

Examining Disparities in Access to Breast and/or Ovarian Cancer Genetic Risk Assessment in “High Incidence Counties” versus “High Usage Counties”

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MPHA Epidemiology Section Conference – April 1, 2016
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BRCA and Hereditary Breast and Ovarian Cancer Syndrome

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- Hereditary Breast and Ovarian Cancer Syndrome (HBOC) will account for ~ 5-10% of all breast and 18-24% of all ovarian cancers
- *BRCA1* & 2: “**BR**east **CA**ncer 1 and 2” Genes
 - Code for tumor suppressor proteins
 - Dominant pattern of inheritance; Affected parent has 50% chance of passing gene mutation onto offspring
- Pathogenic *BRCA* mutations confer increased lifetime risk of certain cancers
 - Breast Cancer = 40-80%
 - Ovarian Cancer = 11-40%
- Estimated 1 in 300 to 1 in 500 individuals will have a pathogenic *BRCA* mutation

NCCN Criteria for HBOC Genetic Risk Assessment (2016)

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- Personal History
 - Known familial mutation in cancer susceptibility gene (i.e. BRCA)
 - Breast cancer at a young age (≤ 50 y.o.)
 - Triple negative breast cancer diagnosed at 60 y.o. or younger
 - Multiple primary HBOC-related cancers
 - Breast cancer at any age, and
 - ≥ 1 close blood relative with breast cancer at young age, or
 - ≥ 1 close blood relative w/ invasive ovarian cancer at any age
 - ≥ 2 close blood relatives w breast cancer and/or pancreatic cancer
 - From a population at increased risk
 - Male breast cancer
 - Ashkenazi Jewish Heritage w/ breast, ovarian, or pancreatic cancer at any age
 - Ovarian Cancer at any age

NCCN Criteria, Cont.

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- Family History; close relative with any of the following:
 - Known familial mutation in cancer susceptibility gene
 - Multiple HBOC primaries in single individual
 - ≥ 2 individuals with breast cancer primaries on same side of family with at least one diagnosed ≤ 50 y.o.
 - Ovarian cancer
 - Male breast cancer
- First or second degree relative with breast cancer ≤ 45 y.o.
- Personal and/or family history of ≥ 3 HBOC-related cancers (e.g. breast, pancreatic, prostate, colon, endometrial, etc.)
- Guidelines are ever-evolving

Data Sources

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- MDHHS *BRCA* Clinical Network Database
 - Funded through multiple CDC cooperative agreements
 - Unique to Michigan
 - Reporting on all *BRCA*-related counseling visits conducted by board-certified genetics professionals in MI
 - 18 contributing institutions – including one telecounseling service (Informed DNA)
 - Visit information consists of de-identified data on patient demographics, *BRCA* genetic testing, surgical procedures, personal and family history of cancer
- Michigan Cancer Surveillance Program (MCSP)
 - MI's cancer registry
 - Contains data on all cancer cases across the state
 - MDHHS Cancer Genomics requests case data for key cancers implicated in hereditary cancer syndromes, including breast and ovarian cancer

Analysis Objectives

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- MDHHS Cancer Genomics aims to increase genetic services utilization for individuals meeting referral criteria
- Examine patient demographics in *BRCA* Clinical Network database
Variables:
 - Year of Initial Visit
 - Dichotomous Race (White/Black only)
 - Ashkenazi Jewish heritage
 - Known Familial Mutation Status
 - Referring Physician
 - Insurance providers
- Compare demographics between two sub-populations:
 - “High Usage Counties” (HUC)
 - “High Incidence Counties” (HIC)

Methods

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High Incidence Counties:

- 12 counties identified by prior analysis of MCSP data:
 - Highest Age-Adjusted Incidence of *Breast Cancer at a young age*, 1998-2007:
 - **Emmet, Manistee, Grand Traverse, Leelanau, & Mason**
 - Highest Age-Adjusted of *Ovarian Cancer*, 2006-2010
 - **Alpena, Huron, Newaygo, Midland, Berrien, Saginaw, & Bay**
- *BRCA* database participants with zip code of residence in these counties included in this group





High Usage Counties:

- Analysis run on initial patient visits in *BRCA* database by zip code
 - High Usage Counties composed of zip codes accounting for top 10% of patient visits by volume
 - **Oakland, Wayne, & Washtenaw**
 - *BRCA* database participants with zip code of residence in these counties included in this group
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- Pearson Chi-square tests used to evaluate statistical dependence between sub-group designation and demographic characteristics
 - Analysis completed using SAS 9.2® software

