



ASCPT 2018

ANNUAL MEETING

MARCH 21 - 24, 2018 • HILTON ORLANDO • ORLANDO, FL

Accommodating substrate-dependence in CYP genotype to activity phenotype translation for pharmacogenetic implementation

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Survey: Challenges to implementing pharmacogenetics in the clinic

What do you think is the most challenging aspect of the implementation of pharmacogenetics into the clinic?

- A. Translation of genetic information into clinical action
- B. Test cost, test reimbursement or other economic issues
- C. Availability of high quality genotyping test (CLIA approved)
- D. Electronic medical record use, such as the application of CDS
- E. Clinician and patient resistance and/or ethical concerns

Survey: top 3 Challenges to implementing pharmacogenetics in the clinic

- 95% of respondents selected: “process required to translate genetic information into clinical actions”
- Next 2 responses
 - Genotype test interpretation (e.g. using genotype information to impute phenotype)
 - Providing recommendations for selecting the drug/gene pairs to implement



- CPIC guidelines are designed to help clinicians understand HOW available genetic test results should be used to optimize drug therapy.
 - Not WHETHER tests should be ordered.
- Key Assumption:
 - Clinical high-throughput and pre-emptive genotyping will become more widespread.
 - Clinicians will be faced with having patients' genotypes available even if they did not order test with drug in mind.



- As of January 2018:
 - 249 Members
 - Clinicians and scientists
 - 178 institutions
 - 28 countries
 - 10 Observers (NIH and FDA)
 - CPIC Informatics
 - >20 members from 12 organizations

2011

- *TPMT* – thiopurines
- *CYP2C19*– clopidogrel
- *CYP2C9, VKORC1* – warfarin

2012

- *CYP2D6* – codeine
- *HLA-B* – abacavir
- *SLCO1B1* – simvastatin

2013

- *HLA-B* – allopurinol
- *CYP2D6, CYP2C19* – TCAs
- *HLA-B* – carbamazepine
- *DPYD* -- 5FU / capecitabine
- *TPMT* – thiopurines—UPDATE
- *CYP2C19* – clopidogrel--UPDATE

2014

- *IL28B* -- PEG interferon α
- *CFTR* -- Ivacaftor
- *G6PD* -- Rasburicase
- *CYP2C9, HLA-B* -- Phenytoin
- *CYP2D6* – codeine--UPDATE
- *HLA-B* – abacavir--UPDATE
- *SLCO1B1* – simvastatin—UPDATE

2015

- *CYP3A5* – tacrolimus
- *CYP2D6, CYP2C19*– SSRIs
- *UGT1A1* – atazanavir
- *HLA-B* – allopurinol—UPDATE

2016

- *CYP2C19* – voriconazole
- *CYP2D6* – ondansetron
- *CYP2C9, VKORC1* – warfarin--
UPDATE
- *CYP2D6, CYP2C19* – TCAs--UPDATE



2017

- *CYP2D6* – tamoxifen
- *HLA-B* – carbamazepine—UPDATE
- *DPYD* -- 5FU / capecitabine—UPDATE-in
review

2018 (in progress)

- *RYR1*– inhaled anesthetics
- *CYP2B6*—efavirenz
- *TPMT/NUDT15* – thiopurines--UPDATE
- *CYP2D6*—atomoxetine
- *CYP2C19*/PPI
- *CYP2C9*/HLA-phenytoin—UPDATE
- *CYP2C9*/celecoxib



CPIC guidelines and list of CPIC genes/drugs



CPIC open meeting on 3/15/2017 in Washington DC - more details on the meetings page

What is CPIC?

The Clinical Pharmacogenetics Implementation Consortium (CPIC) was formed as a shared project between PharmGKB and the Pharmacogenomics

Background

One barrier to clinical implementation of pharmacogenetics is the lack of freely available, peer-reviewed, updatable, and detailed gene/drug clinical

CPIC website: www.cpicpgx.org

Guideline for Voriconazole and CYP2C19

Supplemental Table S1. Evidence linking *CYP2C19* genotype to voriconazole phenotype

[CYP2C19 allele definition table](#) 

[CYP2C19 allele functionality table](#) 

[CYP2C19 frequency table](#) 

[CYP2C19 diplotype-phenotype table](#) 

Gene resource mapping

[CYP2C19 gene resource mappings](#) 

Drug resource mapping

[Voriconazole](#) 

Clinical decision support:^b

[Voriconazole pre- and post-test alerts and flow chart](#) 

tion:

[Pharmacogenetics Implementation Consortium \(CPIC\) Guideline for CYP2C19 and Voriconazole](#)

tions since publication.

manuscript of the guideline:

phenotype based on genotypes

voriconazole based on CYP2C19 phenotype for adult patients

voriconazole based on CYP2C19 phenotype for pediatric patients (children and adolescents <18 years old)

[Pharmacogenetics Implementation Consortium \(CPIC\) Guideline for CYP2C19 and Voriconazole \(October 2016\)](#) 

