#### Check for updates

#### **OPEN ACCESS**

EDITED BY Tiebing Liang, Indiana University Bloomington, United States

REVIEWED BY Miriam Saiz-Rodríguez, Hospital Universitario de Burgos, Spain Tae-Hwi Schwantes-An, Indiana University School of Medicine, United States

\*CORRESPONDENCE Thomas M. Polasek, ☑ tom.polasek@certara.com

RECEIVED 27 January 2023 ACCEPTED 21 April 2023 PUBLISHED 05 May 2023

#### CITATION

Polasek TM (2023), Calculation of the pharmacogenomics benefit score for patients with medicationrelated problems. *Front. Genet.* 14:1152585. doi: 10.3389/fgene.2023.1152585

#### COPYRIGHT

© 2023 Polasek. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

# Calculation of the pharmacogenomics benefit score for patients with medication-related problems

### Thomas M. Polasek<sup>1,2</sup>\*

<sup>1</sup>Certara, Princeton, NJ, United States, <sup>2</sup>Centre for Medicines Use and Safety, Monash University, Melbourne, VIC, Australia

Unexpected poor efficacy and intolerable adverse effects are medication-related problems that may result from genetic variation in genes encoding key proteins involved in pharmacokinetics or pharmacodynamics. Pharmacogenomic (PGx) testing can be used in medical practice "pre-emptively" to avoid future patient harm from medications and "reactively" to diagnose medication-related problems following their occurrence. A structured approach to PGx consulting is proposed to calculate the pharmacogenomics benefit score (PGxBS), a patient-centered objective measure of congruency between medication-related problems and patient genotypes. An example case of poor efficacy with multiple medications is presented, together with comments on the potential benefits and limitations of using the PGxBS in medical practice.

#### KEYWORDS

pharmacogenomics benefit score, pharmacogenomics, precision medicine, precision dosing, adverse drug reactions

## Background

There is growing interest in using pharmacogenomics (PGx) broadly in medical practice to improve the chances of therapeutic success in individual patients by precision dosing (Polasek et al., 2018; Polasek et al., 2019). Clinical guidelines are available to instruct doctors on how to prescribe select medications based on patient genotypes (Relling and Klein, 2011). Ideally, this should be done prior to commencing treatment, which is called "pre-emptive" PGx testing. There are many examples in well-resourced healthcare systems of PGx services being implemented successfully, usually via electronic clinical decision support systems (CDSS) (Dunnenberger et al., 2016); patients are screened and almost all (>95%) are found to have genetic variants with so-called "actionable PGx guideline recommendations" that could influence future prescribing (Mostafa et al., 2019). Less frequently addressed in the PGx literature is the clinical scenario where patients have histories of medication-related problems at standard doses without an obvious explanation, either unexpected poor efficacy or intolerable adverse effects. "Reactive" PGx testing can be used in these patients to diagnose whether PGx is the potential cause. Pharmacogenomic testing is therefore a unique pathology test that has dual clinical utility depending on when the test is ordered and/or reviewed relative to the medication prescribed i.e., a screening test to avoid future patient harm and a diagnostic test in the work-up of differential diagnoses. Whilst there is growing evidence for pre-emptive PGx testing to decrease adverse drug reactions (ADRs), by as much as 30% in some studies (Zhou et al., 2015; Cacabelos et al.,



2019; Swen et al., 2023), the degree to which reactive PGx testing diagnoses the cause of medication-related problems is unclear.

In this report, a structured approach to PGx consulting by a clinical pharmacologist is described based on referrals of patients with current and/or past medication-related problems (Aronson, 2010). The pharmacogenomics benefit score (PGxBS) is proposed as a patient-centered objective measure of congruency between medication-related problems and patient genotypes. An example case of unexpected poor efficacy with multiple medications is presented to show how the PGxBS is calculated. Finally, consideration is given to the potential benefits and limitations of using the PGxBS in medical practice.

# Categories of PGx

There are three categories to consider when diagnosing PGx as the potential cause of medication-related problems.

