

Pharmacogenomics-guided Prescribing and Opportunities for Promoting Health Equity

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No disclosures or conflicts of interest to report

Agenda

Pharmacogenomics in diverse populations

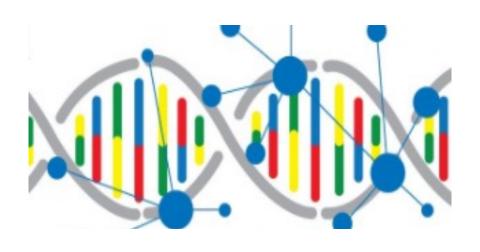
Current and Future Research

Pharmacogenomics (PGx)

Field that examines the relationships between genetic makeup and drug response.



Combines pharmacology and genomics.



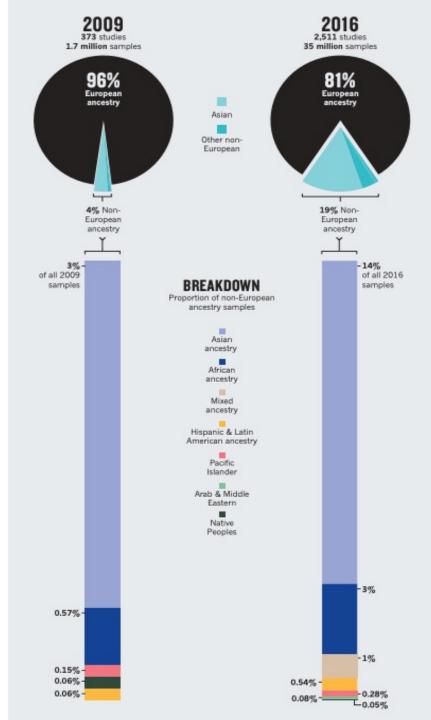
PGx tests for genetic inheritance have <u>lifelong</u> implications for directing drug treatment for a range of health conditions.

Benefits of Personalized Prescribing

- Trial and error prescribing
- Cost of adverse drug events (medical expenditures, morbidity/toxicities, mortality)
- Improve health outcomes



In spite of improvements, the vast majority of data from Genome Wide Association Studies (GWAS) are from European populations



Concern that ethnically diverse populations are being excluded from the benefits of personalized prescribing because:

Dearth of genetic information collected from these populations



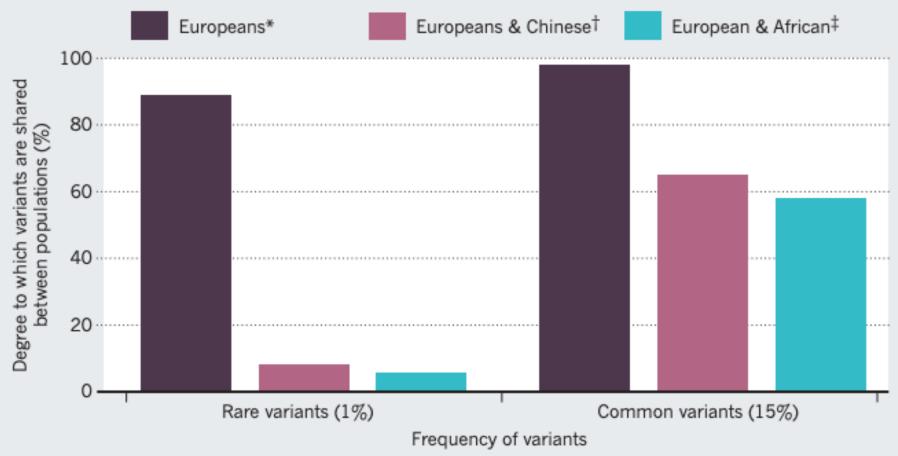
Underrepresentation in PGx studies



Uncertainty of clinical utility in minority populations

COMPARING THE UNCOMPARABLE

The rarer a genetic variant is within a population, the less likely it is to be found in all ethnic groups. One hundred people were sampled from each population.

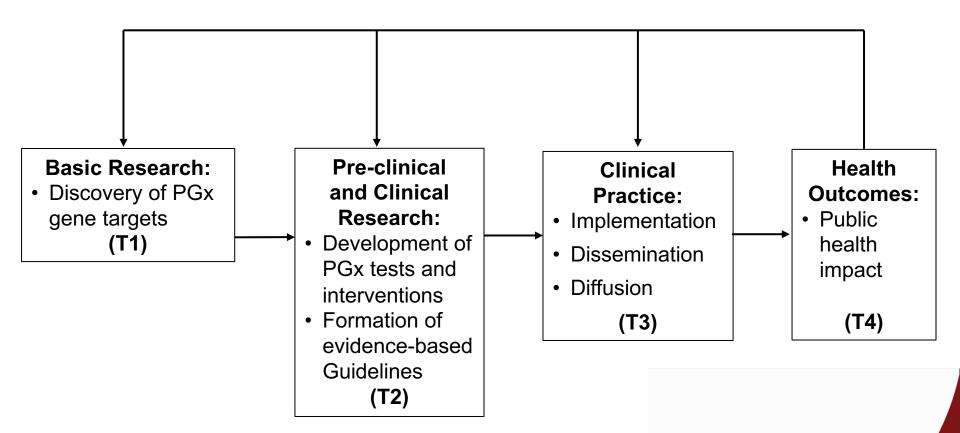


^{*}Comparison of individuals of European descent in Utah and in Tuscany, Italy. † Han Chinese individuals from Beijing compared with Utah sample ‡ Yoruba individuals from Ibadan, Nigeria, compared with Utah sample.