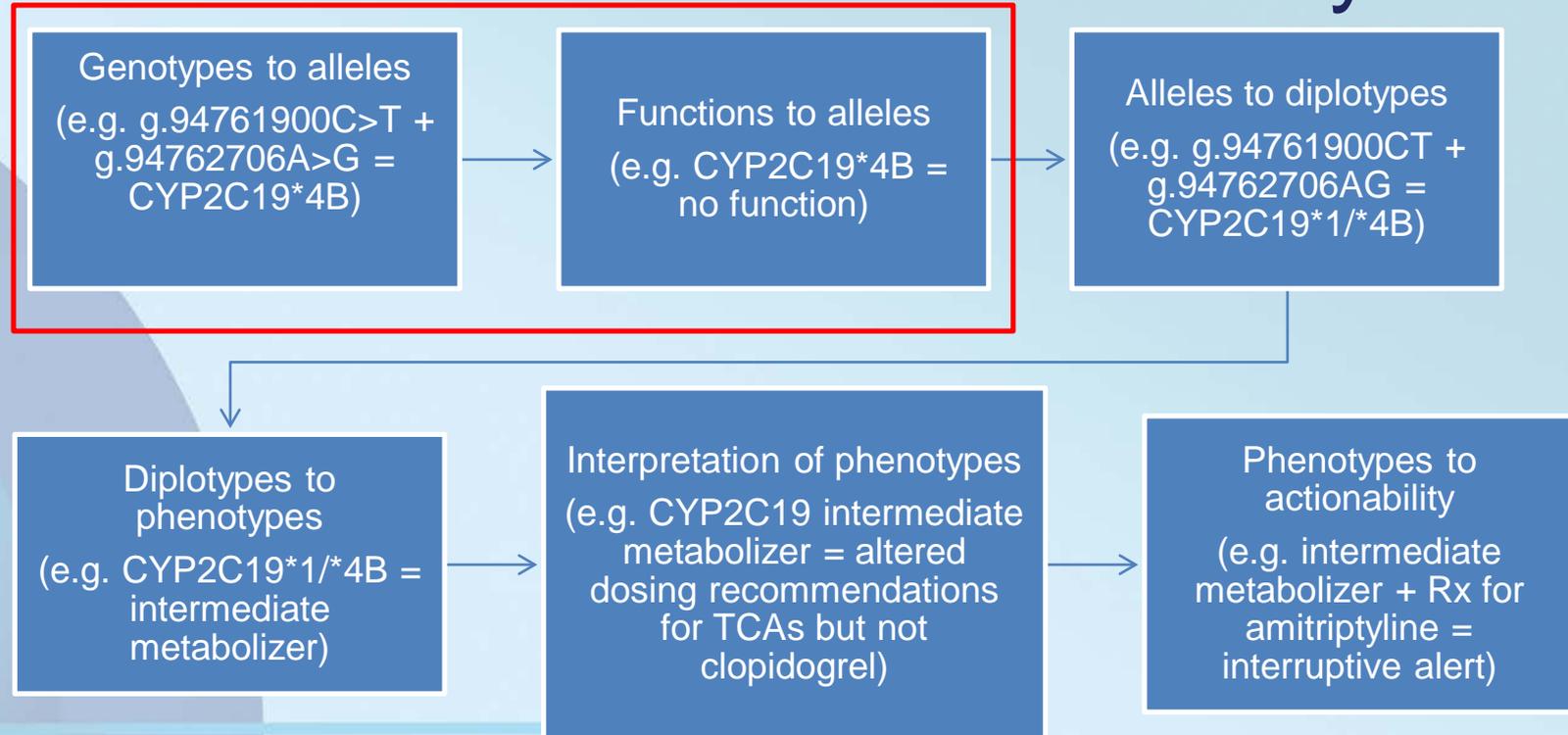
^a in the guideline publication supplement or referenced in the guideline^a:

CYP2C19 genotype to voriconazole phenotype

Gene resource mapping

[CYP2C19 gene resource mappings](#) 

CPIC tables allow translation of genetic test results to actionability



<https://cpicpgx.org/guidelines/>

<https://www.pharmgkb.org/page/cyp2c19RefMaterials>

Allele definition table: genotypes to alleles

CYP2C19_allele_definition_table.xlsx [Protected View] - Excel

PROTECTED VIEW Be careful—files from the Internet can contain viruses. Unless you need to edit, it's safer to stay in Protected View. Enable Editing

B4 : X ✓ fx Position at NC_000010.11 (Homo sapiens chromosome 10, GRCh38.p2)

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
1	GENE: CYP2C19	6/7/2016													
2		Nucleotide change	-2030C>T	-2020C>A	-1439T>C	-1041G>A	-806C>T	-13G>A	1A>G	7C>T	10T>C	50T>C	55A>C	83A>T	151A
3		Effect on protein	5' region	M1V	P3S	F4L	L17P	I19L	K28I	S51G					
4		Position at NC_000010.11	g.94760676C>T	g.94760686C>A	g.94761267T>C	g.94761665G>A	g.94761900C>T	g.94762693G>A	g.94762706A>G	g.94762712C>T	g.94762715T>C	g.94762755T>C	g.94762760A>C	g.94762788A>T	g.94762800G>A
5		Position at NG_000001.1	g.2971C>T	g.2981C>A	g.3562T>C	g.3960G>A	g.4195C>T	g.4988G>A	g.5001A>G	g.5007C>T	g.5010T>C	g.5050T>C	g.5055A>C	g.5083A>T	g.5100G>A
6		rsID	rs113164681	rs111490789	rs17878739	rs7902257	rs12248560	rs367543001	rs28399504	rs367543002	rs367543003	rs55752064	rs17882687		
7	Allele	Allele Functional Status													
8	*1	Normal function	C	C	T	G	C	G	A	C	T	T	A	A	A
9	*2	No function													
10	*3	No function													
11	*4A	No function							G						
12	*4B	No function					T		G						
13	*5	No function													
14	*6	No function													
15	*7	No function													
16	*8	No function													
17	*9	Decreased function													
18	*10	Decreased function													
19	*11	Normal function													
20	*12	Unknown function													
21	*13	Normal function													
22	*14	Unknown function										C			
23	*15	Normal function											C		
24	*16	Decreased function													
25	*17	Increased function					T								
26	*18	Normal function													
27	*19	Decreased function													G
28	*22	No function													
29	*23	Unknown function													
30	*24	No function													
31	*25	Decreased function													

<https://www.pharmgkb.org/page/pgxGeneRef>

<https://cpicpgx.org/guidelines/guideline-for-voriconazole-and-cyp2c19/>



Translating CYP genotypes to allelic functional status

Term/gene category	Final term^a	Functional definition	Genetic definition	Example diplotypes/alleles
Allele functional status: all genes	Increased function	Function greater than normal function	N/A	<i>CYP2C19*17</i>
	Normal function	Fully functional/wild-type	N/A	<i>CYP2C19*1</i>
	Decreased function	Function less than normal function	N/A	<i>CYP2C19*9</i>
	No function	Nonfunctional	N/A	<i>CYP2C19*2</i>
	Unknown function	No literature describing function or the allele is novel	N/A	<i>CYP2C19*29</i>
	Uncertain function	Literature supporting function is conflicting or weak	N/A	<i>CYP2C19*12</i>

