STE0097

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ACE INHIBITORS, PLAIN

ne Royal Dutch Pharmacists Associate eutic dose recommendations for dulc nclude that there are no recommendations	exetine based on C		
Phenotype (Genotype)	Therapeutic Dose Recommendation	Level of Evidence	Clinical Relevance
PM (two inactive (*3-*8, *11-*16, *19-*21, *38, *40, *42) alleles)	No recommendation.	Data on file.	Clinical effect (not statistically significant difference).
IM (two decreased-activity (*9, *10, *17, *29, *36, *41) alleles or carrying one active (*1, *2, *33, *35) and one inactive (*3-*8, *11-*16, *19-*21, *38, *40, *42) allele, or carrying one decreased-activity (*9, *10, *17, *29, *36, *41) allele and one inactive (*3-*8, *11-*16, *19-*21, *38, *40, *42) allele)	No recommendation.	No evidence.	_
UM (a gene duplication in absence of inactive (*3-*8, *11-*16, *19-*21, *38, *40, *42) or decreased-activity (*9, *10, *17, *29, *36, *41) alleles)	No recommendation.	No evidence.	-
	- Clinical Annotati	ons —	
class 3 rs1042713 GG atients with the GG genotype and head spital utilization when treated with G genotype. Other genetic and clinical DRB1	cardiovascular dru	gs as compared to influence efficacy of	patients with the AA

when treated with cardiovascular drugs as compared to patients with the GG genotype. Other genetic

and clinical factors may also influence efficacy of cardiovascular drugs.

ANALGESICS

\mathbf{COMT}	catechol-O-methyltransferase
of headache when discording compared to patients we likelihood of headache in	notype with substance withdrawal syndrome may have an increased likelihood attinuing the use of analgesics (such as opioids, NSAIDs, triptans, ergot) as ith the AA genotype. Other clinical and genetic factors may also influence patients with withdrawal syndrome who discontinue the use of analgesics. Y AGENTS, NON-STEROIDS
COMT	catechol-O-methyltransferase
of headache when discor- compared to patients we likelihood of headache in	notype with substance withdrawal syndrome may have an increased likelihood attinuing the use of analgesics (such as opioids, NSAIDs, triptans, ergot) as ith the AA genotype. Other clinical and genetic factors may also influence patients with withdrawal syndrome who discontinue the use of analgesics. EATMENT OF HIV INFECTIONS, COMBINATIONS
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	- Clinical Annotations —
Patients with the GG georeased risk for hepatoto	GG enotype who are co-infected with HIV and tuberculosis (TB) may have a dexicity when treated with anti-tubercular and antiretroviral drugs as compared genotype. Other genetic and clinical factors may also influence risk of hepato-
CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
	- Clinical Annotations
creased risk for hepatoto	GG enotype who are co-infected with HIV and tuberculosis (TB) may have a dexicity when treated with anti-tubercular and antiretroviral drugs as compared genotype. Other genetic and clinical factors may also influence risk of hepato-

————- Clinical Annotations ———

cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene $1\,$

CYP2A7P1

	Class 3	rs3745274	GG
•	Class o	150140414	GG

Patients with the GG genotype who are co-infected with HIV and tuberculosis (TB) may have a decreased risk for hepatotoxicity when treated with anti-tubercular and antiretroviral drugs as compared to patients with the TT genotype. Other genetic and clinical factors may also influence risk of hepatotoxicity.

BETA BLOCKING AGENTS

ADRB2	adrenoceptor beta 2, surface
Patients with the GG gene hospital utilization when	otype and heart failure may have increased emergency department visits and treated with cardiovascular drugs as compared to patients with the AA or tic and clinical factors may also influence efficacy of cardiovascular drugs. adrenoceptor beta 1
	- Clinical Annotations
	Chinom Thinocomolis
Patients with the CC gend when treated with cardiov and clinical factors may a	otype and heart failure may have increased emergency department utilization rescular drugs as compared to patients with the GG genotype. Other genetic lso influence efficacy of cardiovascular drugs.
DABIGATRAN	
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
People with the GG genore the AA and AG genotype factors may affect exposure. Class 4 rs2032582 Compared the CC genotype with the CC genotype variant at this position, in	type may have decreased exposure to dabigatran compared to patients with es, when also assessed with the rs2032582 allele. Other clinical and genetic re to dabigatran. CC ype may have decreased exposure to dabigatran compared to patients with a cluding genotypes AA, AC, CT, and TT, when assessed in conjunction with 45642. Other clinical and genetic factors may affect exposure to dabigatran.
DRUGS FOR TREATME	ENT OF TUBERCULOSIS
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1

-- Clinical Annotations --

Class 3	rs1045642	GG

Patients with the GG genotype and tuberculosis (TB) may have a decreased risk for hepatotoxicity when treated with anti-TB drugs as compared to patients with the AA genotype. Other genetic and clinical factors may also influence hepatotoxicity.

· Class 3 rs1045642 GG

Patients with the GG genotype who are co-infected with HIV and tuberculosis (TB) may have a decreased risk for hepatotoxicity when treated with anti-tubercular and antiretroviral drugs as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of hepatotoxicity.

CYP2E1	cytochrome P450, family 2, subfamily E, polypeptide 1
when treated with ant majority of studies fin	CC genotype and tuberculosis (TB) may have an increased risk for hepatotoxicity i-TB drugs as compared to patients with the CT or TT genotype. However, the d no association with hepatotoxicity. Other genetic and clinical factors, such as 2 gene, may also influence risk for hepatotoxicity.
DUX1	double homeobox 1
	- Clinical Annotations —
when treated with ant majority of studies fin	CC genotype and tuberculosis (TB) may have an increased risk for hepatotoxicity i-TB drugs as compared to patients with the CT or TT genotype. However, the d no association with hepatotoxicity. Other genetic and clinical factors, such as 2 gene, may also influence risk for hepatotoxicity.
CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
creased risk for hepato	GG genotype who are co-infected with HIV and tuberculosis (TB) may have a detoxicity when treated with anti-tubercular and antiretroviral drugs as compared T genotype. Other genetic and clinical factors may also influence risk of hepato-
CYP2A7P1	cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene 1
	- Clinical Annotations

· Class 3 rs3745274 GG

Patients with the GG genotype who are co-infected with HIV and tuberculosis (TB) may have a decreased risk for hepatotoxicity when treated with anti-tubercular and antiretroviral drugs as compared to patients with the TT genotype. Other genetic and clinical factors may also influence risk of hepatotoxicity.

N-acetyltransferase 2 (arylamine N-acetyltransferase)

 Clinical Annotations

· Class 3 rs1799929 CC

Patients with the CC genotype and tuberculosis (TB) may have a decreased risk for hepatotoxicity when treated with anti-TB drugs as compared to patients with the CT or TT genotype. This SNP is present in a variety of NAT2 * alleles resulting in different NAT2 acetylator phenotypes, and is the signature SNP of NAT2*11. Other genetic and clinical factors may also influence hepatotoxicity.

OPIUM ALKALOIDS AND DERIVATIVES

\cdot Class 3 rs16969968 GG

Patients with the GG genotype who are in chronic pain and receive opioid medications for treatment may be at decreased risk for addiction as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of opiate addiction.

CHRNA5	cholinergic receptor, nicotinic, alpha 5 (neuronal)
	—- Clinical Annotations ————————————————————————————————————

· Class 3 rs16969968 GG

Patients with the GG genotype who are in chronic pain and receive opioid medications for treatment may be at decreased risk for addiction as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of opiate addiction.

PLATINUM COMPOUNDS

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	Clinical Annotations

· Class 3 rs1045642 GG

Patients with the GG genotype and non-small-cell lung cancer may have a better response to platinum-based chemotherapy as compared to patients with the AA or AG genotype. This was only seen in those of Asian ethnicity. Other genetic and clinical factors may also influence response to platinum-based chemotherapy.

· Class 3 rs1128503 GG

Patients with the GG genotype and non-small cell lung cancer may have reduced risk of toxicities when treated with platinum-based chemotherapy compared to patients with the AA genotype. Other clinical and genetic factors may affect risk of toxicities in response to platinum-based chemotherapies.

PYRIMIDINE ANALOGUES

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· Class 1A rs55886062 AA

Patients with the AA genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have a decreased, but not absent, risk for drug toxicity as compared to patients with the AC or CC genotype (DPYD *1/*13 or *13/*13). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 1A rs3918290 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of fluoropyrimidine drugs and 2) decreased, but not non-existent, risk for drug toxicity as compared to patients with the CT or TT genotype (DPYD *1/*2A or *2A/*2A). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine based chemotherapy.

\cdot Class 1A rs67376798 TT

Patients with the TT genotype and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of the drug and 2) decreased, but not absent, risk and reduced severity of drug toxicity as compared to patients with the AT genotype. Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 3 rs1801159 TT

Patients with the TT genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased likelihood of nausea, vomiting, and leukopenia, 2) increased response and 3) increased clearance of fluorouracil as compared to patients with the CT or CC genotype (DPYD *1/*5 or *5/*5). However, other studies find no associations or contradictory associations with fluoropyrimidine-induced drug toxicity or response. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

ACENOCOUMAROL

CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9

· Class 2A rs1057910 AC

Patients with the AC genotype may require decreased dose of acenocoumarol or closer INR monitoring
as compared to patients with the AA genotype. Other genetic and clinical factors may also influence
acenocoumarol dose.

Class 3 rs1799853 CC

Patients with the CC genotype who are taking acenocoumarol may have a decreased risk of a gastrointestinal hemorrhage as compared to patients with the CT or TT genotype. Other genetic and clinical factors may also influence risk of gastrointestinal hemorrhage.

VKORC1	vitamin K epoxide reductase complex, subunit 1
	- Clinical Annotations
0 01	may have decreased dose of acenocoumarol or phenprocoumon as pe GG. Other genetic and clinical factors may also influence the dose non.
PRSS53	protease, serine, 53

Class 2A rs9934438 GA

Patients with the AG genotype may have decreased dose of acenocoumarol or phenprocoumon as compared to patients with genotype GG. Other genetic and clinical factors may also influence the dose of acenocoumarol or phenprocoumon.

ACETAMINOPHEN

UGT1A	UDP glucuronosyltransferase 1 family, polypeptide A complex locus

Class 3 rs8330

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

Class 3 rs10929303

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10	
	—- Clinical Annotations	

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

UGT1A3	UDP glucuronosyltransferase 1 family, polypeptide A3
	— Clinical Annotations —

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

UGT1A4	UDP glucuronosyltransferase 1 family, polypeptide A4
	Clinical Annotations ————————————————————————————————————

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

\cdot Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

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· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

$\mathbf{UGT1A6}$	UDP glucuronosyltransferase 1 family, polypeptide A6

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7	

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

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· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

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UDP glucuronosyltransferase 1 family, polypeptide A9

	Clinical	Annotations	
•	CHITICAL	ATHOGRAFIONS	

· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

UGT1A1

UDP glucuronosyltransferase 1 family, polypeptide A1

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· Class 3 rs8330 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs1042640 GC

Patients with the CG genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

· Class 3 rs10929303 TC

Patients with the CT genotype may have a decreased risk of liver failure due to unintentional acetaminophen overdose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence risk of liver failure due to unintentional acetaminophen overdose.

CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
Women with the TT genot mean arterial pressure of ;= CC genotype. No significan	Type and hypertension may have an increased likelihood of reaching a target = 107 mm Hg when treated with amlodipine as compared to women with the nt associations were seen when considering a target mean arterial pressure of sidering men or men and women together. Other genetic and clinical factors to amlodipine.
CYP3A	cytochrome P450, family 3, subfamily A
	- Clinical Annotations
Women with the TT genot mean arterial pressure of j= CC genotype. No significan j= 92 mm Hg, or when commay also influence response. Class 3 rs776746 CC Healthy males with the CC as compared to healthy materials were seen when consignificance metabolism of an influence metabolism of an influence metabolism.	C (CYP3A5 *3/*3) genotype may have increased metabolism of amlodipine ales with the CT or TT (*3/*1 or *1/*1) genotype. No significant associated idering clearance of amlodipine. Other genetic and clinical factors may also mlodipine.
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
	- Clinical Annotations
as compared to healthy ma	C (CYP3A5 *3/*3) genotype may have increased metabolism of amlodipine ales with the CT or TT (*3/*1 or *1/*1) genotype. No significant associated clearance of amlodipine. Other genetic and clinical factors may also
ZSCAN25	zinc finger and SCAN domain containing 25
	- Clinical Annotations

· Class 3 rs776746 CC

Healthy males with the CC (CYP3A5 *3/*3) genotype may have increased metabolism of amlodipine as compared to healthy males with the CT or TT (*3/*1 or *1/*1) genotype. No significant associations were seen when considering clearance of amlodipine. Other genetic and clinical factors may also influence metabolism of amlodipine.

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
	Clinical Annotations

· Class 4 rs3732219 CC

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

· Class 4 rs3732218 GG

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

UGT1A4	UDP glucuronosyltransferase 1 family, polypeptide A4

· Class 4 rs3732219 CC

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

\cdot Class 4 rs3732218 GG

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

UGT1A5	UDP glucuronosyltransferase 1 family, polypeptide A5

· Class 4 rs3732219 CC

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

\cdot Class 4 rs3732218 GG

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

		— Clinical Annotations –	
Class 4 rs3732219	CC		

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

Class 4 rs3732218

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7
	——- Clinical Annotations ————————————————————————————————————

Class 4 rs3732219 *CC*

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

Class 4 rs3732218

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8

Class 4 rs3732219 *CC*

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

Class 4 rs3732218

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

UGT1A9	UDP glucuronosyltransferase 1 family, polypeptide A9
	— Clinical Annotations

Class 4 rs3732219 *CC*

Patients with the CC genotype may have increased glucuronidation of anastrozole as compared to patients with the CT or TT genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

\cdot Class 4 rs3732218 GG

Patients with the GG genotype may have increased glucuronidation of anastrozole as compared to patients with the AA or AG genotype, as determined by in vitro assays. Glucuronidation allows for the elimination of xenobiotics like anastrozole. Other genetic and clinical factors may also influence glucuronidation of anastrozole.

ANTHRACYCLINES AND RELATED SUBSTANCES

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	- Clinical Annotations —
decreased response to anthrac	ype may have 1) decreased exposure to doxorubicin metabolites and 2) cycline regimens as compared to patients with the AA genotype, however dictory. Other genetic and clinical factors may also influence response to
UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
thracyclines and related subst TT and decreased likelihood	rpe may have increased likelihood of cardiotoxicity when exposed to antances in children with Neoplasms as compared to patients with genotype as compared to patients with the GG genotype, although this has been ther genetic and clinical factors may also influence the risk of toxicity to

· Class 3 rs17863783 GG

anthracyclines and related substances.

Patients with the GG genotype may have decreased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype GT or TT, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

\cdot Class 3 rs6759892 TG

Patients with the GT genotype may have increased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype TT and decreased likelihood as compared to patients with the GG genotype, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

· Class 3 rs17863783 GG

Patients with the GG genotype may have decreased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype GT or TT, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

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UDP glucuronosyltransferase 1 family, polypeptide A7

-- Clinical Annotations -

\cdot Class 3 rs6759892 TG

Patients with the GT genotype may have increased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype TT and decreased likelihood as compared to patients with the GG genotype, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

· Class 3 rs17863783 GG

Patients with the GG genotype may have decreased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype GT or TT, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

UGT1A8

UDP glucuronosyltransferase 1 family, polypeptide A8

————- Clinical Annotations ————

· Class 3 rs6759892 TG

Patients with the GT genotype may have increased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype TT and decreased likelihood as compared to patients with the GG genotype, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

· Class 3 rs17863783 GG

Patients with the GG genotype may have decreased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype GT or TT, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

UGT1A9

UDP glucuronosyltransferase 1 family, polypeptide A9

-- Clinical Annotations

\cdot Class 3 rs6759892 TG

Patients with the GT genotype may have increased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype TT and decreased likelihood as compared to patients with the GG genotype, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

· Class 3 rs17863783 GG

Patients with the GG genotype may have decreased likelihood of cardiotoxicity when exposed to anthracyclines and related substances in children with Neoplasms as compared to patients with genotype GT or TT, although this has been contradicted in one study. Other genetic and clinical factors may also influence the risk of toxicity to anthracyclines and related substances.