Criticisms for the use of race in clinical decision-making

- Scientific basis and definition of race is disputed
 - Arbitrary categorization under influence of social construction and discrimination
 - 'Racial' categories do not apply in a global context, being heavily influenced by national/local sociopolitical factors
 - Genetics may be conflated with race fueling discrimination based in genetic predeterminism
- Self-reported race vs. genetic ancestry

Translational Cycle of Pharmacogenomics (PGx) Research

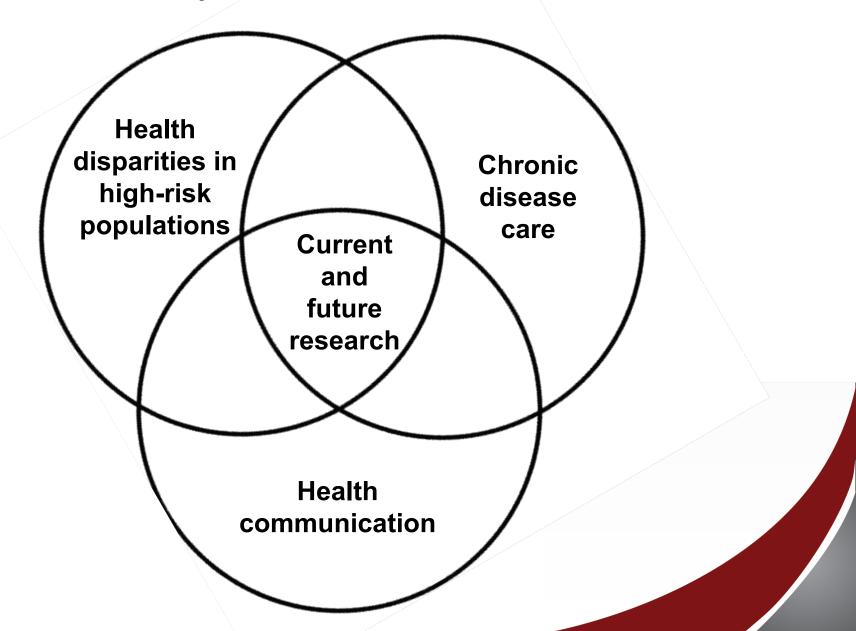


Across translational stages, patients from ethnically diverse populations may experience greater risk for experiencing health disparities.

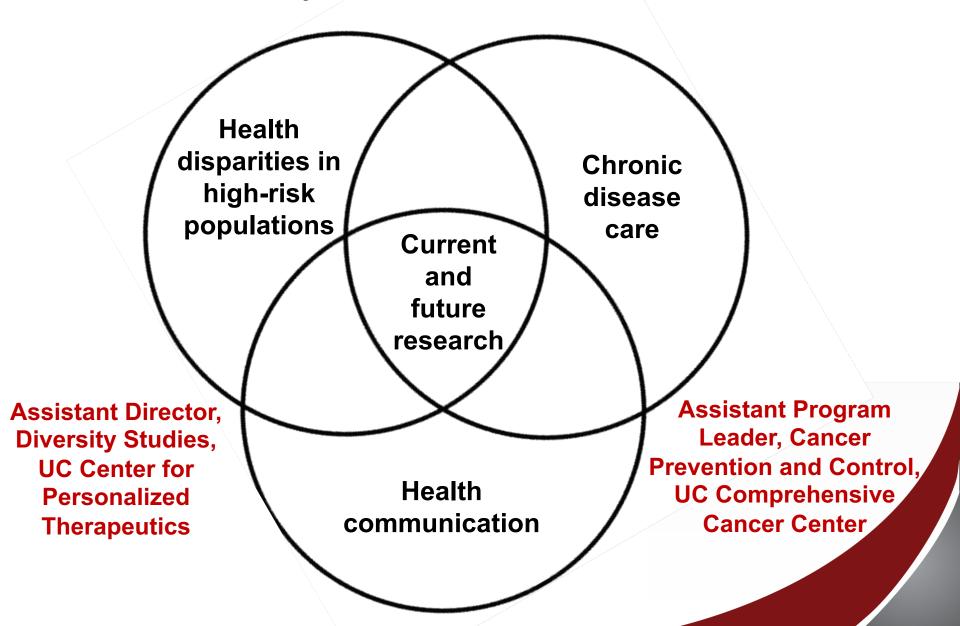
Adapted from Khoury et al Genetics in Medicine 2007

A systematic review of the literature found a documented shortage of evaluations on the impact of pharmacogenomics on health disparities.