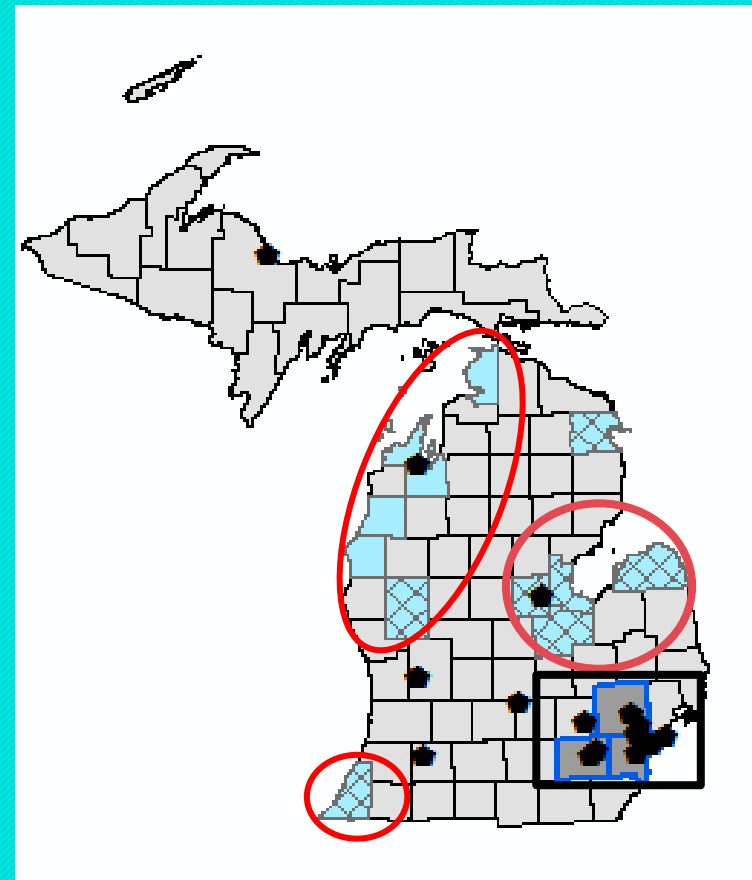
Geographic Distribution of “High Incidence Counties” and “High Usage Counties”

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$N_{\text{total}} = 1,458$
HIC, $N = 380$
HUC, $N = 1,078$

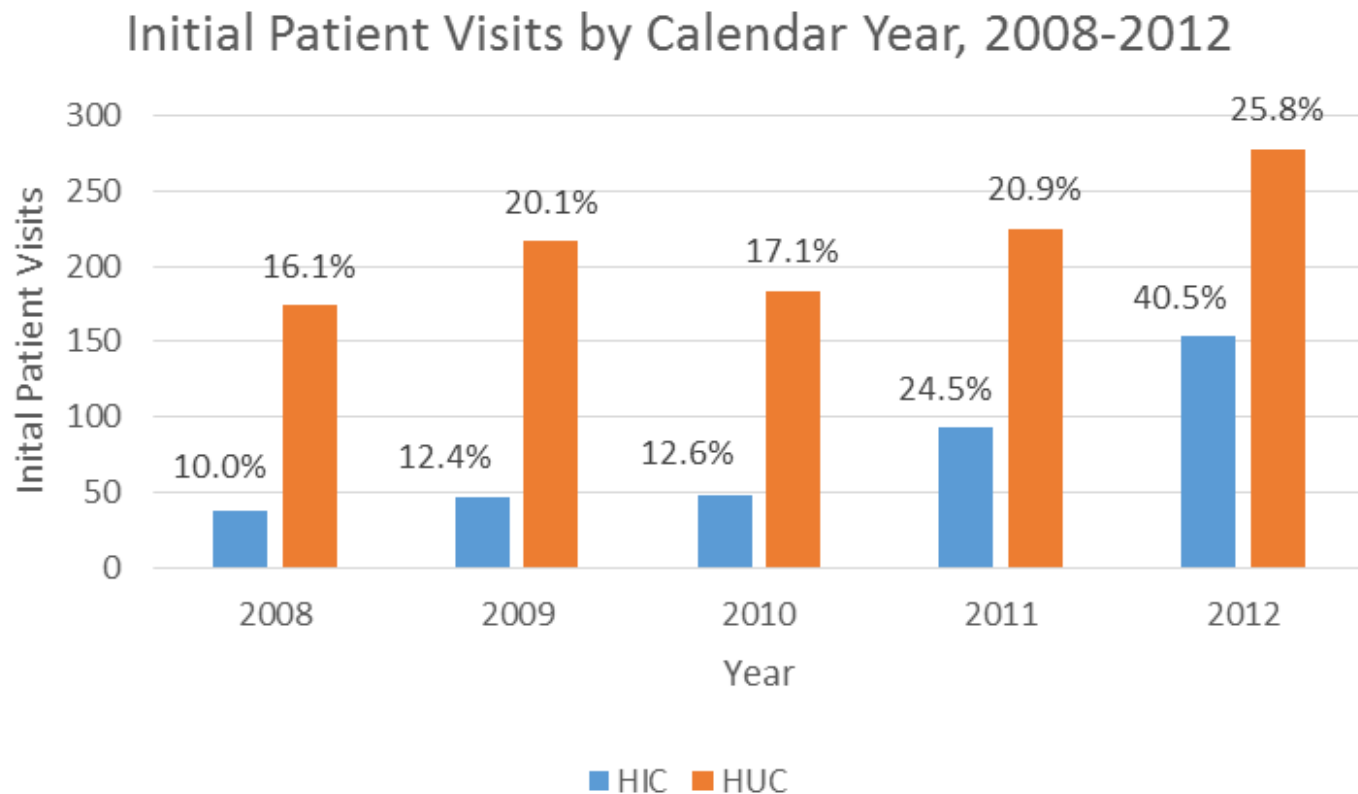
-  = Clinic location
-  = High Incidence Counties:
Breast Cancer at young age
-  = High Incidence Counties:
Ovarian Cancer
-  = High Usage Counties

