Interpretation Guide



Pharmacogenomic (PGx) Report - for your healthcare provider

The following PGx report is a clinical decision support tool based on individual genetic results. It contributes to a better understanding and prediction of medication response and tolerability. This test does not predict the risk of any health problem. Since response to medications is multifactorial, clinical judgment supersedes any recommendations provided.

The report notifies you if the patient carries any genetic variant that can alter the following pharmacological parameters:

- pharmacokinetics: overall **exposure** to a medication depending on metabolic and efflux pump function;
- pharmacodynamics: the potential efficacy of a drug and whether the patient is predisposed to certain atypical effects.

These results do not change with age, but their interpretation can evolve as new data becomes available. Therefore, the Biron PGx reports are updated periodically. These results can also be useful for other medications, not covered by the report.

How to use pharmacogenomic recommendations

- 1. Only medications relevant for your patient need to be consulted.
- 2. Use the **Exposure** column to adjust doses for adequate plasma concentrations.
- 3. Use the Efficacy and Risk of atypical effect columns to choose the most compatible medication.

The **Exposure**, **Efficacy** and **Risk of atypical effect** columns are interpreted independently from each other. Medications are ordered by class with the most compatible options listed first within each class.

Efficacy

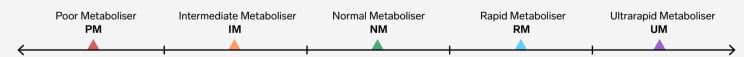
A higher dose may be required to achieve adequate When choosing between multiple clinically appropriate medications, you may give plasma concentrations. preference to a medication in which a lower number of variants have been identified (e.g., 2/2 is better than 4/6), in terms of their association with an increased likelihood of a A lower dose may be required to achieve adequate plasma poorer response or an atypical effect. concentrations. Signifies the presence of a high-impact Signifies the presence of variants Several metabolic pathways are involved, but their gene variant, which increases the associated with an increased risk of particular side effects, compared to capacities are opposed (e.g., PM and UM). Thus, a probability of a poorer response. calculation of dose adjustments is not possible based on non-carriers. current data and closer monitoring is recommended. Signifies that all of the tested variants predict an increased likelihood of a better X Medication not recommended by Drug not recommended by peer-reviewed guidelines due response, compared to non-carriers. This peer-reviewed guidelines due to a to a risk of toxicity or lack of efficacy. risk of severe side effects. medication may be a good option.

The notification "Genetic influence not available" signifies that there is currently no available data allowing for a genetically-based prediction of medication effect.

Nomenclature for enzyme phenotypes

(e.g., cytochrome P450s or CYP)

Exposure



NM is generally used to establish standard doses. This dose may be too high for **PM/IM** or too low for **RM/UM**, warranting a dose adjustments or the consideration of an alternative agent. For a pro-drug (e.g., clopidogrel, tramadol), phenotype variability will have the opposite effect.

<u>Inducible Metaboliser</u> (Ind) - Specific for CYP1A2, which can have increased function in the presence of an inducer, such as tobacco smoke, comparable to RM/UM.

Phone: 1-855-943-6379 Email: genetics@biron.com biron.com/en/genetics/pharmacogenomics



Risk of atypical effect

PHARMACOGENOMIC REPORT



Psychiatry and ADHD

To download the latest version, go here: secur.biron.com/login.

YOUR RESULT'S 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) WIT HOUT 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. For more information, visit biron.com/pgxtest.

| ADMINISTRATIVE DATA | | |
|--|--|--|
| Patient Name Test-Firstname Test-Lastname | Ordering Clinician Meredith Grey | Sample ID: BIO2107270327 Sample Type: test |
| Gender: Female Date of birth: 1999-01-01 Phone Number: (418) 999-9999 Email: test-sample.BIO2107270327@biron.local | Patient Address 1212 some street Ste-foy, Québec G2J4M5 | Date ordered: 2022-08-21 Date of sample reception: 2025-01-14 Date of report: 2025-01-14 |
| Clinical Support Email: genetique@biron.com | Phone: 1-855-943-6379 | Fax: (514) 317-2241 |