1) Exposure PGx. Is the patient at risk of extreme exposure to the medication at standard doses? Pharmacokinetic processes determine "how much" a medication is available at the sites of action, and therefore, assuming typical dose-exposure-response relationships, the magnitude of response. Extremely high medication exposures are associated with an increased risk of adverse effects, whereas persistently low medication exposures may result in subtherapeutic concentrations and poor efficacy. Although many genes influence pharmacokinetics, the cytochrome P450 (CYP) enzymes are the most important for PGx (Doogue and Polasek, 2013).

2) Response PGx. Does the patient have the correct molecular target for the medication? At a given exposure, genetic variability in the molecular target can determine the response (pharmacodynamics). This is best exemplified currently in hematology and oncology; patients are treated with targeted pharmacotherapy based on the results of genetic testing of the molecular targets expressed by cancer cells (Polasek et al., 2016). This category will expand in the future as genomic analyses identify

novel pharmacodynamic biomarkers of response (Dawed et al., 2023).

**3)** Safety PGx. Is the patient at risk of a severe adverse drug reaction to the medication at standard doses? There is some overlap here with category 1 (Exposure PGx) and category 2 (Response PGx) but this category primarily includes rare severe cutaneous adverse drug reactions (SCARs) in patients with certain human leukocyte antigen (HLA) genotypes. In these cases, the patients' immune system carries genetic variants that significantly increase the likelihood of ADRs (Kloypan et al., 2021).

# Structured PGx consult

A structured approach to PGx consulting by a doctor is suggested here because medication-related problems should be considered under differential diagnoses. This requires diagnostic skill and experience, and a broader understanding of the patient beyond simply medications and genotypes (Aronson, 2010) (Figure 1). The doctor may or may not have access to an electronic CDSS with PGx guidance (Wake et al., 2021). The following steps outline the information required to calculate the PGxBS. Binary responses to the main steps are required. A spreadsheet can be used to log answers and calculate scores.

1) List current and past medications. Current medications have priority. Past medications are also important to capture if time permits, since clues on how patients respond to medications more broadly may be garnered, further informing the PGxBS.

2) Determine availability of PGx guidelines. For each medication, determine whether Clinical Pharmacogenomics Implementation Consortium (CPIC<sup>®</sup>) level A or A/B evidence is available (www.cpicpgx.org). Assign one if the answer is "yes" and 0 for "no/unsure". If there are no medications with PGx guidelines, then the PGx consult is complete and the PGxBS for the patient is 0.

**3**) Assess adequacy of therapeutic trials. For each medication with PGx guidelines, determine whether the patient had an adequate therapeutic trial or not. Assign one if the answer is "yes" and 0 for

Medications	PGx Guidances Therapeutic Trials Med		Medication-F	Related Problems	Role of PGx Testing	Outcomes of PGx testing	
	A or B CPIC levels	Good dose and duration	Poor efficacy	Adverse Effect(s)		Congruent	Incongruent
urrent Medications							-
Mirtazapine	0	1	1		PGx unhelpful		
Olanzapine	0	1	1		PGx unhelpful		
Olanzapine	0	1	1		Poxumeipiui		
ast Medications	1						
Quetiapine	0	1		1	PGx unhelpful		
Agomelatine	0	0	1		PGx unhelpful		
	-						
Suvorexant	0	1	1		PGx unhelpful		
Diazepam	0	1	1		PGx unhelpful		
Diazepain	v	1			Poxumeipiu		
Fluoxetine	0	1	1		PGx unhelpful		
Desvenlafaxine	0	1	1		PGx unhelpful		
	-						
Periciazine	0	1	1		PGx unhelpful		
Clomipramine	1	1	1		PGx helpful	1	
ciompianine	-				roxitelpidi	-	
Zopiclone	0	0	1		PGx unhelpful		
					·		
Codeine	1	1	1		PGx helpful		1
Totals	2	10	11	1		1	1
oncidor ordoring PC:	tosting if > 2 "DCh	alaful" roculto		DCy Testing -	Indicated	PGxBS =	0
onsider ordering PGx	testing if 2.2 "PGX he	eipiui results		PGx Testing =	Indicated	PGXB2=	U

"no/unsure". Inadequate therapeutic trials from underdosing or short durations of treatment are common and should be recognised, scoring 0. This section can also be completed for medications without PGx guidelines to improve the medication history, but these responses do not count towards the PGxBS.

