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Pharmacogenomic information from CPIC and DPWG guidelines and its application on drug labels

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ABSTRACT

There are several hurdles to overcome before implementing pharmacogenomics (PGx) in precision medicine. One of the hurdles is unawareness of PGx by clinicians due to insufficient pharmacogenomic information on drug labels. Therefore, it might be important to implement PGx that reflects pharmacogenomic information on drug labels, standard of prescription for clinicians. This study aimed to evaluate the level at which PGx was being used in clinical practice by comparing the Clinical Pharmacogenetics Implementation Consortium and Dutch Pharmacogenetics Working Group guidelines and drug labels of the US Food and Drug Administration (FDA) and the Korea Ministry of Food and Drug Safety (MFDS). Two PGx guidelines and drugs labels were scrutinized, and the concordance of the pharmacogenomic information between guidelines and drug labels was confirmed. The concordance of the label between FDA and MFDS was analyzed. In FDA labels, the number of concordant drug with guidelines was 24, while 13 drugs were concordant with MFDS labels. The number of drugs categorized as contraindication, change dose, and biomarker testing required was 7, 12 and 12 for the FDA and 8, 5 and 4 for the MFDS, respectively. The pharmacogenomic information of 9 drugs approved by both FDA and MFDS was identical. In conclusion, pharmacogenomic information on clinical implementation guidelines was limited on both FDA and MFDS labels because of various reasons including the characteristics of the guidelines and the drug labels. Therefore, more effort from pharmaceutical companies, academia and regulatory affairs needs to be made to implement pharmacogenomic information on drug labels.

Keywords: Pharmacogenomics; Guideline; Drug Labeling; United States Food and Drug Administration; Korea Ministry of Food and Drug Safety

INTRODUCTION

In 2015, President Barack Obama launched the precision medicine initiative, and the United States Food and Drug Administration (FDA) has gathered large-scaled biologic databases such as human genome sequence, metabolomics, and clinical studies for drugs. Like Obama's precision medicine initiative, modern medicine is heading towards precision medicine, a model that pursues personalization of healthcare that covers medical decisions, treatments, practices or products based on individual genotypic, phenotypic or psychosocial characteristics [1].

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Conflict of Interest

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Author Contributions

Conceptualization: Yoon DY, Jang IJ, Lee SH; Data curation: Yoon DY, Lee SH; Formal analysis: Yoon DY, Lee SH; Investigation: Yoon DY, Lee SH; Methodology: Yoon DY, Jang IJ, Lee SH; Project administration: Yoon DY, Lee SH; Supervision: Lee S, Lee SH; Validation: Yoon DY; Visualization: Yoon DY, Lee S; Writing - original draft: Yoon DY; Writing - review & editing: Lee S, Ban MS, Jang IJ, Lee SH. The concept of personalization of healthcare has existed before. For example, therapeutic drug monitoring has been practiced for a long time in clinical fields to adjust the drug dose according to the individual patients' status [2]. By identifying the pharmacokinetics and pharmacodynamics of a drug and the characteristics of the individual variances of a drug, clinicians or pharmacists can adjust the dosage of drugs or change to another drug for an individual patient. Nowadays, precision medicine is also implemented based on the concept of pharmacogenomics (PGx), and crowdsourcing platforms have made it possible for clinicians to access healthcare sources to improve precision medicine [3].

Many organizations associated with PGx such as the Clinical Pharmacogenetics Implementation Consortium (CPIC), Dutch Pharmacogenetics Working Group (DPWG), Pharmacogenomics Research Network and Ubiquitous Pharmacogenomics have been contributing to the implementation of PGx in clinical practice by establishing pharmacogenomic information and developing implementation tools [4,5]. CPIC and DPWG have released public guidelines to implement PGx in clinical practice. These PGx guidelines help clinicians and pharmacists understand how to interpret and apply the results of a genetic test for optimizing pharmacotherapy. Moreover, various tools have been designed like a personalized pocket card that integrates genetic test results into electronic medical records and a clinical decision support system. The card contains personal pharmacogenomic information and a quick response code connected to the individual's personalized dosing recommendations, and it could help clinicians to make a decision for personalized pharmacotherapy [6].