VARIANT(S) OF VERY IMPORTANT PHARMACOGENES (VIPS)

CYP1A2 IND, CYP2C9 IM, CYP2C19 IM, POR IA, CES1 IM, UGT1A1 IM, UGT1A4 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 If patient is carbamazepine-naïve, avoid carbamazepine. If the patient has Risk of hypersensitivity reaction due to the presence of previously used carbamazepine consistently for longer than three months Carbamazepine without incidence of cutaneous adverse reactions, cautiously consider using the HLA-B*15:02 allele. Tegretol® carbamazepine.1 If patient is oxcarbazepine-naïve, avoid oxcarbazepine. If the patient has Oxcarbazepine Risk of hypersensitivity reaction due to the presence of previously used oxcarbazepine consistently for longer than three months without incidence of cutaneous adverse reactions, cautiously consider use of Trileptal® the HLA-B*15:02 allele. oxcarbazepine.1 If patient is phenytoin-naïve, avoid phenytoin. If the patient has previously used Phenytoin Risk of hypersensitivity reaction due to the presence of phenytoin consistently for longer than three months without incidence of the HLA-B*15:02 allele. Dilantin® cutaneous adverse reactions, cautiously consider use of phenytoïn.¹ Clopidogrel Reduced clopid ogrel active metabolite formation; Cardiovascular indications: avoid standard dose (75mg/day) if possible; use increased on-treatment platelet reactivity; increased Plavix® prasugrel or ticagrelor at standard dose if no contraindication. Neurovascular: risk for adverse cardiac and cerebrovascular events (CYP2C19 IM).

CAUTIONARY INFORMATION - MEDICATIONS TO AVOID OR USE WITH CAUTION

| Medication | Identified risk | Recommendation |
|------------|-----------------|---|
| | | consider alternative P2Y12 inhibitor at standard dose if clinically indicated and no contraindication. ² |

PGx RECOMMENDATIONS - PSYCHIATRY

↑ Dose increase may be required.

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Increased probability of a better response.

Dose reduction may be required.

- Greater potential for a poorer response or atypical effect.
- Exposure is difficult to predict, insufficient data to calculate dose adjustments.
- Medication not recommended by peer-reviewed guidelines.

| | ed | | |
|-------------------------------------|--|---|---|
| Medications | Exposure | Efficacy | Risk of atypical effect |
| Antidepressants | | | |
| Selective serotoni | in reuptake inhibitors (SSRIs) | | |
| Fluoxetine (Prozac®) | CYP2D6 NM, CYP2C9 IM - Initiate with recommended dose; a low dose may be adequate. | 4/6 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | Genetic influence not available |
| Fluvoxamine (Luvox®) | CYP2D6 NM, ABCB1 - Initiate with recommended dose but may require a higher dose. | 4/6 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | Genetic influence not available |
| Paroxetine Paxil®) | CYP2D6 NM, ABCB1 - Initiate therapy with recommended starting dose but may require a higher dose. | 4/6 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | Genetic influence not available |
| Citalopram Celexa®) | CYP2C19 IM - Initiate with recommended dose but consider a slower titration schedule and do not exceed the following daily doses: 30mg for adults up to 65 yrs; 15mg for adults 65 yrs or older. ^{3,4} | 4/6 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | 1/1 variant: increased risk of gastrointestinal side effects (HTR2A |
| Escitalopram Cipralex®) | CYP2C19 IM - Initiate with recommended dose but consider a slower titration schedule and do not exceed the following daily doses: 15mg for adults up to 65 yrs; 7.5mg for adults 65 yrs or older. ^{3,4} | 4/6 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | 1/1 variant: increased risk of gastrointestinal side effects (HTR2A) |
| Sertraline Zoloft®) | CYP2B6 NM, CYP2C19 IM - Initiate therapy with recommended starting dose but with a slower titration schedule and a lower maintenance dose. ³ | 4/5 variants: increased likelihood of a poorer response (SLC6A4 - association demonstrated mainly in Caucasians, not observed in Asians, BDNF, FKBP5, HTR7); consider adding L-methylfolate 15mg/day (MTHFR). | 1/1 variant: increased risk of gastrointestinal side effects (HTR2A) |
| Serotonin-norepi | nephrine 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); consider adding L-methylfolate 15mg/day (MTHFR). | Genetic influence not available |
| Levomilnacipran Fetzima®) | CYP3A4 NM, ABCB1 - Initiate therapy with recommended starting dose but may require a higher dose. | 1/2 variants: increased likelihood of a poorer response (FKBP5); consider adding L-methylfolate 15mg/day (MTHFR). | Genetic influence not available |