4) Determine therapeutic outcomes. Two types of medicationrelated problems indicate negative therapeutic outcomes that could be explained by PGx—unexpected poor efficacy or intolerable adverse effects (Polasek et al., 2018). One is chosen here, scoring 1, with the alternative scoring 0. Medications with inadequate therapeutic trials (step 3) are ignored.

5) Determine congruency between therapeutic outcomes and PGx results. Is each medication-related problem consistent with the genotype-predicted phenotype? Again, this is a binary option, with congruent results scoring 1 and incongruent results scoring -1. The same PGx guidelines from step 2 (CPIC<sup>\*</sup>) should be used. An example of a congruent result is a patient who experienced SCAR after starting allopurinol and who was subsequently shown to carry the *HLA-B\*5801* allele (Lucas and Droney, 2022). Alternatively, a chronic pain sufferer with a CYP2D6 poor metabolizer (PM) phenotype who experienced euphoria and intolerable dizziness and nausea with low dose tramadol is an example of an incongruent result (Crews et al., 2021).

6) Calculate the PGxBS. Congruent and incongruent results are added. Scores  $\geq 1$  indicate a possible contribution of PGx to medication-related problems, whereas 0 and negative scores show that PGx is less likely to be important for the patient.

## Example case

A 42-year-old man with a 6-year history of depression, anxiety, insomnia, and chronic lower back pain was referred by his general practitioner to a multi-disciplinary ambulatory care clinic staffed by clinical pharmacologists for "poor responses to psychotropics and pain killers". His mental state had deteriorated over the previous 3 months, and he was awaiting psychiatrist review. Figure 2 shows the spreadsheet used to document the consult and calculate his PGxBS. Since the patient had two medications with PGx guidelines and no previous PGx testing, it was recommended, and the patient accepted the cost (~\$100USD). The PGx results were reconciled with the medication-related problems at the follow-up appointment. His CYP2D6 ultra-rapid (UM) metabolizer phenotype was incongruent with poor analgesic response to codeine (score = -1). However, there was congruency between CYP2D6 UM and CYP2C19 normal metabolizer (NM) phenotypes and no

improvement in mental state with clomipramine (score = 1). The PGxBS was 0. Importantly, this patient held strong beliefs about being "abnormal" and "unable to be helped by drugs". Counselling was provided to explain that no known genetic cause for his poor responses was found. The patient was encouraged to be positive about medications in his overall treatment. The PGx spreadsheet was included in the medical consult note and forwarded to his treating general practitioner and psychiatrist (Figure 2).

# Potential benefits of the PGxBS

The PGxBS is a clinically useful objective measure of congruency between medication-related problems and patient genotypes. The score is patient-centered rather than focused on individual medication-gene pairs ("this is your PGxBS"). The score is easy to understand for patients and non-expert PGx users-positive results indicate a possible role for PGx, whereas zero and negative scores mean that PGx is less likely to be important. The PGxBS may be applied to patients with single or multiple current and/or past medication-related problems. The PGxBS is dynamic and changes with time and changing medication regimens. Calculating a patient's PGxBS requires particular attention to the medication history, which alone has benefits for clinical care. Importantly, the structured PGx consult allows for patient education on the many factors that explain why different patients respond to medications differently, including pharmacokinetic drug-drug interactions that cause CYP phenoconversion (Mostafa et al., 2021; Mostafa et al., 2022). Although the emphasis in this report is on medical practice, pharmacists with expertise in PGx could calculate the PGxBS and integrate it into their clinical practice, ideally in close collaboration with the treating doctor (Polasek et al., 2015).

