



ANTIDEPRESSANTS

USE AS DIRECTED

desvenlafaxine (Pristiq®)
levomilnacipran (Fetzima®)
sertraline (Zoloft®)
vilazodone (Viibryd®)

MODERATE GENE-DRUG INTERACTION

citalopram (Celexa®)	1
escitalopram (Lexapro®)	1
selegiline (Emsam®)	1
trazodone (Desyrel®)	1

SIGNIFICANT GENE-DRUG INTERACTION

bupropion (Wellbutrin®)	1,6
doxepin (Sinequan®)	1,6
fluoxetine (Prozac®)	1,6
mirtazapine (Remeron®)	1,6
venlafaxine (Effexor®)	1,6
amitriptyline (Elavil®)	1,6,8
clomipramine (Anafranil®)	1,6,8
desipramine (Norpramin®)	1,6,8
duloxetine (Cymbalta®)	1,6,8
fluvoxamine (Luvox®)	1,6,8
imipramine (Tofranil®)	1,6,8
nortriptyline (Pamelor®)	1,6,8
vortioxetine (Trintellix®)	1,6,8
paroxetine (Paxil®)	1,4,6,8

CLINICAL CONSIDERATIONS

- 1: Serum level may be too high, lower doses may be required.
- 4: Genotype may impact drug mechanism of action and result in reduced efficacy.
- 6: Use of this drug may increase risk of side effects.
- 8: FDA label identifies a potential gene-drug interaction for this medication.

All psychotropic medications require clinical monitoring.







ANXIOLYTICS AND HYPNOTICS

USE AS DIRECTED

alprazolam (Xanax®)
buspirone (BuSpar®)
chlordiazepoxide (Librium®)
clonazepam (Klonopin®)
clorazepate (Tranxene®)
diazepam (Valium®)
eszopiclone (Lunesta®)
lorazepam (Ativan®)
oxazepam (Serax®)
temazepam (Restoril®)
zolpidem (Ambien®)

MODERATE GENE-DRUG INTERACTION

SIGNIFICANT GENE-DRUG INTERACTION propranolol (Inderal®) 1,6,8

CLINICAL CONSIDERATIONS

- 1: Serum level may be too high, lower doses may be required.
- 6: Use of this drug may increase risk of side effects.
- 8: FDA label identifies a potential gene-drug interaction for this medication.

All psychotropic medications require clinical monitoring.







ANTIPSYCHOTICS

USE AS DIRECTED

asenapine (Saphris®)
cariprazine (Vraylar®)
lurasidone (Latuda®)
paliperidone (Invega®)
thiothixene (Navane®)
ziprasidone (Geodon®)

MODERATE GENE-DRUG INTERACTION

fluphenazine (Prolixin®)	1
quetiapine (Seroquel®)	1
olanzapine (Zyprexa®)	3
clozapine (Clozaril®)	1,8
haloperidol (Haldol®)	1,8

SIGNIFICANT GENE-DRUG INTERACTION

chlorpromazine (Thorazine®)	1,6
aripiprazole (Abilify®)	1,6,8
brexpiprazole (Rexulti®)	1,6,8
iloperidone (Fanapt®)	1,6,8
perphenazine (Trilafon®)	1,6,8
risperidone (Risperdal®)	1,6,8
thioridazine (Mellaril®)	1,6,9

CLINICAL CONSIDERATIONS

- 1: Serum level may be too high, lower doses may be required.
- Difficult to predict dose adjustments due to conflicting variations in metabolism.
- 6: Use of this drug may increase risk of side effects.
- 8: FDA label identifies a potential gene-drug interaction for this medication.
- 9: Per FDA label, this medication is contraindicated for this genotype.

All psychotropic medications require clinical monitoring.







MOOD STABILIZERS

USE AS DIRECTED

carbamazepine (Tegretol®) oxcarbazepine (Trileptal®) valproic acid/divalproex (Depakote®)

MODERATE GENE-DRUG INTERACTION

SIGNIFICANT GENE-DRUG INTERACTION

lamotrigine (Lamictal®)

2

NO PROVEN GENETIC MARKERS						
gabapentin (Neurontin®)	10	topiramate (Topamax®)	10			
lithium (Eskalith®)	10					

CLINICAL CONSIDERATIONS

- 2: Serum level may be too low, higher doses may be required.
- 10: This medication does not have clinically proven genetic markers that allow it to be categorized.