Allele functionality table: alleles to function

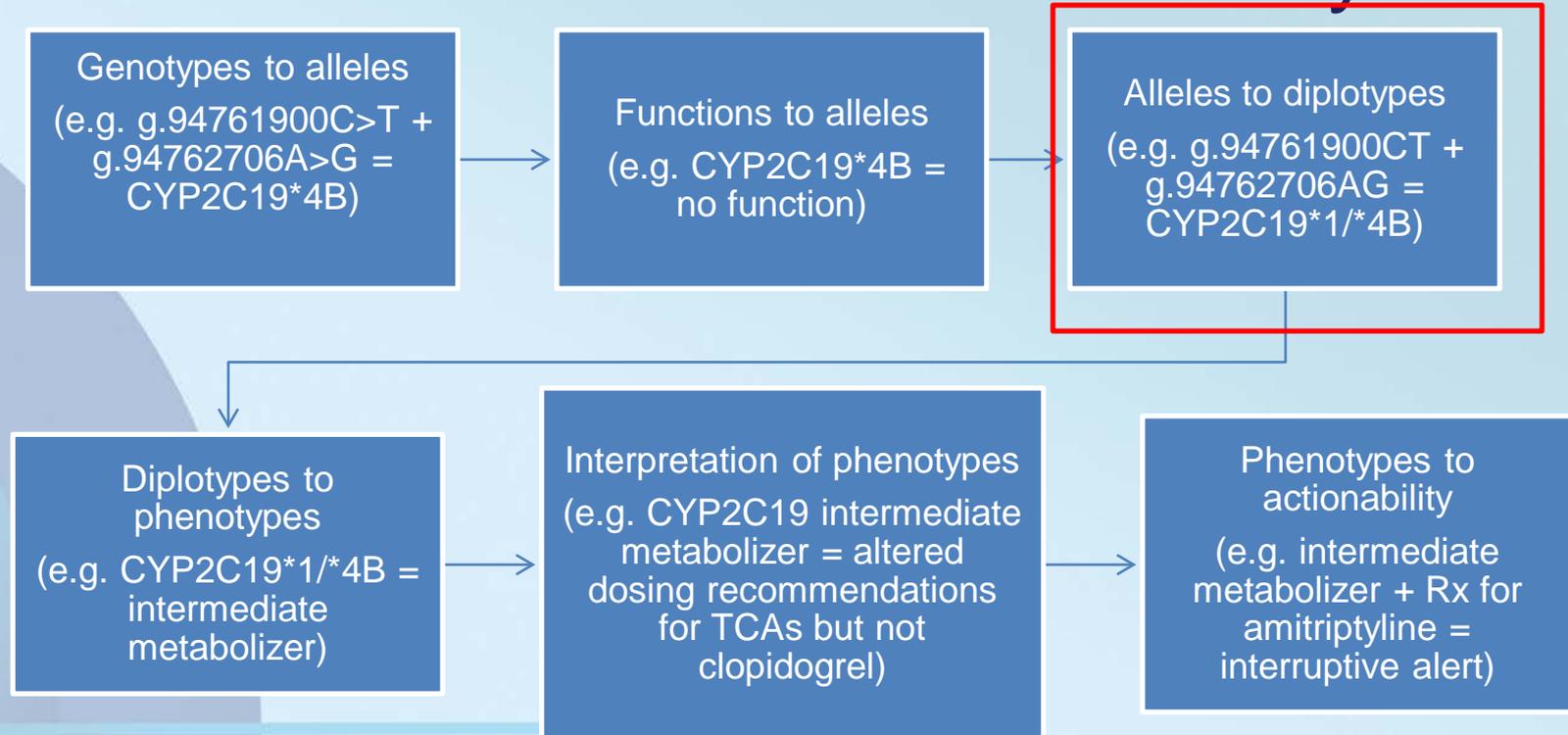
The referenced substrates and cited articles are examples and may not represent all information that may be available for an allele.

GENE: CYP2C19	6/20/2017				
Allele	Allele Functional Status	References	PMID	in vitro	Drug substrate in vivo
*1	Normal function	Romkes 1991	2009263		
		Richardson 1995	7487078	S-mephenytoin, tolbutamide	
		Blaisdell 2002	12464799	S-mephenytoin	
		Hanioka 2007	17455109	S-mephenytoin	
		Hanioka 2008	18312490	omeprazole	
		Wang 2011	21325430	S-mephenytoin, omeprazole	
		Takahashi 2015	25001882	clopidogrel, S-mephenytoin	
*2	No function	de Morais 1994	8195181	S-mephenytoin	
		Ibeanu 1998	9732415		S-mephenytoin
		Lee 2009	19661214		S-mephenytoin, omeprazole
		Xiao 1997	9103550		S-mephenytoin
		de Morais 1994	7969038		S-mephenytoin
*3	No function	Xiao 1997	9103550		S-mephenytoin
		Ferguson 1998	9435198		S-mephenytoin
*4A	No function	Scott 2012	21358751		clopidogrel
*4B	No function	Xiao 1997	9103550		S-mephenytoin
		Ibeanu 1998	10022751	S-mephenytoin, tolbutamide	S-mephenytoin
		Wang 2011	21325430	S-mephenytoin, omeprazole	
		Takahashi 2015	25001882	clopidogrel, S-mephenytoin	
*6	No function	Ibeanu 1998	9732415		S-mephenytoin
		Wang 2011	21325430	S-mephenytoin, omeprazole	
		Takahashi 2015	25001882	clopidogrel, S-mephenytoin	
*7	No function	Ibeanu 1999	10411572		S-mephenytoin
*8	No function	Ibeanu 1999	10411572	S-mephenytoin, tolbutamide	S-mephenytoin
		Wang 2011	21325430	S-mephenytoin, omeprazole	
		Takahashi 2015	25001882	clopidogrel, S-mephenytoin	
*9	Decreased function	Blaisdell 2002	12464799	S-mephenytoin	
		Wang 2011	21325430	S-mephenytoin, omeprazole	
		Takahashi 2015	25001882	clopidogrel, S-mephenytoin	
*10	Decreased function	Blaisdell 2002	12464799	S-mephenytoin	

Supplemental Table S2. Association between allelic variants and CYP2C9 function

Functional Status ^{a, b}	Alleles	In Vitro Activity	
		Substrate	Percent reduction of in vitro metabolism vs CYP2C9*1
Normal Activity	*1		
	*9	S-warfarin Tolbutamide Tolbutamide	82% of Cl _{int} [34] 96% of wild-type activity [34] 93% of Cl _{int} [35]
Decreased Activity	*2	S-warfarin Tolbutamide Phenytoin	32% of Cl _{int} [34] 42% of wild-type activity [34] 71% of Cl _{int} [36]
	*3	S-warfarin Tolbutamide Tolbutamide Phenytoin Phenytoin	21% of wild-type activity [34] 28% of wild-type activity [34] 26% of Cl _{int} [37] 5% of Cl _{int} [36] 7% of Cl _{int} [38]
No Activity	*6	N/A	Frameshift mutation [34]
	*15	N/A Tolbutamide	Nonsense mutation No expression [39]
	*25	N/A	Frameshift mutation [34]
Possible Decreased Activity (no available phenytoin in vitro activity studies)	*4	S-warfarin Tolbutamide	16% of wild-type activity [34] 22% of wild-type activity [34]
	*5	S-warfarin Tolbutamide S-warfarin	19% of wild-type activity [34] 24% of wild-type activity [34] 8% of Cl _{int} [40]