| Cyling (Effexor XR®) Duloxetine (Cymbalta®) Other antidepressants Bupropion (Wellbutrin®) Esketamine (Spravato®) Ketamine (Ketalar®) | P2B6 NM, POR - Initiate with commended dose; a low dose may be adequate. P2B6 NM, POR - Initiate therapy with commended starting dose but may require a higher dose, especially with CYP1A2 inducers, such as smoke. | Efficacy 3/4 variants: increased likelihood of a poorer response (COMT, FKBP5, SLC6A2); consider adding L-methylfolate 15mg/day (MTHFR). 1/3 variants: increased likelihood of a poorer response (FKBP5); consider adding L-methylfolate 15mg/day (MTHFR). 0/1 variant: no increased likelihood of a poorer response for treatment of depressive symptoms (HTR2A). 0/1 variant: no increased likelihood of a poorer response. | Risk of atypical effect Genetic influence not available Genetic influence not available Genetic influence not available 0/1 variant: no increased risk of emergent hypertension. |
|---|--|---|--|
| (Effexor XR®) reching Duloxetine (Cymbalta®) Other antidepressants Bupropion (Wellbutrin®) CYM rechand (Spravato®) rechand (Ketamine (Ketalar®)) | CYP1A2 Ind, CYP2D6 NM - Initiate therapy with recommended starting dose but may require a higher dose, especially with CYP1A2 inducers, such as smoke. YP2B6 NM, POR - Initiate with commended dose; a low dose may be dequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. | poorer response (COMT, FKBP5, SLC6A2); consider adding L-methylfolate 15mg/day (MTHFR). 1/3 variants: increased likelihood of a poorer response (FKBP5); consider adding L-methylfolate 15mg/day (MTHFR). 0/1 variant: no increased likelihood of a poorer response for treatment of depressive symptoms (HTR2A). 0/1 variant: no increased likelihood of a poorer response. | Genetic influence not available Genetic influence not available 0/1 variant: no increased risk of emergent |
| (Cymbalta®) Other antidepressants Bupropion (Wellbutrin®) Esketamine (Spravato®) Ketamine (Ketalar®) | therapy with recommended starting dose but may require a higher dose, especially with CYP1A2 inducers, such as smoke. (P2B6 NM, POR - Initiate with commended dose; a low dose may be lequate. (P2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. (P2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. | poorer response (FKBP5); consider adding L-methylfolate 15mg/day (MTHFR). 0/1 variant: no increased likelihood of a poorer response for treatment of depressive symptoms (HTR2A). 0/1 variant: no increased likelihood of a poorer response. | Genetic influence not available 0/1 variant: no increased risk of emergent |
| Bupropion (Wellbutrin®) Esketamine (Spravato®) Ketamine (Ketalar®) | YP2B6 NM, POR - Initiate with commended dose; a low dose may be dequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. | poorer response for treatment of depressive symptoms (HTR2A). 0/1 variant: no increased likelihood of a poorer response. 0/1 variant: no increased likelihood of a | 0/1 variant: no increased risk of emergent |
| (Wellbutrin®) rectand (Spravato®) Ketamine (Ketalar®) rectand (CYI) rectand (CYI) | commended dose; a low dose may be lequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. | poorer response for treatment of depressive symptoms (HTR2A). 0/1 variant: no increased likelihood of a poorer response. 0/1 variant: no increased likelihood of a | 0/1 variant: no increased risk of emergent |
| (Spravato®) recomma Ketamine (Ketalar®) recomma | commended starting dose; a low dose ay be adequate. YP2B6 NM, POR - Initiate therapy with commended starting dose; a low dose ay be adequate. | poorer response. 0/1 variant: no increased likelihood of a | _ |
| (Ketalar®) rec | commended starting dose; a low dose ay be adequate. | | |
| | /DZ A / NIM Normal accessors | poorer response. | 0/1 variant: no increased risk of emergent hypertension. |
| Trazodone CYI | /P3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available |
| (Viibryd®) rec | (P3A4 NM, ABCB1 - Initiate with commended dose but may require a gher dose. | Genetic influence not available | Genetic influence not available |
| | /P2D6 NM, CYP3A4 NM - Normal posure | Genetic influence not available | Genetic influence not available |
| Mirtazapine (Remeron®) | CYP1A2 Ind, CYP2D6 NM - Initiate with recommended dose but may require a higher dose, especially with CYP1A2 inducers, such as smoke. | 2/2 variants: increased likelihood of a poorer response (FKBP5, TPH2). | Genetic influence not available |
| Tricyclic antidepressan | nts (TCAs) | | |
| Clomipramine No (Anafranil®) | ormal exposure | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| (Norpramin®) rec | YP2D6 NM, ABCB1 - Initiate with commended dose but may require a gher dose. | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| Imipramine No (Tofranil®) | ormal exposure | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| (Aventyl®) rec | YP2D6 NM, ABCB1 - Initiate with commended dose but may require a gher dose. | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| Amitriptyline (Elavil®) | ABCB1, CYP2C19 IM, CYP2D6 NM - Initiate therapy with recommended starting dose but monitor response and tolerance more closely; insufficient data to calculate dose adjustments. | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| Doxepin (Sinequan®) | ABCB1, CYP2C19 IM, CYP2D6 NM - Initiate therapy with recommended starting dose but monitor response and tolerance more closely; insufficient data to calculate dose adjustments. | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |
| | ABCB1, CYP2C19 IM, CYP2D6 NM - Initiate therapy with recommended starting dose but monitor response and tolerance more closely; insufficient data to calculate dose adjustments. | 1/1 variant: increased likelihood of a poorer response (TPH2) | Genetic influence not available |

