Collaboration | The Setup Team

IGL

Trevor McFarland





Sam Medica



Aaron Larson – IGL Project Coordinator (key for organizational support & communications)

HOP researchers & Knight Diagnostics Lab (KDL)

- Sarah McCabe, Manager, Genomics Lab/KDL
- Christopher Corless, Medical Director/KDL
- Paul Spellman, HOP PI/CEDAR & KCI
- Katie Johnson-Camacho, Sr. Human Subjects Research Specialist/CEDAR
- Gregory Goh, Molecular Technologist/KDL
- Travis Hayes, Sr. Specimen Processing Specialist/KDL
- Jane Thanner, Mgr/KDL Support Team/University Applications/ITG
- Other members of KDL

IGL/HOP | Workflow Development

Our starting point –

Platform: QIAGEN QIAsymphony robot Extraction kits: QIAsymphony DNA extraction kits Good partnership with QIAGEN technical and sales staff Issues to address –

- 1. standardized sample intake, storage and transfers
- 2. sample preprocessing for QIAGEN protocol
- 3. saliva volumes needed for required yields
- 4. procedures to insure sample accuracy throughout multi-step process
- 5. information documentation and sharing
- 6. minimizing costs









IGL/HOP | Saliva DNA SOP

Saliva DNA Extraction Workflow (IGL)



- \circ Recommended input = 10 mls saliva in mouthwash
- Turnaround times = 96 to 192 per week (routine; more possible)
- Minimum DNA concentration threshold for library prep = 5 ng/ ul
 - but lower concentrations are working

Sequencing and gene variant analysis/KDL

IGL/HOP | CAP lab set-up

We needed to establish a College of American Pathologists (CAP) accredited laboratory to process saliva samples for genetic testing

Our starting point –

No history of CAP/CLIA lab set-up or operations under these requirements

Partnered with members of the Knight Diagnostic Lab with primary support from Sarah McCabe

Required steps –

- 1. compliant laboratory set-up
- 2. documentation (lots of it!)
- 3. formalized staff training and competency reviews
- 4. College of American Pathologist (CAP) licensing

IGL/HOP | Throughput success !!!

Calendar	Samples
Year	Processed
2019	2880
2020	1056
2021	9120
2022	7392
2023	2784 (thru June)
Total	23232

Samples Processed (Including NTCs)



IGL/HOP | Hiccups & Ongoing Challenges

- •Initial CAP surprise inspection
- •Unexpected staff resignations
 - only one full time HOP technician
- •COVID pandemic

Ongoing Challenges

- Maintaining well-trained and adequate staff
- CAP lab regulatory maintenance with limited staff
- CAP requirements impact on rest of IGL activities/team
- High demand on IGL/HOP staff time during periodic lab inspections
- Adjusting to swings in HOP sample intake numbers

IGL/HOP | Benefits to healthcare mission, research community, and the IGL

Reliable partner for HOP – over 20,000 DNA samples delivered with extremely low failure rates.

✤IGL HOP staff became key contributors to set up of COVID testing lab at OHSU (beginning March , 2020).

IGL staff better informed on issues of working with clinical specimens leading to more insightful support of core projects involving patient samples across campus.

Aspects of CAP compliance requirements translated to better IGL staff training, improved equipment management protocols, and improvements in bench practices and SOPs in other core service areas.

Patient Education Working Group

Sue Williams Rebecca Siego-Coyle

Knight Diagnostics Lab

Chris Corless Sue Richards Gregory Goh Amiee Potter Arpita Kulkarni Sarah McCabe Catherine Driscoll

Genetics

Jone Sampson Kelly Hamman Amelia Mulford Brian O'Roak Andrew Adey

Integrated Genomics Lab Chris Harrington Jacob Buitrago **Britt Daughtry** Tiana Weeks SJ Kim Syber Haverlack Jinah Kim Providence Keri Vartanian JB Rinaldi Lindsey Dickey Bill Wright Aimee Shaykin

Walter Urba

OHSU and University or Oregon Marketing Allen Tomlinson Autumn Shafer Darsen Campbell-Prissle

HOP Staff

Madeleine Mathis

Marit Simmons

Vanessa Serrato

Ryan Lutz

Internal Advisory Board

Lisa Coussens Gordon Mills Rosalie Sears Kerri Winters-Stone Aaron Grossberg Gloria Coronado Emily Ho (Sadik Esener)