Health Policy and Health Services Research



Health Policy and Health Services Research



Translation of pharmacogenomics (PGx) to underserved and underrepresented patient populations at risk for experiencing health disparities

- 1) Patient experience with PGx
- 2) PGx implementation outcomes in minority patient populations
- 3) PGx and social vulnerability (i.e. social determinants of health)

1)Patient experience with PGx

Underrepresented patient views and perceptions of personalized medication treatment through pharmacogenomics

Study Objectives

- Evaluate the reported views and perceptions of care received among genotyped African-American, or Black, and White patients participating in a large institutional pharmacogenomic implementation program.
- 2) Include education as a key covariate because it reflects the availability of resources (e.g. income and health insurance), the ability to process various types of information, and multiple socioeconomic indicators including social and cultural factors

^{***}We particularly focused on self-reported race to define the populations in our study as self-identified race is used to direct pharmacogenomic clinical guidelines as well as clinical decision-making more generally in healthcare settings

Hypothesis

We hypothesized that patients' attitudes and perceptions about pharmacogenomics with respect to their medical care would significantly differ based on self-reported race.

Data and Study Setting

 1200 Patients Project (UChicago Center for Personalized Therapeutics)

Patient surveys completed following outpatient clinical visits

 Surveys provided self-reported demographic information along with patient views/perceptions of care during their most recent health encounter

Methods and Analysis

- Comparison of survey measures between self-reported Black and White study participants
- Multiple surveys collected over time for a single individual
- Statistical analysis involved chi-square tests of independence with the level of statistical significance set at P<0.05 without adjustment for multiple comparisons

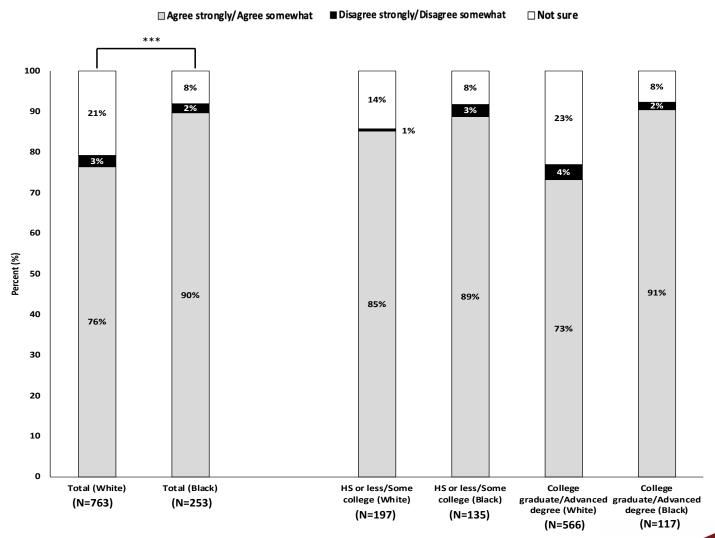
Study Population

			Rac	се				
		White			Black			
Educational attainment		HS or less/ Some college	College graduate/ Advanced degree	Total⁵	HS or less/ Some college	College graduate/ Advanced degree	p-value [†]	
Total survey respondents [N(%)]	332(72)	86(26)	246(74)	131(28)	74(56)	56(43)		
Gender [N(%)]								
Female	138(42)	44(51)	94(38)	94(72)	51(69)	43(77)	<0.0001***	
Age [N(%)]								
Mean (range)	60(19-90)	63(19-90)	59(20-87)	61(19-95)	61(19-89)	61(26-89)		
18-25 years	8(2)	2(2)	6(2)	1(1)	1(1)			
26-39 years	26(8)	3(3)	23(9)	9(7)	6(8)	3(5)		
40-50 years	33(10)	5(6)	28(11)	21(16)	10(14)	11(20)	0.306	
51-64 years	122(37)	33(38)	89(36)	43(33)	22(30)	21(38)		
65+ years	143(43)	43(50)	100(41)	57(44)	35(47)	21(38)		
Total surveys returned evaluating clinical visits (after enrollment) [N(%)]	790(75)	207(26)	583(74)	265(25)	145(55)	119(45)		
Surveys returned per patient (after enrollment) [N(%)]								
Mean (range) *	2(1-16)	2(1-16)	2(1-14)	2(1-7)	2(1-7)	2(1-6)		
1	149(45)	40(47)	109(44)	66(50)	41(55)	24(43)		
2	73(22)	17(20)	56(23)	30(23)	15(20)	15(27)	0.512	
3-4	75(23)	18(21)	57(23)	26(20)	12(16)	14(25)	0.512	
5+	35(11)	11(13)	24(10)	9(7)	6(8)	3(5)		
Self-reported health [N(%)] [‡]	, ,	, ,	, ,	, ,				
Excellent/Very good	479(61)	101(50)	378(65)	105(40)	42(30)	63(53) *		
Good	235(30)	73(36)	162(28)	108(41)	63(44)	45(38)	<0.0001***	
Fair/Poor	68(9)	30(15)	38(7)	48(18)	37(26)	10(8)		
N=	782	204	578	261	142	118		

^{*}*P*≤.05. ***P*≤.01. ****P*≤.001.

