CONFIDENTIAL DOB: 1000-01-01 Date of report: 2023-11-10

PHARMACOGENOMIC REPORT



Psychiatry and ADHD

YOUR RESULTS ARE CONFIDENTIAL. As per the Genetic Non-Discrimination Act (S-201), no person, company or institution, including insurers and employers, can force you to share this report.

DO NOT MAKE ANY CHANGES TO YOUR CURRENT MEDICATION(S) WITHOUT TALKING TO YOUR DOCTOR FIRST. While genetics is important, other factors also contribute to how you react to medications. The final choice of medication used will be based on your health care provider's professional judgement and may be different than what is recommended in this report. This test does not determine your risk of any health problem. It only evaluates select portions of your DNA that help predict how you may react to the medications covered.

ADMINISTRATIVE DATA

 Patient Name
 Ordering Clinician
 Sample ID: T EST 19012314

 First Last Name
 Dr Grey
 Sample Type: Saliva

irst Last Name Dr Grey Sample Type: saliva

Gender: Male Patient Address Date ordered: 2023-11-10

Date of birth: 1000-01-01 N/A Date of sample reception: 2023-11-10

Phone Number: Date of report: 2023-11-10

Email: N/A

Clinical Support

Email: <u>genetics@biron.com</u> Phone: 1-866-923-9222 Fax: (514) 317-2241

VARIANT(S) OF VERY IMPORTANT PHARMACOGENES (VIPS)

CYP1A2 IND, CYP2C9 IM, CYP2D6 UM, POR IA, UGT1A1 IM, UGT2B7 variable, UGT2B15 IM.

Very Important Pharmacogenes (VIPs) - visit pharmgkb.org/vips for more information.

Abbreviations - NM: Normal Metaboliser, IM: Intermediate Metaboliser, PM: Poor Metaboliser, RM: Rapid Metaboliser, UM: Ultrarapid Metaboliser, Ind: Inducible Metaboliser, NA: Normal Activity, IA: Intermediate Activity, PA: Poor Activity.

CAUTIONARY INFORMATION - MEDICATIONS TO AVOID OR USE WITH CAUTION

Medication	Identified risk	Recommendation
Codeine	Potential of serious toxicity due to increased formation of morphine (CYP2D6 UM).	Avoid using codeine. ¹
Tramadol Ultram®	Potential of serious toxicity due to increased formation of active metabolite (CYP2D6 UM).	Avoid using tramadol; if an alternative is not possible, consider reducing the dose by 60% and be alert to side effects. ¹

PGx RECOMMENDATIONS - PSYCHIATRY

Dose increase may be required.

All of the tested variants associate with an increased likelihood of a better response, compared to non-carriers.

Dose reduction may be required.

> 50% of tested variants are associated with an increased likelihood of a poorer response and/or adverse drug reactions, compared to non-carriers.

Exposure is difficult to predict, insufficient data to calculate dose adjustments.

Medication not recommended by peer-reviewed guidelines.