Not only academia but also regulatory affairs have made an effort to encourage PGx implementation. In Unite States, cloud-based next-generation DNA sequencing (NGS) data platform called precision FDA was established. The platform enables researchers to upload and compare data against reference genomes, bioinformatics pipelines, and genomic data [7]. In addition, the FDA released a "Guidance for Industry: Clinical Pharmacogenomics: Premarket Evaluation in Early-Phase Clinical Studies and Recommendations for Labeling" to recommend a prospective collection of samples for PGx research during early phase clinical studies [8]. In the Republic of Korea, the cost of some NGS-based oncology panel tests has been covered by the Korea National Health Insurance Service.

However, there are several huddles for implementing PGx in precision medicine. Cost of genetic tests and reimbursements are common problems facing implementation of PGx in clinical practice around the world [9]. Moreover, lack of education and unawareness of PGx by physicians and pharmacists who actually work with drugs are other obstacles [10,11]. Insufficient pharmacogenomic information on drug labels could be the main reason for the unawareness of PGx by physicians and pharmacists. Therefore, it might be important to implement PGx that reflects evidence-based pharmacogenomic information on drug labels which should be the standard of prescription labels for clinicians.

This study aimed to evaluate the level at which PGx was being used in clinical practice by comparing the PGx guidelines and drug labels of the FDA and Korea Ministry of Food and Drug Safety (MFDS). We reviewed two major public PGx implementation guidelines and identified whether the pharmacogenomic information was reflected on the drug labels. Furthermore, we discussed scientific and regulatory reasons underlying the discordance between the guidelines and drug labels.



METHODS

Data collection

Pharmacogenomic information of drugs listed on the CPIC (https://cpicpgx.org/guidelines/) and DPWG (https://www.knmp.nl/patientenzorg/medicatiebewaking/farmacogenetica/ pharmacogenetics-1/pharmacogenetics) guidelines were reviewed and collected based on the following criteria: generic name of drug, drug related gene, genotype, and phenotype in accordance with genetic information. Based on all the drugs listed in the CPIC and DPWG guidelines, the drug labels of original or reference drugs were reviewed through the FDA (https:// www.accessdata.fda.gov/scripts/cder/daf/) and MFDS (https://nedrug.mfds.go.kr/searchDrug) databases. The following information was collected: contraindication, change of dose (e.g., dose reduction), biomarker testing requirement, and consistent recommendation with the CPIC or DPWG guidelines. All information and data were collected as of September 2020.

Definition

"Concordance" indicated that the pharmacogenomic information between the PGx guidelines and drug labels was identical. When pharmacogenomic information between the guidelines and drug labels was not identical or there was no pharmacogenomic information on the drug labels, it was classified as "discordance." Pharmacogenomic information belonging "concordance" was categorized as "contraindication," "change dose" and "biomarker testing required" depending on the PGx related indication on the drug labels. Indication of both 'changing dosage' or 'not changing dosage' depending on genotypes was included in "change dose."

Data analysis

All drugs listed in either the CPIC or DPWG guidelines were included in the analysis set, except drug categories classified as "This is not a gene-drug interaction" on the DPWG guideline. The drugs included in the analysis set were classified based on the anatomical therapeutic chemical (ATC) code by using the WHO-ATC/DDD Index (https://www.whocc.no/ atc_ddd_index/) and PGx related genes.

To match the pharmacogenomic information in the two guidelines to the drug labels, the PGx related section on the drugs labels of the FDA and MFDS was scrutinized, and the concordance of the pharmacogenomic information between the guidelines and the drug labels was confirmed. The ratio of concordance and discordance, indication category of concordance, and ATC code of concordant drugs were analyzed by FDA and MFDS. Drugs that were not approved by the FDA and MFDS were excluded from the analysis, respectively. Among the pharmacogenomic information of drugs that matched the guidelines to the drug labels, the concordance of the label between the FDA and MFDS was analyzed.

RESULTS

Classification of drugs in the CPIC and DPWG guidelines

A total of 66 and 78 drugs were screened from the CPIC and DPWG guidelines, respectively. Among the 78 drugs listed on the DPWG, 19 drugs were excluded because they were classified as "this is not a gene-drug interaction." The total number of drugs listed on either the CPIC or DPWG were 96 (**Table 1** and **Fig. 1**).