Receptivity to a greater role for personalized prescribing may be more broadly distributed across the Black patient population as Black respondents across education levels indicated similarly high levels of this desire.

I think knowledge of my personal genetic information should have a greater role in my healthcare provider's treatment decisions about me



^{*} 尸፯ !(ዜቭnκ አውው!(ይሳge *ቫታት)/ ውደነው hal genetic information should have a greater role in my healthcare provider's treatment decisions about met Educational attainment

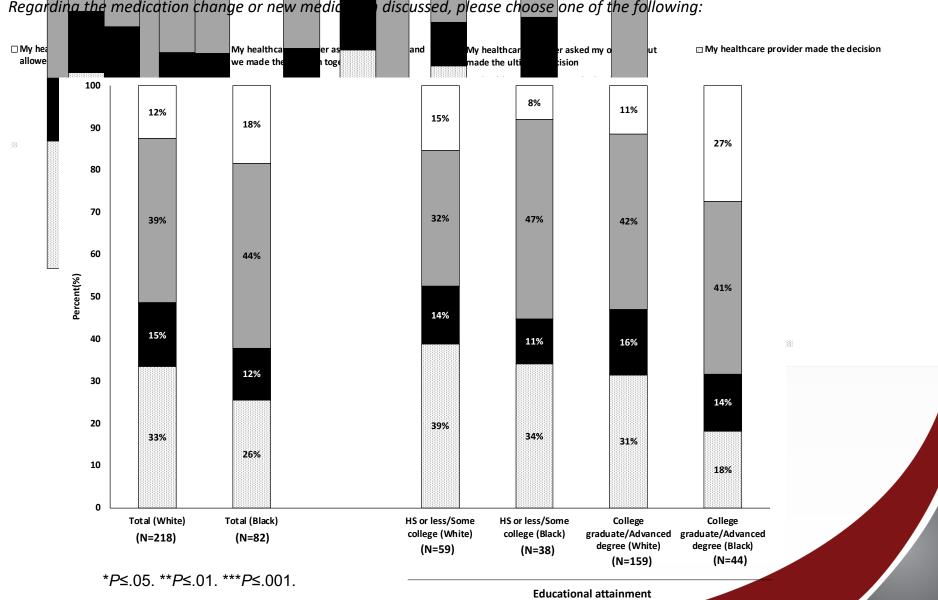
Black patients reported initiating discussions about the impact of personal make-up (genetics) on medication response with their provider far less frequently than White patients.

			Rad	е			
	White			Black			
	Total ^a	HS or less/ Some college	College graduate/ Advanced degree	Total ^b	HS or less/ Some college	College graduate/ Advanced degree	p-value [†]
Survey measure/question	N(%)	N(%)	N(%)	N(%)	N(%)	N(%)	
Did your healthcare provider stop or change one of your medications today, or start a new medication?							
Yes No Unsure N=	237(31) 532(69) 5(1) 774	61(31) 136(69) 1(1) 198	176(31) 396(69) 4(1) 576	87(33) 170(65) 4(2) 261	41(28) 101(70) 2(1) 144	46(40) 68(59) 2(2) 116	0.273
If yes to did your healthcare provider stop or change one of your medications today, or start a new n	nedication						
Did your healthcare provider discuss specific factors about you or your personal make-up which would suggest that you were more likely or less likely than other patients to benefit from the medication change or new medication?							
Yes	128(59)	43(75)	85(53)	43(49)	18(46)	25(52)	
No	77(35)	12(21)	65(41)	29(33)	12(31)	17(35)	0.005**
Unsure	12(6)	2(4)	10(6)	15(17)	9(23)	6(13)	
N=	217	57	160	87	39	48	
If yes, who initiated the discussion about individual factors regarding you and your response to the medication change or new medication?							
I was the one who asked about individual factors My healthcare provider was the one who brought up individual factors Unsure	20(15) 88(67) 23(18)	7(16) 34(76) 4(9)	13(15) 54(63) 19(22)	2(4) 42(86) 5(10)	0(0) 18(86) 3(14)	2(7) 24(86) 2(7)	0.037*
N=	131	45	86	49	21	28	

^{*}*P*≤.05. ***P*≤.01. ****P*≤.001.