CPIC tables allow translation of genetic test results to actionability



<https://cpicpgx.org/guidelines/>

<https://www.pharmgkb.org/page/cyp2c19RefMaterials>

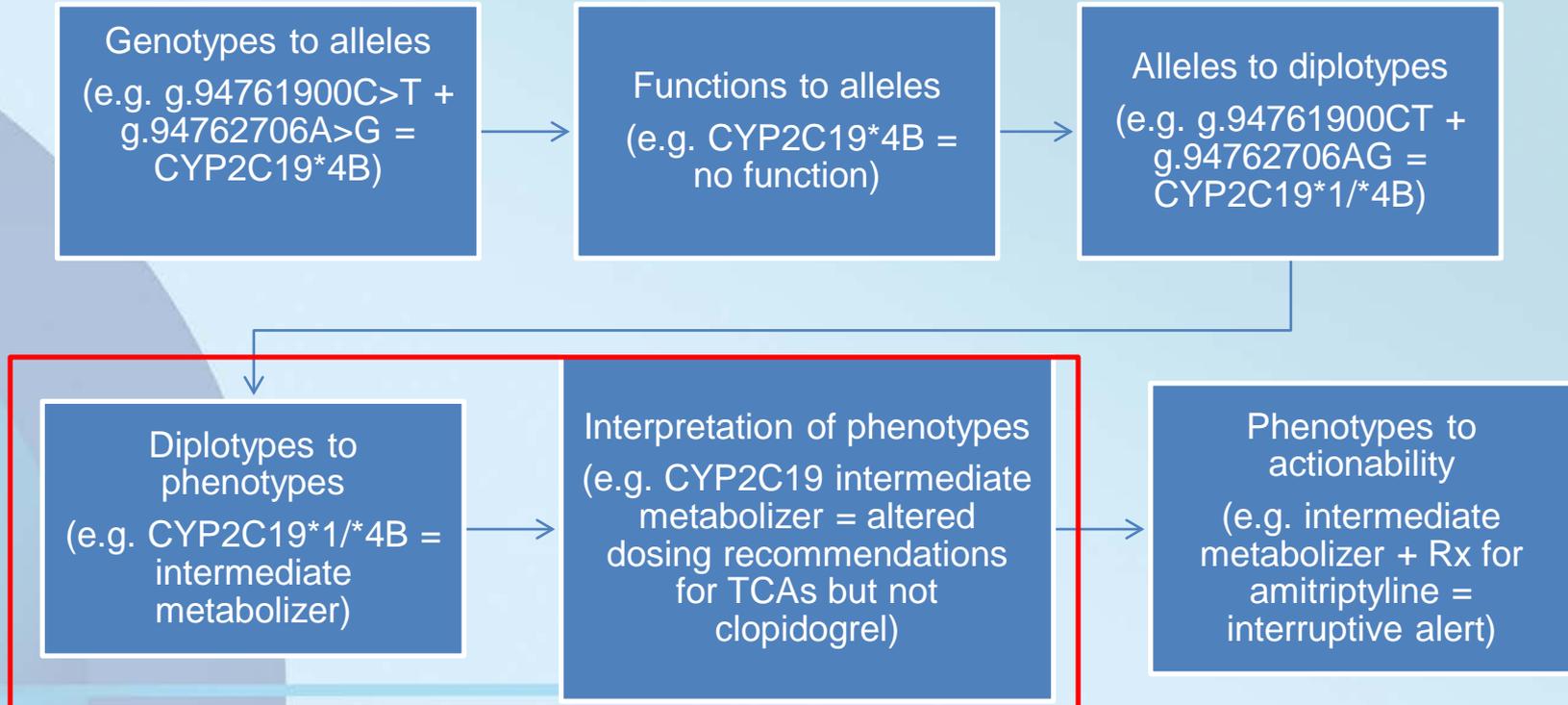
Variants must be phased to assign diplotypes for pharmacogenes

CPIC Gene	Var/var different than var/wt?
<i>TPMT</i>	Yes
<i>CYP2C19</i>	Yes
<i>CYP2D6</i>	Yes
<i>DPYD</i>	Yes
<i>CYP2C9</i>	Yes
<i>SLCO1B1</i>	Yes
<i>HLA-B</i>	No
<i>VKORC1</i>	Yes
<i>IL28-B</i>	Yes
<i>CFTR</i>	No
<i>G6PD</i>	Yes
<i>UGT1A1</i>	Yes
<i>CYP3A5</i>	Yes

CYP2D6: 207 diplotypes observed in first 4046 pts on PG4KDS

TPMT is much simpler

CPIC tables allow translation of genetic test results to actionability



<https://cpicpgx.org/guidelines/>

<https://www.pharmgkb.org/page/cyp2c19RefMaterials>

Genotype to phenotype assignment based on allele function

Table 1 Assignment of likely CYP2C19 phenotypes based on genotypes

Likely phenotype	Genotypes ^a	Examples of CYP2C19 diplotypes
CYP2C19 ultrarapid metabolizer (~2–5% of patients) ^b	An individual carrying two increased function alleles	*17/*17
CYP2C19 rapid metabolizer (~2–30% of patients) ^b	An individual carrying one normal function allele and one increased function allele	*1/*17
CYP2C19 normal metabolizer ^c (~35–50% of patients) ^b	An individual carrying two normal function alleles	*1/*1
CYP2C19 intermediate metabolizer (~18–45% of patients) ^b	An individual carrying one normal function allele and one no function allele or one no function allele and one increased function allele	*1/*2, *1/*3, *2/*17 ^d
CYP2C19 poor metabolizer (~2–15% of patients) ^b	An individual carrying two no function alleles	*2/*2, *2/*3, *3/*3

Clin Pharmacol Ther. 2017 Jul; 102 (1):45-51.

<https://cpicpgx.org/guidelines/guideline-for-voriconazole-and-cyp2c19/>

Genotype to phenotype assignment based on allele function

Table 1 Assignment of likely CYP2D6 phenotypes based on diplotypes

Likely phenotype	Diplotypes
CYP2D6 Ultrarapid (~1–2% of patients)	2xN ^f
CYP2D6 Normal (~77–92% of patients)	/ *5, 41/ *41
CYP2D6 Intermediate (~2–11% of patients)	
CYP2D6 Poor Metabolizer (~5–10% of patients)	/ *6

There are differences in genotype to phenotype assignment between the CPIC and the DPWG guidelines. We are in the process of working together to resolve these discordances.

CYP2C19_Diplotype_Phenotype_Table

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PROTECTED VIEW Be careful—files from the Internet can contain viruses. Unless you need to edit, it's safer to stay in Protected View.