Label indication

Related Indication in the

Drug

	Genes		lelines									
		CPIC	DPWG		oncordance)		MFDS (concordance)					
				Contraindication	Change dose	Biomarker testing required		Contraindication	Change dose	Biomarker testing required	Discordance	
Abacavir	HLA-B	0	0	0	Х	0	Х	0	Х	0	Х	
Aceclofenac	CYP2C9	0	Х			NA		Х	Х	Х	0	
Acenocoumarol*	CYP2C9 VKORC1	Х	0			NA				NA		
Allopurinol	HLA-B	0	Х	Х	Х	Х	0	0	Х	0	Х	
Amitriptyline*	CYP2C9 CYP2C19	0	0	Х	Х	Х	0	Х	Х	Х	0	
Aripiprazole	CYP2D6	Х	0	Х	0	Х	Х	Х	Х	Х	0	
Aspirin	CYP2C9	0	Х	х	Х	Х	0	Х	Х	Х	0	
Atazanavir	UGT1A1	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Atomoxetine	CYP2D6	0	0	Х	0	Х	Х	Х	Х	Х	0	
Atorvastatin	SLCO1B1	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Azathioprine	TPMT	0	0	Х	0	0	х	Х	Х	Х	0	
Brexpiprazole	CYP2D6	Х	0	Х	0	Х	Х			NA		
Carvedilol*	CYP2D6	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Capecitabine	DPYD	0	0	Х	Х	Х	0	0	Х	0	Х	
Carbamazepine	HLA-B HLA-A	0	Х	0	х	0	Х	0	Х	0	Х	
Celecoxib	CYP2C9	0	Х	Х	0	Х	Х	Х	0	Х	Х	
Citalopram	CYP2C19	0	0	Х	0	Х	х	Х	0	Х	х	
Clomipramine*	CYP2D6 CYP2C19	0	0	Х	Х	Х	0	Х	Х	Х	0	
Clopidogrel	CYP2C19	0	0	х	х	0	х	Х	Х	х	0	
Codeine	CYP2D6	0	0	0	Х	Х	Х	0	Х	Х	Х	
Desflurane	RYR1	0	Х	х	х	х	0	Х	Х	х	0	
Desipramine	CYP2C19 CYP2D6	0	Х	Х	х	Х	0			NA		
Dexlansoprazole	CYP2C19	0	Х	х	х	х	0	Х	Х	х	0	
Diclofenac	CYP2C9	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Doxepine	CYP2D6	0	0	Х	Х	Х	0	Х	Х	Х	0	
Efavirenz	CYP2B6	0	0	Х	Х	Х	0	Х	Х	Х	0	
Eliglustat	CYP2D6	Х	0	х	0	0	х			NA		
Enflurane	RYR1	0	Х	Х	Х	Х	0			NA		
Escitalopram	CYP2C19	0	0	Х	Х	Х	0	Х	0	Х	Х	
Esomeprazole*	CYP2C19	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Flecainide	CYP2D6	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Flucloxacillin	HLA-B	Х	0			NA				NA		
Flucytosine	DPYD	Х	0	Х	Х	Х	0			NA		
Fluorouracil	DPYD	0	0	0	Х	Х	Х	0	Х	Х	Х	
Fluoxetine*	CYP2D6	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Flurbiprofen	CYP2C9	0	Х			NA		Х	Х	Х	0	
Fluvoxamine	CYP2D6	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Gefitinib [*]	CYP2D6	Х	0	Х	0	Х	Х	Х	0	Х	Х	
Glibenclamide*	CYP2C9	Х	0			NA		Х	Х	Х	0	
Gliclazide*	CYP2C9	Х	0			NA		Х	Х	Х	0	
Glimepiride*	CYP2C9	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Haloperidol	CYP2D6	Х	0	Х	Х	Х	0	Х	Х	Х	0	
Halothane	RYR1	0	Х			NA				NA		
Ibuprofen	CYP2C9	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Imipramine	CYP2C19 CYP2D6	0	0	Х	Х	Х	0	Х	Х	Х	0	
Indomethacin	CYP2C9	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Irinotecan*	UGT1A1	Х	0	Х	0	0	Х	Х	Х	Х	0	
Isoflurane	RYR1	0	Х	Х	Х	Х	0	Х	Х	Х	0	
Ivacaftor	CFTR	0	Х	0	Х	0	Х			NA		
Lansoprazole*	CYP2C19	0	0	Х	Х	Х	0	Х	Х	Х	0	