More than half of Black and White patients reported being asked their opinion about medication changes during clinical visits, with at least 40% of each patient group reporting being asked their opinion and making the decision together with their provider.

Regarding the medication change or new medic discussed, please choose one of the following:



Conclusions

- Our results suggest an opportunity for enhanced patient-provider communication, especially for self-reported Black patients, around the role of genetic results during prescribing.
- Given the significance of patient-provider communication in influencing patient behaviors and outcomes, tailored communication strategies may need to be developed and employed which address Black patients' perceptions of receiving personalized treatments and help with the interpretation of pharmacogenomic information by providers for minorities and less educated patients.

Health Equity and Risk Communication in Pharmacogenomics

Aim 1: Where might health disparities arise as the use of pharmacogenomics becomes more widespread within the U.S. health system?

Publicly available nationally representative dataset of prescription drug use

Aim 2: What are the views, values, and preferences of minority populations and their providers in relation to pharmacogenomics?

Qualitative interviews

Aim 3: How might communication about PGx impact minority patient views and care preferences?

Survey experiment with hypothetical clinical vignettes

Aim 3: How might communication about PGx impact minority patient views and care preferences?

Survey experiment with hypothetical clinical vignettes (*Health Expect* 2015; *JGIM* 2017)

Exposure to a hypothetical clinical vignettes (Pharmacogenomic risk for adverse drug response)



Responses to measures within a survey instrument

2) PGx implementation outcomes in minority patient populations

Applicability of Pharmacogenomically Guided Medication Treatment during Hospitalization of At-Risk Minority Patients

Study Objectives

- 1) Conduct a multi-site inpatient pharmacogenomic implementation program among self-identified African-Americans (AA; n = 135) with numerous rehospitalizations (n = 341) from 2017 to 2020.
- 2) Evaluate the point-of-care availability of patient pharmacogenomic results to healthcare providers via an electronic clinical decision support tool.

Hypothesis

We hypothesized that evidence-based pharmacogenomic information would be particularly relevant to the medications used in this setting for hospitalized African American patients.

Data and Study Setting

- ACCOuNT (African American Cardiovascular Pharmacogenomics Consortium) Translational Project evaluated the clinical translation of pharmacogenomics to minority patients historically excluded from genomic studies.
- The University of Chicago, the University of Illinois Chicago, and Northwestern University.

ACCOuNT Consortium (Translational Project)

Impact on genetic biomarkers on:

- Drug selection
- Dosing
- Clinical outcomes

ACCOuNT: The African American Cardiovascular Pharmacogenetic Consortium

The University of Chicago

Pre-emptive genotyping of enrolled patients.

Genomic Prescribing System (GPS)- decision support by translating patients' genotypic results into individualized PGx summaries.

Patient PGx information is available and delivered to healthcare providers at the point of the patient encounter to guide prescribing decisions.

Hospitalizations

Medication prescribing frequently occurs during hospitalizations.

With a higher disease burden, AA are a high-risk population with frequent hospitalizations:

Several studies have examined patterns of hospitalization due to ambulatory care-sensitive conditions by race/ethnicity, generally finding higher rates of hospitalization for blacks compared to whites

(Decker et al. 2009; Chang et al. 2008; Howard et al. 2007; Laditka 2003; Oster and Bindman 2003; Pappas et al. 1997; Parket and Schoendorf et al. 2000; Shi et al. 1999)

Acute care setting vs. Routine care settings

 Alongside multiple studies, the lab has demonstrated the critical relevance of PGxguided prescribing for patients enrolled to study through outpatient clinics who were subsequently hospitalized (Lee et al 2019 *Pharmacogenetics and Genomics*).

AA: African-American

Methods and Analysis

- Patients who were hospitalized at any of these 3 institutions were approached for enrollment and collection of a blood sample for broad pharmacogenomic genotyping.
- Then, results were made available for subsequent hospitalizations to enrolled treating providers (physicians, advance practice providers, and pharmacists) via an electronic decision support tool, and prescribing during the hospitalization and at discharge were evaluated.
- Statistical analysis involved chi-square tests of independence with the level of statistical significance set at P<0.05 without adjustment for multiple comparisons