ANTIEPILEPTICS

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	- Clinical Annotations
Class 3 rs1128503	GG
Patients with the GG go decreased risk for resistan However, all other studies	enotype and specifically localization-related epilepsy syndrome may have ance to antiepileptic treatment as compared to patients with the AA genotypes of people with epilepsy have found no association between this variant and other genetic and clinical factors may also influence resistance to antiepileptics
SCN1A	sodium channel, voltage-gated, type I, alpha subunit
particularly carbamazeping factors may also influence	
particularly carbamazepin	ne, as compared to patients with the TT genotype. Other genetic and clinical
particularly carbamazeping factors may also influence NTIPSYCHOTICS	ne, as compared to patients with the TT genotype. Other genetic and clinical eresistance to antiepileptic drugs.
particularly carbamazeping factors may also influence of the section of the secti	ne, as compared to patients with the TT genotype. Other genetic and clinical eresistance to antiepileptic drugs. catechol-O-methyltransferase
particularly carbamazepin factors may also influence of the second of th	catechol-O-methyltransferase ———————————————————————————————————
class 3 rs4680 GA Patients with the AG ger as compared to patients Class 3 rs4680 GA Patients with the AG ger as compared to patients Class 3 rs4680 GA Patients with the AG ger chotics as compared to patients	catechol-O-methyltransferase ———————————————————————————————————
class 3 rs4680 GA Patients with the AG ger as compared to patients Class 3 rs4680 GA Patients with the AG ger as compared to patients Class 3 rs4680 GA Patients with the AG ger chotics as compared to patients	catechol-O-methyltransferase - Clinical Annotations notype may have increased blood pressure when treated with antipsychotics with the GG genotype. Other genetic and clinical factors may also influences receiving antipsychotics. notype may have increased fasting glucose levels when treated with antipsycatients with the GG genotype. Other genetic and clinical factors may also

Class 3 rs2032582 CC

Patients with the CC genotype and schizophrenia who responded to treatment with antipsychotics may require an increased dose of antipsychotics as compared to patients with the AA genotype. Other genetic and clinical factors may also influence dose of antipsychotics.

· Class 3 rs1045642 GG

Patients with the GG genotype and schizophrenia who responded to treatment with antipsychotics may require an increased dose of antipsychotics as compared to patients with the AA genotype. Other genetic and clinical factors may also influence dose of antipsychotics.

ATA	7	TAT	A T	7 TD
A I A		N I	ΑI	/ I K

UGT1A UDP glucuronosyltransferase 1 family, polypeptide A complex locus - Clinical Annotations

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
	- Clinical Annotations

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

|--|

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

$\mathbf{UGT1A4}$	UDP glucuronosyltransferase 1 family, polypeptide A4

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A5	UDP glucuronosyltransferase 1 family, polypeptide A5

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A6

UDP glucuronosyltransferase 1 family, polypeptide A6

— Clinical Annotations	
- Chinca Annotations	

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

\cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGTIA7	UDP glucuronosyltransferase I family, polypeptide A7

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

\cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

$\mathbf{UGT1A9}$	UDP glucuronosyltransferase 1 family, polypeptide A9

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A1	UDP glucuronosyltransferase 1 family, polypeptide A1 $$
	— Dosing Guideline —

UGT1A1:*1/*80 Strong

There is no need to avoid prescribing of atazanavir based on UGT1A1 genetic test result. Inform the patient that some patients stop atazanavir because of jaundice (yellow eyes and skin), but that this

	Clinical Approximations
Class 3 rs10929303	3 TC
Patients with the CT with atazanavir and rit nephrolithiasis as comp	genotype and HIV may have a decreased risk of nephrolithiasis when treated to navir as compared to patients with the TT genotype and an increased risk opered to people with the CC genotype. Other genetic and clinical factors may rolithiasis in patients with HIV who are taking atazanavir and ritonavir.
with atazanavir and rit nephrolithiasis as comp	genotype and HIV may have a decreased risk of nephrolithiasis when treated to a compared to patients with the GG genotype and an increased risk of patients with the CC genotype. Other genetic and clinical factors may relithiasis in people with HIV who are taking atazanavir and ritonavir. C
with atazanavir and rit nephrolithiasis as comp	genotype and HIV may have a decreased risk of nephrolithiasis when treated to navir as compared to patients with the GG genotype and an increased risk of pared to people with the CC genotype. Other genetic and clinical factors may rolithiasis in patients with HIV who are taking atazanavir and ritonavir.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	- Clinical Annotations
to patients with the A	genotype and HIV may have increased concentrations of atazanavir as compared A genotypes, although this is contradicted in one study. There is no evidence in the little of the contradicted in the contradict
Other clinical and general HIV.	
Other clinical and general HIV.	
Other clinical and general HIV.	etic factors may also influence the concentrations of atazanavir in patients wit
Other clinical and general HIV. FENOLOL	etic factors may also influence the concentrations of atazanavir in patients wit
Other clinical and general HIV. FENOLOL ADRB2	adrenoceptor beta 2, surface ———————————————————————————————————
Other clinical and general HIV. FENOLOL ADRB2 Class 3 rs1042714 Patients with the GG triglyceridemia when the general triglyceridemia when triglyceri	adrenoceptor beta 2, surface ———————————————————————————————————
Other clinical and general HIV. FENOLOL ADRB2 Class 3 rs1042714 Patients with the GG triglyceridemia when the general triglyceridemia when triglyceri	GG genotype and hypertension may have an increased risk of developing hypereated with atenolol or metoprolol as compared to patients with the CC or CC
Other clinical and general HIV. FENOLOL ADRB2 Class 3 rs1042714 Patients with the GG triglyceridemia when the genotype. Other genetic	adrenoceptor beta 2, surface ———————————————————————————————————

CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
ent drug when treated with a	pe may be more likely to require a decrease in dose or switch to a differ- atorvastatin or simvastatin as compared to patients with the CC or CT clinical factors may also influence dose of simvastatin or atorvastatin, or ifferent drug.
CYP3A	cytochrome P450, family 3, subfamily A
ent drug when treated with a	pe may be more likely to require a decrease in dose or switch to a differ- atorvastatin or simvastatin as compared to patients with the CC or CT clinical factors may also influence dose of simvastatin or atorvastatin, or ifferent drug.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
· Class 3 rs2032582 CC	

· Class 3 rs1042713 GG

Patients with the GG genotype and hypertension may have a greater decrease in diastolic blood pressure when treated with benazepril as compared to patients with the AA genotype. No significant results have been seen for systolic blood pressure. Additionally, the same study reported no significant differences in systolic or diastolic blood pressure between genotypes in a different cohort. Other genetic and clinical factors may also influence change in diastolic or systolic blood pressure.

-- Clinical Annotations

BEVACIZUMAB

ADRB2

adrenoceptor beta 2, surface

	——————————————————————————————————————
· Class 3 rs2010963 CG	
Patients with the CG genotype	e and choroidal neovascularization may have a better response to anti- l to patients with the CC genotype. Other genetic and clinical factors anti-VEGF treatment.
capecitabine, fluorouracil, irino	and the AC genotype may have a reduced response to bevacizumab, stecan, leucovorin, or oxaliplatin as compared to patients with the CC enetic factors may also affect response to chemotherapy in people with
BUPROPION	
CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
· Class 3 rs3211371 CC	
Patients with the CC genotype treated with bupropion as com	who are smokers may have a lower chance of smoking cessation when apared to patients with the CT or TT genotype, although this is congenetic and clinical factors may also influence likelihood of smoking
· Class 3 rs2279343 AA	
pion as compared to individua	sorder and the AA genotype may have an improved response to buprols with the AG and GG genotypes. Other clinical and genetic factors propion in individuals with tobacco use disorder.
BUSULFAN	
CYP2C19	cytochrome P450, family 2, subfamily C, polypeptide 19

metabolism of busulfan as compared to patients with the CT (*1/*17) or TT (*17/*17) genotype. However, some contradictory evidence exists for this association. Other genetic and clinical factors may also influence metabolism of busulfan.

CYP2C9 cytochrome P450, family 2, subfamily C, polypeptide 9 — Clinical Annotations —

Class 3 rs1799853 CC

Patients with the CC (CYP2C9 *1/*1) genotype undergoing hemopoietic stem cell transplant may have increased metabolism of busulfan as compared to patients with the CT ($^*1/^*2$) or TT ($^*2/^*2$) genotype. Other genetic and clinical factors may also influence metabolism of busulfan.

CAPECITABIN	$\mathbf{C}\mathbf{A}$	ΛPE	\mathbf{CI}	TA	В	IN	١E
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ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
Class 3 rs1045642 GG	
in people with Colorectal Ne are not associated with decre epirubicin and gemcitabine i	hay have increased risk of hand-foot syndrome when treated with capecitabin oplasms as compared to patients with genotype AA. Genotypes AG + GG based clinical outcome when treated with capecitabine, cisplatin, docetaxel, in people with Pancreatic Neoplasms as compared to genotype AA. Other hay influence the response to capecitabine.
in people with Colorectal N	hay have increased risk of hand-foot syndrome when treated with capecitabin eoplasms as compared to patients with genotype AA. Other genetic and
	nence the response to capecitabine.
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to	
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to Other genetic and clinical fa	type and colorectal cancer may have an increased risk of neutropenia or reated with capecitabine as compared to patients with the AA genotype.
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to	type and colorectal cancer may have an increased risk of neutropenia or reated with capecitabine as compared to patients with the AA genotype. ctors may also influence risk of neutropenia or hand-foot syndrome.
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to Other genetic and clinical fa	type and colorectal cancer may have an increased risk of neutropenia or reated with capecitabine as compared to patients with the AA genotype. ctors may also influence risk of neutropenia or hand-foot syndrome. cytidine deaminase
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to Other genetic and clinical factors CDA Class 3 rs602950 AA Cancer patients with the A treated with capecitabine-ba	type and colorectal cancer may have an increased risk of neutropenia or reated with capecitabine as compared to patients with the AA genotype. ctors may also influence risk of neutropenia or hand-foot syndrome. cytidine deaminase
Class 3 rs1128503 GG Patients with the GG genot hand-foot syndrome when to Other genetic and clinical factors CDA Class 3 rs602950 AA Cancer patients with the A treated with capecitabine-ba	cytidine deaminase A genotype may have a decreased risk of neutropenia or cytidine deaminase A genotype may have a decreased risk of diarrhea or dehydration when sed therapy as compared to patients with the AG or GG genotype.

Class 1A rs55886062 AA

Use label-recommended dosage and administration.

Patients with the AA genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidinebased chemotherapy may have a decreased, but not absent, risk for drug toxicity as compared to patients with the AC or CC genotype (DPYD *1/*13 or *13/*13). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or

———- Clinical Annotations –

with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

Class 1A rs3918290 *CC*

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of fluoropyrimidine drugs and 2) decreased, but not non-existent, risk for drug toxicity as compared to patients with the CT or TT genotype (DPYD *1/*2A or *2A/*2A). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine based chemotherapy.

\cdot Class 1A rs67376798 TT

Patients with the TT genotype and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of the drug and 2) decreased, but not absent, risk and reduced severity of drug toxicity as compared to patients with the AT genotype. Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 3 rs1801160 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased metabolism of fluorouracil and 2) decreased risk for drug toxicities as compared to patients with the CT or TT genotype (DPYD *1/*6 or *6/*6). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin) or with other drugs such as bevacizumab, cetuximab, raltitrexed. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

Class 3 rs1801159 TT

Patients with the TT genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased likelihood of nausea, vomiting, and leukopenia, 2) increased response and 3) increased clearance of fluorouracil as compared to patients with the CT or CC genotype (DPYD *1/*5 or *5/*5). However, other studies find no associations or contradictory associations with fluoropyrimidine-induced drug toxicity or response. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 3 rs1801158 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased but not absent risk of toxicity and 2) increased DPYD activity as compared to patients with the CT genotype (DPYD *1/*4). However, some studies find no association with drug toxicity or DPYD activity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

VEGFA	vascular endotnellal growth factor A
	- Chinical Affilotations

· Class 3 rs699947 AC

Patients with colorectal cancer and the AC genotype may have a reduced response to bevacizumab, capecitabine, fluorouracil, irinotecan, leucovorin, or oxaliplatin as compared to patients with the CC

genotype. Other clinical and genetic factors may also affect response to chemotherapy in people with colorectal cancer.

· Class 3 rs2010963 CG

Patients with the CG genotype and colorectal cancer may have a poorer response when treated with capecitabine and oxaliplatin (XELOX) as compared to patients with the CC or GG genotype. Other genetic and clinical factors may also influence response to XELOX treatment.

CARBAMAZEPINE

SCN1A	sodium channel, voltage-gated, type I, alpha subunit

\cdot Class 2B rs3812718 CT

Patients with the CT genotype who are treated with carbamazepine may require a higher dose as compared to patients with the CC genotype. Other genetic and clinical factors may also influence dose of carbamazepine.

· Class 2B rs3812718 CT

Patients with the CT genotype and epilepsy may be less likely to be resistant to antiepileptic treatment, particularly carbamazepine, as compared to patients with the TT genotype. Other genetic and clinical factors may also influence resistance to antiepileptic drugs.

 \cdot Class 3 rs3812718 CT

Patients with epilepsy and the CT genotype may have decreased metabolism of carbamazepine, resulting in increased exposure as compared to patients with the TT genotype.

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1

· Class 3 rs1045642 GG

Patient with genotype GG may have decreased likelihood of drug resistance when treated with antiepileptics and carbamazepine in people with Epilepsy as compared to patients with genotype AA. However, contradictory findings have been reported. Other genetic and clinical factors may also influence response to carbamazepine.

· Class 3 rs1128503 GG

African American and white patients with the GG genotype and epilepsy may have decreased clearance of carbamazepine as compared to patients with the AA or AG genotype. This association was not found in Chinese patients. Other genetic and clinical factors may also influence clearance of carbamazepine.

 \cdot Class 3 rs1045642 GG

Patients with the GG genotype and epilepsy may have decreased metabolism of carbamazepine and may need a decreased dose as compared to patients with the AG genotype. However, multiple studies have shown no association with dose or concentrations of carbamazepine. Other genetic and clinical factors may also influence concentrations of carbamazepine.

ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2

Patients with the CT genotype may have decreased metabolism of carbamazepine in men with Epilepsy as compared to patients with genotype CC. This association was only significant in male patients. Other genetic and clinical factors may also influence the metabolism of carbamazepine.

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cytochrome P450, family 3, subfamily A, polypeptide 5

-- Clinical Annotations

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and epilepsy may have decreased clearance and increased concentrations of carbamazepine, and require lower doses of the drug, as compared to patients with the CT (*1/*3) or TT (*1/*1) genotype. Other genetic and clinical factors may also influence dose or concentrations of carbamazepine.

· Class 3 rs15524 AA

Patients with the AA genotype and epilepsy may have increased concentrations of carbamazepine compared to patients with the AG and GG genotypes when patients were also taking phenytoin or phenobarbital. Other clinical and genetic factors may affect concentrations of carbamazepine.

ZSCAN25

zinc finger and SCAN domain containing 25

— Clinical Annotations	
- Chincal Alliforations	

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and epilepsy may have decreased clearance and increased concentrations of carbamazepine, and require lower doses of the drug, as compared to patients with the CT (*1/*3) or TT (*1/*1) genotype. Other genetic and clinical factors may also influence dose or concentrations of carbamazepine.

· Class 3 rs15524 AA

Patients with the AA genotype and epilepsy may have increased concentrations of carbamazepine compared to patients with the AG and GG genotypes when patients were also taking phenytoin or phenobarbital. Other clinical and genetic factors may affect concentrations of carbamazepine.

CYP3A

cytochrome P450, family 3, subfamily A

- Clinical Annotations
- Chincal Annotations

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and epilepsy may have decreased clearance and increased concentrations of carbamazepine, and require lower doses of the drug, as compared to patients with the CT (*1/*3) or TT (*1/*1) genotype. Other genetic and clinical factors may also influence dose or concentrations of carbamazepine.

· Class 3 rs2740574 TT

Patients with the TT genotype and epilepsy may have increased clearance of carbamazepine as compared to patients with the CC or CT genotype. Other genetic and clinical factors may also influence clearance of carbamazepine.

CYP3A4

cytochrome P450, family 3, subfamily A, polypeptide 4

carbamazepine as compared studies conflict. Other generations of the conflict	otype (CYP3A4 *1/*1) and epilepsy may have increased concentrations of d to patients with the CT (*1/*1G) or TT (*1G/*1G) genotype. However, etic and clinical factors may also influence concentrations of carbamazepine.
CYP1A2	cytochrome P450, family 1, subfamily A, polypeptide 2
	- Clinical Annotations —
as compared to pediatric pa	lepsy and the AA genotype may have increased clearance of carbamazepine atients with epilepsy and the AC or CC genotypes. Other clinical and genetic clearance of carbamazepine in pediatric patients with epilepsy.
UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
glucuronidation of carvedil responsible for the glucuron	G UGT1A1 *1/*1) genotype and angina or heart failure may have increased lol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is nidation of target substrates, rendering them water soluble and allowing for nation. Other genetic and clinical factors may also influence metabolism of
UGT1A3	UDP glucuronosyltransferase 1 family, polypeptide A3
glucuronidation of carvedil responsible for the glucuron	G UGT1A1 *1/*1) genotype and angina or heart failure may have increased lol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is nidation of target substrates, rendering them water soluble and allowing for nation. Other genetic and clinical factors may also influence metabolism of
UGT1A4	UDP glucuronosyltransferase 1 family, polypeptide A4

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

UGT1A5	UDP glucuronosyltransferase 1 family, polypeptide A5

· Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

· Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

· Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

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UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
	- Clinical Annotations —

· Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

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UDP glucuronosyltransferase 1 family, polypeptide A9

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· Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

\cdot Class 3 rs4148323 GG

Patients with the GG (i.e. UGT1A1 *1/*1) genotype and angina or heart failure may have increased glucuronidation of carvedilol as compared to patients with the AA (*6/*6) genotype. UGT1A1 is responsible for the glucuronidation of target substrates, rendering them water soluble and allowing for their biliary or renal elimination. Other genetic and clinical factors may also influence metabolism of carvedilol.