	Genetic Associations Identified					
Medications	Exposure	Efficacy	Adverse Drug Reactions			
Antidepressants						
Selective seroton	in reuptake inhibitors (SSRIs)					
Citalopram (Celexa®)	CYP2C19 NM, CYP3A4 NM, ABCB1 - Initiate with recommended dose; a low dose may be adequate.	3/6 variants: increased likelihood of a poorer response (FKBP5, HTR2A, BDNF).	0/3 variants: no increased risk of gastrointestinal or SSRI-induced sexual side effects.			
Escitalopram (Cipralex®)	CYP2C19 NM, CYP3A4 NM, ABCB1 - Initiate with recommended dose; a low dose may be adequate.	3/6 variants: increased likelihood of a poorer response (FKBP5, HTR2A, BDNF).	0/3 variants: no increased risk of gastrointestinal or SSRI-induced sexual side effects.			
F luoxetine (Prozac®)	CYP2D6 UM - Initiate with recommended dose but monitor more closely for response; may require a higher dose.	3/6 variants: increased likelihood of a poorer response (FKBP5, HTR2A, BDNF).	no marker available			
Sertraline (Zoloft®)	CYP2B6 NM, CYP2C19 NM, ABCB1 - Initiate with recommended dose; a low dose may be adequate.	2/5 variants: increased likelihood of a poorer response (BDNF, FKBP5).	0/3 variants: no increased risk of gastrointestinal or SSRI-induced sexual side effects.			
Fluvoxamine (Luvox®)	CYP2D6 UM - Initiate with recommended starting dose but may require a higher dose. If patient does not respond to recommended maintenance dosing, consider an alternative drug not predominantly metabolized by CYP2D6.	3/6 variants: increased likelihood of a poorer response (FKBP5, HTR2A, BDNF).	no marker available			
Paroxetine (Paxil®)	CYP2D6 UM - Consider an alternative drug not predominantly metabolized by CYP2D6 due to potential lack of efficacy. ²	3/6 variants: increased likelihood of a poorer response (FKBP5, HTR2A, BDNF).	no marker available			
Serotonin-norep	inephrine reuptake inhibitors (SNRIs)					
Desvenlafaxine (Pristiq®)	UGT1A1 IM, UGT2B15 IM - Initiate with recommended dose; a low dose may be adequate.	1/2 variants: increased likelihood of a poorer response (FKBP5).	no marker available			
_evomilnacipran [Fetzima®)	CYP3A4 NM, ABCB1 - Initiate with recommended dose; a low dose may be adequate.	1/2 variants: increased likelihood of a poorer response (FKBP5).	no marker available			
Duloxetine (Cymbalta®)	CYP1A2 Ind, CYP2D6 UM - Initiate therapy with recommended starting dose but may require a higher dose, especially with CYP1A2 inducers, such as smoke.	1/3 variants: increased likelihood of a poorer response (FKBP5).	no marker available			
/enlafaxine Effexor XR®)	CYP2D6 UM - Consider a dose increase up to 150% of the standard dose. If dose adjustment does not result in efficacy, consider an alternative drug not predominantly metabolized by CYP2D6. ³	3/4 variants: increased likelihood of a poorer response (COMT, FKBP5, SLC6A2).	no marker available			

inducers, such as smoke.

PGx RECOMMENDATIONS - COMPLEMENTARY TREATMENTS

Dose increase may be required.

All of the tested variants associate with an increased likelihood of a better response, compared to non-carriers.

Dose reduction may be required.

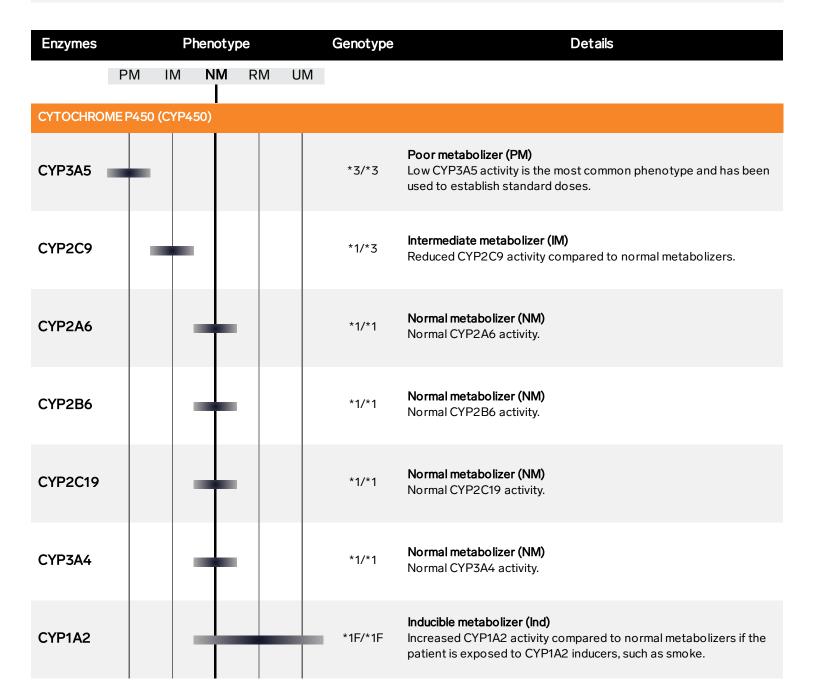
> 50% of tested variants are associated with an increased likelihood of a poorer response and/or adverse drug reactions, compared to non-carriers.