**Italics denote high OC incidence vs. BC at young age*



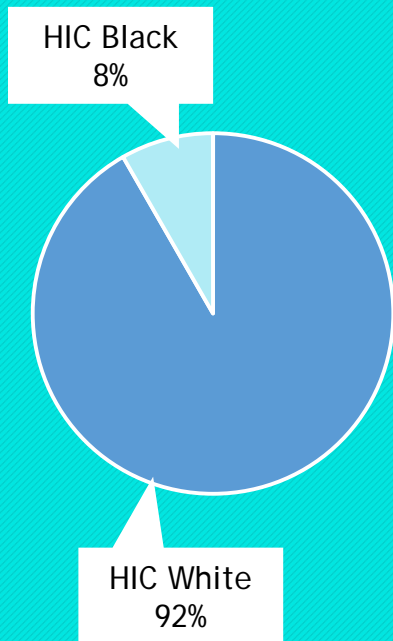
Visit Year Distribution, 2008-2012

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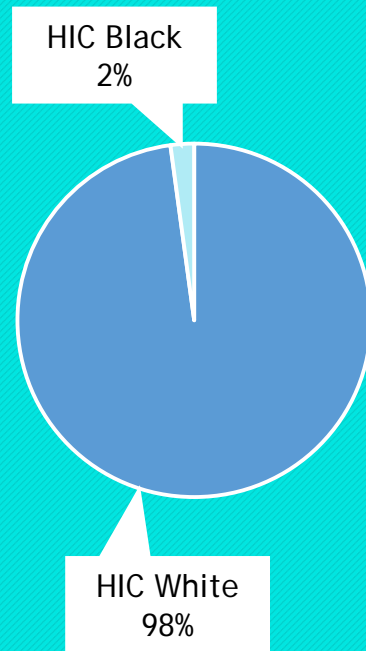


Racial Differences in Genetic Counseling Visits

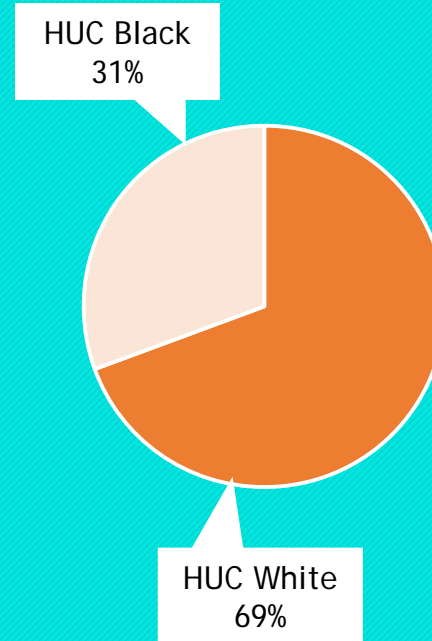
HIC Census Population (2010)