All psychotropic medications require clinical monitoring.







PATIENT GENOTYPES AND PHENOTYPES



PHARMACODYNAMIC GENES



Lower Risk

SLC6A4 Normal Response

L/L

This patient is homozygous for the long promoter polymorphism of the serotonin transporter gene. The long promoter allele is reported to express normal levels of the serotonin transporter. The patient is predicted to have a normal response to selective serotonin reuptake inhibitors.

HTR2A

Increased Sensitivity

G/G

This individual is homozygous variant for the G allele of the -1438G>A polymorphism for the Serotonin Receptor Type 2A. They carry two copies of the G allele. This genotype has been associated with an increased risk of adverse drug reactions with certain selective serotonin reuptake inhibitors.

HLA-B*1502 Not Present

This patient does not carry the HLA-B*1502 allele or a closely related *15 allele. Absence of HLA-B*1502 and the closely related *15 alleles suggests lower risk of serious dermatologic reactions including toxic epidermal necrolysis (TEN) and Stevens-Johnson syndrome (SJS) when taking certain mood stabilizers.

HLA-A*3101 Lower Risk

A/A

This patient is homozygous for the A allele of the rs1061235 A>T polymorphism indicating absence of the HLA-A*3101 allele. This genotype suggests a lower risk of serious hypersensitivity reactions, including Stevens-Johnson syndrome (SJS), toxic epidermal necrolysis (TEN), maculopapular eruptions, and Drug Reaction with Eosinophilia and Systemic Symptoms when taking certain mood stabilizers.







PATIENT GENOTYPES AND PHENOTYPES



PHARMACOKINETIC GENES



CYP1A2

Extensive (Normal) Metabolizer

-163C>A - C/A, 5347C>T - C/T

This genotype is most consistent with the extensive (normal) metabolizer phenotype.

CYP2B6

Intermediate Metabolizer

*1/*6

CYP2B6*1 allele enzyme activity: Normal CYP2B6*6 allele enzyme activity: Reduced

This genotype is most consistent with the intermediate metabolizer phenotype. This patient may have reduced enzyme activity as compared to individuals with the normal phenotype.

CYP2C19

Extensive (Normal) Metabolizer

*1/*17

CYP2C19*1 allele enzyme activity: Normal CYP2C19*17 allele enzyme activity: Increased

This genotype is most consistent with the extensive (normal) metabolizer phenotype.

CYP2C9

Extensive (Normal) Metabolizer

*1/*1

CYP2C9*1 allele enzyme activity: Normal CYP2C9*1 allele enzyme activity: Normal

This genotype is most consistent with the extensive (normal) metabolizer phenotype.

CYP2D6 *4/*41 **Poor Metabolizer**

CYP2D6*4 allele enzyme activity: None CYP2D6*41 allele enzyme activity: Reduced

This genotype is most consistent with the poor metabolizer phenotype. This patient may have reduced enzyme activity as compared to individuals with the normal phenotype.

CYP3A4

Extensive (Normal) Metabolizer

*1/*1

CYP3A4*1 allele enzyme activity: Normal CYP3A4*1 allele enzyme activity: Normal

This genotype is most consistent with the extensive (normal) metabolizer phenotype.

UGT1A4

Ultrarapid Metabolizer

*1/*3

UGT1A4*1 allele enzyme activity: Normal UGT1A4*3 allele enzyme activity: Increased

This genotype is most consistent with the ultrarapid metabolizer phenotype. This patient may have increased enzyme activity as compared to individuals with the normal phenotype.

UGT2B15

Extensive (Normal) Metabolizer

*1/*2

UGT2B15*1 allele enzyme activity: Normal UGT2B15*2 allele enzyme activity: Reduced

This genotype is most consistent with the extensive (normal) metabolizer phenotype. The patient is expected to have normal enzyme activity.