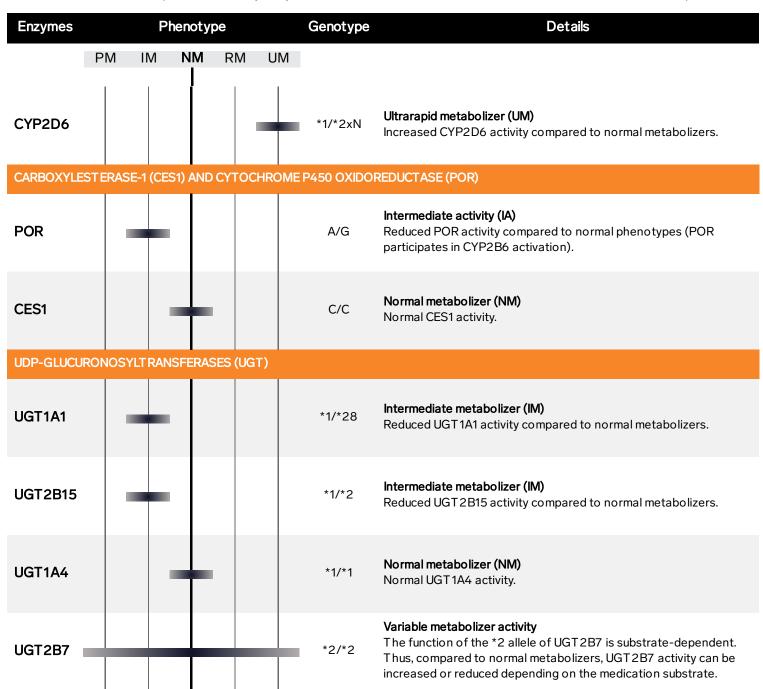
Exposure is difficult to predict, insufficient data to calculate dose adjustments.

Medication not recommended by peer-reviewed guidelines.

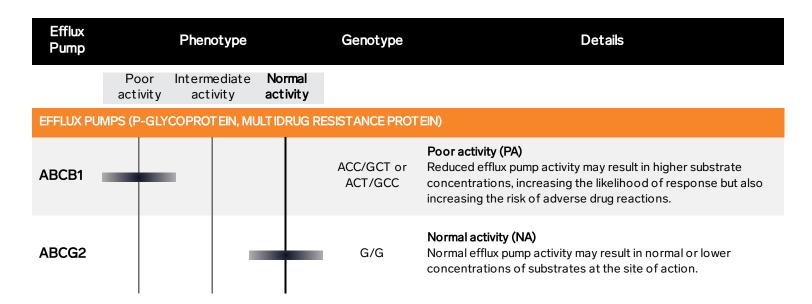
	Genetic Associations Identified					
Medications	Exposure	Efficacy	Adverse Drug Reactions			
Antiemetics						
Dimenhydrinate (Gravol®)	no marker available	no marker available	no marker available			
Granisetron (Kytril®)	CYP3A4 NM, CYP3A5 PM - Normal exposure	no marker available	no marker available			
Palonosetron (Aloxi®)	CYP2D6 UM - Initiate with recommended dose but may require a higher dose.	no marker available	no marker available			
Ondansetron (Zofran®)	CYP2D6 UM - Consider an alternative drug not predominantly metabolized by CYP2D6 (i.e. granisetron) due to decreased response. ⁸	no marker available	no marker available			
Proton pump inhib	oitors (PPI)					
Esomeprazole (Nexium®)	no marker available	no marker available	no marker available			
Dexlansoprazole (Dexilant®)	CYP2C19 NM - Initiate therapy with standard dose but consider increasing the dose by 50-100% for the treatment of H. pylori infection and erosive esophagitis. Daily dose may be given in divided doses. Monitor for efficacy.	no marker available	no marker available			
Lansoprazole (Prevacid®)	CYP2C19 NM - Initiate therapy with standard dose but consider increasing the dose by 50-100% for the treatment of H. pylori infection and erosive esophagitis. Daily dose may be given in divided doses. Monitor for efficacy.	no marker available	no marker available			
Omeprazole (Losec®)	CYP2C19 NM - Initiate therapy with standard dose but consider increasing the dose by 50-100% for the treatment of H. pylori infection and erosive esophagitis. Daily dose may be given in divided doses. Monitor for efficacy.	no marker available	no marker available			
Pantoprazole (Pantoloc®)	CYP2C19 NM - Initiate therapy with standard dose but consider increasing the dose by 50-100% for the treatment of H. pylori infection and erosive esophagitis. Daily dose may be given in divided doses. Monitor for efficacy.	no marker available	no marker available			

METABOLISM	The level of activity	The level of activity of the enzymes listed below influence your exposure to active metabolite(s).			
	CYP450	POR	CES	UGT	
Variants detected / tested	3/8	1/1	0/1	3/4	





DISTRIBUTION	The level of activity of the efflux pumps listed below influence your exposure to medications at the site of action (e.g., central nervous system).	
	ABCB1	ABCG2
Variant(s) detected / tested	1/1	0/1



The variants listed below have been associated with a higher likelihood of a poorer response to medications, compared to non-carriers.

