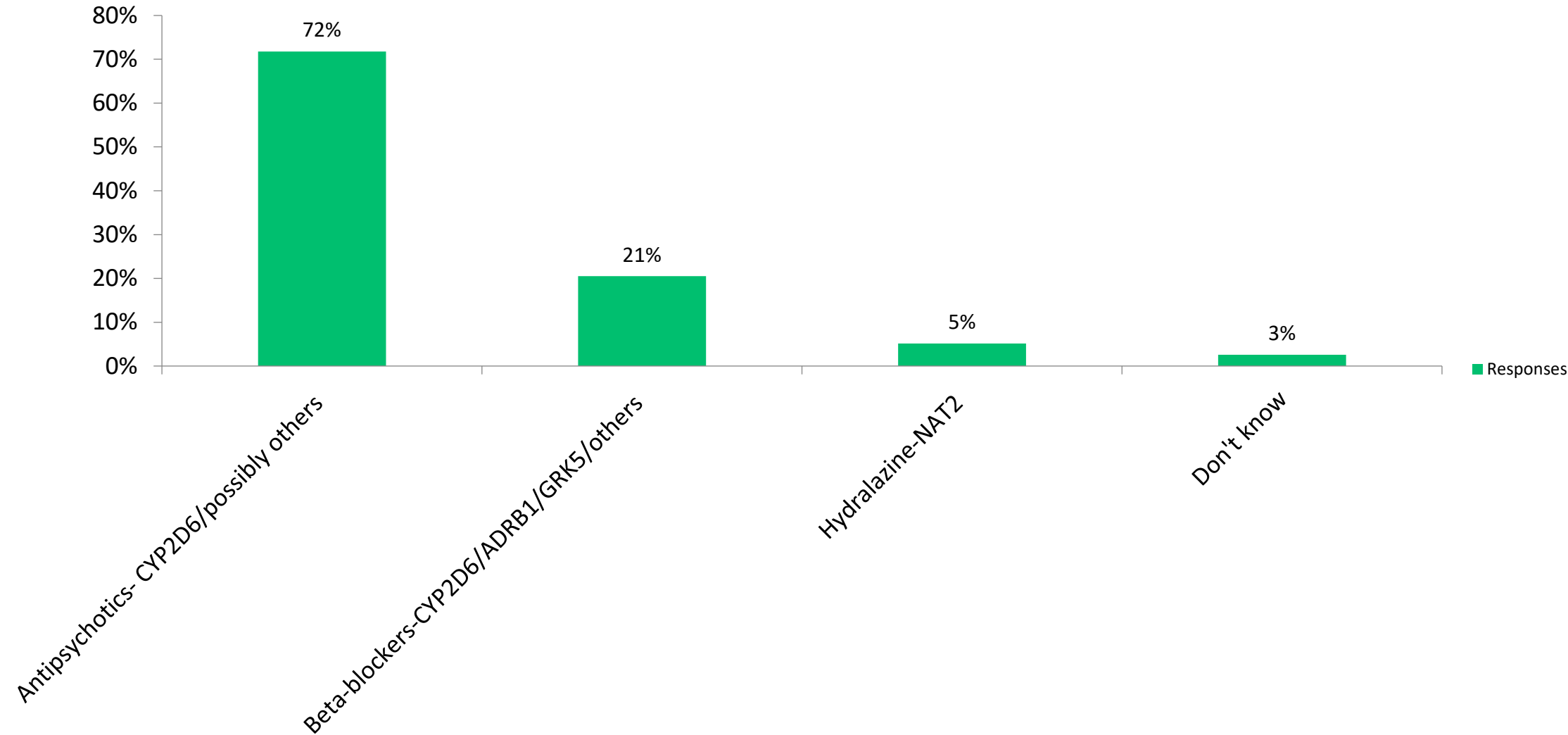


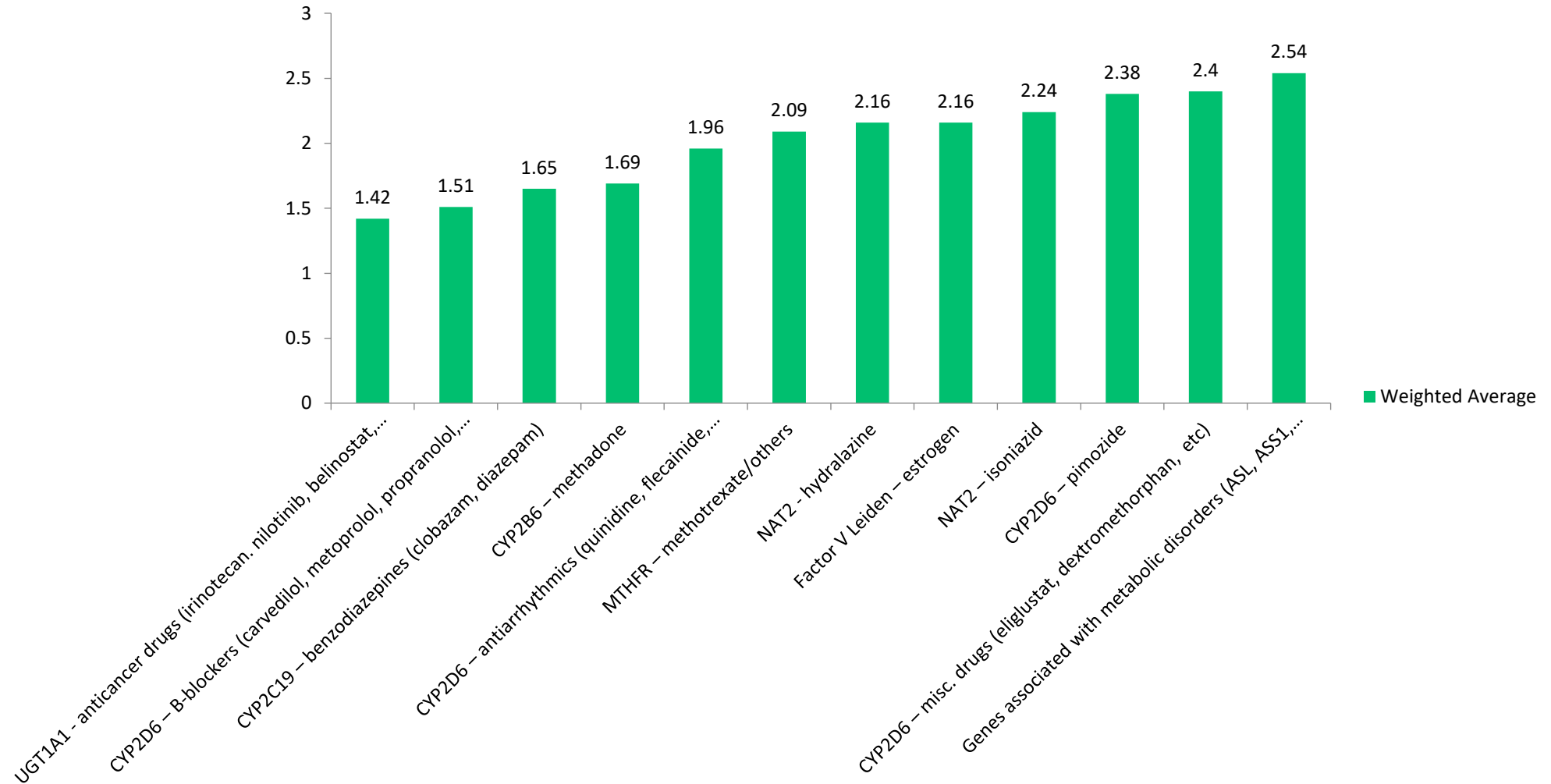
Member survey for guideline  
prioritization

Of these 3, which do you consider the highest priority or respond “don’t know”