C23 : X ✓ fx None

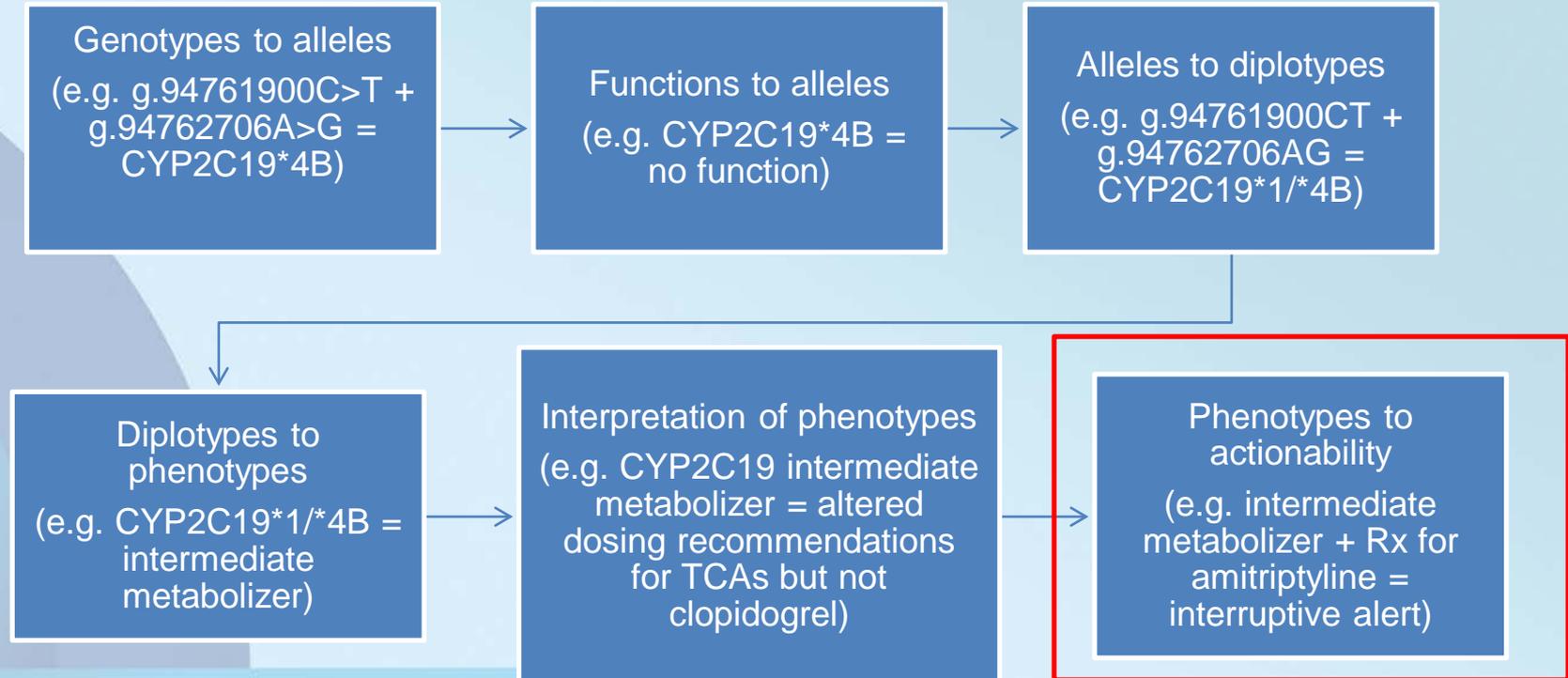
	A	B	C	D	E
1	CYP2C19 Diplotype	Coded Diplotype/Phenotype Summary^a	EHR Priority Result Notation^b		
2	*1/*1	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
3	*1/*2	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
4	*1/*3	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
5	*1/*4A	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
6	*1/*4B	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
7	*1/*5	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
8	*1/*6	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
9	*1/*7	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
10	*1/*8	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
11	*1/*9	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
12	*1/*10	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
13	*1/*11	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
14	*1/*12	Indeterminate	None		
15	*1/*13	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
16	*1/*14	Indeterminate	None		
17	*1/*15	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
18	*1/*16	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
19	*1/*17	CYP2C19 Rapid Metabolizer	Abnormal/Priority/High Risk		
20	*1/*18	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
21	*1/*19	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
22	*1/*22	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
23	*1/*23	Indeterminate	None		
24	*1/*24	CYP2C19 Intermediate Metabolizer	Abnormal/Priority/High Risk		
25	*1/*25	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
26	*1/*26	CYP2C19 Likely Intermediate Metabolizer	Abnormal/Priority/High Risk		
27	*1/*27	Indeterminate	None		
28	*1/*28	CYP2C19 Normal Metabolizer	Normal/Routine/Low Risk		
29	*1/*29	Indeterminate	None		
30	*1/*30	Indeterminate	None		

Possible CYP2C19 Diplotype | 2C19 Interpretation consult note | CYP2C19 Implementation work ...

<https://www.pharmgkb.org/page/cyp2c19RefMaterials>
<https://cpicpgx.org/guidelines/guideline-for-voriconazole-and-cyp2c19/>



CPIC tables allow translation of genetic test results to actionability



<https://cpicpgx.org/guidelines/>

<https://www.pharmgkb.org/page/cyp2c19RefMaterials>

Problem List

Management Discipline View All Problems Change View

Qualifier	Name of Problem	Onset Date	Classification
<input type="checkbox"/>	All Problems		
	ACUTE LYMPHOCYTIC LEUKEMIA	5/2/2011	HIMS Sum...
	ALL (acute lymphoblastic leukemia)	5/11/2011	HIMS Sur
	Consented to all optional research testing...	6/14/2011	Medical
	CYP2D6 POOR METABOLIZER	5/25/2011	Medical
	LOW RISK CONSOL T16	6/23/2011	Medical
	Peg Asp 2500 u/m2/IV randomized	2011	Medical
	PT. HAS HICKMAN LINE SINGLE LUMEN	5/2/2011	Medi
<input type="checkbox"/>	PT. HAS SUBQPORT SINGLE	12/17/2013	Medical
	TPMT INTERMEDIATE METABOLIZER	2/15/2012	Medical

Drive CDS off of problem list entry

Discern (2 of 2)

Cerner

WARNING

Based on the genotype result, this patient is predicted to be a TPMT-INTERMEDIATE METABOLIZER. The patient is at risk for myelosuppression with normal doses of 6-mercaptopurine. Consider starting 6-mercaptopurine doses at 30 - 70% of the normal dose. Please consult a clinical pharmacist or review the pharmacogenetics tab for more information.

Alert Action

Cancel entry
 Dose altered accordingly
 Modify

History Add Info OK

Post-test alert can incorporate non-genetic info too: based on CYP2C19 phenotype, route of administration, age

Discern: (2 of 2)

 **POOR METABOLIZER**

Based on the genotype result, this patient is predicted to be a **CYP2C19 POOR METABOLIZER**. If voriconazole is prescribed to a CYP2C19 poor metabolizer adverse events are likely. **For a patient 12 years of age or older and a CYP2C19 PM phenotype**, initiate voriconazole at a reduced dose of **200 mg PO Q12H** and follow up with therapeutic drug monitoring. Please consult a clinical pharmacist, review the pharmacogenetics tab or click on the link below for more information.