CATECHOLAMINES

ADRB1 adrenoceptor beta 1 — Clinical Annotations

· Class 3 rs1801253 GG

Patients with the GG genotype and coronary artery disease may require an increased dose of catecholamines as compared to patients with the CC or CG genotype. Other genetic and clinical factors may also influence required dose of catecholamines.

CELECOXIB

CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9
	- Clinical Annotations -

\cdot Class 2A rs1057910 AC

Patients with the AC (CYP2C9 *1/*3) genotype may have reduced metabolism of celecoxib as compared to patients with the AA (*1/*1) genotype, and increased metabolism as compared to patients with the CC (*3/*3) genotype. Other genetic and clinical factors may also influence metabolism of celecoxib.

CLOPIDOGREL

MED12L	mediator complex subunit 12-like
Class 3 rs2046934 A	
_	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also lopidogrel.
P2RY12	purinergic receptor P2Y, G-protein coupled, 12
Class 3 rs2046934 A	A
Class 3 rs2046934 A	
Class 3 rs2046934 A. Patients with the AA geno	type may have increased risk of adverse cardiac events when treated with
Class 3 rs2046934 A. Patients with the AA gene clopidogrel as compared to	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also
Class 3 rs2046934 A. Patients with the AA gene clopidogrel as compared to	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also
Class 3 rs2046934 A. Patients with the AA geno	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to clopidographic description.	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also lopidogrel.
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to clopidographic description.	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to clopidographic description.	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also lopidogrel.
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to clopidographic description.	otype may have increased risk of adverse cardiac events when treated with patients with genotype GG. Other genetic and clinical factors may also lopidogrel. catechol-O-methyltransferase
Class 3 rs2046934 A. Patients with the AA gend clopidogrel as compared to influence the response to clopidographic communication. COMT Class 3 rs4680 GA	catechol-O-methyltransferase ———————————————————————————————————
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to class 3 rs4680 GA Patients with the AG gence clopidogrel as compared to influence the response to class 3 rs4680 GA	catechol-O-methyltransferase — Clinical Annotations cotype and schizophrenia may have a poorer response when treated with
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to class 3 rs4680 GA Patients with the AG gence clopidogrel as compared to influence the response to class 3 rs4680 GA Patients with the AG gence clopidogrel as compared to clopidogrel as compa	catechol-O-methyltransferase ———————————————————————————————————
Class 3 rs2046934 A. Patients with the AA gence clopidogrel as compared to influence the response to class 3 rs4680 GA Patients with the AG gence clozapine as compared to present the compared to pre	catechol-O-methyltransferase ———————————————————————————————————

AA genotype. Other genetic and clinical factors may also influence concentrations and risk of clozapine-

- Clinical Annotations -

People with GG genotype may have decreased, but not absent, risk of major adverse cardiovascular

Class 3 rs1045642 GG

induced toxicity.

CODEINE

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
risk for CNS depression	GG chose mothers have the GG genotype and are taking codeine may be at decreased as compared to those whose mothers have the AA genotype. Other genetic also influence the risk of CNS depression in breast-feeding infants.
COTININE	
CHRNA3	cholinergic receptor, nicotinic, alpha 3 (neuronal)
	- Clinical Annotations
cotinine, a metabolite o	co Use Disorder and the GG genotype may have decreased concentrations of of nicotine, as compared to individuals with the AG or AA genotype. Other ors may also contribute to cotinine concentrations in individuals with Tobacco cholinergic receptor, nicotinic, alpha 5 (neuronal)
	- Clinical Annotations
cotinine, a metabolite o	co Use Disorder and the GG genotype may have decreased concentrations of of nicotine, as compared to individuals with the AG or AA genotype. Other ors may also contribute to cotinine concentrations in individuals with Tobacco
CYP2A6	cytochrome P450, family 2, subfamily A, polypeptide 6
	- Clinical Annotations
_	AT enotype may have increased 7-hydroxylation of coumarin compared to patients Other genetic and clinical factors may also influence metabolism of coumarin.

CYCLOPHOSPHAMIDE

phamide may have a shorter pe	the TT genotype and breast cancer who are treated with cyclophoseriod of time before chemotherapy-induced ovarian failure compared to enotype. Other genetic and clinical factors may also influence time to
chemotherapy-induced ovarian	failure.
CYP3A	cytochrome P450, family 3, subfamily A
Class 3 rs2740574 TT	
Premenopausal patients with a phamide may have a shorter po	the TT genotype and breast cancer who are treated with cyclophoseriod of time before chemotherapy-induced ovarian failure compared to enotype. Other genetic and clinical factors may also influence time to failure.
VEGFA	vascular endothelial growth factor A
and clinical factors may also in Class 3 $\mathbf{rs1570360}$ AG Patients with the AG genotyp when treated with docetaxel place.	namide as compared to patients with the CC genotype. Other genetic fluence length of progression-free survival. e and prostate cancer may have longer progression-free survival time as oral metronomic cyclophosphamide as compared to patients with the ad clinical factors may also influence progression-free survival time.
CYP1B1	cytochrome P450, family 1, subfamily B, polypeptide 1
cyclophosphamide, epirubicin a genetic and clinical factors may	be and breast cancer may have a better response when treated with and fluorouracil as compared to patients with the CC genotype. Other also influence response to treatment with cyclophosphamide, epirubicin C/G variant, particularly in a gene on the minus chromosomal strand,
CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
Class 3 rs4802101 TT	

 \cdot Class 3 rs2740574 TT

Patients with the TT genotype may have increased metabolism of cyclophosphamide, resulting in increased concentrations of active cyclophosphamide metabolites, and increased risk of gastrointestinal toxicity, or leukopenia, as compared to patients with the CT or CC genotypes. Other clinical and genetic factors may also influence metabolism of cyclophosphamide, as well as risk of toxicity in patients with lupus.

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ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1

· Class 3 rs1128503 GG

Patients with the GG genotype and myasthenia gravis or organ transplantation may have increased clearance of cyclosporine and therefore may require an increased dose of cyclosporine, compared to patients with the AA genotype. Patients with the GG genotype may also have a decreased risk of infection as compared to those with the AA or AG genotype. Other genetic and clinical factors may also influence clearance and dose of cyclosporine.

· Class 3 rs2032582 CC

Patients with the CC genotype may have lower blood trough concentrations of cyclosporine compared to patients with the AA genotype, and may require dose adjustments. Other genetic and clinical factors may also influence cyclosporine blood concentrations.

· Class 3 rs1045642 GG

Patients with genotype GG may have decreased intracellular and blood concentrations of cyclosporine in people with Transplantation as compared to patients with genotype AA or AG. However, contradictory findings have been reported. Other genetic and clinical factors may also influence the concentration of cyclosporine.

· Class 3 rs2032582 CC

Patients with the CC genotype and cystic fibrosis may have increased clearance of dicloxacillin, when it is coadministered with cyclosporine, as compared to patients with the AA genotype. Other genetic and clinical factors may also influence clearance of dicloxacillin.

and clinical factors may also	influence clearance of dicloxacillin.
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
	- Clinical Annotations
· Class 2B rs776746 CC	one (CVP3A5 *3/*3) may require a lower dose of cyclosporine to reach

Patients with the CC genotype (CYP3A5 *3/*3) may require a lower dose of cyclosporine to reach target blood concentration as compared to patients with the CT (CYP3A5 *1/*3) or TT (CYP3A5 *1/*1) genotype, although this is contradicted in some studies. Other genetic and clinical factors may also influence dose of cyclosporine.

· Class 2B rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) may require a lower dose of cyclosporine to reach target blood concentration as compared to patients with the CT (CYP3A5 *1/*3) or TT (CYP3A5 *1/*1) genotype, although this is contradicted in some studies. Other genetic and clinical factors may also influence dose of cyclosporine.

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cytochrome P450, family 3, subfamily A

-- Clinical Annotations -

· Class 2B rs776746 *CC*

Patients with the CC genotype (CYP3A5 *3/*3) may require a lower dose of cyclosporine to reach target blood concentration as compared to patients with the CT (CYP3A5 *1/*3) or TT (CYP3A5 *1/*1) genotype, although this is contradicted in some studies. Other genetic and clinical factors may also influence dose of cyclosporine.

Class 3 rs35599367 *GG*

Patients with the GG genotype and organ transplantation administered cyclosporine may have a 1) increased metabolism of cyclosporine 2) increased clearance of cyclosporine and 3) a decreased risk in adverse events (e.g. graft rejection or kidney function) as compared to patients with the AA genotype. Other clinical and genetic factors may also affect metabolism and incidence of adverse events in organ transplant patients administered cyclosporine.

CYP3A4

cytochrome P450, family 3, subfamily A, polypeptide 4

—- Clinical Annotations —

· Class 3 rs28371759 AA

Patients with the AA genotype (CYP3A4 *1/*1) who underwent kidney transplantation may have decreased metabolism of cyclosporine as compared to patients with the GG genotype (*18B/*18B). Other genetic and clinical factors may also influence metabolism of cyclosporine.

· Class 3 rs35599367 GG

Patients with the GG genotype and organ transplantation administered cyclosporine may have a 1) increased metabolism of cyclosporine 2) increased clearance of cyclosporine and 3) a decreased risk in adverse events (e.g. graft rejection or kidney function) as compared to patients with the AA genotype. Other clinical and genetic factors may also affect metabolism and incidence of adverse events in organ transplant patients administered cyclosporine.

CYTARABINE

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	—- Clinical Annotations —

· Class 3 rs1045642 GG

Patients with the GG genotype may have 1) decreased exposure to doxorubicin metabolites and 2) decreased response to anthracycline regimens as compared to patients with the AA genotype, however the evidence is highly contradictory. Other genetic and clinical factors may also influence response to anthracycline regimens.

· Class 3 rs1128503 GG

Patients with the GG genotype and acute myeloid leukemia may have a poorer response when treated with cytarabine, alone or in combination with daunorubicin, or dexrazoxane as compared to patients with the AA or AG genotype, however some evidence contradicts this. Other genetic and clinical factors may also influence response to cytarabine.

SLCO1B1	solute carrier organic anion transporter family, member 1B1
	Clinical Annotations —

· Class 3 rs2291075 CT

Patients with the CT genotype may have more favorable event-free and overall survival in children with de novo acute myeloid leukemia (AML) treated with cytarabine, daunorubicin, etoposide and mitoxantrone as compared to patients with genotype CC. Other genetic and clinical factors may also influence the treatment outcome in acute myeloid leukemia.

DAUNORUBICIN

SLCO1B1	solute carrier organic anion transporter family, member 1B1

· Class 3 rs2291075 CT

Patients with the CT genotype may have more favorable event-free and overall survival in children with de novo acute myeloid leukemia (AML) treated with cytarabine, daunorubicin, etoposide and mitoxantrone as compared to patients with genotype CC. Other genetic and clinical factors may also influence the treatment outcome in acute myeloid leukemia.

DEFERASIROX

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10

· Class 3 rs887829 CT

Patients with the CT genotype and beta-thalassemia may have decreased concentrations of deferasirox as compared to patients with the TT genotype. Other genetic and clinical factors may also influence concentrations of deferasirox.

· Class 3 rs3806596 TC

Patients with the CT genotype and beta-thalassemia may have worse response to, and decreased concentrations of deferasirox as compared to patients with the CC genotype. Other clinical and genetic factors may also influence concentrations of and response to deferasirox in patients with beta-thalassemia.

· Class 4 rs4124874 TG

Pediatric patients with major thalassemia and the GT genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the TT genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype GT, therefore there is no information for patients with the GG or TT genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

Class 4 rs10929302 GA

Pediatric patients with major thalassemia and the AG genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the GG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the AA or GG genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

\cdot Class 4 rs4148323 GG

Pediatric patients with major thalassemia and the GG genotype may have a decreased risk of adverse reactions when administered deferasirox as compared to patients with the AA or AG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the GG or AA genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

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UDP glucuronosyltransferase 1 family, polypeptide A3

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· Class 3 rs887829 CT

Patients with the CT genotype and beta-thalassemia may have decreased concentrations of deferasirox as compared to patients with the TT genotype. Other genetic and clinical factors may also influence concentrations of deferasirox.

\cdot Class 3 rs3806596 TC

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UGT1A5

UDP glucuronosyltransferase 1 family, polypeptide A5

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· Class 3 rs887829 CT

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Pediatric patients with major thalassemia and the GT genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the TT genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype GT, therefore there is no information for patients with the

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UGT1A6	UDP glucuronosyltransferase 1 family, polypeptic	de A6
	- Clinical Annotations	

· Class 3 rs887829 CT

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· Class 3 rs3806596 TC

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Class 3 rs887829 CT

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· Class 3 rs887829 CT

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\cdot Class 3 rs3806596 TC

Patients with the CT genotype and beta-thalassemia may have worse response to, and decreased concentrations of deferasirox as compared to patients with the CC genotype. Other clinical and genetic factors may also influence concentrations of and response to deferasirox in patients with beta-thalassemia.

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patient with major thalassemia of genotype GT, therefore there is no information for patients with the GG or TT genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

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· Class 4 rs4148323 GG

Pediatric patients with major thalassemia and the GG genotype may have a decreased risk of adverse reactions when administered deferasirox as compared to patients with the AA or AG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the GG or AA genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

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UDP glucuronosyltransferase 1 family, polypeptide A9

Clinical Annotations	
Chincal Allifotations	

· Class 3 rs887829 CT

Patients with the CT genotype and beta-thalassemia may have decreased concentrations of deferasirox as compared to patients with the TT genotype. Other genetic and clinical factors may also influence concentrations of deferasirox.

· Class 3 rs3806596 TC

Patients with the CT genotype and beta-thalassemia may have worse response to, and decreased concentrations of deferasirox as compared to patients with the CC genotype. Other clinical and genetic factors may also influence concentrations of and response to deferasirox in patients with beta-thalassemia.

· Class 4 rs4124874 TG

Pediatric patients with major thalassemia and the GT genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the TT genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype GT, therefore there is no information for patients with the GG or TT genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

· Class 4 rs10929302 GA

Pediatric patients with major thalassemia and the AG genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the GG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the AA or GG genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

· Class 4 rs4148323 GG

Pediatric patients with major thalassemia and the GG genotype may have a decreased risk of adverse reactions when administered deferasirox as compared to patients with the AA or AG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the

GG or AA genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

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UDP glucuronosyltransferase 1 family, polypeptide A1

-- Clinical Annotations ----

· Class 3 rs887829 CT

Patients with the CT genotype and beta-thalassemia may have decreased concentrations of deferasirox as compared to patients with the TT genotype. Other genetic and clinical factors may also influence concentrations of deferasirox.

\cdot Class 4 rs4124874 TG

Pediatric patients with major thalassemia and the GT genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the TT genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype GT, therefore there is no information for patients with the GG or TT genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

· Class 4 rs10929302 GA

Pediatric patients with major thalassemia and the AG genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the GG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the AA or GG genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

· Class 4 rs4148323 GG

Pediatric patients with major thalassemia and the GG genotype may have a decreased risk of adverse reactions when administered deferasirox as compared to patients with the AA or AG genotype. Please note, the evidence comes solely from a single case study report of a single individual, a 3 year old female patient with major thalassemia of genotype AG, therefore there is no information for patients with the GG or AA genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2
	—- Clinical Annotations ————

\cdot Class 3 rs2273697 GG

Patients with the GG genotype and beta-thalassemia may have decreased concentrations of deferasirox as compared to patients with the AG genotype. Other genetic and clinical factors may also influence concentrations of deferasirox.

· Class 4 rs717620 CT

Pediatric patients with major thalassemia and the CT genotype may have an increased risk of adverse reactions when administered deferasirox as compared to patients with the CC genotype. Please note, the evidence comes solely from a single case study report of a 3 year old female patient with major thalassemia of genotype CT, therefore there is no information for patients with the CC or TT genotypes. Other clinical and genetic factors may also influence risk of adverse reactions in patients with major thalassemia who are administered deferasirox.