Table 1. Summary of concordant drugs listed in the PGx guidelines and on the drug labels

(continued to the next page)

Drug	Related Genes		on in the elines	he Label indication									
		CPIC	DPWG		FDA (c	oncol	rdance)			MFDS (c	oncordance)		
				Contraindication	Change dose	l tes	Biomarker ting required		Contraindication	Change dose	Biomarker testing required	Discordance	
Lornoxicam	CYP2C9	0	Х			NA			Х	Х	X	0	
Lumiracoxib	CYP2C9	0	Х			NA					NA		
Meloxicam	CYP2C9	0	Х	Х	Х		Х	0	Х	Х	Х	0	
Methoxyflurane	RYR1	0	Х			NA					NA		
Metamizole	CYP2C9	0	Х			NA					NA		
Mercaptopurine	TPMT	0	0	Х	Х		0	Х	0	Х	Х	Х	
Metoprolol	CYP2D6	Х	0	Х	Х		Х	0	Х	Х	Х	0	
Moclobemide*	CYP2C19	Х	0			NA			Х	Х	Х	0	
Nabumetone	CYP2C9	0	Х	Х	Х		Х	0	Х	Х	Х	0	
Naproxen	CYP2C9	0	Х	Х	Х		Х	0	Х	Х	Х	0	
Nortriptyline	CYP2C19 CYP2D6	0	0	Х	Х		Х	0	Х	Х	Х	0	
Omeprazole*	CYP2C19	0	0	Х	Х		Х	0	Х	Х	Х	0	
Ondansetron	CYP2D6	0	х	х	Х		Х	0	Х	х	Х	0	
Oxcarbazepine	HLA-B HLA-A	0	Х	Х	Х		0	Х	Х	Х	Х	0	
Oxycodone*	CYP2D6	х	0	х	х		Х	0	х	х	х	0	
Pantoprazole*	CYP2C19		0	X	X		X	0	X	X	X	0	
Paroxetine [*]	CYP2D6	0	0	X	X		X	0	X	X	X	0	
Peginterferon alfa-2a	IFNL3	0	X	X	X		X	0	X	X	X	0	
Peginterferon alfa-2b	IFNL3	0	X	X	X		X	0			NA	-	
Phenprocoumon*	VKORC1	X	0			NA		-			NA		
Phenytoin	CYP2C9	0	0	х	Х		Х	0	х	Х	Х	0	
Pimozide	CYP2D6	X	0	X	Х		0	X	X	Х	X	0	
Piroxicam	CYP2C9	0	X	X	X		X	0	X	X	X	0	
Propafenone	CYP2D6	Х	0	Х	Х		Х	0	Х	Х	Х	0	
Rabeprazole*	CYP2C19		0	х	Х		Х	0	х	Х	х	0	
Rasburicase	G6PD	0	Х	0	Х		0	Х	0	Х	Х	Х	
Ribavirin	IFNL3	0	Х	Х	Х		Х	0	х	Х	Х	0	
Risperidone*	CYP2D6	Х	0	Х	Х		Х	0	Х	Х	Х	0	
Sertraline	CYP2C19	0	0	Х	Х		Х	0	х	Х	Х	0	
Sevoflurane	RYR1	0	Х	Х	Х		Х	0	Х	Х	Х	0	
Simvastatin	SLCO1B1	0	0	Х	Х		Х	0	х	Х	Х	0	
Siponimod	CYP2C9	Х	0	Х	0		Х	Х			NA		
Succinylcholine	RYR1	0	Х	Х	Х		Х	0	х	Х	Х	0	
Tacrolimus	CYP3A5	0	0	Х	Х		Х	0	Х	Х	Х	0	
Tamoxifen	CYP2D6	0	0	Х	Х		Х	0	Х	Х	Х	0	
Tegafur	DPYD	0	0			NA			Х	Х	Х	0	
Tenoxicam	CYP2C9	0	Х			NA					NA		
Thioguanine	TPMT	0	0	Х	0		0	Х			NA		
Tolbutamide*	CYP2C9	Х	0			NA					NA		
Tramadol	CYP2D6	Х	0	0	Х		Х	Х	Х	Х	Х	0	
Trimipramine	CYP2D6 CYP2C19	0	Х			NA					NA		
Tropisetron	CYP2D6	0	Х			NA					NA		
Venlafaxine	CYP2D6	X	0	Х	х		Х	0	х	Х	X	0	
Voriconazole	CYP2C19		0	X	X		X	0	X	X	X	0	
Warfarin	CYP2C9 VKORC1	0	0	X	0		X	x	x	0	x	x	
Zuclopenthixol	CYP2D6	Х	0			NA					NA		
Total No.	0200	66	59	7 (9.0%)	12 (15.4%		12 (15.4%)	54 (69.2%)	8 (10.7%)	5 (6.7%)	4 (5.3%)	62 (82.7%)	