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Questions? Call 855.891.9415 or email medinfo@assurexhealth.com

GENE-DRUG INTERACTIONS

		USE AS	DIRECTED					
	CYP1A2	CYP2B6	CYP2C19	CYP2C9	CYP3A4	CYP2D6	UGT1A4	UGT2B15
ANTIDEPRESSANTS								
desvenlafaxine (Pristiq®)			0		0			
levomilnacipran (Fetzima®)			0		0	•		
sertraline (Zoloft®)		•	0	0	0	•		
vilazodone (Viibryd®)			0		0	•		
ANXIOLYTICS AND HYPNOTICS								
alprazolam (Xanax®)					0			
buspirone (BuSpar®)					0	•		
chlordiazepoxide (Librium®)	0				0			0
clonazepam (Klonopin®)					0			
clorazepate (Tranxene®)	0				0			0
diazepam (Valium®)	0	•	0	0	0			0
eszopiclone (Lunesta®)				0	0			
lorazepam (Ativan®)								0
oxazepam (Serax®)								0
temazepam (Restoril®)		•		0	0			0
zolpidem (Ambien®)	0		0	0	0	•		
ANTIPSYCHOTICS								
asenapine (Saphris®)	0				0	•	•	
cariprazine (Vraylar®)					0	•		
lurasidone (Latuda®)					0			
paliperidone (Invega®)					0	•		
thiothixene (Navane®)	0							
ziprasidone (Geodon®)	0				0			
MOOD STABILIZERS								
carbamazepine (Tegretol®)		•			0			
oxcarbazepine (Trileptal®)								
valproic acid/divalproex (Depakote®)		•		0			•	
	MODE	RATE GENE	DRUG INTER	ACTION				
	CYP1A2	CYP2B6	CYP2C19	CYP2C9	CYP3A4	CYP2D6	UGT1A4	UGT2B15
ANTIDEPRESSANTS								
citalopram (Celexa®)			0		0	•		
escitalopram (Lexapro®)			0		0	•		
selegiline (Emsam®)	0	•	0		0			
trazodone (Desyrel®)	0				0	•		
ANTIPSYCHOTICS								
clozapine (Clozaril®)	0				0	•	•	

Variation was found in patient genotype that may impact medication response.



O - This gene is associated with medication response, but patient genotype is normal.



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GENE-DRUG INTERACTIONS

MODERATE GENE-DRUG INTERACTION								
	CYP1A2	CYP2B6	CYP2C19	CYP2C9	CYP3A4	CYP2D6	UGT1A4	UGT2B15
ANTIPSYCHOTICS								
fluphenazine (Prolixin®)	0		0	0	0	•		
haloperidol (Haldol®)	0				0	•	•	
olanzapine (Zyprexa®)	0				0	•	•	
quetiapine (Seroquel®)					0	•		
	SIGNIFI	CANT GENE	-DRUG INTER	RACTION				
	CYP1A2	CYP2B6	CYP2C19	CYP2C9	CYP3A4	CYP2D6	UGT1A4	UGT2B15
ANTIDEPRESSANTS								
amitriptyline (Elavil®)	0		0	0	0	•	•	
bupropion (Wellbutrin®)		•			0	•		
clomipramine (Anafranil®)	0		0		0	•		
desipramine (Norpramin®)						•		
doxepin (Sinequan®)	0		0	0	0	•	•	
duloxetine (Cymbalta®)	0					•		
fluoxetine (Prozac®)			0	0	0	•		
fluvoxamine (Luvox®)	0					•		
imipramine (Tofranil®)	0		0		0	•		
mirtazapine (Remeron®)	0			0	0	•		
nortriptyline (Pamelor®)						•		
paroxetine (Paxil®)					0	•		
venlafaxine (Effexor®)			0	0	0	•		
vortioxetine (Trintellix®)		•	0	0	0	•		
ANXIOLYTICS AND HYPNOTICS								
propranolol (Inderal®)	0					•		
ANTIPSYCHOTICS								
aripiprazole (Abilify®)					0	•		
brexpiprazole (Rexulti®)					0	•		
chlorpromazine (Thorazine®)	0				0	•		
iloperidone (Fanapt®)					0	•		
perphenazine (Trilafon®)	0		0		0	•		
risperidone (Risperdal®)					0	•		
thioridazine (Mellaril®)	0		0		0	•		
MOOD STABILIZERS								
lamotrigine (Lamictal®)								



Variation was found in patient genotype that may impact medication response.