| | G | ed | |
|---|--|---|--|
| Medications | Exposure | Efficacy | Risk of atypical effect |
| Guanfacine (Intuniv XR®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available |
| Mood Stabilizers | | | |
| Gabapentin (Neurontin®) | ABCB1 - Normal exposure | Genetic influence not available | Genetic influence not available |
| Levetiracetam (Keppra®) | ABCB1 - Initiate with recommended dose; a low dose may be adequate. | Genetic influence not available | Genetic influence not available |
| Lithium (Carbolith®) | Genetic influence not available | 0/1 variant: no increased likelihood of a poorer response; 0/2 variants: normal likelihood of relapse after successful lithium therapy (IncRNA) - for bipolar disorder. | Genetic influence not available |
| Pregabalin (Lyrica®) | Genetic influence not available | Genetic influence not available | Genetic influence not available |
| Topiramate (Topamax®) | Genetic influence not available | 0/1 variant: no increased likelihood of a poorer response for treatment of alcohol-related disorders. | Genetic influence not available |
| Valproic acid, Divalproex (Depakene®, Epival®) | CYP2A6 NM, CYP2C9 IM - Normal exposure | Genetic influence not available | 1/1 variant: increased likelihood of weight gain (ANKK1). |
| Lamotrigine (Lamictal®) | ABCG2, UGT2B7 IM - Initiate with recommended dose; a low dose may be adequate. | Genetic influence not available | HLA-B*15:02 positive, the risk of hypersensitivity reaction is increased by a factor of 3.6 and estimated to be 0.4%. Avoid lamotrigine if an alternative is available. If not, advise the patient to report any rash immediately. ⁵ |
| Carbamazepine Tegretol®) UGT2B7 RM - Initiate with recommended dose but may require a higher dose. | | Genetic influence not available | HLA-B*15:02 positive,HLA-A*31:01 negative, increased risk of hypersensitivity reaction - If patient is carbamazepine-naïve, avoid carbamazepine. If the patient has previously used carbamazepine consistently for longer than three months without incidence of cutaneous adverse reactions, cautiously consider using carbamazepine. 1 |
| Oxcarbazepine (Trileptal®) UGT2B7 RM - Initiate with recommended dose but may require a higher dose. | | Genetic influence not available | HLA-B*15:02 positive, increased risk of hypersensitivity reaction - If patient is oxcarbazepine-naïve, avoid oxcarbazepine. If the patient has previously used oxcarbazepine consistently for longer than three months without incidence of cutaneous adverse reactions, cautiously consider use of oxcarbazepine. ¹ |
| Phenytoin (Dilantin®) CYP2C9 IM - Initiate with recommended starting dose but consider using 25% less than the typical maintenance dose based on response and tolerance. | | Genetic influence not available | HLA-B*15:02 positive - If patient is phenytoin-naïve, avoid phenytoin. If the patient has previously used phenytoin consistently for longer than three months without incidence of cutaneous adverse reactions, cautiously consider use of phenytoin. 6 |
| Norepinephrine Re | uptake Inhibitor | | |
| Atomoxetine | CYP2D6 NM - Normal exposure ^{5,7} | Genetic influence not available | Genetic influence not available |
| Strattera®) | CTI ZDO INIVI - NOTITIAI exposure | | |