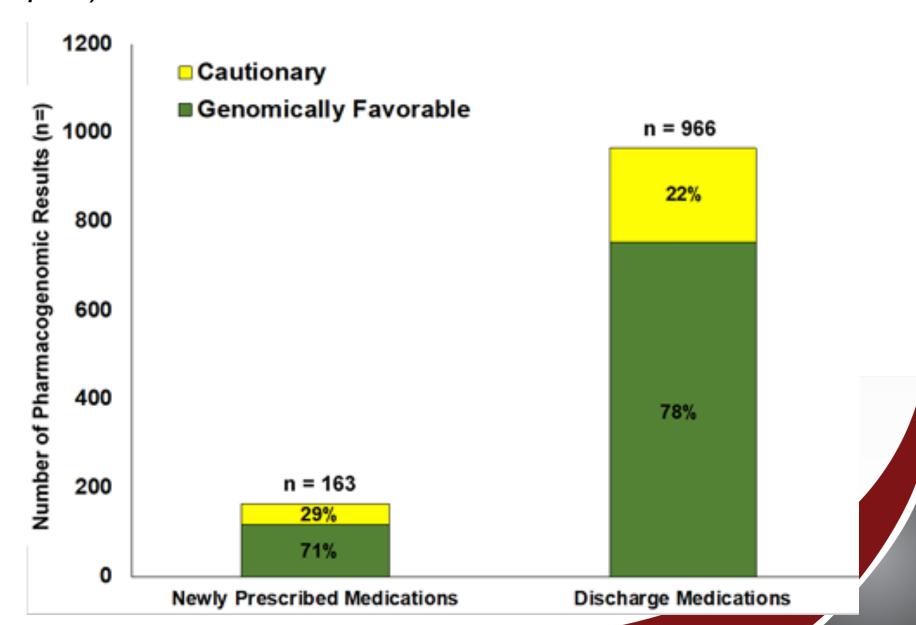
Patient Study Population

- The population was predominantly female (61%) with a mean age of 53 years (range 19-86).
- The majority of the population (80%) had an educational attainment of HS or less/some college.
- Mean number of comorbidities reported was 7 (range of 1-19).
- Mean number of prescription drugs was 12 (range of 1-28)
- On average, six medications were newly prescribed during each individual hospital admission.

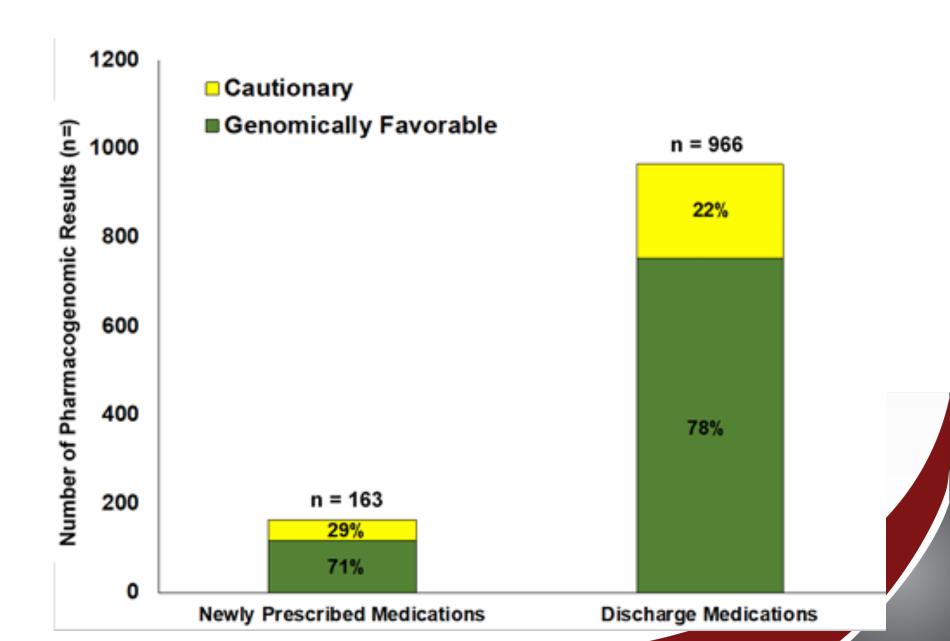
ACCOuNT Study Population: Providers

Number of Providers (n=2		
Characteristic		
Provider Type	n (%)	
MD (Hospitalist Physician + Primary Care)	158 (61)	
PA	14 (5)	
APN	16 (6)	
PharmD (Hospitalist Pharmacist + PP_Pharm	nacist) 73 (28)	
Years in practice (mean ± SD)	8 ± 7	
Institution (all providers)	n (%)	
UChicago	128 (49)	
Northwestern	89 (34)	
UIC	44 (17)	