Table 1. (Continued) Summary of concordant drugs listed in the PGx guidelines and on the drug labels Delated Indiantian in the

Data are presented as the total number and ratio (%) of drugs and total number of drugs with pharmacogenetic information in the CPIC and/or DPWG guideline except for not approved drugs (total FDA = 78, MFDS = 75).

CPIC, Clinical Pharmacogenetics Implementation Consortium; DPWG, Dutch Pharmacogenetics Working Group; FDA, US Food and Drug Administration; MFDS, Korea Ministry of Food and Drug Safety.

*Drug represents that no action is required for this gene-drug interaction. NA indicates that the drug is not approved in United States and/or Republic of Korea.

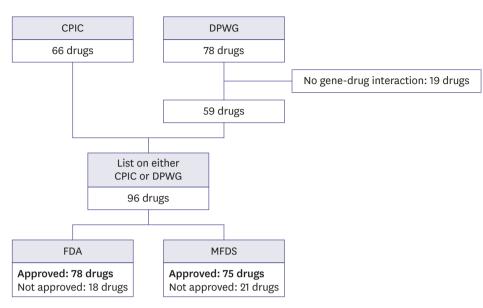


Figure 1. Selection of drugs for comparison between the pharmacogenomics guidelines and drug labels of the regulatory affairs.

CPIC, Clinical Pharmacogenetics Implementation Consortium; DPWG, Dutch Pharmacogenetics Working Group; FDA, US Food and Drug Administration; MFDS, Korea Ministry of Food and Drug Safety.

The most common type of drug classified by ATC code was nervous system in both the CPIC (33.3%) and DPWG (37.3%) guidelines (**Fig. 2A**). In the case of PGx related genes, CYP2D6 and CYP2C19 were dominant genes in both guidelines (**Fig. 2B**).

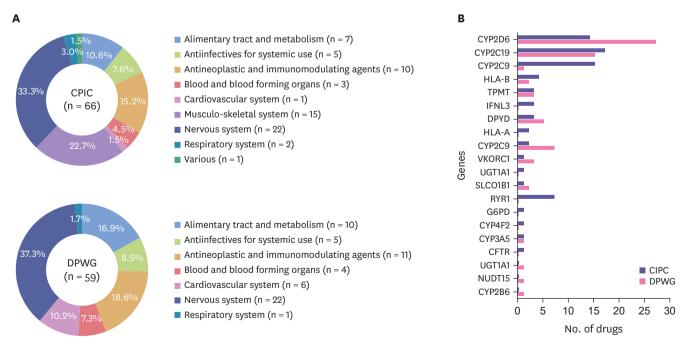


Figure 2. Classification of drugs listed in the CPIC and DPWG guidelines by (A) the anatomical therapeutic chemical code and (B) pharmacogenomics related genes. Drug categories classified as "This is not a gene-drug interaction" on the DPWG guideline.

CPIC, Clinical Pharmacogenetics Implementation Consortium; DPWG, Dutch Pharmacogenetics Working Group.