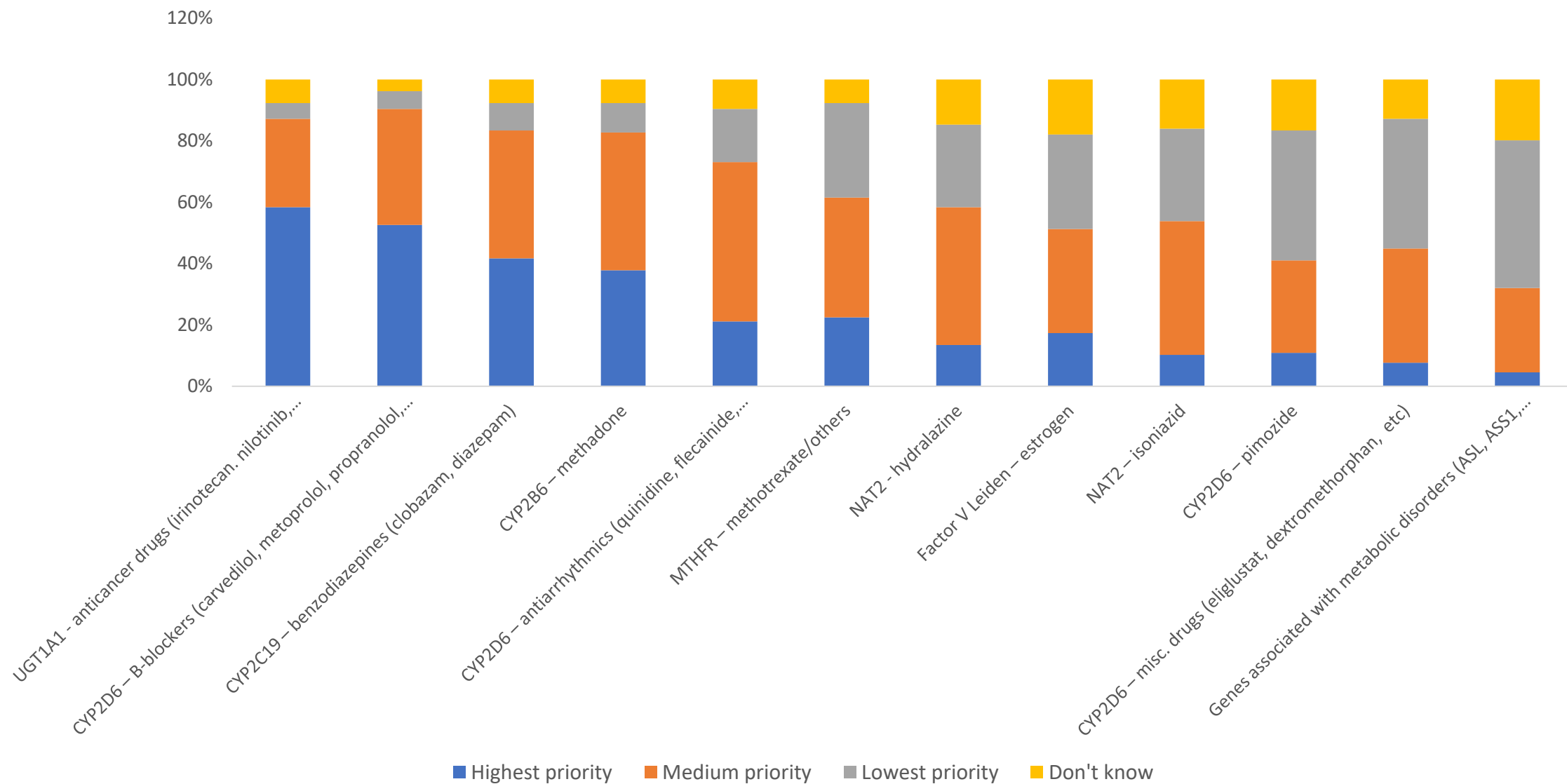


n=156

Rank the following gene/drug groups for their priority as the subject of a new CPIC guideline (highest priority, medium priority, lowest priority,



n=156



# Guideline prioritization

Antipsychotics- CYP2D6/possibly others

CYP2D6 – B-blockers (carvedilol, metoprolol, propranolol, timolol)