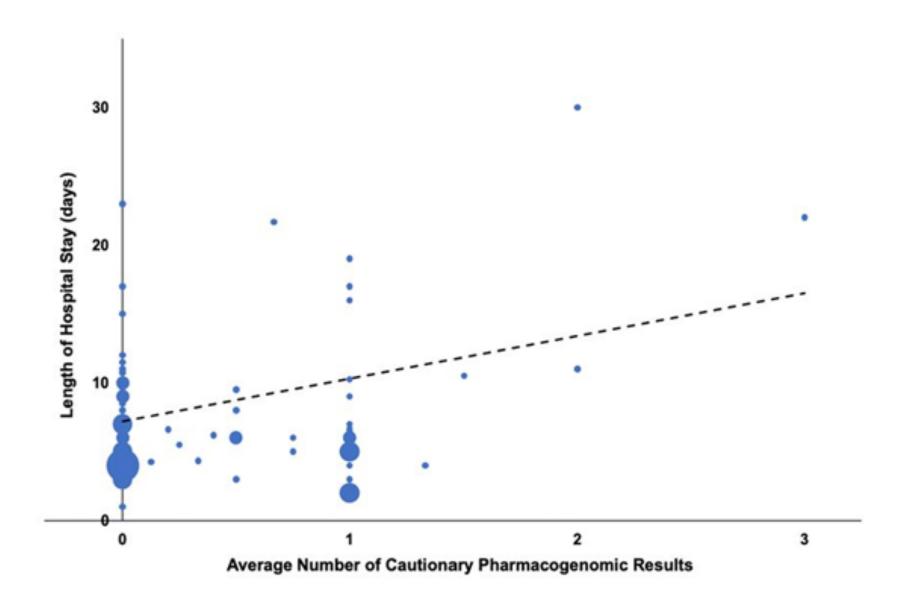
Most results indicated genomic favorability, although nearly 29% of newly prescribed medications indicated increased genomic caution (increase in toxicity risk/suboptimal response).



More than one of every five medications prescribed to AA patients at hospital discharge were associated with cautionary pharmacogenomic results.



Positive relationship between the per-patient average number of cautionary pharmacogenomic results associated with newly prescribed medications during hospitalizations and length of hospital stay.



Conclusions

- Our results indicated that high-risk pharmacogenomic results (genomic contraindication) were exceedingly rare in our hospitalized AA study population.
- We conclude that the applicability of pharmacogenomic information during hospitalizations for vulnerable populations at-risk for experiencing health disparities is substantial and warrants continued prospective investigation.

3) PGx and social vulnerability (i.e. social determinants of health)

Effect Modification by Social Determinants of Pharmacogenetic Medication Interactions on 90-Day Hospital Readmissions within an Integrated U.S. Healthcare System

Study Objective

Investigate the association between pharmacogenetic interactions and 90-day readmission in a diverse study population.

Hypothesis

We hypothesized that social determinants were important contributors to readmission and that there would be effect modification of the gene-x-drug interactions on risk of 90-day readmission, particularly by medical complexity.

Data and Study Setting

- Evaluated 90-day hospital readmission (primary outcome) for 19,999
 adults from 2010 through 2020 who underwent testing with a 13-gene
 pharmacogenetic panel.
- For genes with evidence of variation in drug response, genetic indicators (genotype-based phenotypes) were collected and annotated for:
 - CYP2B6, CYP2C19, CYP2C9, CYP2D6, CYP3A5, CYP4F2, DPYD, IFNL3, NUDT15, SLCO1B1, TPMT, UGT1A1, VKORC1.
- The NorthShore University HealthSystem

Methods and Analysis

Univariate logistic regression analyses demonstrated that strongest associations with 90-day hospital readmissions were the number of medications prescribed within 30 days of a first hospital admission that had Clinical Pharmacogenomics Implementation Consortium (CPIC) guidance.

Study Population

Table 1. Overall Study Population Sociodemographic and Clinical Characteristics (n = 19,999).

	Inpatient Admissions 2010–2020 a						
	Overall No				90-Day Hospital Readmission ^b		
n (%) √		Yes	n = 4740				
	19,999 (100)	15,259 (76)	4740 (24)		No 4120 (87)	Yes 620 (13)	
Age, median (IQR)	53 (42-64)	53 (42-63)	54 (41-68)	*	52 (41-67)	62 (46-72)	*
Age group (years)							
18-39	4097 (20.5)	3128 (76.3)	969 (23.7)		877 (90.5)	92 (9.5)	
40-49	4275 (21.4)	3208 (75)	1067 (25)		976 (91.5)	91 (8.5)	
50-64	6849 (34.2)	5679 (82.9)	1170 (17.1)		1011 (86.4)	159 (13.6)	
65 or above	4778 (23.9)	3244 (67.9)	1534 (32.1)		1256 (81.9)	278 (18.1)	
Gender	, ,	, ,	, ,	*			*
Female	12,852 (64.3)	9324 (72.5)	3528 (27.5)		3106 (88)	422 (12)	
Male	7147 (35.7)	5935 (83)	1212 (17)		1014 (83.7)	198 (16.3)	
Race	, ,	, ,	, ,	*			*
White	13,961 (69.9)	10,292 (73.7)	3669 (26.3)		3172 (86.5)	497 (13.5)	
Black/African American	612 (3.1)	437 (71.4)	175 (28.6)		131 (74.9)	44 (25.1)	
Asian	1293 (6.5)	1020 (78.9)	273 (21.1)		254 (93)	19 (7)	
American Indian/Alaska Native	37 (0.2)	28 (75.7)	9 (24.3)		8 (88.9)	1 (11.1)	
Pacific Islander/Hawaiian Native	18 (0.1)	16 (88.9)	2 (11.1)		2 (100)	0 (0)	
Other	3751 (18.8)	3157 (84.2)	594 (15.8)		537 (90.4)	57 (9.6)	

Health insurance was not associated with risk of 90-day readmission.