Alert Action

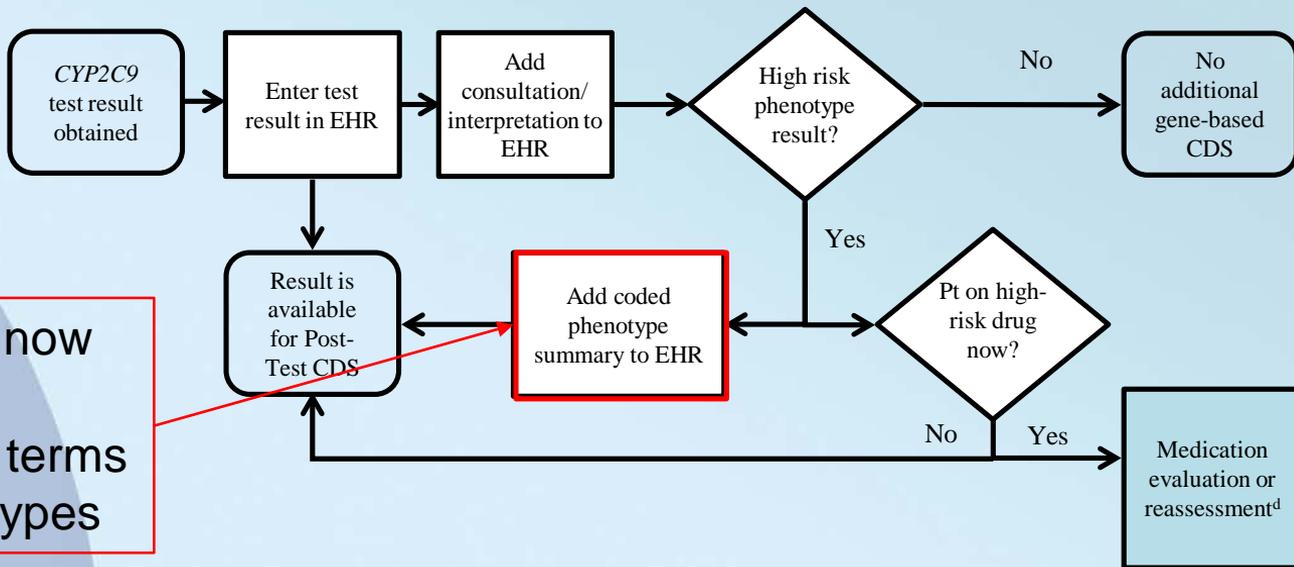
Check BELOW for age and phenotype adjusted dose

Continue with different dose

Add Order for:

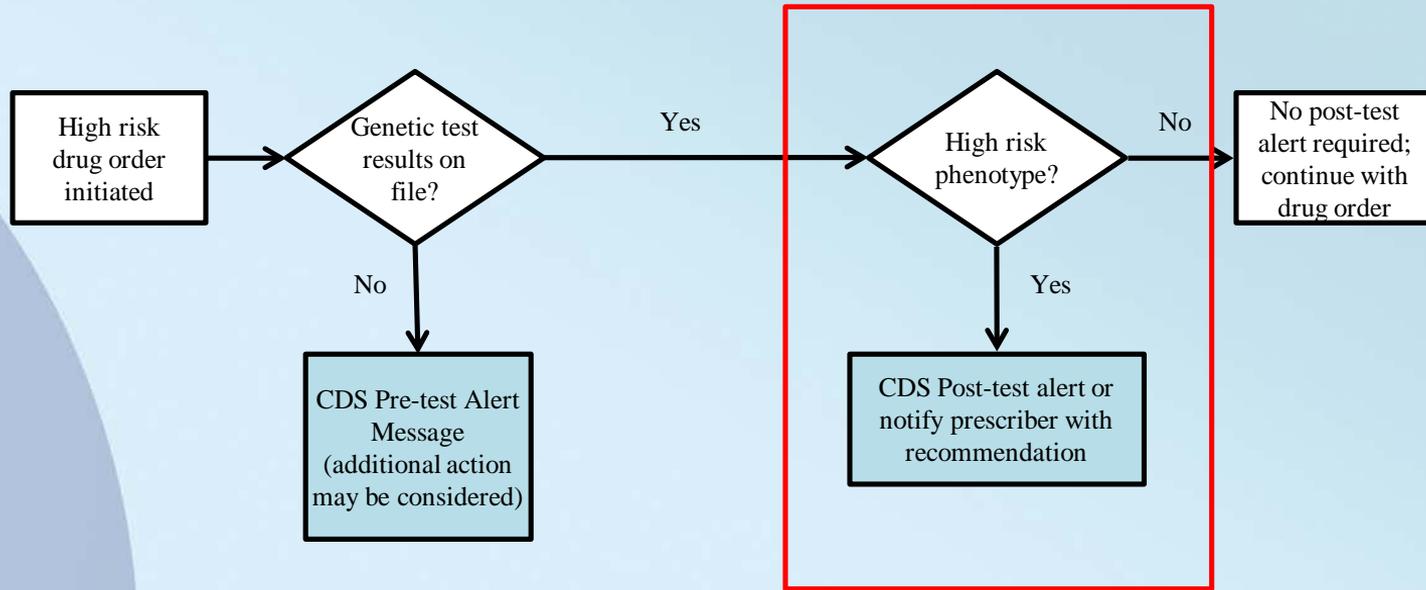
Voriconazole oral -> 200 mg = PO Q12H, Routine, CYP2C19 POOR METABOLIZER. Age 12 years or above

Integrating substrate-dependence into genotype-to-phenotype translation



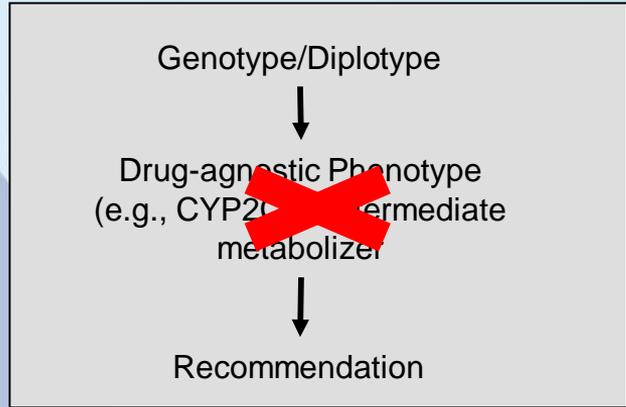
Blue shading indicates interaction with provider