UGT1A1 - anticancer drugs (irinotecan, nilotinib, belinostat, others)

CYP2C19 – benzodiazepines (clobazam, diazepam)

CYP2B6 – methadone

CYP2D6 – antiarrhythmics (quinidine, flecainide, propafenone)

MTHFR – methotrexate/others

NAT2 - hydralazine

Factor V Leiden – estrogen

NAT2 – isoniazid

CYP2D6 – pimozide

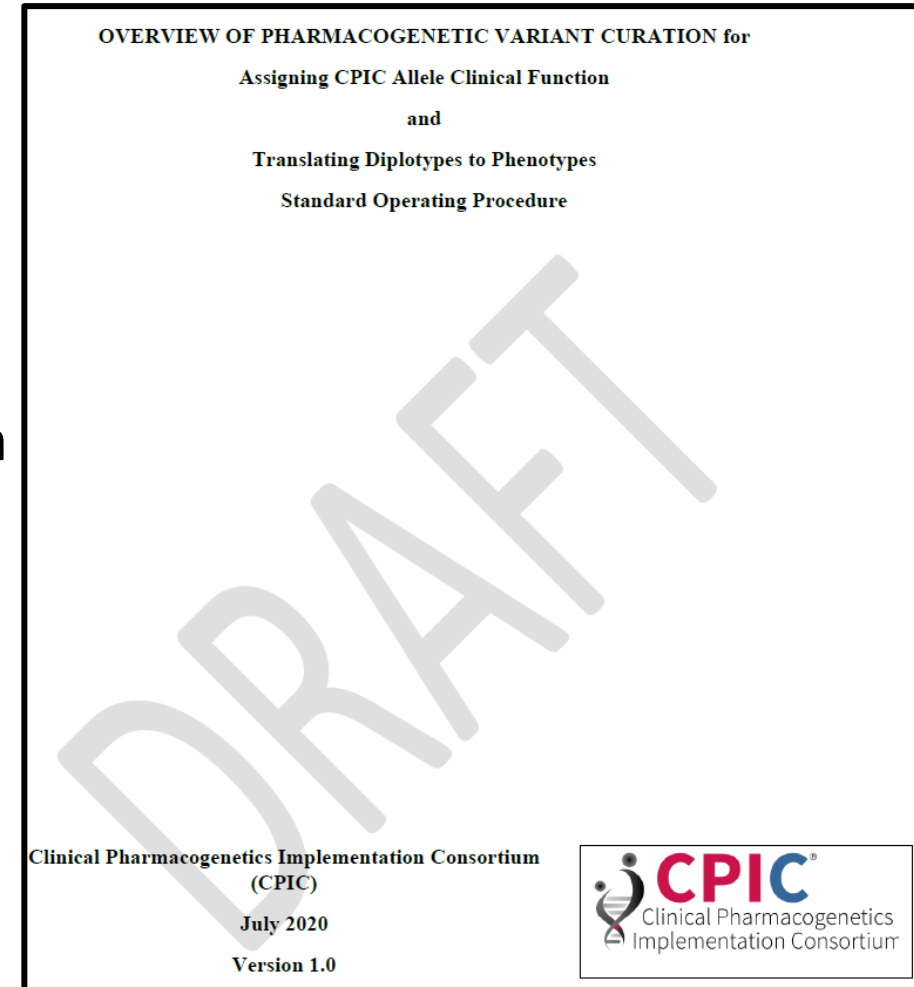
CYP2D6 – misc. drugs (eliglustat, dextromethorphan, etc)