Pharmacogenomic information from PGx guidelines and drug labels

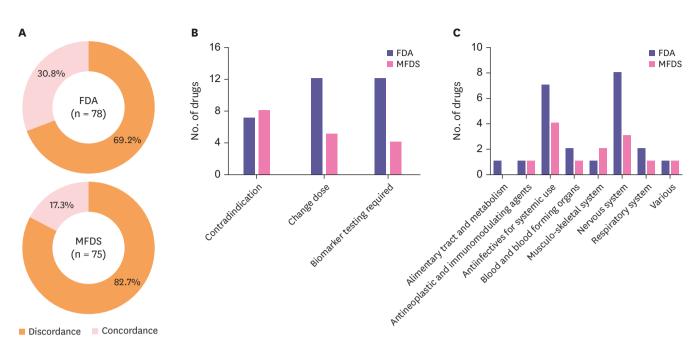


Figure 3. Concordance between the pharmacogenomic information in the guidelines and on the drug labels (A) pie chart showing the ratio of the concordance and discordance of the FDA and MFDS (B) bar chart representing the number of concordant drugs categorized by indication (C) bar chart representing the number of concordant drugs categorized by the anatomical therapeutic chemical code. FDA, US Food and Drug Administration; MFDS, Korea Ministry of Food and Drug Safety.

Concordance of the pharmacogenomic information between the guidelines and drug labels

Among the 96 drugs listed on the PGx guidelines, 78 and 75 drugs were approved by the FDA and MFDS, respectively. In the FDA labels, the number of concordant drugs was 24 (30.8%), while 13 drugs (17.3%) were concordant to the MFDS labels (**Fig. 3A**). In detail, the number of drugs categorized as contraindication, change dose and biomarker testing required was 7, 12 and 12 in the FDA and 8, 5 and 4 in the MFDS, respectively. Nervous system and antineoplastic and immunomodulation agents were dominant drugs among concordant drugs.

Concordance of pharmacogenomic information between the FDA and MFDS labels

The drugs that had the same pharmacogenomic information between the FDA and MFDS were as follow: abacavir, carbamazepine, celecoxib, citalopram, codeine, fluorouracil, geftinib, rasburicase and warfarin (**Table 2**). Abacavir, carbamazepine, codeine, fluorouracil and rasburicase were prohibited to people with certain genotypes. In the case of carbamazepine, biomarker test for HLA-B*1502 should be conducted prior to treatment. The dosage of celecoxib, citalopram and warfarin should be changed depending on the genotype, while the dosage of geftinib did not need to be adjusted regardless of the genotype.

DISCUSSION

The comparison of pharmacogenomic information between the guidelines and drug labels was conducted to identify whether the pharmacogenomic information in guidelines was well reflected on the drug labels which was easily accessible information for physicians and pharmacists. As a result of the study, a total of 96 drugs listed in either the CPIC or DPWG guidelines were screened, and 78 and 75 drug labels of the FDA and MFDS, respectively, were

Drug	Gene	CPIC	DPWG	FDA/MFDS	Indication
Abacavir	HLA-B	Abacavir is not recommended for carrier of HLA-B*57:01 because of significantly increased risk of abacavir hypersensitivity	Abacavir is contra- indicated for HLA-B*5701- positive patients	Patients who carry the HLA-B*5701 allele are at a higher risk of experiencing a hypersensitivity reaction to abacavir	Contraindication
Carbamazepine	HLA-B	If patient is carbamazepine- naive and alternative agents are available, do not use carbamazepine	NA	Patients with ancestry in genetically at-risk populations should be screened for the presence of HLA-B*1502 prior to initiating treatment with carbamazepine. Patients testing positive for the allele should not be treated with carbamazepine unless the benefit clearly outweighs the risk	Contraindication, Biomarker testing required
Celecoxib	CYP2C9	Initiate therapy with 25–50% of the lowest recommended starting dose	NA	In patients who are known or suspected to be poor CYP2C9 metabolizers (i.e., CYP2C9*3/*3), based on genotype or previous history/experience with other CYP2C9 substrates administer celecoxib starting with half the lowest recommended dose	Change dose
Citalopram	CYP2C19	Consider a 50% reduction of the recommended starting dose. Utilize therapeutic drug monitoring to guide dose adjustments	Do not exceed the following daily doses (50% of the standard maximum dose)	Dosage adjustment is recommended in CYP2C19 poor metabolizers	Change dose
Codeine	CYP2D6	Avoid codeine use due to potential for toxicity	Codeine is contra- indicated	Ultra-rapid metabolism of codeine and other risk factors for life-threatening respiratory depression in children	Contraindication
Fluorouracil	DYPD	Avoid use of 5-fluorouracil or 5-fluorouracil prodrug-based regimens	Start with 50% of the standard dose or avoid fluorouracil	Increased risk of serious or fatal adverse reactions in patients with low or absent dipyrimidine dehydrogenase activity	Contraindication
Geftinib	CYP2D6	NA	NO action is needed for this gene-drug interaction	No dose adjustment is recommended in patients with a known CYP2D6 poor metabolizer genotype, but these patients should be closely monitored for adverse reactions	Change dose
Rasburicase	G6PD	Rasburicase is contraindicated	NA	Do not administer rasburicase to patients with glucose-6phosphate dehydrogenase (G6PD) deficiency	Contraindication
Warfarin	CYP2C9	Dosing Recommendations with Consideration of Genotype	Dosing Recommendations with Consideration of Genotype	Dosing Recommendations with Consideration of Genotype	Change dose