# Limitations of the PGxBS

The PGxBS does not apply to pre-emptive PGx testing, where, at least in principle, almost all patients will benefit i.e., >95% have genetic variants with so-called "actionable PGx guideline recommendations" (Mostafa et al., 2019; Swen et al., 2023). In patients with medication-related problems who have not been tested, two or more medications with PGx guidelines is the suggested cut-off for reactive PGx testing. This is only a guide since the clinical need (indication) for reactive PGx testing depends on many factors, including disease status, differential diagnoses, severity of treatment outcomes, treatment alternatives, and test affordability. The PGxBS is not validated for clinical decision-making, including prescribing. To date, the score has not been applied beyond one clinical pharmacology referral stream in Australia. Whether a patient's present score reflects the future clinical utility of PGx for that patient is

## References

Aronson, J. K. (2010). A manifesto for clinical pharmacology from principles to practice. *Br. J. Clin. Pharmacol.* 70, 3–13. doi:10.1111/j.1365-2125. 2010.03699.x

unknown. The PGxBS often depends on the recollection of subjective past experiences with medications, occurring years previously in some cases, and there may be intrinsic biases. Finally, there are nuances to the PGxBS that are debatable, such as the PGx guidelines and levels of evidence chosen (step 2).

#### Conclusion

Despite the promise of superior patient care and considerable academic and commercial interests, adoption of PGx in routine medical practice has been limited (Pearce et al., 2022). Whilst there is growing evidence for pre-emptive PGx testing to avoid ADRs, the degree to which reactive PGx testing diagnoses the cause of medication-related problems is less clear. Rather than details about individual medication-gene pairs, patients with histories of medication-related problems and their doctors are often more interested in whether PGx is "the answer". In such cases, a structured approach to PGx consulting is recommended to generate the PGxBS, a patient-centered objective measure of congruency between medication-related problems and patient genotypes.

## Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

## Author contributions

The author confirms being the sole contributor of this work and has approved it for publication.

# Conflict of interest

TP provides a pharmacogenomics consulting service for Sonic Healthcare Ltd.

# Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Cacabelos, R., Cacabelos, N., and Carril, J. C. (2019). The role of pharmacogenomics in adverse drug reactions. *Expert Rev. Clin. Pharmacol.* 12, 407–442. doi:10.1080/ 17512433.2019.1597706 Crews, K. R., Monte, A. A., Huddart, R., Caudle, K. E., Kharasch, E. D., Gaedigk, A., et al. (2021). Clinical pharmacogenetics implementation Consortium guideline for CYP2D6, OPRM1, and COMT genotypes and select opioid therapy. *Clin. Pharmacol. Ther.* 110, 888–896. doi:10.1002/cpt.2149

Dawed, A. Y., Mari, A., Brown, A., McDonald, T. J., Li, L., Wang, S., et al. (2023). Pharmacogenomics of GLP-1 receptor agonists: A genome-wide analysis of observational data and large randomised controlled trials. *Lancet Diabetes Endocrinol.* 11, 33–41. doi:10.1016/S2213-8587(22)00340-0

Doogue, M. P., and Polasek, T. M. (2013). The ABCD of clinical pharmacokinetics. *Ther. Adv. Drug Saf.* 4, 5–7. doi:10.1177/2042098612469335

Dunnenberger, H. M., Biszewski, M., Bell, G. C., Sereika, A., May, H., Johnson, S. G., et al. (2016). Implementation of a multidisciplinary pharmacogenomics clinic in a community health system. *Am. J. Health-Syst Pharm.* 73, 1956–1966. doi:10.2146/ ajhp160072

Kloypan, C., Koomdee, N., Satapornpong, P., Tempark, T., Biswas, M., and Sukasem, C. (2021). A comprehensive review of HLA and severe cutaneous adverse drug reactions: Implication for clinical pharmacogenomics and precision medicine. *Pharm. (Basel)* 14, 1077. doi:10.3390/ph14111077