Genes associated with metabolic disorders (ASL, ASS1, CPS1, NAGS, OTC, GBA, HPRT1, NAGS, POLG, etc) – various drugs (valproic acid, velagluceras alfa, mycophenolic acid, carglumic acid, etc)

Updated allele function SOP

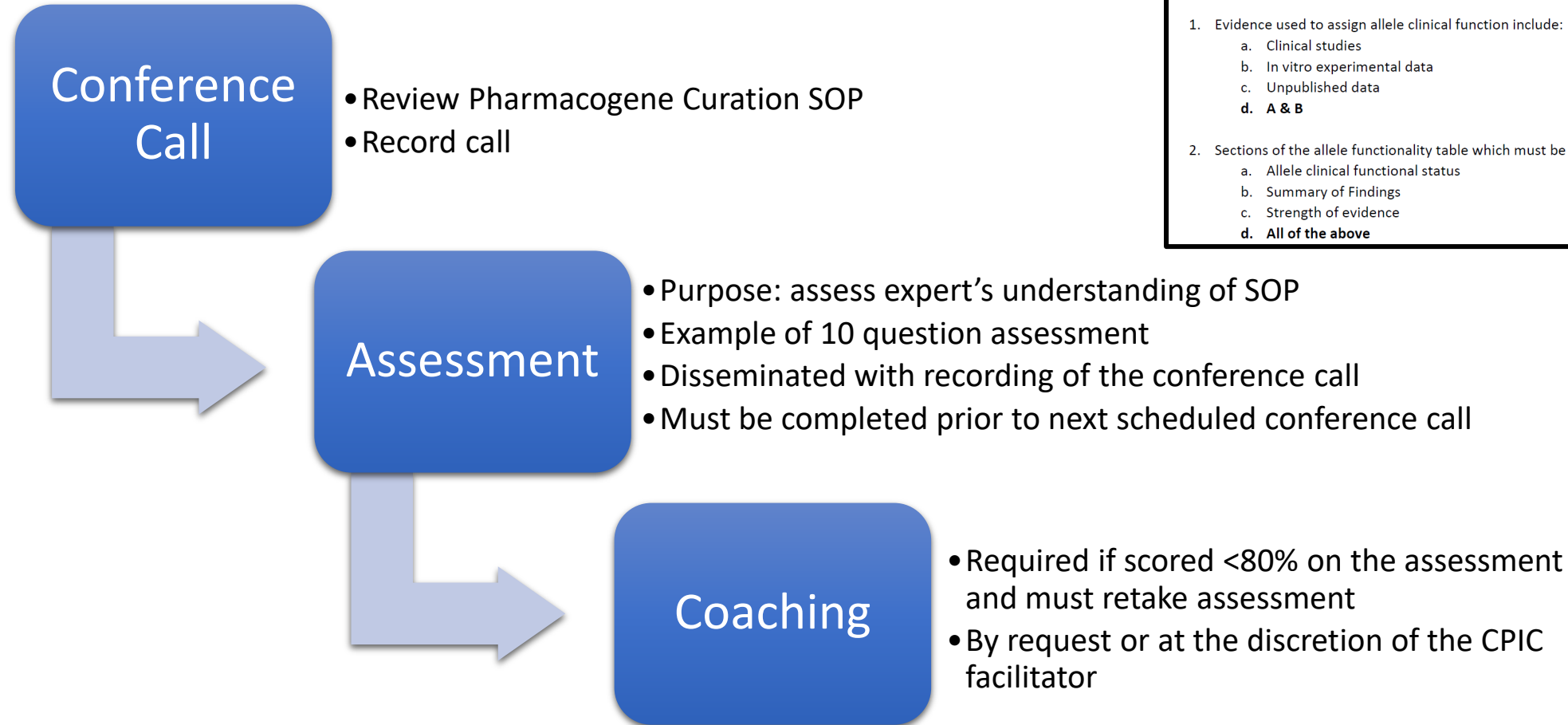
# Pharmacogene Curation SOP

- Describes procedures for
  - expert selection and training
  - evidence collection and inclusion of alleles
  - assigning strength of evidence and allele function
  - summarizing the evidence
  - translating diplotypes to phenotypes
  - re-evaluations and updates



# Professional Training

- Required for all experts



**CPIC Assessment of Expert Understanding of Pharmacogene Curation SOP**

Author Expert: \_\_\_\_\_ Date: \_\_\_\_\_

Please mark the correct answers and submit.