O - This gene is associated with medication response, but patient genotype is normal.





TEST INFORMATION

The buccal swab sample was collected on 2/13/2020 and received in the laboratory on 2/14/2020. Genomic DNA was isolated and the relevant genomic regions were amplified by polymerase chain reaction (PCR). Analysis of CYP2D6 deletion and duplication, HLA-B*1502 and SLC6A4 was completed by electrophoresis of PCR products. Analysis of rs1061235 (indicating presence of the HLA-A*3101 allele or certain HLA-A*33 alleles), CYP1A2, CYP2B6, CYP2C19, CYP2C9, CYP2D6, CYP3A4, HTR2A, UGT1A4 and UGT2B15 was completed by using iPLEX MassARRAY® technology (Agena Bioscience). The following genetic variants may be detected in the assay: CYP1A2 -3860G>A (NG_008431.1:g.28338G>A), -2467T>delT (NM_000761.4:c.-1635delT), -739T>G (NM_000761.4:c.-10+103T>G), -729C>T (NM_000761.4:c.-10+113C>T), -163C>A (NM_000761.4:c.-9-154C>A), 125C>G (NM_000761.4:c.125C>G), 558C>A (NM_000761.4:c.558C>A), 2116G>A (NM_000761.4:c.1042G>A), 2473G>A (NM_000761.4:c.1130G>A), 2499A>T (NM_000761.4:c.1156A>T), 3497G>A (NM_000761.4:c.1217G>A), 3533G>A (NM 000761.4:c.1253+1G>A), 5090C>T (NM 000761.4:c.1291C>T), 5166G>A (NM 000761.4:c.1367G>A), 5347C>T (NM 000761.4:c.1548C>T); CYP2B6 *1, *4 (NM_000767.4:c.785A>G), *6 (NM_000767.4:c.516G>T; c.785A>G), *9 (NM_000767.4:c.516G>T); CYP2C19 *1, *2 (NM_000769.2:c.681G>A), *3 (NM_000769.2:c.636G>A), *4 (NM_000769.2:c.1A>G), *5 (NM_000769.2:c.1297C>T),*6 (NM_000769.2:c.395G>A), *7 (NM_000769.2:c.819+2T>A), *8 (NM 000769.2:c.358T>C), *17 (NM 000769.2:c.-806C>T); CYP2C9 *1, *2 (NM 000771.3:c.430C>T), *3 (NM 000771.3:c.1075A>C), *4 (NM_000771.3:c.1076T>C), *5 (NM_000771.3:c.1080C>G), *6 (NM_000771.3:c.817deIA); CYP2D6 *1, *2 (NM_000106.5:c.886C>T; c.1457G>C), *2A (NM_000106.5:c.-1584C>G; c.886C>T; c.1457G>C), *3 (NM_000106.5:c.775delA), *4 (NM_000106.5:c.506-1G>A; c.100C>T; c.1457G>C), *5 (CYP2D6 Deletion), *6 (NM_000106.5:c.454delT), *7 (NM_000106.5:c.971A>C), *8 (NM_000106.5:c.505G>T; c.886C>T; c.1457G>C), *9 (NM_000106.5:c.841_843delAAG), *10 (NM_000106.5:c.100C>T; c.1457G>C), *11, *12 (NM_000106.5:c.124G>A; c.886C>T; c.1457G>C), *14 (NM_000106.5:c.505G>A; c.886C>T; c.1457G>C), *15, *17 (NM_000106.5:c.320C>T; c.886C>T; c.1457G>C), *41 (NM_000106.5:c.985+39G>A; c.886C>T; c.1457G>C), gene duplication; CYP3A4 *1, *13 (NM_017460.5:c.1247C>T), *15A (NM_017460.5:c.485G>A), *22 (NM_017460.5:c.522-191C>T); HLA-B*1502; rs1061235 (NM_002116.7:c.*66A>T); HTR2A -1438G>A (NM_000621.4:c.-998G>A); SLC6A4 L, S; UGT1A4 *1, *3 (NM_007120.2:c.142T>G); UGT2B15 *1, *2 (NM_001076.3:c.253G>T). The following rare genetic variants have not been observed by the Assurex Health, Inc. laboratory: CYP1A2 125C>G, 558C>A; CYP2C19 *7.