	- Clinical Annotations —
 as compared to patients with the AC influence concentrations of deferasirox. Class 3 rs2470890 TT Patients with the TT genotype and beta possibly to levels below therapeutic eff 	a-thalassemia may have decreased concentrations of deferasirox or CC genotype. Other genetic and clinical factors may also a-thalassemia may have decreased concentrations of deferasirox, ficacy, as compared to patients with the CC or CT genotype. also influence concentrations of deferasirox.
DIURETICS	
ADRB2	adrenoceptor beta 2, surface
	- Clinical Annotations ————————————————————————————————————
hospital utilization when treated with	art failure may have increased emergency department visits and cardiovascular drugs as compared to patients with the AA or al factors may also influence efficacy of cardiovascular drugs. adrenoceptor beta 1
	- Clinical Annotations ————————————————————————————————————
	rt failure may have increased emergency department utilization as compared to patients with the GG genotype. Other genetic efficacy of cardiovascular drugs.
DOBUTAMINE	
ADRB1	adrenoceptor beta 1
	- Clinical Annotations ————————————————————————————————————

· Class 3 rs1801253 GG

Healthy males with the GG genotype may have smaller increases in fractional shortening and systolic blood pressure when given dobutamine, as compared to healthy males with the CC genotype. No significant differences were seen for heart rate. Other genetic and clinical factors may also influence fractional shortening and systolic blood pressure.

DOCETAXEL

Class 3 rs2740574 TT Patients with the TT genotype m	hay have decreased clearance of docetaxel and a decreased risk of an
infusion-related reaction as comp experience a decreased risk of ne	ared to patients with the CC or CT genotype. These patients may protoxicity with docetaxel treatment, though reports conflict. Other lso influence clearance of and reactions to docetaxel.
CYP3A	cytochrome P450, family 3, subfamily A
infusion-related reaction as comp experience a decreased risk of neu	hay have decreased clearance of docetaxel and a decreased risk of an ared to patients with the CC or CT genotype. These patients may protoxicity with docetaxel treatment, though reports conflict. Other lso influence clearance of and reactions to docetaxel.
CYP4B1	cytochrome P450, family 4, subfamily B, polypeptide 1
9 11	ay have a decreased but not absent risk of toxicity with docetaxel and nts with the CT or TT genotypes. Other genetic and clinical factors conse.
NAT2	$\hbox{N-acetyltransferase 2 (arylamine N-acetyltransferase)}$
	- Clinical Annotations
	hay have an increased risk of toxicity with docetaxel and thalidomide AA genotype. Other genetic and clinical factors may also influence
VEGFA	vascular endothelial growth factor A
Class 3 rs1570360 AG	
compared to patients with the GO	d breast cancer may have a better response to docetaxel treatment as G genotype. However, contradictory evidence exists when considering enetic and clinical factors may also influence response to docetaxel.

Current literature evidence finds no significant effect of the AC genotype on progression-free survival

time in patients taking docetaxel.

· Class 3 rs1570360 AG

—- Clinical Annotations —

Patients with the AG genotype and prostate cancer may have longer progression-free survival time when treated with docetaxel plus oral metronomic cyclophosphamide as compared to patients with the AA genotype. Other genetic and clinical factors may also influence progression-free survival time.

HNF4A her	patocyte nuclear factor 4, alpha

\cdot Class 3 rs2273618 TC

Patients with nasopharyngeal cancer and the CT genotype may have more severe anemia when treated with docetaxel as compared to patients with the TT genotype. Other clinical and genetic factors may also influence severity of anemia in patients with nasopharyngeal cancer who are treated with docetaxel.

· Class 3 rs3746574 TC

Patients with nasopharyngeal cancer and the CT genotype may have more severe anemia when treated with docetaxel as compared to patients with the TT genotype. Other clinical and genetic factors may also influence severity of anemia in patients with nasopharyngeal cancer who are treated with docetaxel.

DOXORUBICIN

ABCB1 ATP-binding cassette, sub-family B (MDR/TAP), member 1 - Clinical Annotations

· Class 3 rs1045642 GG

Patients with the GG genotype may have 1) decreased exposure to doxorubicin metabolites and 2) decreased response to anthracycline regimens as compared to patients with the AA genotype, however the evidence is highly contradictory. Other genetic and clinical factors may also influence response to anthracycline regimens.

· Class 3 rs2032582 CC

Patients with the CC genotype may have increased metabolism of doxorubicin in people with Breast Neoplasms as compared to patients with genotype AA. Other genetic and clinical factors may also influence the metabolism of doxorubicin.

EFAVIRENZ

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$

· Class 3 rs1045642 GG

Patients with the GG genotype and HIV infection who are treated with efavirenz may have reduced clearance of efavirenz as compared to patients with the AG genotype. Some studies have shown no association between this polymorphism and efavirenz clearance, plasma concentrations or exposure, or PBMC concentrations. Other genetic and clinical factors may also influence efavirenz pharmacokinetics.

Class 3 rs2032582 *CC*

Patients with the CC genotype may have increased likelihood of emerging viral drug resistance when exposed to efavirenz in people with HIV Infections as compared to patients with the AA genotype. This varaint is not associated with plasma exposure of efavirenz. Other genetic and clinical factors may also influence the response to efavirenz

\cdot Class 4 rs1128503 GG

Patients with GG genotype and HIV may have increased concentrations of efavirenz in plasma compared to patients with AA genotype. However, this association was not significant and was not found in another study of plasma and PBMCs. Other clinical and genetic factors may affect efavirenz concentrations.

CYP2A6	cytochrome P450, family 2, subfamily A, polypeptide 6

· Class 3 rs28399433 AA

Patients with the AA genotype and HIV may have decreased plasma concentrations of efavirenz as compared to patients with the AC or CC genotype. Other genetic and clinical factors may also influence plasma concentrations of efavirenz.

CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
	- Clinical Annotations

· Class 2B rs4803419 CC

Patients with HIV and the CC genotype may have lower plasma concentrations of efavirenz as compared to patients with the TT genotype. Other clinical and genetic factors may also influence plasma concentrations of efavirenz in patients with HIV.

· Class 3 rs8192709 CC

Patients with genotype CC may have decreased metabolism of efavirenz in people with HIV Infections as compared to patients with genotype CT. Other genetic and clinical factors may also influence the metabolism of efavirenz.

· Class 3 rs8192719 CC

Patients with the CC genotype and HIV may have decreased concentrations of efavirenz as compared to patients with the CT or TT genotype. Other genetic and clinical factors, such as rs3745274, may also influence concentrations of efavirenz.

\cdot Class 3 rs1042714 GG

Patients with the GG genotype and left ventricular hypertrophy may have a greater percent reduction in left ventricular mass index when treated with enalapril as compared to patients with the CC genotype. Other genetic and clinical factors may also influence reduction in left ventricular mass index.

	——————————————————————————————————————
Class 3 rs699947 AC Patients with hypertension and the AC genotype may have an improved response to enalapril as compared to patients with the CC genotype. Other clinical and genetic factors may also influence respons to enalapril in patients with hypertension.	
ERLOTINIB	
CYP1A2	cytochrome P450, family 1, subfamily A, polypeptide 2
_	$\cal A$ stype may have increased concentrations of erlotinib as compared to patients her genetic and clinical factors may also influence concentrations of erlotinib.
ESCITALOPRAM	
CYP2D6	cytochrome P450, family 2, subfamily D, polypeptide 6
Patients with the GG genotreated with escitalopram factors may also effect pat	TG by type and depression may have a increased response and remission rate when as compared to patients with the AA genotype. Other genetic and clinical ients response.
ETHAMBUTOL	
NAT2	N-acetyltransferase 2 (arylamine N-acetyltransferase)
_	TT otype and tuberculosis (TB) may have an increased risk for hepatotoxicity drugs as compared to patients with the CC genotype. Other genetic and

clinical factors may also influence risk for hepatotoxicity.

Class 2A rs1799930 AA

Patients with the AA genotype and tuberculosis (TB) may have an increased risk of hepatotoxicity when treated with anti-TB drugs as compared to patients with the GG genotype. They also may have decreased clearance of isoniazid as compared to those with the AG or GG genotype. Other genetic and clinical factors may also influence risk for hepatotoxicity and clearance of isoniazid.

Class 3 rs1799931 *GG*

Patients with the GG genotype and tuberculosis (TB) may have a decreased risk of hepatotoxicity when treated with anti-TB drugs as compared to patients with the AA or AG genotype. However, some studies find no association with hepatotoxicity. Other genetic and clinical factors may also influence risk of hepatotoxicity.

CHRNA3	cholinergic receptor, nicotinic, alpha 3 (neuronal)
_	GG otype may have an increased risk for alcoholism as compared to patients with genetic and clinical factors may also influence risk of alcoholism.
CHRNA5	cholinergic receptor, nicotinic, alpha 5 (neuronal)
	- Clinical Annotations
9	GG otype may have an increased risk for alcoholism as compared to patients with genetic and clinical factors may also influence risk of alcoholism.
SLCO1B1	solute carrier organic anion transporter family, member 1B1
	- Clinical Annotations —
Patients with the CT gen with de novo acute myel- mitoxantrone as compared influence the treatment of	notype may have more favorable event-free and overall survival in children oid leukemia (AML) treated with cytarabine, daunorubicin, etoposide and d to patients with genotype CC. Other genetic and clinical factors may also atcome in acute myeloid leukemia.
FEXOFENADINE	
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1

ciation with fexofenadine plasma concentrations. Other genetic and clinical factors may also influence

plasma concentrations of fexofenadine and dose requirements.

FLUOROURACIL

	ave decreased risk of diarrhea when treated with fluorouracil in people pared to patients with genotype AA. Other genetic and clinical factors are to fluorouracil.
ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2 $$
Class 3 rs717620 CT	
treated with FOLFOX (fluoroura	d colon cancer may have a decreased risk of thrombocytopenia when acil, leucovorin, oxaliplatin) as compared to patients with the CC cal factors may also influence risk of thrombocytopenia.
syndromes when treated with FO	and colorectal cancer may have decreased severity of neurotoxicity LFOX (fluorouracil, leucovorin, oxaliplatin) as compared to patients netic and clinical factors may also influence severity of neurotoxicity
CYP1B1	cytochrome P450, family 1, subfamily B, polypeptide 1
	- Clinical Annotations
cyclophosphamide, epirubicin and genetic and clinical factors may al	and breast cancer may have a better response when treated with I fluorouracil as compared to patients with the CC genotype. Other so influence response to treatment with cyclophosphamide, epirubicin /G variant, particularly in a gene on the minus chromosomal strand,
DPYD	dihydropyrimidine dehydrogenase
	——————————————————————————————————————
DPYD:*1/*1 Moderate	
Use label-recommended dosage ar	
	(DPYD *1/*1) and cancer who are treated with fluoropyrimidine- decreased, but not absent, risk for drug toxicity as compared to

patients with the AC or CC genotype (DPYD *1/*13 or *13/*13). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to

fluoropyrimidine-based chemotherapy.

- Clinical Annotations -

Class 1A rs3918290 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of fluoropyrimidine drugs and 2) decreased, but not non-existent, risk for drug toxicity as compared to patients with the CT or TT genotype (DPYD *1/*2A or *2A/*2A). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine based chemotherapy.

\cdot Class 1A rs67376798 TT

Patients with the TT genotype and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of the drug and 2) decreased, but not absent, risk and reduced severity of drug toxicity as compared to patients with the AT genotype. Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 3 rs1801160 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased metabolism of fluorouracil and 2) decreased risk for drug toxicities as compared to patients with the CT or TT genotype (DPYD *1/*6 or *6/*6). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin) or with other drugs such as bevacizumab, cetuximab, raltitrexed. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

Class 3 rs1801159 TT

Patients with the TT genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased likelihood of nausea, vomiting, and leukopenia, 2) increased response and 3) increased clearance of fluorouracil as compared to patients with the CT or CC genotype (DPYD *1/*5 or *5/*5). However, other studies find no associations or contradictory associations with fluoropyrimidine-induced drug toxicity or response. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 3 rs1801265 GA

Patients with the AG genotype (DPYD *1/*9A) and cancer who are treated with fluorouracil may have 1) an increased risk for drug toxicities, 2) decreased response and 3) increased DPYD activity as compared to patients with the AA genotype (DPYD *1/*1). Patients with the AG genotype were also found to have increased clearance of fluorouracil as compared to those with the GG genotype (DPYD *9A/*9A). However, multiple studies find contradictory or negative evidence for drug toxicities, fluorouracil clearance, and DPYD activity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

Class 3 rs1801158 CC

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased but not absent risk of toxicity and 2) increased DPYD activity as compared to patients with the CT genotype (DPYD *1/*4). However, some studies find no association with drug toxicity or DPYD activity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

\cdot Class 4 rs1801266 GG

Patients with the GG genotype (DPYD *1/*1) may have increased DPYD activity as compared to those with the AA or AG genotype (DPYD *8/*8 or *1/*8). Other genetic and clinical factors may also affect DPYD activity.

Class 4	rs1801268	CC

Patients with the CC genotype (DPYD *1/*1) may have increased DPYD activity as compared to those with the AC or AA genotype (DPYD *1/*10 or *10/*10). Other genetic and clinical factors may also affect DPYD activity.

· Class 4 rs72549306 CC

Patients with the CC genotype (DPYD $^*1/^*1$) may have increased DPYD activity as compared to those with the AA or AC genotype (DPYD $^*11/^*11$ or $^*1/^*11$). Other genetic and clinical factors may also affect DPYD activity.

VEGFA	vascular endothelial growth factor A
· Class 3 rs699947 AC	
Patients with colorectal cancerage capecitabine, fluorouracil, iri	er and the AC genotype may have a reduced response to bevacizumab notecan, leucovorin, or oxaliplatin as compared to patients with the CC genetic factors may also affect response to chemotherapy in people with
IGFBP3	insulin-like growth factor binding protein 3
	——————————————————————————————————————
9 1	pe and stomach cancer may have a poorer survival outcomes when treated d to patients with the GT or TT genotype. Other genetic and clinical revival outcome.
FLUVASTATIN	
SLCO1B1	solute carrier organic anion transporter family, member 1B1
	- Clinical Annotations -
· Class 3 rs11045819 CC Patients with the CC genotype as compared to patients with	pe who are treated with fluvastatin may have a lesser reduction in LDL-C

· Class 3 rs4986910 AA

Patients with the AA genotype may have a smaller increase in HDL cholesterol when treated with fluvastatin as compared to patients with the AG genotype. Other genetic and clinical factors may also influence HDL cholesterol response.

- Clinical Annotations -

CYP3A4

cytochrome P450, family 3, subfamily A, polypeptide 4

· Class 3 rs4986910 AA Patients with the AA genotype may have a smaller increase in HDL cholesterol when treated wit fluvastatin as compared to patients with the AG genotype. Other genetic and clinical factors may als influence HDL cholesterol response.	
GEFITINIB	
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	- Clinical Annotations —
and skin rash when treate	GG notype and non-small cell lung cancer may have a decreased risk for diarrhead with gefitinib as compared to patients with the AA genotype. Other geneticalso influence drug toxicity risk in patients receiving gefitinib.
GEMCHABINE	
CDA	cytidine deaminase
to patients with the TT g	the CT genotype may have increased metabolism of gemcitabine as compared enotype. However, this has been contradicted by some studies. Other genetic also influence metabolism of gemcitabine.
HALOPERIDOL	
COMT	catechol-O-methyltransferase
· Class 3 rs4680 GA Patients with the AG gen	otype and schizophrenia may have an increased risk for developing extrapyra-

midal symptoms when treated with haloperidol as compared to patients with the AA or GG genotype. Other genetic and clinical factors may also influence risk for extrapyramidal symptoms when taking haloperidol.

HMG COA REDUCTASE INHIBITORS

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\cdot Class 3 rs1045642 GG

Patients with the GG genotype may have decreased serum creatine kinase levels when treated with hmg CoA reductase inhibitors as compared to patients with the AA genotype. Other genetic and clinical factors may also influence serum creatine kinase levels.

· Class 3 rs1128503 GG

Patients with the GG genotype may have decreased serum creatine kinase levels when treated with hmg CoA reductase inhibitors as compared to patients with the AA genotypes. Other genetic and clinical factors may also influence serum creatine kinase levels.

ILOPERIDONE

· Class 3 rs1065852 GG

Patients with the GG genotype and schizophrenia may have an increased QTc interval when treated with iloperidone as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence QTc interval.

IMATINIB

· Class 3 rs1045642 GG

Patients with the GG genotype and chronic myeloid leukemia may have an increased likelihood of achieving complete molecular response when treated with imatinib, as compared to patients with the AA or AG genotype. However, this was only significant in an exclusively Caucasian population. Additionally, no significant results were seen when considering major molecular response. Other genetic and clinical factors may also influence likelihood of achieving complete molecular response.

· Class 3 rs1128503 GG

Patients with the GG genotype and chronic myeloid leukemia may have a better response to imatinib treatment as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence response to imatinib.