Poorer response more likely

Variant(s) detected / tested

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15/23

Poorer response more likely				
Gene	Reference SNP cluster ID	Variant detected	Associated medication(s)	
ADRA2A	rs1800544	yes	methylphenidate	
ANKK1	rs1800497	yes	aripiprazole, risperidone	
BDNF	rs6265	yes	citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine	
CACNG2	rs2283967	yes	lithium	
COMT	rs4680	yes	venlafaxine (AA), methylphenidate (GG)	
FKBP5	rs4713916	yes	citalopram, desvenlafaxine, duloxetine, escitalopram, levomilnacipran, sertraline, fluoxetine, fluoxamine, mirtazapine, paroxetine, venlafaxine	
GNB3	rs5443	yes	clonidine	
GRIK1	rs2832407	yes	topiramate	
HTR2A	rs2770296	yes	bupropion	
HTR2A	rs6311	yes	citalopram, escitalopram, fluoxetine, fluvoxamine, paroxetine	
IncRNA	rs74795342	yes	lithium	
IncRNA	rs75222709	yes	lithium	
SLC6A2	rs2242446	yes	venlafaxine	
SLC6A2	rs28386840	yes	methylphenidate	
TPH2	rs1487278	yes	amitriptyline, clomipramine, desipramine, doxepin, imipramine, mirtazapine, nortriptyline, trimipramine	
ANKK1	rs1800497	no	bupropion	
BDNF	rs6265	no	esketamine, ketamine	
DRD3	rs963468	no	duloxetine	

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Gene	Reference SNP cluster ID	Variant detected	Associated medication(s)
GRIK4	rs1954787	no	citalopram, desvenlafaxine, duloxetine, escitalopram, fluoxetine, fluvoxamine, levomilnacipran, paroxetine, sertraline, venlafaxine
HTR7	rs7905446	no	citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine
SLC6A2	rs5569	no	methylphenidate
SLC6A4	5-HTTLPR	no	citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine
ТН	rs2070762	no	methylphenidate

PGx ASSOCIATIONS - ADVERSE DRUG REACTIONS

The variants listed below have been associated with a higher likelihood of specific adverse drug reactions, compared to non-carriers.

Increased risk of adverse drug reactions

Variant(s) detected / tested

4/13

Increased risk of adverse drug reactions				
Gene	Reference SNP cluster ID	Variant detected	Associated medication(s)	
ANKK1	rs1800497	yes	weight gain: valproic acid	
INSIG2	rs17047764	yes	antipsychotic-induced weight gain: clozapine, olanzapine, risperidone	
MC4R	rs489693	yes	antipsychotic-induced weight gain: clozapine, olanzapine, quetiapine, risperidone	
MC4R	rs17782313	yes	antipsychotic-induced weight gain: quetiapine	
ADRA2A	rs1800544	no	antipsychotic-induced weight gain: clozapine, olanzapine, risperidone	
DRD2	rs6275	no	antipsychotic-induced weight gain: clozapine, olanzapine, risperidone	
GNB3	rs5443	no	antipsychotic-induced weight gain: clozapine, olanzapine, risperidone	
HLA-A	*31:01	no	cutaneous adverse reactions: carbamazepine	
HLA-B	*15:02	no	cutaneous adverse reactions: carbamazepine, oxcarbazepine, phenytoin	
HTR2A	rs6311	no	gastrointestinal side effects (nausea, vomiting): citalopram, escitalopram, sertraline	
HTR2C	rs3813929	no	antipsychotic-induced weight gain: clozapine, olanzapine, risperidone	
SLC6A2	rs28386840	no	hypertension: esketamine	
SLC6A5	rs2298826	no	extrapyramidal side effects: haloperidol	

ANALYTICAL RESULTS

The following analytical results were used to generate the pharmacogenomic interpretations found in this report. Technical limitations inherent with the methods used to produce these results may hinder the attribution of a definitive phenotype (see "TEST METHODOLOGY AND LIMITATIONS").