1. Evidence used to assign allele clinical function include:

- Clinical studies
- In vitro experimental data
- Unpublished data
- A & B**

2. Sections of the allele functionality table which must be completed for every allele include:

- Allele clinical functional status
- Summary of Findings
- Strength of evidence
- All of the above**



# Scoring the Evidence Process Overview

- Two authors will independently
  - assign strength of evidence
  - assign allele clinical function
  - assign activity value for genes using activity scores
  - summarize the evidence
- Independent evaluations are compiled into a working draft of the Allele functionality table
- Table is disseminated to all other authors for review

Allele/cDNA/rsID	Activity Value (Optional)	Allele Biochemical Functional Status (Optional)	Allele Clinical Functional Status (Required)	Allele Clinical Function Substrate Specificity (Optional)	PMID (Required)	Strength of Evidence (Required)	Summary of Findings (Required)
*1	1	Normal Function	Normal Function				
*2	0.5	Decreased function	Decreased function		29283396; 20150829; 15637526; 12742136; 10413320; 27179628; 25775139; 19082874;	Definitive	CYP2C9*2 is assigned decreased function based on definitive evidence in homozygous and heterozygous patients and in vitro experimental data. CYP2C9*2 has been studied for more than 20 years and its association with impaired function compared to wildtype is well-established (29283396, 20150829, 15637526). The phenotype for patients homozygous for the CYP2C9*2 variant was updated from poor metabolizer to intermediate metabolizer after reevaluation of prior and new evidence demonstrating CYP2C9*2 has more enzyme activity than CYP2C9*3, which results in a similar phenotype for