CYP2B6 cytochrome P450, family 2, subfamily B, polypeptide 6 — Clinical Annotations

· Class 3 rs3745274 GG

Patients with the GG genotype and chronic myeloid leukemia may have a 1) a better response to treatment with imatinib as compared to patients with the TT genotype, 2) an increased risk of developing cytogenetic resistance to imatinib as compared to patients with the GT genotype, and 3) a greater risk

CYP2A7P1	cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene 1
Class 3 rs3745274	
ment with imatinib as cytogenetic resistance t for side effects as comp	genotype and chronic myeloid leukemia may have a 1) a better response to treat- compared to patients with the TT genotype, 2) an increased risk of developing to imatinib as compared to patients with the GT genotype, and 3) a greater risk ared to patients with the GT or TT genotype. Other genetic and clinical factors conse, resistance and risk of side effects in patients taking imatinib.
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
	- Clinical Annotations
Class 3 rs776746	CC
Patients with the CC g	enotype and chronic myeloid leukemia have have increased trough concentrations to patients with the CT and TT genotypes. Other genetic and clinical factors
ZSCAN25	zinc finger and SCAN domain containing 25
Class 3 rs776746	CC
Patients with the CC g	enotype and chronic myeloid leukemia have have increased trough concentrations to patients with the CT and TT genotypes. Other genetic and clinical factors
CYP3A	cytochrome P450, family 3, subfamily A
	- Clinical Annotations
Class 3 rs776746	CC
Patients with the CC g	enotype and chronic myeloid leukemia have have increased trough concentrations to patients with the CT and TT genotypes. Other genetic and clinical factors
BESARTAN	
CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9

Patients with the AC genotype and essential hypertension may have decreased metabolism or clearance of irbesartan as compared to patients with the AA genotype, but may have no difference in response. Other clinical or genetic factors may also influence concentrations of irbesartan in patients with essential hypertension.

· Class 3 rs72558187 TT

Individuals with the TT genotype may have increased metabolism and clearance of irbesartan which may result may in decreased exposure of irbesartan as compared to patients with the CT genotype. Other clinical and genetic factors may also influence metabolism of irbesartan.

IRINOTECAN

SLCO1B1	solute carrier organic anion transporter family, member 1B1
	- Clinical Annotations —

· Class 3 rs4149056 TT

Patients with the TT genotype and cancer may have a decreased risk of neutropenia when treated with irinotecan or irinotecan-based regimens, as compared to patients with the CC or CT genotype. However, a different study of similar size found no association between the TT genotype and neutropenia. No significant results have been seen for diarrhea. Other genetic and clinical factors may also influence risk of neutropenia or diarrhea.

· Class 3 rs4149015 GG

Patients with the GG genotype and non-small cell lung cancer may have a decreased risk of neutropenia when treated with irinotecan as compared to patients with the AG or GG genotype. No association has been seen for diarrhea. Other genetic and clinical factors may also influence risk of neutropenia.

· Class 3 rs2306283 AG

Patients with the AG genotype and solid tumors may experience increased risk of neutropenia compared to patients with the AA genotype. However, studies conflict as to this association. Other clinical and genetic factors may affect risk of neutropenia with irinotecan therapy.

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10

· Class 3 rs3832043 T/del

Patients with the T/del genotype and non-small cell lung cancer may have a decreased risk of diarrhea when treated with irinotecan or irinotecan-based regimens as compared to patients with the del/del genotype. However, a different study of similar size found no association between this genotype and diarrhea. No significant results have been seen when considering neutropenia or tumor response. Other genetic and clinical factors may also influence risk of diarrhea.

· Class 3 rs11692021 TC

Patients with the CT genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the TT genotype, or a decreased risk as compared to patients with the CC genotype. This may be due to decreased enzymatic activity toward SN-38, the active metabolite of irinotecan, found in cells with the C allele as compared to those with the T allele. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

 \cdot Class 3 rs2070959 AG

Patients with the AG genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the GG genotype, or a decreased risk as compared to patients with the AA genotype. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

· Class 3 rs10929302 GA

The AG genotype may be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

UGT1A8 UDP glucuronosyltransferase 1 family, polypeptide A8

Clinical Annotations -

· Class 3 rs3832043 T/del

Patients with the T/del genotype and non-small cell lung cancer may have a decreased risk of diarrhea when treated with irinotecan or irinotecan-based regimens as compared to patients with the del/del genotype. However, a different study of similar size found no association between this genotype and diarrhea. No significant results have been seen when considering neutropenia or tumor response. Other genetic and clinical factors may also influence risk of diarrhea.

· Class 3 rs11692021 TC

Patients with the CT genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the TT genotype, or a decreased risk as compared to patients with the CC genotype. This may be due to decreased enzymatic activity toward SN-38, the active metabolite of irinotecan, found in cells with the C allele as compared to those with the T allele. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

\cdot Class 3 rs2070959 AG

Patients with the AG genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the GG genotype, or a decreased risk as compared to patients with the AA genotype. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

· Class 3 rs10929302 GA

The AG genotype may be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

Class 3 rs3832043 T/del

Patients with the T/del genotype and non-small cell lung cancer may have a decreased risk of diarrhea when treated with irinotecan or irinotecan-based regimens as compared to patients with the del/del genotype. However, a different study of similar size found no association between this genotype and diarrhea. No significant results have been seen when considering neutropenia or tumor response. Other genetic and clinical factors may also influence risk of diarrhea.

· Class 3 rs11692021 TC

Patients with the CT genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the TT genotype, or a decreased risk as compared to patients with the CC genotype. This may be due to decreased enzymatic activity toward

SN-38, the active metabolite of irinotecan, found in cells with the C allele as compared to those with the T allele. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

· Class 3 rs2070959 AG

Patients with the AG genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the GG genotype, or a decreased risk as compared to patients with the AA genotype. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

· Class 3 rs10929302 GA

The AG genotype may be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2

\cdot Class 3 rs2273697 GG

Patients with the GG genotype and colorectal cancer may have decreased metabolism of irinotecan as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence metabolism of irinotecan.

· Class 3 rs3740066 CT

Patients with the CT genotype and non-small cell lung cancer may have a decreased risk of diarrhea when treated with irinotecan as compared to patients with the CC genotype. No association has been seen for neutropenia. Other genetic and clinical factors may also influence risk of diarrhea.

UGT1A6	UDP glucuronosyltransferase 1 family, polypeptide A6
	- Chinical Affiliations

· Class 3 rs11692021 TC

Patients with the CT genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the TT genotype, or a decreased risk as compared to patients with the CC genotype. This may be due to decreased enzymatic activity toward SN-38, the active metabolite of irinotecan, found in cells with the C allele as compared to those with the T allele. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

\cdot Class 3 rs2070959 AG

Patients with the AG genotype and colorectal cancer may have an increased risk for severe neutropenia when treated with irinotecan as compared to patients with the GG genotype, or a decreased risk as compared to patients with the AA genotype. Other genetic and clinical factors, such as UGT1A1*28, may also influence the risk for neutropenia in patients taking irinotecan.

· Class 3 rs10929302 GA

The AG genotype may be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

when treated with irinotecan as compared to patients with the CG SN-38, the active metabolite of it the T allele. Other genetic and concurred in patients taking irinotecan as rs2070959 AG Patients with the AG genotype are when treated with irinotecan as compared to patients with the Acmay also influence the risk for ne Class 3 rs10929302 GA The AG genotype may be associated to patients with the Acmay also influence the risk for ne Class 3 rs10929302 GA	and colorectal cancer may have an increased risk for severe neutropenia compared to patients with the TT genotype, or a decreased risk as C genotype. This may be due to decreased enzymatic activity toward rinotecan, found in cells with the C allele as compared to those with clinical factors, such as UGT1A1*28, may also influence the risk for notecan. Independent concert may have an increased risk for severe neutropenia compared to patients with the GG genotype, or a decreased risk as A genotype. Other genetic and clinical factors, such as UGT1A1*28, utropenia in patients taking irinotecan. Independent concert in the concert of
UGT1A3	UDP glucuronosyltransferase 1 family, polypeptide A3
0 01	ated with a reduced risk for irinotecan-induced grade 3 or 4 hema- icities, including neutropenia and diarrhea, as compared to the AA
The AG genotype may be associtological and gastrointestinal tox	

tological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

— Clinical Annotations –

Class 3 rs10929302 GA

The AG genotype may be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematological and gastrointestinal toxicities, including neutropenia and diarrhea, as compared to the AA genotype.

UGT1A1

UGT1A5

UDP glucuronosyltransferase 1 family, polypeptide A1

UDP glucuronosyltransferase 1 family, polypeptide A5

	GA be associated with a reduced risk for irinotecan-induced grade 3 or 4 hematinal toxicities, including neutropenia and diarrhea, as compared to the AA
VEGFA	vascular endothelial growth factor A
	Clinical Annotations
capecitabine, fluorouraci	cancer and the AC genotype may have a reduced response to bevacizumab, il, irinotecan, leucovorin, or oxaliplatin as compared to patients with the CC and genetic factors may also affect response to chemotherapy in people with
ISONIAZID	
NAT2	N-acetyltransferase 2 (arylamine N-acetyltransferase)
	Clinical Annotations
when treated with anti-Clinical factors may also Class 2A rs1799930 Patients with the AA go when treated with anti-T decreased clearance of isc clinical factors may also Class 3 rs1799931 Patients with the GG gen treated with anti-TB dr	enotype and tuberculosis (TB) may have an increased risk for hepatotoxicity TB drugs as compared to patients with the CC genotype. Other genetic and influence risk for hepatotoxicity. AA enotype and tuberculosis (TB) may have an increased risk of hepatotoxicity TB drugs as compared to patients with the GG genotype. They also may have oniazid as compared to those with the AG or GG genotype. Other genetic and influence risk for hepatotoxicity and clearance of isoniazid.
IVACAFTOR	
	is transmembrane conductance regulator (ATP-binding cassette sub-family C,
	Clinical Annotations
· Class 1A rs7865542	$oldsymbol{1} GG$

Patients with the GG genotype and cystic fibrosis may not respond when treated with ivacaftor as compared to patients with the AA and AG genotypes. Other genetic and clinical factors may also influence the efficacy of ivacaftor.

\mathbf{L}	A	Λ	Λī	7	7 T	T	\Box	T	V	E
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ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	Clinical Annotations

· Class 3 rs1045642 GG

Patients with the GG genotype and HIV may have an increased risk of virological failure when receiving highly active antiretroviral therapy (HAART), as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of virological failure on HAART.

LAMOTRIGINE

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10 $$

\cdot Class 2B rs2011425 TT

Patients with the TT genotype and epilepsy who are administered lamotrigine may have increased serum concentrations of lamotrigine, as well as improved response to lamotrigine, and may need a higher dose as compared to patients with the GG genotype. Other clinical and genetic factors may also influence metabolism, response, and dose of lamotrigine.

\cdot Class 2B rs2011425 TT

Patients with the TT genotype and epilepsy who are administered lamotrigine may have increased serum concentrations of lamotrigine, as well as improved response to lamotrigine, and may need a higher dose as compared to patients with the GG genotype. Other clinical and genetic factors may also influence metabolism, response, and dose of lamotrigine.

\cdot Class 2B rs2011425 TT

Patients with the TT genotype and epilepsy who are administered lamotrigine may have increased serum concentrations of lamotrigine, as well as improved response to lamotrigine, and may need a higher dose as compared to patients with the GG genotype. Other clinical and genetic factors may also influence metabolism, response, and dose of lamotrigine.

	Class 2B rs2011425 T	
	serum concentrations of lam	ype and epilepsy who are administered lamotrigine may have increased notrigine, as well as improved response to lamotrigine, and may need a
	influence metabolism, respon	patients with the GG genotype. Other clinical and genetic factors may also use, and dose of lamotrigine.
	UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7
٠	serum concentrations of lam	ype and epilepsy who are administered lamotrigine may have increased notrigine, as well as improved response to lamotrigine, and may need a patients with the GG genotype. Other clinical and genetic factors may also
	UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
		——————————————————————————————————————
	CI	an and a second a second and a second and a second and a second and a second a second and a second a second and a second a second a second a second and a second a second a second a second a second and a second and
•		ype and epilepsy who are administered lamotrigine may have increased
		notrigine, as well as improved response to lamotrigine, and may need a patients with the GG genotype. Other clinical and genetic factors may also use, and dose of lamotrigine.
	UGT1A9	UDP glucuronosyltransferase 1 family, polypeptide A9
	Class 2B rs2011425 T	T
	Patients with the TT genot serum concentrations of lam	ype and epilepsy who are administered lamotrigine may have increased notrigine, as well as improved response to lamotrigine, and may need a patients with the GG genotype. Other clinical and genetic factors may also
L(ORAZEPAM	
	UGT2B15	UDP glucuronosyltransferase 2 family, polypeptide B15
	Class 2B rs1902023 A	C
•		pe may have decreased clearance of oxazepam or lorazepam as compared to
		pe, or increased clearance as compared to subjects with the AA genotype.

Other genetic and clinical factors may also influence the oral clearance of oxazepam or lorazepam.

— Clinical Annotations —

CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9
0 02	who are treated with losartan may have decreased metabolism of s with the AA genotype. Other genetic and clinical factors may also
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	may have poorer response to losartan in people with hypertension as A or AG genotype. Other genetic and clinical factors may also influence
MERCAPTOPURINE	
$ ext{TPMT}$	thiopurine S-methyltransferase
	——————————————————————————————————————

TPMT:*1/*1 Strong

Start with normal starting dose (e.g., 75 mg/m2/d or 1.5 mg/kg/d) and adjust doses of mercaptopurine (and of any other myelosuppressive therapy) without any special emphasis on mercaptopurine compared to other agents. Allow 2 weeks to reach steady state after each dose adjustment.

\cdot Class 3 rs1142345 TT

Pediatric patients with the TT genotype and Precursor Cell Lymphoblastic Leukemia-Lymphoma may experience decreased GI toxicity when treated with mercaptopurine and may require an increased dose as compared to patients with the CT or CC genotypes. Other genetic and clinical factors may also influence the likelihood of GI toxicity and dose of mercaptopurine in pediatric patients with Precursor Cell Lymphoblastic Leukemia-Lymphoma.

\cdot Class 4 rs3931660 AA

Patients with the AA genotype may have increased TPMT activity toward mercaptopurine as compared to patients with the AT genotype. Other genetic and clinical factors may also influence TPMT activity.

METHADONE

an increased dose of the drug as factors may also influence dose Class 3 rs2279343 AA Patients with the AA genotype an increased dose of the drug	who are being treated with methadone for heroin addiction may require as compared to patients with the GG genotype. However, one study his variant and methadone dose. Other genetic and clinical factors may
CYP2A7P1	cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene 1
	who are being treated with methadone for heroin addiction may require compared to patients with the TT genotype. Other genetic and clinical of methadone.
CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
when treated with methadone in TT or CT. Other genetic and collection of Class 3 rs3735451 TT Patients with the TT genotype withdrawal symptoms when tree	who are heroin dependent may have less severe side effects and opioid eated with methadone as compared to patients with genotype eated with methadone as compared to patients with genotype eated with methadone as compared to patients with the CC genotype.

METHOTREXATE

SLCO1B1 solute carrier organic anion transporter family, member 1B1

— Clinical Annotations

· Class 3 rs4149056 TT

Pediatric patients with the TT genotype and acute lymphoblastic leukemia may have increased clearance of methotrexate as compared to patients with the CC or CT genotype. Other genetic and clinical factors may also influence clearance of methotrexate.

\cdot Class 3 rs2306283 AG

Pediatric patients with the AG genotype and acute lymphoblastic leukemia may have increased clearance of methotrexate as compared to patients with the GG genotype. Other genetic and clinical factors may also influence clearance of methotrexate.

ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2 $$
	- Clinical Annotations
as compared to patients study done in an Asian por decreased clearance of me distribution of volume of Class 3 rs717620 C Patients with lymphoblas drug toxicity and decreas to patients with the CC a French population with	otype may have increased clearance and distribution of volume of methotrexate with the CC genotype. Please note: the opposite effect was observed in a oppulation with various types of lymphomas. The T allele was associated with ethotrexate. Other clinical and genetic factors may also affect clearance and methotrexate. The ticleukemia-lymphoma and the CT genotype may have an increased risk of sed clearance of methotrexate when treated with methotrexate as compared genotype. Please note: the opposite effect was observed in a study done in a various types of lymphomas. The C allele was associated with decreased et. Other clinical and genetic factors may also influence risk of drug toxicity
METOPROLOL	
ADRB2	adrenoceptor beta 2, surface
	- Clinical Annotations
triglyceridemia when trea	enotype and hypertension may have an increased risk of developing hyperted with atenolol or metoprolol as compared to patients with the CC or CG and clinical factors may also influence risk of hypertriglyceridemia.
MIDAZOLAM	
CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
to patients with the AG g midazolam. • Class 4 rs12721627	otype and tumors may have increased metabolism of midazolam as compared genotype. Other genetic and clinical factors may also influence metabolism of GG ruct caring the G variant is not associated with decreased clearance of mida-

-- Clinical Annotations

cytochrome P450, family 3, subfamily A

CYP3A

Patients with the GG genotype and tumors may have increased metabolism of midazolam as compared to patients with the AG genotype. Other genetic and clinical factors may also influence metabolism of midazolam.

Class 4 rs12721627 GG

The expression of a construct caring the G variant is not associated with decreased clearance of midazolam in transfected cells.