Genes	Variant Detail	ls (GRCH38.p12)	Result
ABCB1	rs1045642	chr7:87509329	A G
	rs2032582	chr7:87531302	C C
	rs2032583	chr7:87531245	C T
ABCG2	rs2231142	chr4:88131171	G G
ADRA2A	rs1800544	chr10:111076745	C C
ANKK1	rs1800497	chr11:113400106	GIG
BDNF	rs6265	chr11:27658369	CIC
CACNG2	rs2283967	chr22:36567486	CIT
CES1	rs71647871	chr16:55823658	CIC
CNR1	rs806380	chr6:88154934	AlG
COMT	rs4680	chr22:19963748	AIG
CYP1A2	rs762551	chr15:74749576	AlA
CIFIAZ	rs2069514	chr15:74745879	GG
CYP2A6	rs1801272	chr19:40848628	AlA
CITZAO	rs5031017	chr19:40843845	GIG
	rs28399433	chr19:40850474	AIA
CYP2B6	rs2279343	chr19:41009358	AlA
011 250	rs3745274	chr19:41006936	GIG
	rs28399499	chr19:41012316	TIT
CYP2C9	rs1057910	chr10:94981296	CIA
	rs1799853	chr10:94942290	cic
	rs7900194	chr10:94942309	G G
	rs9332131	chr10:94949282-94949283	AļA
	rs9332239	chr10:94989020	CIC
	rs28371685	chr10:94981224	CIC
	rs28371686	chr10:94981301 chr10:94941958	C C
	rs72558187 rs72558190	chr10:94947782	T T C C
CYP2C19	rs4244285	chr10:94781859	GIG
CITZCIS	rs4986893	chr10:94781653	G G
	rs6413438	chr10:94781858	CIC
	rs12248560	chr10:94761900	cic
	rs12769205	chr10:94775367	AA
	rs17884712	chr10:94775489	GIG
	rs28399504	chr10:94762706	AlA
	rs41291556	chr10:94775416	T T
	rs56337013	chr10:94852738	CIC
	rs72552267 rs72558186	chr10:94775453 chr10:94781999	G G T T
CYP2D6	rs16947	chr22:42127941	GIA
CIPZDO	rs1065852	chr22:42130692	GIG
	rs1135840	chr22:42126611	CIG
	rs3892097	chr22:42128945	CIC
	rs5030655	chr22:42129084	A A
	rs5030656	chr22:42128174-42128178	ΑİΑ
	rs5030862	chr22:42130668	CIC
	rs5030865	chr22:42129033	CIC
	rs5030867	chr22:42127856	T T
	rs28371725	chr22:42127803	C C
	rs28371706 rs35742686	chr22:42129770 chr22:42128242	G G T T
	rs59421388	chr22:42127608	CIC
	rs72549354	chr22:42128815-42128817	
	rs201377835	chr22:42129910	CIC
	Gene Deletion	n/a	Not Detecte
	Gene Duplication	n/a	Detected

Genes	Variant Deta	nils (GRCH38.p12)	Result
CYP3A4	rs4986907	chr7:99769804	C C
	rs35599367	chr7:99768693	GIG
	rs55785340	chr7:99768360	A A
	rs67666821 rs72552799	chr7:99758184-99758188 chr7:99770165	D D C C
CYP3A5	rs776746	chr7:99672916	CIC
077 0710	rs10264272	chr7:99665212	CIC
	rs28365083	chr7:99652613	GİG
	rs28383479	chr7:99660516	CIC
	rs41303343	chr7:99652771	DID
	rs55817950	chr7:99676198	GIG
	rs56411402	chr7:99665237	T T
DRD2	rs6275	chr11:113412755	G G
DRD3	rs963468	chr3:114144040	AIG
FAAH	rs324420	chr1:46405089	C C
FKBP5	rs4713916	chr6:35702206	G G
GNB3	rs5443	chr12:6845711	C C
GRIK1	rs2832407	chr21:29595188	A C
GRIK4	rs1954787	chr11:120792654	C C
HLA- A*31:01	rs1061235	chr6:29945521	A A
HLA- B*15:02	rs144012689	chr6:31355003	T T
HTR2A	rs6311	chr13:46897343	C T
	rs6313	chr13:46895805	AIG
	rs2770296	chr13:46866425	T T
HTR2C	rs3813929	chrX:114584047	C T
HTR7	rs7905446	chr10:90859404	G G
INSIG2	rs17047764	chr2:118111006	C C
long non-	rs74795342	chr21:18954018	GIG
coding (Inc) RNA	rs75222709	chr21:18955109	T T
MC4R	rs489693	chr18:60215554	AA
	rs17782313	chr18:60183864	CIC
MTHFR	rs1801131	chr1:11794419	TIT
000144	rs1801133	chr1:11796321	GIG
OPRM1	rs1799971	chr6:154039662	AIA
POR	rs2868177	chr7:75960585	AIG
SLC6A2	rs5569	chr16:55697923	GIG
	rs2242446 rs28386840	chr16:55656513 chr16:55652906	C T A A
SLC6A4	5-HTTLPR	chr17:30190154-30240133	SIL
SLC6A5	rs2298826	chr11:20638211	GIG
TH	rs2070762	chr11:2165105	A A
TPH2	rs1487278	chr12:72007071	TIT
UGT1A1	rs4148323	chr2:233760498	G G
JGTIAI	rs34815109	chr2:233760234-233760248	6 7
UGT1A4	rs2011425	chr2:233718962	TIT
UGT2B7	rs7439366	chr4:69098620	C C
UGT2B15	rs1902023	chr4:68670366	CIA