# Evidence Summary

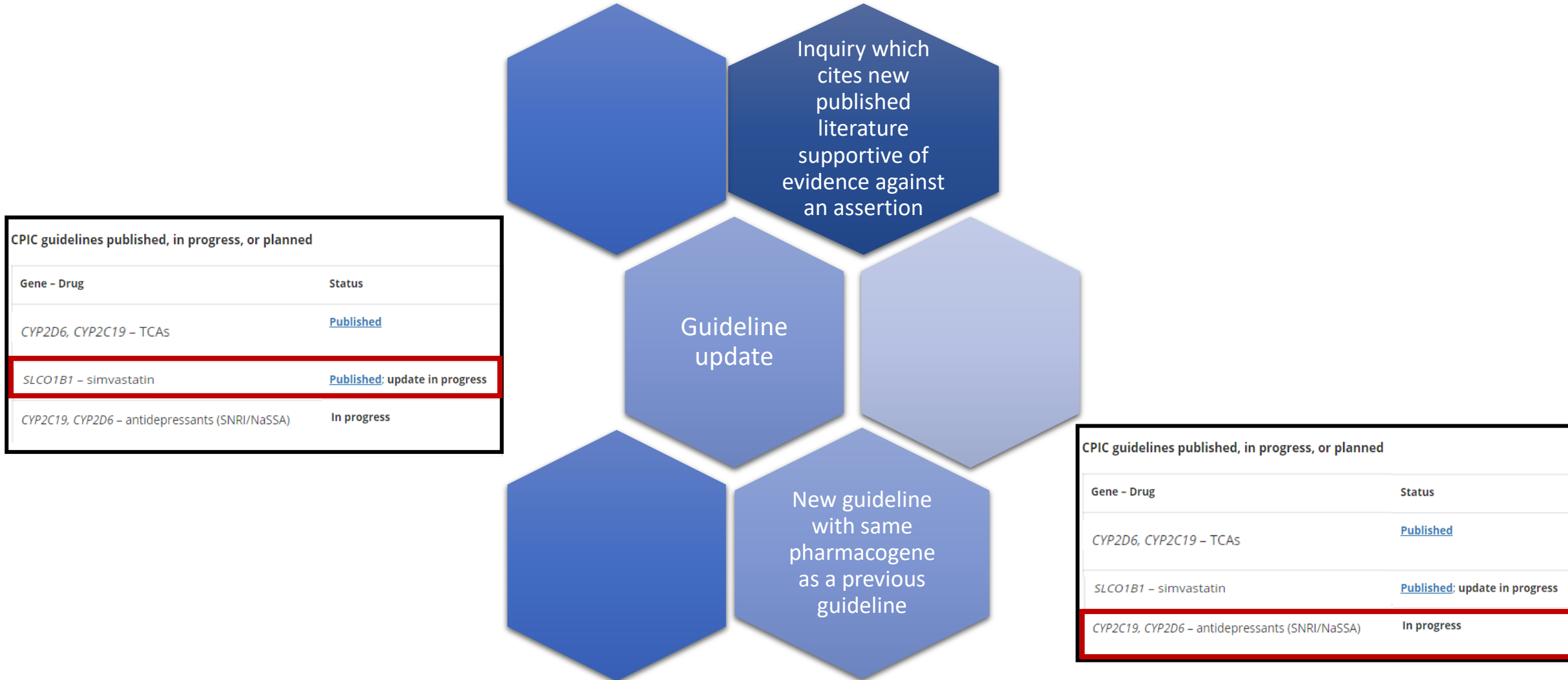
- (a) summarizes the evidence supporting the assigned allele function
  - must note if it is a well-established association that did not require a primary literature review
  - may point to an authoritative resource for allele function
- (b) notes conflicting evidence and summarizes the experts' assessment of this evidence
  - includes consensus and rationale for final assertion if there was disagreement among experts
- (c) notes if an assertion was modified based on clinical expertise
  - includes the original assertion and the experts' rationale for modification

# Expert Attestation

- Attestation of expert review and agreement with the assertions in the evidence table is required
- Documents consensus achieved among at least 70% of experts
- Ensures evidence summaries adequately capture consensus, rationale, and acknowledgment of disagreements

Expert Attestation of Review and Agreement with CPIC assertions	
Gene table update review for _____ (gene)	Date gene table circulated for review: _____
Author Expert: _____	Date of attestation: _____
Please review and submit.	
The Pharmacogene Curation SOP was followed when creating the allele functionality table, phenotype mapping table, and diplotype to phenotype table.	
<input type="checkbox"/> I agree	
<input type="checkbox"/> I do not agree	
I have reviewed the allele functionality table and agree with the assertions made for allele clinical function, strength of evidence, and summary of findings for every allele.	
<input type="checkbox"/> I agree	
<input type="checkbox"/> I do not agree	
I have reviewed the phenotype mapping table and diplotype to phenotype table and agree with the documented conventions and translations of the diplotypes to phenotypes.	
<input type="checkbox"/> I agree	
<input type="checkbox"/> I do not agree	
For any statement not agreed to please specify below which aspects of the reviewed material you do not agree with. Please identify specific assertions and provide rationale for disagreement.	

# Three ways to initiate re-evaluations and updates



# Genes with updated tables

- CYP2C9
  - Sent to CYP2C9 authors for review
  - Authors need to take the assessment and attestation
- CYP2C19
  - Authors of current CYP2C19 guidelines (clopidogrel and SSRI) will go through process of re-evaluations
  - Once final, will send to other CYP2C19 guideline authors (e.g., voriconazole, TCAs, PPIs, etc) for approval
  - All authors need to take the assessment and attestation
- CYP2D6-underway
- SLCO1B1-guideline being updated now

# Process for re-evaluations due to updated guideline or guideline with same pharmacogene as a previous guideline