Integrating substrate-dependence into genotype-to-phenotype translation

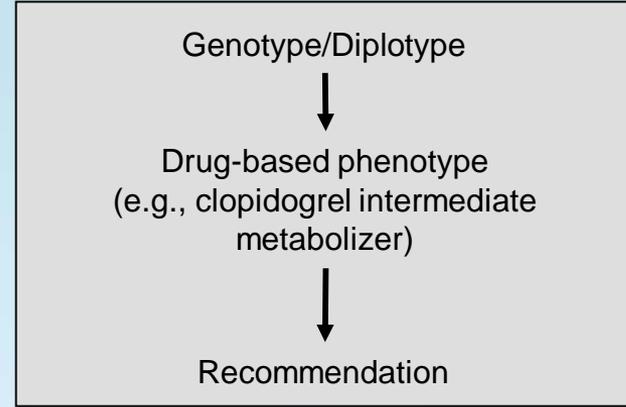


Integrating substrate-dependence into genotype-to-phenotype translation

Option 1:



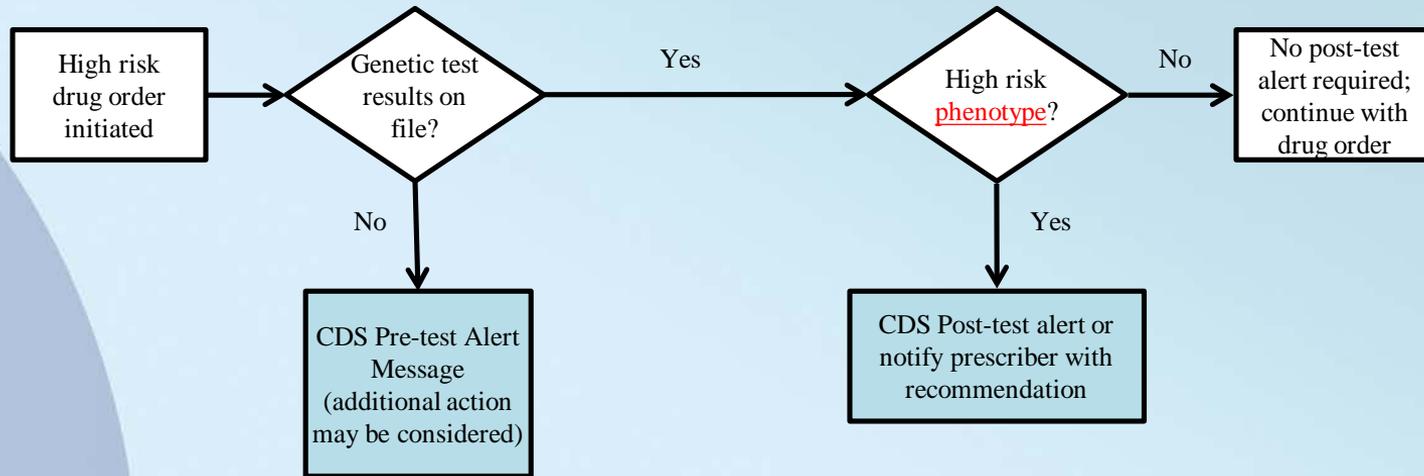
Option 2:



- Is there enough clinical evidence to allow for clinical recommendations based on genotype/diplotype?
- How would this affect current system/implementation?

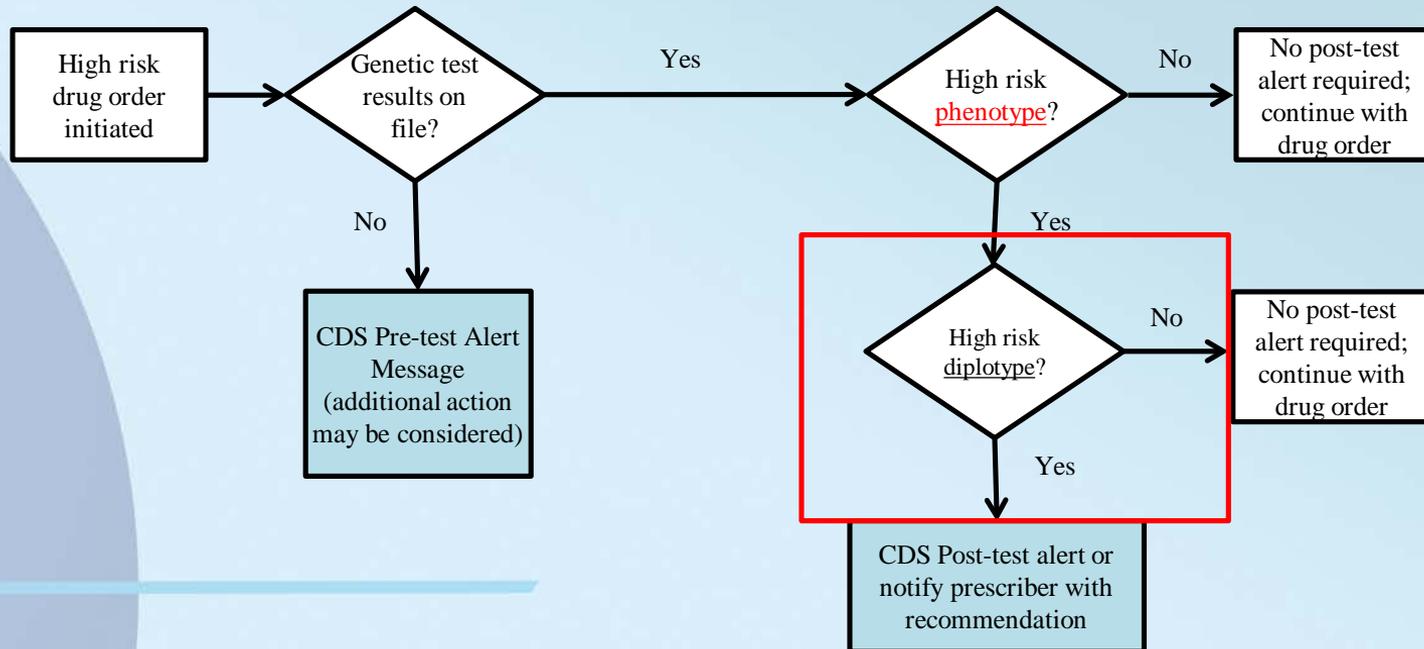
Integrating substrate-dependence into genotype-to-phenotype translation

Change to diplotype or drug-based phenotype?



How would this impact problem list entry or use of standardized terminology such SNOMED?