т.	T (\mathbf{T}	P	TTI	гт т	\mathbf{T}
IV	41	JК	P	н	IIV	H

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	— Clinical Annotations —

· Class 3 rs1045642 GG

Patients with the GG genotype may have decreased pain reduction when treated with morphine in cancer patients as compared to patients with genotype AA. Other genetic and clinical factors may also influence response to morphine.

NEVIRAPINE

CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6

· Class 2A rs3745274 GG

Patients with the GG genotype and HIV infection may have increased clearance of and decreased exposure to nevirapine as compared to patients with the TT or GT genotype. Other genetic and clinical factors may also influence clearance of nevirapine and exposure to drug.

· Class 3 rs28399499 TT

Patients with the TT genotype and HIV may have a decreased risk for Stevens-Johnson Syndrome/toxic epidermal necrolysis (SJS/TEN) when treated with nevirapine as compared to patients with the CC or CT genotype. Other genetic and clinical factors may also influence risk for developing SJS/TEN when receiving nevirapine.

CYP2A7P1	cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene 1
	- Clinical Annotations

Class 2A rs3745274 GG

Class 3 rs776746

CC

Patients with the GG genotype and HIV infection may have increased clearance of and decreased exposure to nevirapine as compared to patients with the TT or GT genotype. Other genetic and clinical factors may also influence clearance of nevirapine and exposure to drug.

clinical factors may also influence	clearance of nevirapine and exposure to drug.
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
	——- Clinical Annotations ————————————————————————————————————

Patients with the CC genotype and HIV infection who are treated with nevirapine may have increased clearance of the drug as compared to patients with the CT and TT genotype. Association with clearance was not found in a larger cohort in a separate study. Patients may also have differences in alanine aminotransferase levels, but association with toxicity has not been reported. Other genetic and clinical factors may also influence clearance of nevirapine.

ZSCAN25	zinc finger and SCAN domain containing 25
- Clin	nical Annotations
Class 3 rs776746 CC	

Patients with the CC genotype and HIV infection who are treated with nevirapine may have increased clearance of the drug as compared to patients with the CT and TT genotype. Association with clearance was not found in a larger cohort in a separate study. Patients may also have differences in alanine aminotransferase levels, but association with toxicity has not been reported. Other genetic and clinical factors may also influence clearance of nevirapine.

CYP3A cytochrome P450, family 3, subfamily A -- Clinical Annotations -

Class 3 rs776746 CC

Patients with the CC genotype and HIV infection who are treated with nevirapine may have increased clearance of the drug as compared to patients with the CT and TT genotype. Association with clearance was not found in a larger cohort in a separate study. Patients may also have differences in alanine aminotransferase levels, but association with toxicity has not been reported. Other genetic and clinical factors may also influence clearance of nevirapine.

ABCC10 ATP-binding cassette, sub-family C (CFTR/MRP), member 10 -- Clinical Annotations

Class 3 rs2125739 TT

Patients with the TT genotype and HIV may have increased concentrations of nevirapine as compared to patients with the CC genotype. Other genetic and clinical factors may also influence concentrations of nevirapine.

CYP2D6 cytochrome P450, family 2, subfamily D, polypeptide 6 -- Clinical Annotations

Class 3 rs28371706 GG

Pediatric patients with the GG genotype and HIV may have increased clearance of nevirapine as compared to pediatric patients with the AA or AG genotype. No significant association was seen in adults. Other genetic and clinical factors may also influence clearance of nevirapine.

NICOTINE

Patients with with the GG of this SNP	the AA genotype may have increased metabolism of nicotine as compared to patients or AG genotype. Other variants within the CYP2A6 gene should be considered - allele is part of the $*7$, $*10$, $*19$, $*36$, $*37$ CYP2A6 alleles. Other genetic and clinical factors ence metabolism of nicotine.
DDC	dopa decarboxylase (aromatic L-amino acid decarboxylase)
	2718541 $$ GG the GG genotype who smoke to bacco may have a decreased risk of addiction as compared to the AA genotype. Other genetic and clinical factors may also influence risk of smoking
NIFEDIPINE	
CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
· Class 4 rs4 In vitro, the c	987161 AA onstruct expressing the wild type allelic protein has average nifedipine metabolism.
CYP3A	cytochrome P450, family 3, subfamily A
	Clinical Annotations —
women with to of nifedipine. Class 4 rs4	nen with the CC genotype may have decreased clearance of nifedipine as compared to the CT or TT genotype. Other genetic and clinical factors may also influence clearance
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
	- Clinical Annotations
_	76746 CC nen with the CC genotype may have decreased clearance of nifedipine as compared to the CT or TT genotype. Other genetic and clinical factors may also influence clearance

——- Clinical Annotations —

_	CC genotype may have decreased clearance of nifedipine as compared to genotype. Other genetic and clinical factors may also influence clearance
ONDANSETRON	
CYP3A5	cytochrome P450, family 3, subfamily A, polypeptide 5
with the TT genotype. Oth	type may have increased metabolism of ondansetron as compared to patients are genetic and clinical factors may also influence metabolism of ondansetron.
ZSCAN25	zinc finger and SCAN domain containing 25
_	type may have increased metabolism of ondansetron as compared to patients are genetic and clinical factors may also influence metabolism of ondansetron.
CYP3A	cytochrome P450, family 3, subfamily A
with the TT genotype. Oth	type may have increased metabolism of ondansetron as compared to patients her genetic and clinical factors may also influence metabolism of ondansetron.
OPIOIDS	
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	type may have a decreased risk of opioid dependence when exposed to opioids ith the AG genotype. Other clinical and genetic factors may also influence

cate chol-O-methyl transfer as e

 \mathbf{COMT}

—- Clinical Annotations ————

of headache when disconting compared to patients with	ype with substance withdrawal syndrome may have an increased likelihood nuing the use of analgesics (such as opioids, NSAIDs, triptans, ergot) as the AA genotype. Other clinical and genetic factors may also influence atients with withdrawal syndrome who discontinue the use of analgesics.
PACLITAXEL	
CYP2C8	cytochrome P450, family 2, subfamily C, polypeptide 8
Patients with the TT genoty the AA or AT genotypes, h may also influence clearance Class 4 rs11572080 C Patients with the CC genoty the CT or TT genotypes, h may also influence clearance Class 4 rs10509681 T Patients with the TT genotypes	ype may have increased clearance of paclitaxel as compared to patients with lowever this has not been shown in vivo. Other genetic and clinical factors e of paclitaxel. The type may have increased metabolism of paclitaxel as compared to patients pes, however this has not been shown in vivo. Other genetic and clinical
CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
Patients with the GG gen	GG otype may have increased metabolism of paclitaxel as compared to pagenotypes. Other genetic and clinical factors may also influence paclitaxel

· Class 3 rs12721627 GG

CYP3A

Patients with the GG genotype may have increased metabolism of paclitaxel as compared to patients with the CC or CG genotypes. Other genetic and clinical factors may also influence paclitaxel metabolism.

——- Clinical Annotations —

cytochrome P450, family 3, subfamily A

Class 3 rs776746 CC

Patients with the CC genotype may have decreased but not absent risk of neurotoxicity when treated with paclitaxel as compared to patients with the TT genotype. Other genetic and clinical factors may also influence risk of toxicity with paclitaxel.

Class 3 rs776746 CC	
Patients with the CC genotyp	pe may have decreased but not absent risk of neurotoxicity when treated to patients with the TT genotype. Other genetic and clinical factors may with paclitaxel.
ZSCAN25	zinc finger and SCAN domain containing 25
Class 3 rs776746 CC	
Patients with the CC genotyp	pe may have decreased but not absent risk of neurotoxicity when treated to patients with the TT genotype. Other genetic and clinical factors may with paclitaxel.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
rate when treated with paclif AG. However, contradictory Caucasians. Other genetic an Class 3 rs1045642 GG	iated with increased disease control rate and increased overall survival taxel in Asians with metastatic breast cancer as compared to genotype findings have been reported and no association have been reported for a clinical factors may influence the response to paclitaxel.
when treated with paclitaxel	pe may have decreased risk of Neutropenia and Neurotoxicity Syndromes in cancer patients as compared to patients with genotype AA. Other ay influence the risk of adverse events to paclitaxel. 2A
when treated with paclitaxel genetic and clinical factors m	l in cancer patients as compared to patients with genotype AA. Other ay influence the risk of adverse events to paclitaxel.
when treated with paclitaxel genetic and clinical factors m CGINTERFERON ALFA-2 IFNL3	l in cancer patients as compared to patients with genotype AA. Other ay influence the risk of adverse events to paclitaxel. 2A
when treated with paclitaxel genetic and clinical factors m CGINTERFERON ALFA-2 IFNL3	l in cancer patients as compared to patients with genotype AA. Other ay influence the risk of adverse events to paclitaxel. 2A interferon, lambda 3
when treated with paclitaxel genetic and clinical factors m CGINTERFERON ALFA-2 IFNL3	in cancer patients as compared to patients with genotype AA. Other ay influence the risk of adverse events to paclitaxel. 2A interferon, lambda 3 — Dosing Guideline —

Patients with genotype CT may have decreased response to daclatasvir, peginterferon alfa-2a, peginterferon alfa-2b and ribavirin in people with Hepatitis C, Chronic as compared to genotypes CC. SVR24 rates are higher in patients treated with the combination of daclatasvir and pegIFN-alfa/RBV than those receiving pegIFN-alfa/RBV alone across all IL28B genotypes (CC, CT, or TT) regardless of viral subtypes. Other genetic and clinical factors may also influence the response to daclatasvir therapy.

	Clinical Apparations
feron alfa-2b and ribavirin in rates are higher in patients those receiving pegIFN-alfa/	may have decreased response to daclatasvir, peginterferon alfa-2a, pegintern people with Hepatitis C, Chronic as compared to genotypes CC. SVR24 treated with the combination of daclatasvir and pegIFN-alfa/RBV than RBV alone across all IL28B genotypes (CC, CT, or TT) regardless of viral clinical factors may also influence the response to daclatasvir therapy.
LDLR	low density lipoprotein receptor
HIV may have an increased	type who are co-infected with chronic hepatitis C, genotype 1 or 4, and likelihood of sustained virological response when treated with pegylated compared to patients with the AA or AG genotype. Other genetic and
	uence likelihood of sustained virological response.
IFNL3	interferon, lambda 3
	——————————————————————————————————————
IFNL3:rs12979860C/rs1 Phenotype (Genotype)	2979860C Strong
	- Clinical Annotations
· Class 3 rs12979860 C	T nay have decreased response to daclatasvir, peginterferon alfa-2a, peginter-

feron alfa-2b and ribavirin in people with Hepatitis C, Chronic as compared to genotypes CC. SVR24 rates are higher in patients treated with the combination of daclatasvir and pegIFN-alfa/RBV than those receiving pegIFN-alfa/RBV alone across all IL28B genotypes (CC, CT, or TT) regardless of viral subtypes. Other genetic and clinical factors may also influence the response to daclatasvir therapy.

IFNL4

IFNL4

interferon, lambda 4 (gene/pseudogene)

· Class 3 rs12979860 CT Patients with genotype CT ma	y have decreased response to daclatasvir, peginterferon alfa-2a, peginter-
feron alfa-2b and ribavirin in prates are higher in patients truthose receiving pegIFN-alfa/Rl	people with Hepatitis C, Chronic as compared to genotypes CC. SVR24 eated with the combination of daclatasvir and pegIFN-alfa/RBV than BV alone across all IL28B genotypes (CC, CT, or TT) regardless of viral linical factors may also influence the response to daclatasvir therapy.
LDLR	low density lipoprotein receptor
· Class 3 rs14158 GG	
HIV may have an increased linterferon and ribavirin as con	be who are co-infected with chronic hepatitis C, genotype 1 or 4, and kelihood of sustained virological response when treated with pegylated impared to patients with the AA or AG genotype. Other genetic and ince likelihood of sustained virological response.
PHENPROCOUMON	
VKORC1	vitamin K epoxide reductase complex, subunit 1
· Class 2A rs9934438 GA	
Patients with the AG genoty	pe may have decreased dose of acenocoumarol or phenprocoumon as otype GG. Other genetic and clinical factors may also influence the dose oumon.
PRSS53	protease, serine, 53
· Class 2A rs9934438 GA	
Patients with the AG genoty	pe may have decreased dose of acenocoumarol or phenprocoumon as otype GG. Other genetic and clinical factors may also influence the dose oumon.
PHENYTOIN	
CON1 A	
SCN1A	sodium channel, voltage-gated, type I, alpha subunit
· Class 2B rs3812718 CT	
Patients with the CT genotype	who are treated with phenytoin may require a higher dose as compared to . Other genetic and clinical factors may also influence dose of phenytoin.

LDLR	low density lipoprotein receptor
	- Clinical Annotations
ment as compared to patient	ype and vascular diseases may have a poorer response to pravastatin treats with the TT genotype, or a better response as compared to patients with netic and clinical factors may also influence pravastatin response.
ROPOFOL	
UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
treated with propofol as comfactors may also influence re	ype may have decreased but not non-existent risk of adverse effects when apared to patients with the AA or AG genotype. Other genetic and clinical esponse to propofol.
UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
	- Clinical Annotations
	ype may have decreased but not non-existent risk of adverse effects when pared to patients with the AA or AG genotype. Other genetic and clinical
UGT1A9	UDP glucuronosyltransferase 1 family, polypeptide A9
	- Clinical Annotations
	ype may have decreased but not non-existent risk of adverse effects when apared to patients with the AA or AG genotype. Other genetic and clinical
CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6

Class 3 rs3745274 GG

Patients under general anaesthesia with genotypes GG may need increased dose of propofol as compared to patients with genotype TT or GT. Other genetic and clinical factors may also influence the dose of propofol.

CYP2A7P1	cytochrome P450, family 2, subfamily A, polypeptide 7 pseudogene 1
	Clinical Annotations
· Class 3 rs3745274 (GG
_	esthesia with genotypes GG may need increased dose of propofol as compared TT or GT. Other genetic and clinical factors may also influence the dose of
QUETIAPINE	
COMT	catechol-O-methyltransferase
CI	
quetiapine as compared to	notype and schizophrenia may have a poorer response to treatment with patients with the GG genotype, or a better response as compared to patients ther genetic and clinical factors may also influence quetiapine response.
Patients with the CG generated to	notype and schizophrenia may have a poorer response to treatment with patients with the GG genotype, or a better response as compared to patients ther genetic and clinical factors may also influence quetiapine response.
RALOXIFENE	
UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
· Class 3 rs1042597	CC
	with the CC genotype and schizophrenia may have increased response to

Post menopausal women with the CC genotype and schizophrenia may have increased response to raloxifene compared to patients with the CG genotype. Other genetic and clinical factors may affect response to raloxifene.

RANIBIZUMAB

VEGFA	vascular endothelial growth factor A
	annotations —

· Class 3 rs2010963 CG

Patients with the CG genotype and choroidal neovascularization may have a better response to anti-VEGF treatment, as compared to patients with the CC genotype. Other genetic and clinical factors may also influence response to anti-VEGF treatment.

REPAGLINIDE

SLCO1B1	solute carrier organic anion transporter family, member 1B1
· Class 3 rs2306283 A While the GG genotype is shown for the GA genotype	associated with reduced plasma concentrations of repaglinide, no results are
CYP2C8	cytochrome P450, family 2, subfamily C, polypeptide 8
compared to patients with in blood glucose lowering ϵ	(CYP2C8*1/*1) genotype may have decreased metabolism of repaglinide the CT genotype (CYP2C8*3/*1). No association was found with differences efficacy. Please note, the study supporting this annotation was carried out in genetic and clinical factors may also influence metabolism of repaglinide.
NAT2	N-acetyltransferase 2 (arylamine N-acetyltransferase)
Class 2A rs1041983	TT
when treated with anti-TI clinical factors may also in	otype and tuberculosis (TB) may have an increased risk for hepatotoxicity drugs as compared to patients with the CC genotype. Other genetic and fluence risk for hepatotoxicity.
Patients with the AA gen when treated with anti-TE decreased clearance of ison clinical factors may also in	AA otype and tuberculosis (TB) may have an increased risk of hepatotoxicity drugs as compared to patients with the GG genotype. They also may have liazed as compared to those with the AG or GG genotype. Other genetic and affluence risk for hepatotoxicity and clearance of isoniazid.
Patients with the GG geno treated with anti-TB drug	type and tuberculosis (TB) may have a decreased risk of hepatotoxicity when gs as compared to patients with the AA or AG genotype. However, some with hepatotoxicity. Other genetic and clinical factors may also influence
SLCO1B1	solute carrier organic anion transporter family, member 1B1

· Class 3 rs11045819 CC

Patients with the CC genotype may have decreased clearance of rifampin as compared to patients with the AC genotype. Other genetic and clinical factors may also influence rifampin clearance.

	~1 -		. ~
•	Class 3	rs2306283	AG

Patients with the AG genotype may have increased clearance of rifampin as compared to patients with the GG genotype. Other genetic and clinical factors may also influence rifampin clearance.