TEST METHODOLOGY AND LIMITATIONS

The Biron pharmacogenomic test for psychiatry and pain management is a MALDI-TOF-based single nucleotide primer extension genotyping test; laboratory developed and validated test (LDT), not approved by Health Canada. Nucleic acid amplification techniques may be subject to general interference by factors such as reaction inhibitors and low quality or quantity of extracted DNA. Factors influencing the amount and quality of extracted DNA include but are not limited to patient oral hygiene, collection technique and presence of dietary or microbial source of nucleic acids and nuclease. When present, these interferents typically yield no result rather than an inaccurate one. Risk of suboptimal DNA quantity or quality is significantly reduced by automated DNA extraction which uses chemistry without PCR inhibitors (magnetic beads) and systematic dilution, quantitation and normalization of DNA before nucleic acid amplification. Very infrequent variants or polymorphisms occurring in primer-binding regions may also affect testing and could produce an erroneous result or assay failure. The test does not detect all known and unknown variations in the genes tested, nor does absence of a detectable variant (typically reported as *1 for metabolic enzymes) rule out the presence of other, non-detected variants. The test detects CYP2D6 deletion and duplication but cannot differentiate duplication in the presence of deletion. CYP2D6 deletion and duplication assays can translate into equivocal phenotype results where a range of enzyme activity level must be reported. Test results and clinical interpretation may be inaccurate for individuals who have undergone or are receiving non-autologous blood transfusions, tissue, and/or organ transplant therapies.

DISCLAIMER

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Biron Health Group developed this pharmacogenomic report. This test does not diagnose any disorder, condition or disease. The interpretations and recommendations provided in this report are intended as a clinical support tool (DST) to be used solely by a healthcare professional. Treatment decisions for the patient remain the sole responsibility of the treating healthcare provider. The interpretations of the results provided by this report were determined by Biron's data curation protocol, which were established as per the current available scientific evidence available at the time this report version was created. As more evidence becomes available in the future, these interpretations may change. Some variants tested may not be used to provide report interpretations due to a lack of clear gene-drug association as determined by Biron's data curation protocol. The presence of a notification within the "Exposure", "Efficacy" or "Adverse Drug Reactions" categories for a given drug indicates that an associated genetic variant was detected. The lack of a notification within these categories for a given drug does not eliminate the requirement for dose adjustments for optimal dosage, does not guarantee effective drug therapy and does not eliminate the risks of adverse drug reactions. Commercial names are indicated as examples and do not consist an exhaustive list.

REFERENCES

For the full list of references, contact pgxinfo@biron.com

Reference(s) cited in this report:

- 1. Crews KR, et al. Clinical Pharmacogenetics Implementation Consortium Guideline for CYP2D6, OPRM1, and COMT Genotypes and Select Opioid Therapy. Clin Pharmacol Ther (2021).
- 2. Bousman CA, et al. Clinical Pharmacogenetics Implementation Consortium (CPIC) Guideline for CYP2D6, CYP2C19, CYP2B6, SLC6A4, and HTR2A Genotypes and Serotonin Reuptake Inhibitor Antidepressants. Clin Pharmacol Ther (2023).
- Dutch Pharmacogenetics Working Group (DPWG) 2021 update (https://api.pharmgkb.org/v1/download/file/attachment/DPWG_May_2021.pdf)
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