Lucas, G., and Droney, L. (2022). Severe adverse drug reaction to allopurinol. Aust. Prescr. 45, 130–131. doi:10.18773/austprescr.2022.032

Mostafa, S., Kirkpatrick, C. M. J., Byron, K., and Sheffield, L. (2019). An analysis of allele, genotype and phenotype frequencies, actionable pharmacogenomic (PGx) variants and phenoconversion in 5408 Australian patients genotyped for CYP2D6, CYP2C19, CYP2C9 and VKORC1 genes. *J. Neural Transm. (Vienna)* 126, 5–18. doi:10. 1007/s00702-018-1922-0

Mostafa, S., Polasek, T. M., Bousman, C. A., Müeller, D. J., Sheffield, L. J., Rembach, J., et al. (2022). Pharmacogenomics in psychiatry - the challenge of cytochrome P450 enzyme phenoconversion and solutions to assist precision dosing. *Pharmacogenomics* 23, 857–867. doi:10.2217/pgs-2022-0104

Mostafa, S., Polasek, T. M., Sheffield, L. J., Huppert, D., and Kirkpatrick, C. M. J. (2021). Quantifying the impact of phenoconversion on medications with actionable

pharmacogenomic guideline recommendations in an acute aged persons mental health setting. *Front. Psychiatry* 12, 724170. doi:10.3389/fpsyt.2021.724170

Pearce, A., Terrill, B., Alffenaar, J. W., Patanwala, A. E., Kummerfeld, S., Day, R., et al. (2022). Pharmacogenomic testing: Perception of clinical utility, enablers and barriers to adoption in Australian hospitals. *Intern Med. J.* 52, 1135–1143. doi:10.1111/imj.15719

Polasek, T. M., Ambler, K., Scott, H. S., Sorich, M. J., Kaub, P. A., Rowland, A., et al. (2016). Targeted pharmacotherapy after somatic cancer mutation screening. *F1000Res* 5, 1551. doi:10.12688/f1000research.9040.2

Polasek, T. M., Mina, K., and Suthers, G. (2019). Pharmacogenomics in general practice - the time has come. *Aust. J. General Pract.* 48, 100–105. doi:10.31128/AJGP-10-18-4733

Polasek, T. M., Rowland, A., Wiese, M. D., and Sorich, M. J. (2015). Pharmacists in Australian general practice: An opportunity for expertise in precision medicine. *Ther. Adv. Drug Saf.* 6, 186–188. doi:10.1177/2042098615599947

Polasek, T. M., Shakib, S., and Rostami-Hodjegan, A. (2018). Precision dosing in clinical medicine: Present and future. *Expert Rev. Clin. Pharmacol.* 11, 743–746. doi:10. 1080/17512433.2018.1501271

Relling, M. V., and Klein, T. E. (2011). CPIC: Clinical pharmacogenetics implementation Consortium of the pharmacogenomics Research network. *Clin. Pharmacol. Ther.* 89, 464–467. doi:10.1038/clpt.2010.279

Swen, J. J., van der Wouden, C. H., Manson, L. E., Abdullah-Koolmees, H., Blagec, K., Blagus, T., et al. (2023). A 12-gene pharmacogenetic panel to prevent adverse drug reactions: An open-label, multicentre, controlled, cluster-randomised crossover implementation study. *Lancet* 401, 347–356. doi:10.1016/S0140-6736(22)01841-4

Wake, D. T., Smith, D. M., Kazi, S., and Dunnenberger, H. M. (2021). Pharmacogenomic clinical decision support: A review, how-to guide, and future vision. *Clin. Pharmacol. Ther.* 112, 44–57. doi:10.1002/cpt.2387

Zhou, Z. W., Chen, X. W., Sneed, K. B., Yang, Y. X., Zhang, X., He, Z. X., et al. (2015). Clinical association between pharmacogenomics and adverse drug reactions. *Drugs* 75, 589–631. doi:10.1007/s40265-015-0375-0