RISPERIDONE

ABCB1 ATP-binding cassette, sub-family B (MDR/TAP), member 1 — Clinical Annotations

· Class 3 rs1045642 GG

Patients with the GG genotype and schizophrenia may have a shorter QTc interval when treated with risperidone as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence QTc interval in patients taking risperidone.

CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4

\cdot Class 3 rs35599367 GG

Patients with the GG genotype and psychiatric disorders may have increased clearance of risperidone compared to patients with the AG genotype. Other clinical and genetic factors likely affect risperidone pharmacokinetics.

CYP3A	cytochrome P450, family 3, subtamily A

· Class 3 rs35599367 GG

Patients with the GG genotype and psychiatric disorders may have increased clearance of risperidone compared to patients with the AG genotype. Other clinical and genetic factors likely affect risperidone pharmacokinetics.

RITONAVIR.

UGT1A	UDP glucuronosyltransferase 1 family, polypeptide A complex locus

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

Class 3 rs8330 *GC*

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A3

UDP glucuronosyltransferase 1 family, polypeptide A3

— Clinical Annotations

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

|--|

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A5	UDP glucuronosyltransferase 1 family, polypeptide $A5$

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A6	UDP glucuronosyltransferase 1 family, polypeptide A6

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A7

UDP glucuronosyltransferase 1 family, polypeptide A7

— Clinical Annotations	
- Chinca Annotations	

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGTIA8	UDP glucuronosyltransferase I family, polypeptide A8

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

· Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

· Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

UGT1A1	UDP glucuronosyltransferase 1 family, polypeptide A			

· Class 3 rs10929303 TC

Patients with the CT genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the TT genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs1042640 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to patients with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in people with HIV who are taking atazanavir and ritonavir.

 \cdot Class 3 rs8330 GC

Patients with the CG genotype and HIV may have a decreased risk of nephrolithiasis when treated with atazanavir and ritonavir as compared to patients with the GG genotype and an increased risk of nephrolithiasis as compared to people with the CC genotype. Other genetic and clinical factors may also affect risk of nephrolithiasis in patients with HIV who are taking atazanavir and ritonavir.

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1

· Class 3 rs1045642 GG

Patients with the GG genotype and HIV may have an increased risk of virological failure when receiving highly active antiretroviral therapy (HAART), as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of virological failure on HAART.

 \cdot Class 3 rs1045642 GG

Patients with the GG genotype and HIV may have increased concentrations of atazanavir as compared to patients with the AA genotypes, although this is contradicted in one study. There is no evidence that the GG genotype is associated with hyperbilirubinemia, drug discontinuation, or nephrolithiasis. Other clinical and genetic factors may also influence the concentrations of atazanavir in patients with HIV.

ROSIGLITAZONE

CYP2C8	cytochrome P450, family 2, subfamily C, polypeptide 8
a larger change in HbA1c, an (CYP2C8*3/*3) or CT (CYP2c	C8*1/*1) genotype may have decreased metabolism of rosiglitazone, d an increased risk of edema as compared to patients with the CC C8*3/*1) genotype. One study found no association with blood glucose al factors may also influence metabolism of rosiglitazone, risk of edema
SLCO1B1	solute carrier organic anion transporter family, member 1B1
0 11	e may have decreased response to rosiglitazone in people with type II to patients with genotype CC or CT. Other genetic and clinical factors e to rosiglitazone.
VEGFA	vascular endothelial growth factor A
	may have decreased response to sildenafil in men with Erectile Dysfuncith genotype CC. Other genetic and clinical factors may also influence
SIMVASTATIN	
HMGCR	3-hydroxy-3-methylglutaryl-CoA reductase
	- Clinical Annotations
Class 4 mg2946669 CC	

The GG genotype may be associated with decreased induction of full-length transcripts and increased expression of spliced HMGCRv1 transcript as compared to AA genotype. ABCC2 ATP-binding cassette, sub-family C (CFTR/MRP), member 2 -- Clinical Annotations --Class 3 rs717620 Patients with the CT genotype may have decreased dose of simvastatin and atorvastatin as compared to patients with genotype CC. Other genetic and clinical factors may also influence the dose of simvastatin. ABCB1 ATP-binding cassette, sub-family B (MDR/TAP), member 1 -- Clinical Annotations --Class 3 rs1128503 GG Patients with the GG genotype and hypercholesterolemia may lesser reduction in LDL and total cholesterol when treated with simvastatin as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence cholesterol levels. Class 3 rs1128503 GG Patients with the GG genotype and hypercholesterolemia may have an increased risk for myalgia when treated with simvastatin as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk for myalgia. CYP3A4 cytochrome P450, family 3, subfamily A, polypeptide 4 -- Clinical Annotations Class 3 rs2740574 TTPatients with the TT genotype may be more likely to require a decrease in dose or switch to a different drug when treated with atorvastatin or simvastatin as compared to patients with the CC or CT genotype. Other genetic and clinical factors may also influence dose of simvastatin or atorvastatin, or likelihood of switching to a different drug. CYP3A cytochrome P450, family 3, subfamily A —- Clinical Annotations -Class 3 rs2740574

Patients with the TT genotype may be more likely to require a decrease in dose or switch to a different drug when treated with atorvastatin or simvastatin as compared to patients with the CC or CT genotype. Other genetic and clinical factors may also influence dose of simvastatin or atorvastatin, or likelihood of switching to a different drug.

SORAFENIB

		Clinical Annotations –	
Class 3 rs2010963	CG		

Patients with the CG genotype may have increased risk of hand-foot syndrome when treated with sorafenib in people with Carcinoma, Renal Cell as compared to patients with genotype CC. Other genetic and clinical factors may also influence the toxicity to sorafenib.

· Class 3 rs1570360 AG

Patients with the AG genotype may have unfavorable progression-free survival when treated with sorafenib in people with Carcinoma, Renal Cell as compared to patients with genotype GG. Other genetic and clinical factors may also influence the response to sorafenib.

· Class 3 rs2010963 CG

Patients with the CG genotype may have increased progression-free survival and increased overall survival when treated with sorafenib in people with Hepatocellular Carcinoma as compared to patients with genotype GG. Other genetic and clinical factors may also influence the response to sorafenib.

SLCO1B1	solute carrier organic anion transporter family, member 11		
	- Clinical Annotations —		

\cdot Class 3 rs4149056 TT

Patients with the TT genotype may have increased likelihood of developing Thrombocytopenia when treated with sorafenib as compared to patients with genotype CC. Other genetic and clinical factors may also influence the response to sorafenib.

 \cdot Class 3 rs2306283 AG

Patients with the AG genotype may have decreased likelihood of developing Diarrhea when treated with sorafenib as compared to patients with genotype AA. Other genetic and clinical factors may also influence the response to sorafenib.

ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), mem		
	—- Clinical Annotations ————		

\cdot Class 3 rs717620 CT

Patients with the CT genotype may have decreased risk of skin rash when treated with sorafenib in people with Carcinoma, Renal Cell as compared to patients with genotype CC. Other genetic and clinical factors may also influence the toxicity to sorafenib.

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	- Clinical Annotations —

· Class 3 rs1045642 GG

Patients with the GG genotype may have decreased risk of hypertension when treated with sorafenib in people with Carcinoma, Renal Cell as compared to patients with genotype AA or AG. Other genetic and clinical factors may also influence the toxicity to sorafenib.

SULFONAMIDES, UREA DERIVATIVES

	AC rith the AC genotype were not statistically significant.
SUNITINIB	
VEGFA	vascular endothelial growth factor A
of developing grade 3 hy	AC enotype may have higher increase in systolic blood pressure and increased risk ypertension when treated with sunitinib as compared to patients with genotype clinical factors may also influence the response to sunitinib.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
	- Clinical Annotations —
· Class 3 rs1045642	GG
when treated with suni no association between Other genetic and clinic	enotype and renal cell carcinoma may have an increased risk for adverse effects tinib as compared to patients with the AA or AG genotype. One study found this SNP and thrombocytopenia, neutropenia, anemia or hand-food syndrome. cal factors may also influence risk for sunitinib toxicities. CC
when treated with sunit	carcinoma and the CC genotypes may have an increased risk of neutropenia inib as compared to patients with any of the following genotypes: AA, AC, AT netic factors may also influence risk of neutropenia in patients with renal cell ted with sunitinib. GG
Patients with renal cell increased risk of neutronalthough this has been	carcinoma and the GG genotype who are treated with sunitinib may have an penia, leukopenia, and diarrhea as compared to patients with the AA genotypes, a contradicted by some studies. Other clinical and genetic factors may also a patients with renal cell carcinoma who are administered sunitinib. CC

Patients with renal cell carcinoma and the CC genotype may have an incressed response to sunitinib as compared to patients with the AA genotypes. There is no association between this SNP and overall or progression free survival. Response here refers to stable disease or partial response and non-response to progressive disease. Other clinical and genetic factors may also influence response to sunitinib in patients with renal cell carcinoma.

TACROLIMUS

· Class 3 rs1045642 GG

Patients with the GG genotype who are undergoing organ transplantation may have increased clearance and dose requirements of tacrolimus, as compared to patients with the AA or AG genotype. However, the vast majority of studies find no association between this SNP and clearance or dose of tacrolimus. Other genetic and clinical factors, such as CYP3A5*3, may also influence clearance and dose of tacrolimus.

· Class 3 rs1045642 GG

Patients with the GG genotype who are CYP2C19 extensive metabolizers and are receiving tacrolimus after renal transplantation may have increased plasma concentrations of (R)-lansoprazole but no significant differences in the frequency of gastroesophageal symptoms as compared to patients with the AA genotype. Other genetic and clinical factors may also influence lansoprazole clearance.

· Class 3 rs1045642 GG

Pediatric patients with the GG genotype who are treated with prednisone and tacrolimus may have an increased risk of remaining on steroids 1 year after heart transplantation compared to patients with the AA or AG genotype. Other genetic and clinical factors may also influence risk of remaining on steroids 1 year after transplantation.

\cdot Class 3 rs1045642 GG

Patients who receive a kidney with the GG genotype may have increased estimated glomerular filtration rate (eGFR) when treated with tacrolimus as compared to patients with the AA or AG genotype. No significant results were seen when recipient genotype was considered. Other genetic and clinical factors may also influence eGFR.

· Class 3 rs2032582 CC

Patients with CC genotype may have lower success rate in achieving short-term remission when treated with tacrolimus in people with Colitis, Ulcerative as compared to patients with the AA genotype. The majority of studies find no association with dose of tacrolimus in people with transplantations as compared and genotypes of this SNP. Other genetic or clinical factors may influence response and dose of tacrolimus.

\cdot Class 3 rs1128503 GG

Patients with the GG genotype who are undergoing organ transplantation may have decreased concentrations of tacrolimus as compared to patients with the AA or AG genotype. However, the majority of the literature evidence shows no association between this variant and tacrolimus concentrations, clearance or dose. Other genetic and clinical factors may also influence concentrations of tacrolimus.

· Class 3 rs2032582 CC

Patients with the CC genotype who are undergoing organ transplantation may have increased metabolism and dose requirements of tacrolimus, as compared to patients with the AA, AC, CT or TT genotypes. However, the majority of studies have found no association between this polymorphism and metabolism or dose of tacrolimus. Other genetic and clinical factors, such as CYP3A5*3, may also influence metabolism and dose of tacrolimus.

\cdot Class 3 rs1045642 GG

Patients with the GG genotype and ulcerative colitis may have a poorer chance at achieving remission when treated with tacrolimus as compared to patients with the AA genotype. Other genetic and clinical factors may also influence likelihood of ulcerative colitis remission.

\cdot Class 3 rs1045642 GG

Patients with the GG genotype who are undergoing kidney transplantation and are treated with tacrolimus may have decreased risk of experiencing transplant rejection as compared to patients with the AG genotype. However, the majority of studies find no association between this polymorphism and risk for transplant rejection. Other genetic and clinical factors may also influence risk of transplant rejection.

Patients with the GG genotype who are undergoing kidney transplantation may have a decreased risk of hypokalemia when treated with tacrolimus as compared to patients with the AG genotype. Other genetic and clinical factors may also influence risk of hypokalemia.

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cytochrome P450, family 3, subfamily A, polypeptide 5

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Dosing Guider	.1116		

CYP3A5:*1A/*1A Strong

Increase starting dose 1.5 to 2 times recommended starting dose. Total starting dose should not exceed 0.3mg/kg/day. Use therapeutic drug monitoring to guide dose adjustments.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing liver transplantation may have an increased risk for renal dysfunction when treated with tacrolimus as compared to patients with the CT or TT genotype (*1/*3 or *1/*1). Other genetic and clinical factors may also influence risk for renal dysfunction.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for neurotoxicity when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence risk for neurotoxicity in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing kidney transplantation may have decreased systolic and diastolic blood pressure when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. However, the majority of studies show no association between the CC genotype and blood pressure. Other genetic and clinical factors may also influence changes in blood pressure in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for infections when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. Other genetic and clinical factors may also influence risk for infections in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and ulcerative colitis may have an increased chance of achieving remission when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence chance of remission from ulcerative colitis.

ZSCAN25	zinc finger and SCAN domain containing 25
	nnotations —

Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing liver transplantation may have an increased risk for renal dysfunction when treated with tacrolimus as compared to patients with the CT

or TT genotype ($^*1/^*3$ or $^*1/^*1$). Other genetic and clinical factors may also influence risk for renal dysfunction.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for neurotoxicity when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence risk for neurotoxicity in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing kidney transplantation may have decreased systolic and diastolic blood pressure when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. However, the majority of studies show no association between the CC genotype and blood pressure. Other genetic and clinical factors may also influence changes in blood pressure in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for infections when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. Other genetic and clinical factors may also influence risk for infections in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and ulcerative colitis may have an increased chance of achieving remission when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence chance of remission from ulcerative colitis.

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cytochrome P450, family 3, subfamily A

— Clinical Ann	notations	
—- Unincai Ani	notations —	

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing liver transplantation may have an increased risk for renal dysfunction when treated with tacrolimus as compared to patients with the CT or TT genotype (*1/*3 or *1/*1). Other genetic and clinical factors may also influence risk for renal dysfunction.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for neurotoxicity when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence risk for neurotoxicity in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing kidney transplantation may have decreased systolic and diastolic blood pressure when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. However, the majority of studies show no association between the CC genotype and blood pressure. Other genetic and clinical factors may also influence changes in blood pressure in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) undergoing organ transplantation may have a decreased risk for infections when treated with tacrolimus as compared to patients with the CT or TT (*1/*3 or *1/*1) genotype. Other genetic and clinical factors may also influence risk for infections in patients receiving tacrolimus.

· Class 3 rs776746 CC

Patients with the CC genotype (CYP3A5 *3/*3) and ulcerative colitis may have an increased chance of achieving remission when treated with tacrolimus as compared to patients with the CT (*1/*3) genotype. Other genetic and clinical factors may also influence chance of remission from ulcerative colitis.

· Class 3 rs35599367 GG

Transplant recipients with the GG genotype (also known as CYP3A4 *1/*1) may have increased metabolism of tacrolimus, resulting in decreased exposure and a higher dose requirement (among other pharmacokinetic parameters), as compared to patients with the AG or AA genotype (CYP3A4 *22/*1 and *22/*22). Around half of studies report an association, and half report no association. One study found an association with liver transplant donor genotype but not recipient genotype. Other genetic and clinical factors, such as CYP3A5 *3 (rs776746), may also influence metabolism of tacrolimus.

· Class 4 rs4986910 AA

Patients with the AA genotype who are undergoing kidney transplantation may require an increased dose of tacrolimus as compared to patients with the GG genotype. Other genetic and clinical factors, such as CYP3A5*3, may also influence dose of tacrolimus.

CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
	- Clinical Annotations
	- Chincal Affilotations

· Class 3 rs28371759 AA

Patients with the AA genotype (CYP3A4 *1/*1) who underwent kidney transplantation may have decreased metabolism of tacrolimus as compared to patients with the AG genotype (*1/*18B). Other genetic and clinical factors, such as rs776746 (CYP3A5*3), may also influence metabolism of tacrolimus.

· Class 3 rs35599367 GG

Transplant recipients with the GG genotype (also known as CYP3A4 *1/*1) may have increased metabolism of tacrolimus, resulting in decreased exposure and a higher dose requirement (among other pharmacokinetic parameters), as compared to patients with the AG or AA genotype (CYP3A4 *22/*1 and *22/*22). Around half of studies report an association, and half report no association. One study found an association with liver transplant donor genotype but not recipient genotype. Other genetic and clinical factors, such as CYP3A5 *3 (rs776746), may also influence metabolism of tacrolimus.

· Class 4 rs4986910 AA

Patients with the AA genotype who are undergoing kidney transplantation may require an increased dose of tacrolimus as compared to patients with the GG genotype. Other genetic and clinical factors, such as CYP3A5*3, may also influence dose of tacrolimus.