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|--|--|---|--|--|--|--|
| | G | enetic Associations Identific | Associations Identified | | | |
| Medications | Exposure | Efficacy | Risk of atypical effect | | | |
| Psychostimulants | | | | | | |
| Amphetamine / Dextroamphetamine (Adderall XR®) | CYP2D6 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Dextroamphetamine (Dexed rine®) | CYP2D6 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Lisdexamfetamine (Vyvanse®) | CYP2D6 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Methylphenidate - Biphentin® | CES1 IM - Consider using a lower dose. | 2/5 variants: increased likelihood of a poorer response (COMT, SLC6A2). | Genetic influence not available | | | |
| Methylphenidate - Concerta® | CES1 IM - Consider using a lower dose. | 2/5 variants: increased likelihood of a poorer response (COMT, SLC6A2). | Genetic influence not available | | | |
| Methylphenidate - Foquest® | CES1 IM - Consider using a lower dose. | 2/5 variants: increased likelihood of a poorer response (COMT, SLC6A2). | Genetic influence not available | | | |
| Methylphenidate - Ritalin® | CES1 IM - Consider using a lower dose. | 2/5 variants: increased likelihood of a poorer response (COMT, SLC6A2). | Genetic influence not available | | | |
| Sedative-Hypnotic | cs . | | | | | |
| Daridorexant (Quviviq®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Diphenydramine (Benadryl®) | Genetic influence not available | Genetic influence not available | Genetic influence not available | | | |
| Eszopicione (Lunesta®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Lemborexant (Dayvigo®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Melatonin | Genetic influence not available | Genetic influence not available | Genetic influence not available | | | |
| Phenobarbital (Phenobarb®) | CYP2C9 IM - Initiate therapy with recommended starting dose; a low dose may be adequate. | Genetic influence not available | Genetic influence not available | | | |
| Triazolam (Halcion®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Zolpidem (Sublinox®) | CYP3A4 NM, CYP2C9 IM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Zopiclone (Imovane®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Wakefulness-prom | noting agents | | | | | |
| Modafinil (Alertec®) | CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Pitolisant (Wakix®) | CYP2D6 NM, CYP3A4 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | | |
| Sodium oxybate (Xyrem®) | Genetic influence not available | Genetic influence not available | Genetic influence not available | | | |
| Solriam fetol (Sunosi®) | Genetic influence not available | Genetic influence not available | Genetic influence not available | | | |

PGx RECOMMENDATIONS - COMPLEMENTARY TREATMENTS

Dose increase may be required.