This test was developed and its performance characteristics determined by Assurex Health. It has not been cleared or approved by the U.S. Food and Drug Administration.

These interpretations are based upon data available in scientific literature and prescribing information for the relevant drugs. Interpretations are, in some instances, based on data regarding the pharmacokinetic, pharmacodynamic and pharmacogenomics properties of a drug derived from non-clinical studies (e.g. *in vitro* studies). Findings from studies performed in a non-clinical setting or clinical studies involving healthy subjects are not necessarily indicative of clinical performance in a particular patient.

This report was reviewed and verified on 2/18/2020 by:

Disclaimer of Liability

The information contained in this report is provided as a service and does not constitute medical advice. At the time of report generation this information is believed to be current and is based upon published research; however, research data evolves and amendments to the prescribing information of the drugs listed will change over time. While this report is believed to be accurate and complete as of the date issued, THE DATA IS PROVIDED "AS IS", WITHOUT WARRANTIES OF ANY KIND, EXPRESS OR IMPLIED, INCLUDING WITHOUT LIMITATION, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS FOR A PARTICULAR PURPOSE. As medical advice must be tailored to the specific circumstances of each case, the treating healthcare provider has ultimate responsibility for all treatment decisions made with regard to a patient including any made on the basis of a patient's genotype.

GeneSight Psychotropic is covered by U.S. Patent No. 9,111,028

Genetic testing was completed by a CLIA and CAP accredited laboratory in the United States located at:

6000 Mason-Montgomery Road

Mason, OH 45040

Customer Service

Please contact 855.891.9415 or medinfo@assurexhealth.com for assistance with report interpretation. For all other inquires please contact 866.757.9204 or support@assurexhealth.com.

GeneSight Psychotropic Test Version: 3.0.2





GeneSight® Order Medical Necessity Documentation

Patient Nar	me	Patient Date of Birth	Order Number
GeneSi	ght Psychotropic		
ICD-10 Co	de(s)		
F41.9	Anxiety disorder, unspecified		
F43.9	Reaction to severe stress, unspecified		
F33.1	Major depressive disorder, recurrent, moderate		
	to DSM-5 criteria, does the patient suffer from ressive Disorder (MDD)?	_	e the patient's depressive symptoms: n excess of the five required to make the diagnoses and the principle impairment
Depress Diminisi Significa Insomni Psychol Fatigue Feelings Diminisi	sed mood hed interest in activities ant weight gain/loss ia or hypersomnia motor agitation/retardation	✓ Moderate: symptoms or fund	tional impairment are between 'mild' and 'severe' I the symptoms markedly interfere with functioning; can
Medical No	ecessity ntemplating an alteration* in a neuropsychiatric medication treatment? No	Medication that the patient has Prozac® (fluoxetine)	failed or is currently failing:
Has the pa	tient failed or are they currently failing at least one niatric medication?		
✓ Yes	□No		
Additional	Information (optional)		
None prov	ided.		
The undersi consent for	include medication elimination, switching, augmentation, and/or dose adjustments gned attests that he/she is licensed to order the selected test(s). I acknowledge tha genetic testing from the patient or his/her legal authorized representative. I attest t it and treatment decisions for the above referenced patient and agree to provide an	hat the selected genetic test(s) are medical	ly necessary and that these results will be used in the medical
Insurers req in your patie	uire that you maintain documentation supporting the medical necessity for GeneSigent's medical record. In addition to the information provided for this order, please e	ght tests in the patient's medical record. Ple nsure that the patient's medical record is u	ease verify that the order information above is correct and include p-to-date with DSM-5-based diagnostic information.
Healthcare	Provider Information		
Name ₋			
Signatu	re		Date