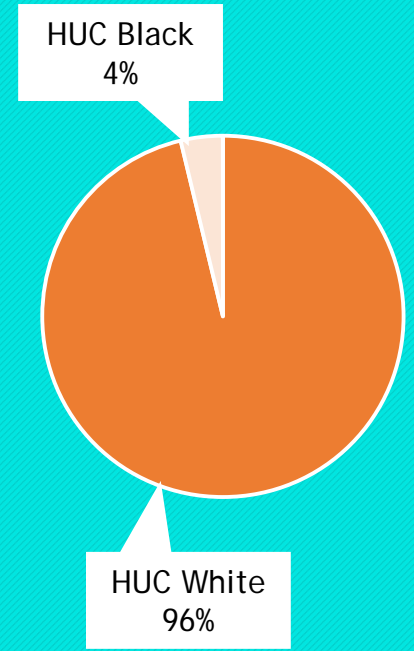
HIC Clinical Population



HUC Census Population (2010)



HUC Clinical Population



Ashkenazi Jewish Heritage and Known Familial Mutation Status

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	High Incidence Counties		High Usage Counties		X ²	p
	N	%	N	%		
Ashkenazi Jewish (AJ)					95.19	<.0001
No	374	98.42	819	75.79		
Yes	6	1.58	259	24.03		
Known Familial mutation (KFM)					6.48	.011
No	317	83.42	954	88.50		
Yes	63	16.58	124	11.50		

- Core Jewish population in MI exists in Wayne, Oakland, and Macomb counties
 - AJ = 1.9% of overall population in Metro Detroit
 - AJ = 7.5% in BRCA Clinical Network Database
- Overall proportion of KFM in BRCA Clinical Network Database = 12.03%

Results: Insurance Providers

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	High Incidence Counties		High Usage Counties		X ²	p
	N	%	N	%		
Top 5 Insurers					104.74	<.0001
Aetna	30	12.71	30	3.56		
Blue Care Network	12	5.08	133	15.78		
Blue Cross Blue Shield	153	64.83	526	62.40		
HAP			105	12.46		
Priority Health	31	13.14	42	4.98		
Uninsured	10	4.24	7	0.83		
Medicare					0.11	0.74
No	327	86.05	935	86.73		
Yes	53	13.95	143	13.27		
Medicaid					20.40	<.0001
No	361	95.00	1,066	98.89		
Yes	19	5.00	12	1.11		

Conclusions

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- Results of the current analysis:
 - show an increase in BRCA-related genetic counseling over time for HIC and HUC
 - Aligned with HealthyPeople2020 Genomics Objective
 - Highlight importance of determining barriers and facilitators to access to genetic counseling for populations at highest risk
- Other data sources are being used to conduct additional surveillance activities
- Long-term CDC project objective: reduce the incidence and mortality of hereditary cancers, including breast and ovarian cancer

Acknowledgements

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Clinical Partner Institutions: Beaumont Cancer Genetics Program, Henry Ford Health System Cancer Genetics Program, InformedDNA, Karmanos Cancer Genetic Counseling Service, Lacks Cancer Center Genetics at St. Mary's Healthcare, Marquette General Hereditary Cancer Program, Michigan State University Division of Clinical Genetics, MidMichigan Health Cancer Genetics Program, Munson Healthcare, Beaumont Hospital - Dearborn Genetics Risk Assessment for Cancer Clinic, Providence Hospital Medical Genetics, St. John Providence Cancer Genetics Program, Cancer Genetics Program at St. Joseph Mercy Hospital, Spectrum Health Cancer Genetics, University of Michigan Breast and Ovarian Cancer Risk and Evaluation Program, University of Michigan Cancer Genetics Clinic, and West Michigan Cancer Center

MDHHS Staff: Debra Duquette, MS, CGC, Robert Wahl, DVM, MS, Maricar Macalincag, MS, Sarah Lyon-Callo, MA, MS

This presentation was supported by the Grant or Cooperative Agreement Number, DP5357-02, funded by the Centers for Disease Control and Prevention. Its contents are solely the responsibility of the authors and do not necessarily represent the official views of the Centers for Disease Control and Prevention or the Department of Health and Human Services.

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