Integrating substrate-dependence into genotype-to-phenotype translation



Integrating substrate-dependence into genotype-to-phenotype translation

Table 1 Assignment of likely CYP2C19 phenotypes based on genotypes

Likely phenotype	Genotypes	Examples of diplotypes
Ultrarapid metabolizer: normal or increased activity (~5–30% of patients)	An individual carrying two increased activity alleles (*17) or one functional allele (*1) plus one increased-activity allele (*17)	*1/*17, *17/*17
Extensive metabolizer: homozygous wild-type or normal activity (~35–50% of patients)	An individual carrying two functional (*1) alleles	*1/*1
Intermediate metabolizer: heterozygote or intermediate activity (~18–45% of patients)	An individual carrying one functional allele (*1) plus one loss-of-function allele (*2–*8) or one loss-of-function allele (*2–*8) plus one increased-activity allele (*17)	*1/*2, *1/*3, *2/*17
Poor metabolizer: homozygous variant, mutant, low, or deficient activity (~2–15% of patients)	An individual carrying two loss-of-function alleles (*2–*8)	*2/*2, *2/*3, *3/*3

Some rare genotype combinations have unclear predicted metabolic phenotypes; see **Supplementary Table S5** online.

Integrating substrate-dependence into genotype-to-phenotype translation

Table 2 Antiplatelet therapy recommendations based on CYP2C19 status when considering clopidogrel for ACS/PCI patients

Phenotype (genotype)	Implications for clopidogrel	Therapeutic recommendations	Classification of recommendations ^a
Ultrarapid metabolizer (UM) (*1/*17, *17/*17) and extensive metabolizer (EM) (*1/*1)	Normal (EM) or increased (UM) platelet inhibition; normal (EM) or decreased (UM) residual platelet aggregation ^b	Clopidogrel: label-recommended dosage and administration	Strong
Intermediate metabolizer (*1/*2, *1/*3, *2/*17)	Reduced platelet inhibition; increased residual platelet aggregation; increased risk for adverse cardiovascular events	Alternative antiplatelet therapy (if no contraindication), e.g., prasugrel, ticagrelor	Moderate
Poor metabolizer (*2/*2, *2/*3, *3/*3)	Significantly reduced platelet inhibition; increased residual platelet aggregation; increased risk for adverse cardiovascular events	Alternative antiplatelet therapy (if no contraindication), e.g., prasugrel, ticagrelor	Strong

Integrating substrate-dependence into genotype-to-phenotype translation

Table 3 Dosing recommendations for voriconazole treatment based on CYP2C19 phenotype for pediatric patients (children and adolescents <18 years old)

CYP2C19 phenotype	Implications for voriconazole pharmacologic measures	Therapeutic recommendations	Classification of recommendations ^a
CYP2C19 ultrarapid metabolizer (*17/*17)	In patients for whom an ultrarapid metabolizer genotype (*17/*17) is	Choose an alternative agent that is not dependent on CYP2C19 metabo-	Moderate

BUT BEFORE THIS CHANGE....

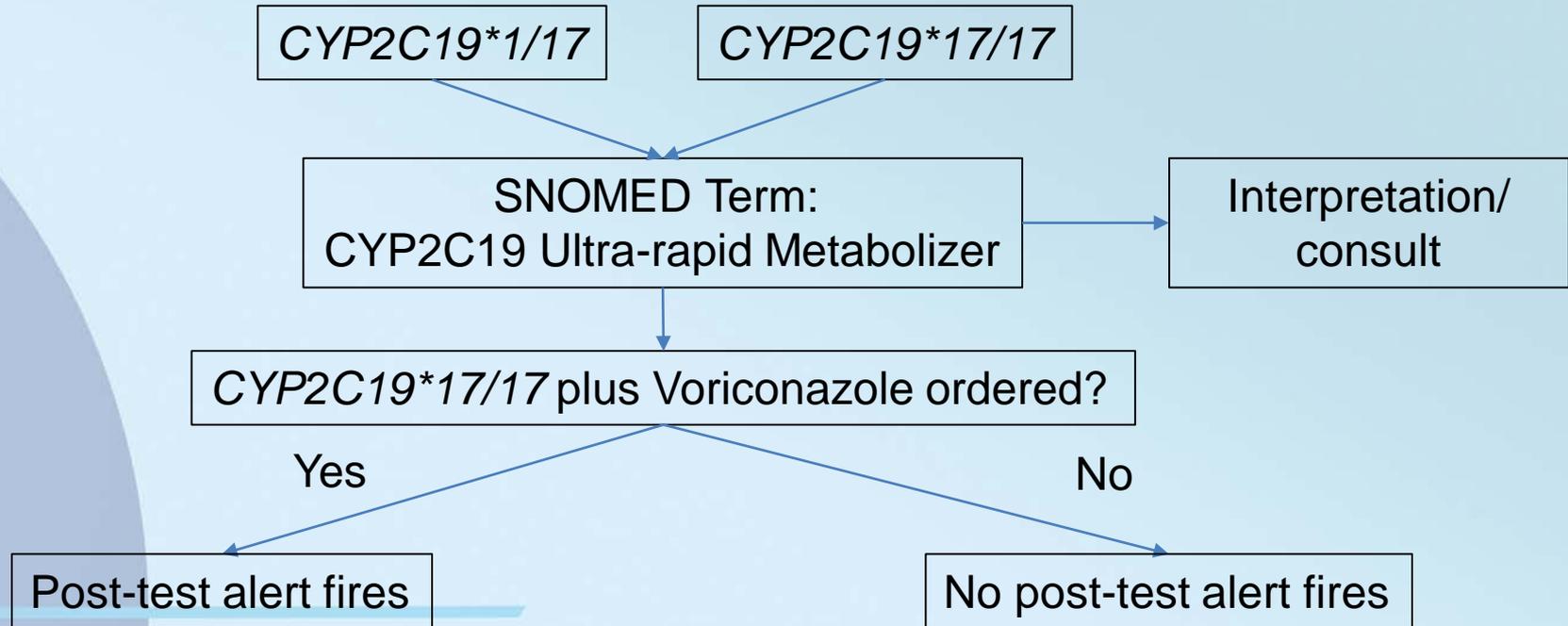
CYP2C19 poor metabolizer

Higher dose required to reach concentrations of voriconazole and may increase probability of adverse events

Choose an alternative agent that is not dependent on CYP2C19 metabolism as primary therapy in lieu of voriconazole. Such agents include liposomal amphotericin B and posaconazole.^{b,e}
In the event that voriconazole is considered to be the most appropriate agent, based on clinical advice, for a patient with poor metabolizer genotype, voriconazole should be administered at a preferably lower than standard dosage with careful therapeutic drug monitoring.

Moderate

Integrating substrate-dependence into genotype-to-phenotype translation



Why not so straightforward?

- SNOMED CT terms
- Clinical evidence
- Scalability/Shareable
- Diplotype not always a discrete field in EHR

PharmCAT

To automate the annotation of .vcf files with the appropriate haplotypes or diplotypes from the CPIC guideline genes, and generate a report with the corresponding CPIC guideline prescribing recommendations



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