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Increased probability of a better response.

Dose reduction may be required.

Greater potential for a poorer response or atypical effect.

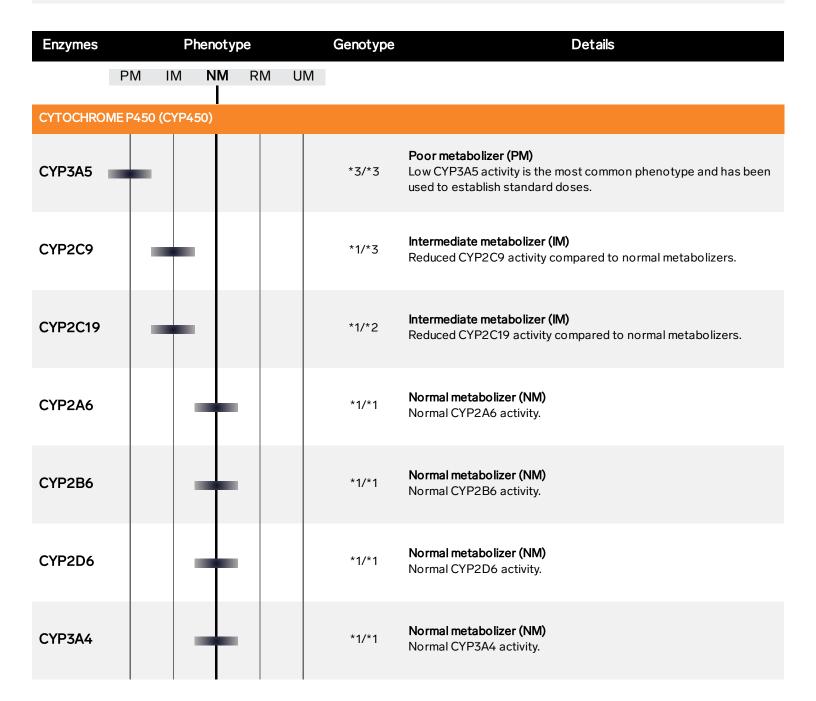
Exposure is difficult to predict, insufficient data to calculate dose adjustments.

Medication not recommended by peer-reviewed guidelines.

| | G | ienetic Associations Identified | | | |
|------------------------------------|---|---------------------------------|---------------------------------|--|--|
| Medications | Exposure | Efficacy | Risk of atypical effect | | |
| Antiemetics | | | | | |
| Dimenhydrinate (Gravol®) | Genetic influence not available | Genetic influence not available | Genetic influence not available | | |
| Granisetron (Kytril®) | CYP3A4 NM, CYP3A5 PM - Normal exposure | Genetic influence not available | Genetic influence not available | | |
| Ondansetron (Zofran®) | CYP2D6 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | |
| Palonosetron (Aloxi®) | CYP2D6 NM - Normal exposure | Genetic influence not available | Genetic influence not available | | |
| Proton pump inhib | itors (PPI) | | | | |
| Esomeprazole (Nexium®) | Genetic influence not available | Genetic influence not available | Genetic influence not available | | |
| Dexiansoprazole (Dexilant®) | CYP2C19 IM - Initiate therapy with standard dose but for chronic therapy (> 12 weeks) and efficacy achieved, consider reducing the daily dose by 50% and monitor for continued efficacy. ⁸ | Genetic influence not available | Genetic influence not available | | |
| Lansoprazole (Prevacid®) | CYP2C19 IM - Initiate therapy with standard dose but for chronic therapy (> 12 weeks) and efficacy achieved, consider reducing the daily dose by 50% and monitor for continued efficacy. ⁸ | Genetic influence not available | Genetic influence not available | | |
| Omeprazole (Losec®) | CYP2C19 IM - Initiate therapy with standard dose but for chronic therapy (> 12 weeks) and efficacy achieved, consider reducing the daily dose by 50% and monitor for continued efficacy. ⁸ | Genetic influence not available | Genetic influence not available | | |
| Pantoprazole (Pantoloc®) | CYP2C19 IM - Initiate therapy with standard dose but for chronic therapy (> 12 weeks) and efficacy achieved, consider reducing the daily dose by 50% and monitor for continued efficacy. ⁸ | Genetic influence not available | Genetic influence not available | | |