Table 4. Unadjusted and Adjusted Logistic Regression Analysis of 90-day Hospital Readmissions Among Patients Prescribed a CPIC Medication Within 30 days of Inpatient Admission (n = 4404).

uOR (95% CI)		<i>p</i> -Value	aOR (95% CI)	p-Value
Age Group				
18-39	Reference	59/0c0/A/0	Reference	3343490
40-49	1.05 (0.75-1.47)	0.001	1.04 (0.74-1.47)	0.529
50-64	1.73 (1.27-2.35)	0.026	1.28 (0.93-1.78)	0.104
65 or above	2.35 (1.77-3.12)	< 0.0001	1.14 (0.80-1.65)	0.783
Gender				
Female	Reference		Reference	
Male	1.47 (1.21-1.80)	0.0001	1.11 (0.89-1.38)	0.371
Race *				
White	Reference		Reference	
Black/African American	2.12 (1.47-3.07)	< 0.0001	2.12 (1.42-3.17)	< 0.0001
Asian	0.43 (0.26-0.70)	< 0.0001	0.62 (0.37-1.03)	0.002
Ethnicity				
Non-Hispanic	Reference			
Hispanic/Latino	1.10 (0.54-2.26)	0.975		
Marital status				
Unmarried	Reference		20	-
Married	0.60 (0.49-0.72)	0.441		
Employment status **	,			
Employed	Reference		Reference	
Unemployed	2.19 (1.82-2.64)	< 0.0001	1.74 (1.39-2.18)	0.007
Insurance status	10.0000		10. 800 0 10.000	
Commercial	Reference		Reference	
Government	1.81 (1.50-2.20)	0.307	1.04 (0.81-1.34)	0.6605
Out-of-pocket (self-pay)	1.67 (0.47-5.90)	0.736	1.48 (0.41-5.32)	0.5742
		0.737.77		

The odds of 90-day readmission for patients with one or more identified genex-drug interactions after adjustment for these covariates was attenuated by 10%

Table 4. Unadjusted and Adjusted Logistic Regression Analysis of 90-day Hospital Readmissions Among Patients Prescribed a CPIC Medication Within 30 days of Inpatient Admission (n = 4404).

uOR (95% CI)		p-Value	aOR (95% CI)	p-Value		
Median household income in relation to US median household income ***						
Less than \$64,994	Reference		Reference			
\$64,994 or more	1.33 (0.86-2.05)	0.203	1.63 (1.03-2.58)	0.035		
BMI, range 13.3-79.2	1.03 (1.02-1.04)	< 0.0001	1.01 (1.00-1.03)	0.050		
Smoking status	10000					
No	Reference		Reference			
Yes	1.38 (0.87-2.18)	0.176	1.26 (0.78-2.04)	0.347		
COVID status						
Yes vs. No	0.56 (0.17-1.84)	0.341				
		morbidities ****				
0	Reference		Reference			
1	1.57 (1.24-1.98)	0.018	1.28 (0.99-1.64)	0.044		
2	2.54 (1.91-3.39)	0.006	1.83 (1.33-2.52)	0.070		
3 or more	3.36 (2.61-4.32)	< 0.0001	2.23 (1.66-3.02)	< 0.0001		
Number	of CPIC medications prescribed	within 30 days prior to i				
1	Reference					
2	1.14 (0.82-1.58)	< 0.0001	į-	-		
3	2.50 (1.77-3.53)	0.417		-		
4	3.07 (2.13-4.41)	0.008	(-	-		
5 or more	7.66 (5.45-10.77)	<0.0001				
G	ene-x-drug interactions within 3	30 days prior to inpatien	t admission			
Absent	Reference		Reference			
Present	1.41 (1.18-1.70)	0.0002	1.31 (1.08-1.59)	0.006		

Conclusions

Our results highlight the major contribution of social determinants and medical complexity to risk for hospital readmission, and that these determinants may modify the effect of gene-x-drug interactions on rehospitalization risk.

Future Research



Health Services Research

RESEARCH ARTICLE 🙃 Open Access 💿 🛈 😑 💲

The social vulnerability metric (SVM) as a new tool for public health

First published: 19 November 2022 | https://doi.org/10.1111/1475-6773.14102

Funding information: National Human Genome Research Institute, Grant/Award Number: K08 HG011505; National Institute on Aging, Grant/Award Number: R56 AG066127

Future Research (continued)

Improve health outcomes for underserved and underrepresented populations through precision prescribing

- Tailoring PGx implementation across clinical settings and therapeutic areas
- Evolve PGx clinical decision support tools at the point-of-care to best support targeted care delivery (e.g. patient-provider communication on PGx results and prescribing decisions)
- Aligning PGx care to patient preferences

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Thank you!

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