Table 2. List of concordant drugs for which the pharmacogenetic information on the FDA and MFDS labels was identical

NA indicates that the drug is not approved in United States and/or Republic of Korea.

FDA, US Food and Drug Administration; MFDS, Korea Ministry of Food and Drug Safety; CPIC, Clinical Pharmacogenetics Implementation Consortium; DPWG, Dutch Pharmacogenetics Working Group.

scrutinized. Less than half of the pharmacogenomic information from the approved drugs was consistent with the PGx guidelines.

Inconsistency between the CPIC and DPWG guidelines could be one of the factors why PGx related recommendations by guidelines were not reflected on the drug labels. Bank et al. reported that CPIC and DPWG showed a high level of consistency; however, some differences between them existed such as methodology and terminology [12]. Methodology for the scaling level of evidence, utilizing the source of information and processing establishment of dose recommendation between the two PGx guidelines was inconsistent. Like the example that "increased function" referred to "greater than normal function" in the CPIC guideline had a similar meaning of "gain-of-function" in the DPWG guideline; thus, there was subtle gap for the terminology between the two guidelines. These methodological and terminological differences possibly hindered the integration of pharmacogenomic information, leading to inconsistency between the guidelines and drug labels.

Above all, the main reason of discordance of the pharmacogenomic information between the guidelines and labels is different characteristics as informative tools. PGx guidelines are generally established and updated by various experts in the field based on the latest knowledge. The underlying assumption for the guidelines is that genotyping will be common, and establishing guidelines enables clinicians to translate genotyping results into actionable clinical decisions [5]. Drug labels are released after permission from the reviewers of regulatory affairs under a comprehensive and meticulous regulatory principle. The drug label can be revised if strong evidence and the potential impact of the gene-drug response are supported [13]. Therefore, PGx guidelines tend to reflect up-to-date knowledge, whereas drug labels are relatively slowly updated.

Among approved drugs from both the FDA and MFDS, the pharmacogenomic information of only 9 drugs was identical. One of the reasons for the inconsistent information between the FDA and MFDS might be the ethnic characteristics of US and South Korean citizens in that South Korea is close to a mono-ethnicity, while the United States is a poly-ethnicity with a small percentage of Asians. Ethnicity is one of the factors that affect the pharmacokinetics, pharmacodynamics and PGx of a drug, resulting in variability of the drug response, and a difference in the frequency of genetic polymorphisms of the drug-metabolizing enzymes between Caucasian and East Asians were reported [11,14]. In addition, a difference in the system of establishing or revising drug labeling between the two agencies possibly triggered a gap between the FDA and MFDS drug labels.

One of the limitations of this study was that the labels of the European Medicines Agency and Pharmaceuticals and Medical Devices Agency were not considered. Also, the other limitation is that some labeling information between original and generic drugs containing same active pharmaceutical ingredient is inconsistent [15]. To minimize bias, the original and reference drug were selected in the process of collecting the pharmacogenomic information of the drug labels in this study.