PGx ASSOCIATIONS - EXPOSURE

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



| 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 | 1/1 |

| Efflux Pump | | Phenotype | | Genotype | Details |
|----------------|------------------|-----------------------|-----------------|--------------|--|
| | Poor activity | Intermediate activity | Normal activity | | |
| EFFLUX PUM | IPS (P-GLY | COPROTEIN, MU | JLT IDRUG RI | ESISTANCE PF | ROTEIN) |
| ABCB1 ■ | + | - | | AAT/ACT | Poor activity (PA) Reduced efflux pump activity may result in higher substrate concentrations, increasing the likelihood of response but also increasing the risk of adverse drug reactions. |
| ABCG2 | | + | | G/T | Intermediate activity (IA) Reduced efflux pump activity may result in higher substrate concentrations, increasing the likelihood of response but also increasing the risk of adverse drug reactions. |

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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10/23

| Poorei | response m | ore likely | |
|--------|--------------------------------|---------------------|---|
| Gene | Reference SNP cluster ID | Variant detected | Associated medication(s) |
| ANKK1 | rs1800497 | yes | aripiprazole, risperidone |
| BDNF | rs6265 | yes | citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine |
| 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 |
| HTR7 | rs7905446 | yes | citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine |
| SLC6A2 | rs2242446 | yes | venlafaxine |
| SLC6A2 | rs28386840 | yes | methylphenidate |
| SLC6A4 | 5-HTTLPR | yes | citalopram, escitalopram, sertraline, fluoxetine, fluvoxamine, paroxetine |
| TPH2 | rs1487278 | yes | amitriptyline, clomipramine, desipramine, doxepin, imipramine, mirtazapine, nortriptyline, trimipramine |
| ADRA2A | rs1800544 | no | methylphenidate |
| ANKK1 | rs1800497 | no | bupropion |
| BDNF | rs6265 | no | esketamine, ketamine |
| CACNG2 | rs2283967 | no | lithium |
| DRD3 | rs963468 | no | duloxetine |
| GRIK1 | rs2832407 | no | topiramate |
| GRIK4 | rs1954787 | no | citalopram, desvenlafaxine, duloxetine, escitalopram, fluoxetine, fluvoxamine, levomilnacipran, paroxetine, sertraline, venlafaxine |
| HTR2A | rs2770296 | no | bupropion |