TAMOXIFEN

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	- Clinical Annotations

\cdot Class 3 rs1045642 GG

Women with the GG genotype and breast cancer may have a decreased chance of disease recurrence when treated with tamoxifen as compared to patients with the AG genotype. Other genetic and clinical factors may also influence breast cancer recurrence.

	also influence risk of endometrial cancer.
CYP3A	cytochrome P450, family 3, subfamily A
Class 3 rs2740574	TT
cancer following tamoxid	notype and breast cancer may have a decreased risk of developing endometria fen treatment as compared to patients with the CT genotype. Other geneticalso influence risk of endometrial cancer.
ABCC2	ATP-binding cassette, sub-family C (CFTR/MRP), member 2 $$
	- Clinical Annotations —
Class 4 rs717620 (CT
treated with tamoxifen factors may also influence	enotype and breast neoplasms may have increased disease-free survival whereast compared to patients with the CC genotype. Other genetic and clinicate disease-free survival with tamoxifen treatment.
UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
	obi graculonosymianolerase i lammy, polypepulae ilio
	- Clinical Annotations
Class 3 rs2011425 Patients with the TT gn-glucuronide when taki	——————————————————————————————————————
Class 3 rs2011425 Patients with the TT g n-glucuronide when taki clinical and genetic factor	——————————————————————————————————————
Class 3 rs2011425 Patients with the TT gn-glucuronide when taki	——————————————————————————————————————
Class 3 rs2011425 Patients with the TT g n-glucuronide when taki clinical and genetic factor	——————————————————————————————————————
Class 3 rs2011425 Patients with the TT g n-glucuronide when taki clinical and genetic facto UGT1A4 Class 3 rs2011425 Patients with the TT g n-glucuronide when taki	——————————————————————————————————————
Class 3 rs2011425 Patients with the TT g n-glucuronide when taki clinical and genetic facto UGT1A4 Class 3 rs2011425 Patients with the TT g n-glucuronide when taki	— Clinical Annotations TT enotype and breast cancer may have decreased concentrations of tamoxifering tamoxifen compared to patients with the GG and GT genotypes. Other ors may affect the metabolism of tamoxifen. UDP glucuronosyltransferase 1 family, polypeptide A4 — Clinical Annotations TT enotype and breast cancer may have decreased concentrations of tamoxifering tamoxifen compared to patients with the GG and GT genotypes. Other

	UDP glucuronosyltransferase 1 family, polypeptide A6
	- Clinical Annotations —
n-glucuronide when taking tak	pe and breast cancer may have decreased concentrations of tamoxifen moxifen compared to patients with the GG and GT genotypes. Other by affect the metabolism of tamoxifen.
UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7
	- Clinical Annotations —
n-glucuronide when taking tak	pe and breast cancer may have decreased concentrations of tamoxifen moxifen compared to patients with the GG and GT genotypes. Other by affect the metabolism of tamoxifen.
UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
	pe and breast cancer may have decreased concentrations of tamoxifen moxifen compared to patients with the GG and GT genotypes. Other
_	y affect the metabolism of tamoxifen.
_	
clinical and genetic factors ma	y affect the metabolism of tamoxifen.
Class 3 rs2011425 TT Patients with the TT genotypen-glucuronide when taking tak	UDP glucuronosyltransferase 1 family, polypeptide A9 ———————————————————————————————————
Class 3 rs2011425 TT Patients with the TT genotypen-glucuronide when taking tak	UDP glucuronosyltransferase 1 family, polypeptide A9 — Clinical Annotations — ce and breast cancer may have decreased concentrations of tamoxifen moxifen compared to patients with the GG and GT genotypes. Other

Patients with the TT genotype and breast cancer may have decreased concentrations of tamoxifenn-glucuronide when taking tamoxifen compared to patients with the GG and GT genotypes. Other

	Clin	ical	Anı	nota	tions
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· Class 1A rs55886062 AA

Patients with the AA genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have a decreased, but not absent, risk for drug toxicity as compared to patients with the AC or CC genotype (DPYD *1/*13 or *13/*13). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

· Class 1A rs3918290 *CC*

Patients with the CC genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of fluoropyrimidine drugs and 2) decreased, but not non-existent, risk for drug toxicity as compared to patients with the CT or TT genotype (DPYD *1/*2A or *2A/*2A). Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine based chemotherapy.

\cdot Class 1A rs67376798 TT

Patients with the TT genotype and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) increased clearance of the drug and 2) decreased, but not absent, risk and reduced severity of drug toxicity as compared to patients with the AT genotype. Fluoropyrimidines are often used in combination chemotherapy such as FOLFOX (fluorouracil, leucovorin and oxaliplatin), FOLFIRI (fluorouracil, leucovorin and irinotecan) or FEC (fluorouracil, epirubicin and cyclophosphamide) or with other drugs such as bevacizumab, cetuximab, raltitrexed. The combination and delivery of the drug may influence risk for toxicity. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

\cdot Class 3 rs1801159 TT

Patients with the TT genotype (DPYD *1/*1) and cancer who are treated with fluoropyrimidine-based chemotherapy may have 1) a decreased likelihood of nausea, vomiting, and leukopenia, 2) increased response and 3) increased clearance of fluorouracil as compared to patients with the CT or CC genotype (DPYD *1/*5 or *5/*5). However, other studies find no associations or contradictory associations with fluoropyrimidine-induced drug toxicity or response. Other genetic and clinical factors may also influence response to fluoropyrimidine-based chemotherapy.

CYP2A6	cytochrome P450, family 2, subfamily A, polypeptide 6

· Class 4 rs28399433 AA

Hepatic cells with the AA genotype may have increased expression of the CYP2A6 gene, resulting in increased metabolism of tegafur, as compared to those with the AC or CC genotype. Other genetic and clinical factors may also influence CYP2A6 expression and tegafur metabolism.

THALIDOMIDE

	- Clinical Annotations —
0 0.	e may have a decreased but not absent risk of toxicity with docetaxel and atients with the CT or TT genotypes. Other genetic and clinical factors response.
NAT2	N-acetyltransferase 2 (arylamine N -acetyltransferase)
	- Clinical Annotations
	e may have an increased risk of toxicity with docetaxel and thalidomide the AA genotype. Other genetic and clinical factors may also influence
ΓICAGRELOR	
CYP3A4	cytochrome P450, family 3, subfamily A, polypeptide 4
	be and acute coronary syndrome may have decreased concentrations of ts with the CT genotype. Other factors may affect concentrations of
SLCO1B1	solute carrier organic anion transporter family, member 1B1
	be and acute coronary syndrome may have decreased concentrations of swith the CC and CT genotypes. Other factors may affect concentrations
FOLBUTAMIDE	
CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9
	e may have increased metabolism of tolbutamide as compared to patients. Other genetic and clinical factors may also influence tolbutamide

metabolism.

 \cdot Class 3 rs2070959 AG

SLC22A1	solute carrier family 22 (organic cation transporter), member 1
	- Clinical Annotations
exposed to tramadol in healt genetic or clinical factors may Class 3 rs34130495 GG Patients with the GG genoty when exposed to tramadol in	pe may have lower plasma concentrations of O-desmethyltramadol when thy individuals as compared to patients with the TT genotype. Other y influence the response to tramadol.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
response to tramadol. VALPROIC ACID	
UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
	e may require an increased dose of valproic acid compared to patients with etic and clinical factors may also influence a patients dose requirements.
UGT1A6	UDP glucuronosyltransferase 1 family, polypeptide A6
0 0.	e may require an increased dose of valproic acid compared to patients with etic and clinical factors may also influence a patients dose requirements.
UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7

the AA genotype. Other	er genetic and clinical factors may also influence a patients dose requirements.
UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8
_	AG enotype may require an increased dose of valproic acid compared to patients with er genetic and clinical factors may also influence a patients dose requirements.
UGT1A9	UDP glucuronosyltransferase 1 family, polypeptide A9
	- Clinical Annotations
Patients with the AG ge	AG enotype may require an increased dose of valproic acid compared to patients with α genetic and clinical factors may also influence a patients dose requirements.
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
Patients with genotype pared to patients with g to have different respons	GG and depressive disorder may have increased response to venlafaxine compensional entry and are also as to venlafaxine compared to patients with other genotypes. Other clinical and affect response to venlafaxine.
CYP2D6	cytochrome P450, family 2, subfamily D, polypeptide 6
	- Clinical Annotations —
· Class 4 rs36754300 Patients with the GG g	GG senotype were not studied.
/ERAPAMIL	
ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1
· Class 3 rs2032582	CC

Patients with the AG genotype may require an increased dose of valproic acid compared to patients with

Patients with the CC genotype may have decreased metabolism of verapamil as compared to patients with the AA or AC genotype. Other genetic and clinical factors may also impact the metabolism of verapamil.

· Class 3 rs1045642 GG

Patients with the GG genotype may have decreased metabolism of verapamil as compared to patients with the AA or AG genotype. Other genetic and clinical factors may also impact the metabolism of verapamil.

T 7	-	-	-	45.7	- T	-
V	11	ľΑ	ιN	/	IIN	H)

CYP4F2	cytochrome P450, family 4, subfamily F, polypeptide 2

· Class 3 rs2108622 CC

Patients with the CC genotype may have decreased steady-state levels of vitamin E when taking vitamin E supplements as compared to patients with the CT and TT genotypes. Other clinical and genetic factors may also influence steady-state levels of vitamin E in patients taking vitamin E supplements.

· Class 4 rs3093105 AA

The AA genotype may be associated with decreased CYP4F2 activity and decreased vitamin e metabolism as compared to the AC or CC genotype. This is based solely on an in vitro study in a haploid heterologous cell system. Other clinical and genetic factors may also influence metabolism of vitamin e.

WARFARIN

CYP2C9	cytochrome P450, family 2, subfamily C, polypeptide 9
	— Dosing Guideline ————————————————————————————————————

CYP2C9:*1/*1 N/A

Estimate the anticipated stable dose of warfarin using the algorithms available on http://www.warfarindosing.org, the IWPC Pharmacogenetic Dosing Algorithm or the FDA-approved drug label

—- Clinical Annotations —

Class 1A rs1057910 AC

Patients with the AC genotype: 1) may require a decreased dose of warfarin as compared to patients with the AA genotype 2) may have an increased risk for adverse events as compared to patients with the AA genotype.

 \cdot Class 2A rs7900194 GG

Patients with the GG genotype who are treated with warfarin may require a higher maintenance dose as compared to patients with the AG or GG genotype. Other clinical or genetic factors may also influence warfarin dose.

 \cdot Class 2A rs56165452 TT

Patients with the TT genotype may required higher dose of warfarin as compared to patients with the CT or CC genotype. Other clinical or genetic factors may also influence warfarin dose. This variant rs56165452 defines CYP2C9*4.

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CYP2C9:*1/*1 N/A

Estimate the anticipated stable dose of warfarin using the algorithms available on http://www.warfarindosing.org, the IWPC Pharmacogenetic Dosing Algorithm or the FDA-approved drug label

-- Clinical Annotations ----

· Class 1B rs9934438 GA

Patients with the AG genotype who are treated with warfarin may require a lower dose as compared to patients with the GG genotype, and a higher dose as compared to patients with the AA genotype. Other clinical and genetic factors may also influence a patients required dose of warfarin.

 \cdot Class 2A rs9923231 CT

Patients with genotype CT may require shorter time to therapeutic INR when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the response to warfarin.

· Class 2A rs9923231 CT

Patients with the CT genotype may have increased risk of over-anticoagulation when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the toxicity to warfarin.

 \cdot Class 2B rs7196161 GA

Patients with the AG genotype may require an increased dose of warfarin as compared to patients with the GG genotype and a decreased dose of warfarin as compared to patients with the AA genotype. Other clinical and genetic factors may also influence the dose of warfarin.

 \cdot Class 3 rs17880887 GG

Patients with the GG genotype may require lower dose of warfarin as compared to patients with the GT or TT genotype. Other genetic and clinical factors may also influence warfarin dose. This variant rs17880887 is part of VKORC1 H8 and H9 haplotypes.

 \cdot Class 3 rs9923231 CT

Patients with the CT genotype may spent less time in INR therapeutic range (TTR) when treated with warfarin as compared with patients with genotype CC. Contradictory findings have also been reported. Other genetic and clinical factors may also influence the response to warfarin.

· Class 3 rs9934438 GA

Patients with the AG genotype may have decreased time in the rapeutic range of INR (TTR) when treated with warfarin as compared to genotype GG. Other genetic and clinical factors may also influence the response to warfarin.

PRSS53 protease, serine, 53

- Clinical Annotations

· Class 1B rs9934438 GA

Patients with the AG genotype who are treated with warfarin may require a lower dose as compared to patients with the GG genotype, and a higher dose as compared to patients with the AA genotype. Other clinical and genetic factors may also influence a patients required dose of warfarin.

 \cdot Class 2A rs9923231 CT

Patients with genotype CT may require shorter time to the rapeutic INR when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the response to warfarin.

Class 2A rs9923231 CT

Patients with the CT genotype may have increased risk of over-anticoagulation when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the toxicity to warfarin.

· Class 3 rs9923231 CT

Patients with the CT genotype may spent less time in INR therapeutic range (TTR) when treated with warfarin as compared with patients with genotype CC. Contradictory findings have also been reported. Other genetic and clinical factors may also influence the response to warfarin.

· Class 3 rs9934438 GA

Patients with the AG genotype may have decreased time in the rapeutic range of INR (TTR) when treated with warfarin as compared to genotype GG. Other genetic and clinical factors may also influence the response to warfarin.

BCKDK	branched chain ketoacid dehydrogenase kinase
	- Clinical Annotations

Class 2A rs9923231 CT

Patients with genotype CT may require shorter time to the rapeutic INR when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the response to warfarin.

 \cdot Class 2A rs9923231 CT

Patients with the CT genotype may have increased risk of over-anticoagulation when treated with warfarin as compared with patients with genotype CC. Other genetic and clinical factors may also influence the toxicity to warfarin.

· Class 3 rs9923231 CT

Patients with the CT genotype may spent less time in INR therapeutic range (TTR) when treated with warfarin as compared with patients with genotype CC. Contradictory findings have also been reported. Other genetic and clinical factors may also influence the response to warfarin.

CYP4F2	cytochrome P450, family 4, subfamily F, polypeptide 2		

· Class 3 rs2108622 CC

Patients with the CC genotype may have increased international normalized ratio variability (INR-var) when treated with warfarin as compared to patients with genotype TT or CT in European-Americans after the warfarin dose-titration phase. Other genetic and clinical factors may also influence the response to warfarin.

UGT1A10	UDP glucuronosyltransferase 1 family, polypeptide A10
	- Clinical Annotations
	- Chinical Affilotations

· Class 3 rs887829 CT

Patients with the CT genotype and heart valve replacement may require a larger stable dose of warfarin compared to patients with the CC genotypes. Other clinical and genetic factors affect stable dose of warfarin.

	rpe and heart valve replacement may require a larger stable dose of warfaring the CC genotypes. Other clinical and genetic factors affect stable dose of
UGT1A4	UDP glucuronosyltransferase 1 family, polypeptide A4
	pe and heart valve replacement may require a larger stable dose of warfaring the CC genotypes. Other clinical and genetic factors affect stable dose of
UGT1A5	UDP glucuronosyltransferase 1 family, polypeptide A5
UGT1A6	UDP glucuronosyltransferase 1 family, polypeptide A6 ———————————————————————————————————
	ope and heart valve replacement may require a larger stable dose of warfaring the CC genotypes. Other clinical and genetic factors affect stable dose of
UGT1A7	UDP glucuronosyltransferase 1 family, polypeptide A7
· · ·	the CC genotypes. Other clinical and genetic factors affect stable dose of
UGT1A8	UDP glucuronosyltransferase 1 family, polypeptide A8

Class 3 rs887829 CT

Patients with the CT genotype and heart valve replacement may require a larger stable dose of warfarin compared to patients with the CC genotypes. Other clinical and genetic factors affect stable dose of warfarin.

UGT1A9	UDP glucuronosyltransferase 1 family, polypeptide A9
	- Clinical Annotations
Class 3 rs887829	CT

Patients with the CT genotype and heart valve replacement may require a larger stable dose of warfarin compared to patients with the CC genotypes. Other clinical and genetic factors affect stable dose of warfarin.

UGT1A1 UDP glucuronosyltransferase 1 family, polypeptide A1

— Clinical Annotations —

· Class 3 rs887829 CT

Patients with the CT genotype and heart valve replacement may require a larger stable dose of warfarin compared to patients with the CC genotypes. Other clinical and genetic factors affect stable dose of warfarin.

ZIDOVUDINE

ABCB1	ATP-binding cassette, sub-family B (MDR/TAP), member 1 $$
	- Clinical Annotations

· Class 3 rs1045642 GG

Patients with the GG genotype and HIV may have an increased risk of virological failure when receiving highly active antiretroviral therapy (HAART), as compared to patients with the AA genotype. Other genetic and clinical factors may also influence risk of virological failure on HAART.