GeneSight® Order Medical Necessity Documentation

Patient Na	me	Patient Date of Birth	Order Number				
GeneSight Analgesic ICD-10 Code(s)		Medical Necessity Patient has failed or is currently failing at least one analgesic medication and I am contemplating an alteration in an analgesic medication treatment.					
		adverse drug events. Patient suspected of abuse	ations for his/her condition which increases the risk for and/or diversion with current medication regimen. atient with no prior pharmacological treatment history				
GeneSi	ight ADHD ide(s)	I am contemplating an alter. Patient is on multiple medic adverse drug events. Patient suspected of abuse	ently failing at least one neuropsychiatric medication and ation in a neuropsychiatric medication treatment. ations for his/her condition which increases the risk for and/or diversion with current medication regimen. atient with no prior pharmacological treatment history				
GeneSi	ight MTHFR						
ICD-10 Co F41.9	de(s) Anxiety disorder, unspecified	Medical Necessity I am considering folate supp Patient has low serum folate					
F43.9	Reaction to severe stress, unspecified	✓ Other TBD					

Ordered by:





NORMAL FOLIC ACID CONVERSION

REDUCED FOLIC ACID CONVERSION



SIGNIFICANTLY REDUCED **FOLIC ACID CONVERSION**

Note: Serum levels of folate may be too low. Folate supplementation or higher daily intake of folic acid may be required.

PATIENT GENOTYPE AND PHENOTYPE

MTHFR

Intermediate Activity

C/T

This individual is heterozygous for the C677T polymorphism in the MTHFR gene. This genotype is associated with reduced folic acid metabolism, moderately decreased serum folate levels, and moderately increased homocysteine levels.

TEST INFORMATION

The buccal swab sample was collected on 2/13/2020 and received in the laboratory on 2/14/2020. Genomic DNA was isolated and the relevant genomic regions were amplified by polymerase chain reaction (PCR). Analysis of MTHFR was completed by using iPLEX MassARRAY® technology (Agena Bioscience). The following genetic variant may be detected in the assay: MTHFR 677C>T (NM_005957.4:c.665C>T).

This test was developed and its performance characteristics determined by Assurex Health. It has not been cleared or approved by the U.S. Food and Drug Administration.

These interpretations are based upon data available in scientific literature and prescribing information for the relevant drugs. Interpretations are, in some instances, based on data regarding the pharmacokinetic, pharmacodynamic and pharmacogenomics properties of a drug derived from non-clinical studies (e.g. in vitro studies). Findings from studies performed in a non-clinical setting or clinical studies involving healthy subjects are not necessarily indicative of clinical performance in a particular patient.

This report was reviewed and verified on 2/18/2020 by:

Disclaimer of Liability

The information contained in this report is provided as a service and does not constitute medical advice. At the time of report generation this information is believed to be current and is based upon published research; however, research data evolves and amendments to the prescribing information of the drugs listed will change over time. While this report is believed to be accurate and complete as of the date issued, THE DATA IS PROVIDED "AS IS", WITHOUT WARRANTIES OF ANY KIND, EXPRESS OR IMPLIED, INCLUDING WITHOUT LIMITATION, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS FOR A PARTICULAR PURPOSE. As medical advice must be tailored to the specific circumstances of each case, the treating healthcare provider has ultimate responsibility for all treatment decisions made with regard to a patient including any made on the basis of a patient's genotype.

Genetic testing was completed by a CLIA and CAP accredited laboratory in the United States located at:

6000 Mason-Montgomery Road

Mason, OH 45040

Customer Service

Please contact 855.891.9415 or medinfo@assurexhealth.com for assistance with report interpretation. For all other inquires please contact 866.757.9204 or support@assurexhealth.com.

GeneSight MTHFR Test Version: 1.0