DOB: 1999-01-01

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

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5/13

| Increa | Increased risk of adverse drug reactions | | | | | |
|--------|--|---------------------|--|--|--|--|
| Gene | Reference SNP cluster ID | Variant detected | Associated medication(s) | | | |
| ANKK1 | rs1800497 | yes | weight gain: valproic acid | | | |
| HLA-B | *15:02 | yes | cutaneous adverse reactions: carbamazepine, oxcarbazepine, phenytoin | | | |
| HTR2A | rs6311 | yes | gastrointestinal side effects (nausea, vomiting): citalopram, escitalopram, sertraline | | | |
| HTR2C | rs3813929 | yes | antipsychotic-induced weight gain: clozapine, olanzapine, risperidone | | | |
| MC4R | rs489693 | yes | antipsychotic-induced weight gain: clozapine, olanzapine, quetiapine, risperidone | | | |
| 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 | | | |
| INSIG2 | rs17047764 | no | antipsychotic-induced weight gain: clozapine, olanzapine, risperidone | | | |
| MC4R | rs17782313 | no | antipsychotic-induced weight gain: quetiapine | | | |
| 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 Detai | ls (GRCH38.p12) | Result |
|---------|----------------------------|---|-------------|
| ABCB1 | rs1045642 | chr7:87509329 | A A |
| | rs2032582 | chr7:87531302 | AIC |
| | rs2032583 | chr7:87531245 | T T |
| ABCG2 | rs2231142 | chr4:88131171 | G T |
| ADRA2A | rs1800544 | chr10:111076745 | C G |
| ANKK1 | rs1800497 | chr11:113400106 | GIG |
| BDNF | rs6265 | chr11:27658369 | CIC |
| CACNG2 | rs2283967 | chr22:36567486 | CIC |
| CES1 | rs71647871 | chr16:55823658 | CIT |
| CNR1 | rs806380 | chr6:88154934 | AlA |
| | | | AIG |
| COMT | rs4680 | chr22:19963748 | |
| CYP1A2 | rs762551 rs2069514 | chr15:74749576 chr15:74745879 | C A G G |
| CVD247 | | | |
| CYP2A6 | rs1801272 | chr19:40848628 | A A |
| | rs5031017 rs28399433 | chr19:40843845 chr19:40850474 | G G A A |
| CYP2B6 | rs2279343 | chr19:41009358 | <u> </u> |
| C1P2B0 | rs3745274 | chr19:41009358 | G G G G |
| | rs28399499 | chr19:41012316 | T T |
| CYP2C9 | rs1057910 | chr10:94981296 | CIA |
| CIFZC9 | rs1799853 | chr10:94942290 | CIC |
| | rs7900194 | chr10:94942309 | GIG |
| | rs9332131 | chr10:94949282-94949283 | AA |
| | rs9332239 | chr10:94989020 | cic |
| | rs28371685 | chr10:94981224 | CIC |
| | rs28371686 | chr10:94981301 | CIC |
| | rs72558187 | chr10:94941958 | T T |
| | rs72558190 | chr10:94947782 | CIC |
| CYP2C19 | rs4244285 | chr10:94781859 | GA |
| | rs4986893 | chr10:94780653 | G G |
| | rs6413438 rs12248560 | chr10:94781858 chr10:94761900 | C C C C |
| | rs12769205 | chr10:94761900 | GIA |
| | rs17884712 | chr10:94775489 | GIG |
| | rs28399504 | chr10:94762706 | AlA |
| | rs41291556 | chr10:94775416 | ΤİΤ |
| | rs56337013 | chr10:94852738 | cic |
| | rs72552267 | chr10:94775453 | GIG |
| | rs72558186 | chr10:94781999 | T T |
| CYP2D6 | rs16947 | chr22:42127941 | GIG |
| | rs1065852 | chr22:42130692 | G G |
| | rs1135840 | chr22:42126611 | CIC |
| | rs3892097 rs5030655 | chr22:42128945 chr22:42129084 | CIC |
| | rs5030656 | chr22:42128174-42128178 | A A A A |
| | rs5030862 | chr22:42130668 | CIC |
| | rs5030865 | chr22:42129033 | CIC |
| | rs5030867 | chr22:42127856 | T T |
| | rs28371725 | chr22:42127803 | cic |
| | rs28371706 | chr22:42129770 | GIG |
| | rs35742686 | chr22:42128242 | T T |
| | rs59421388 | chr22:42127608 | C C |
| | rs72549354 | chr22:42128815-42128817 | |
| | rs774671100 rs201377835 | chr22:42130555-42130755 chr22:42129910 | G G C C |
| | Gene Deletion | n/a | Not Detecte |
| | Gene Duplication | n/a | Not Detecte |
| | 2:2:: 2 = 2p::000:0:: | 2 | 0.0000 |

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

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

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