In conclusion, pharmacogenomic information on clinical implementation guidelines was limited in both FDA and MFDS labels because of various reasons including the characteristics of the guidelines and drug labels. Therefore, more effort from pharmaceutical companies, academia and regulatory affairs needs to be made to implement pharmacogenomic information onto drug labels.

REFERENCES

- Jameson JL, Longo DL. Precision medicine--personalized, problematic, and promising. N Engl J Med 2015;372:2229-2234.
 PUBMED | CROSSREF
- Cremers S, Guha N, Shine B. Therapeutic drug monitoring in the era of precision medicine: opportunities! Br J Clin Pharmacol 2016;82:900-902.
 PUBMED | CROSSREF
- Weinshilboum RM, Wang L. Pharmacogenomics: precision medicine and drug response. Mayo Clin Proc 2017;92:1711-1722.
 - PUBMED | CROSSREF
- van der Wouden CH, Cambon-Thomsen A, Cecchin E, Cheung KC, Dávila-Fajardo CL, Deneer VH, et al. Implementing pharmacogenomics in Europe: design and implementation strategy of the ubiquitous pharmacogenomics consortium. Clin Pharmacol Ther 2017;101:341-358.
 PUBMED | CROSSREF
- Caudle KE, Klein TE, Hoffman JM, Muller DJ, Whirl-Carrillo M, Gong L, et al. Incorporation of pharmacogenomics into routine clinical practice: the Clinical Pharmacogenetics Implementation Consortium (CPIC) guideline development process. Curr Drug Metab 2014;15:209-217.
 PUBMED | CROSSREF

- Blagec K, Romagnoli KM, Boyce RD, Samwald M. Examining perceptions of the usefulness and usability of a mobile-based system for pharmacogenomics clinical decision support: a mixed methods study. PeerJ 2016;4:e1671.
 PUBMED | CROSSREF
- Altman RB, Prabhu S, Sidow A, Zook JM, Goldfeder R, Litwack D, et al. A research roadmap for nextgeneration sequencing informatics. Sci Transl Med 2016;8:335ps10.
- 8. US Food and Drug Administration. Guidance for industry: clinical pharmacogenomics: premarket evaluation in early-phase clinical studies and recommendations for labeling. January 2013. https://www.fda.gov/media/84923/download. Accessed August 8, 2020.
- Abou Diwan E, Zeitoun RI, Abou Haidar L, Cascorbi I, Khoueiry Zgheib N. Implementation and obstacles of pharmacogenetics in clinical practice: An international survey. Br J Clin Pharmacol 2019;85:2076-2088. PUBMED | CROSSREF
- Klein ME, Parvez MM, Shin JG. Clinical implementation of pharmacogenomics for personalized precision medicine: barriers and solutions. J Pharm Sci 2017;106:2368-2379.
 PUBMED | CROSSREF
- Yasuda SU, Zhang L, Huang SM. The role of ethnicity in variability in response to drugs: focus on clinical pharmacology studies. Clin Pharmacol Ther 2008;84:417-423.
 PUBMED | CROSSREF
- Bank PC, Caudle KE, Swen JJ, Gammal RS, Whirl-Carrillo M, Klein TE, et al. Comparison of the guidelines of the Clinical Pharmacogenetics Implementation Consortium and the Dutch Pharmacogenetics Working Group. Clin Pharmacol Ther 2018;103:599-618.
 PUBMED | CROSSREF
- Drozda K, Pacanowski MA, Grimstein C, Zineh I. Pharmacogenetic labeling of FDA-approved drugs: a regulatory retrospective. JACC Basic Transl Sci 2018;3:545-549.
 PUBMED | CROSSREF
- 14. Kim K, Johnson JA, Derendorf H. Differences in drug pharmacokinetics between East Asians and Caucasians and the role of genetic polymorphisms. J Clin Pharmacol 2004;44:1083-1105. PUBMED | CROSSREF
- Pfistermeister B, Saß A, Criegee-Rieck M, Bürkle T, Fromm MF, Maas R. Inconsistencies and misleading information in officially approved prescribing information from three major drug markets. Clin Pharmacol Ther 2014;96:616-624.
 PUBMED | CROSSREF