## **Contents**

Chapter 1	2
Chapter 2	2
Table 1: PharmGKB 'Very Important Pharmacogenes' as of 2020	2
Chapter 3	7
Table 2: Full data extraction of trials included in Chapter 3 biomarker review	7
Table 3: Full list of evidence cited by TARGET trial in biomarker review	8
Table 4: Full list of evidence cited by EU-PACT trial in biomarker review	g
Table 5: Full list of evidence cited by SHIVA trial in biomarker review	10
Table 6: Full list of evidence cited by GIST trial in biomarker review	11
Table 7: Full list of evidence cited by Precision Medicine Guided Treatment for Cancer Pain trial in biomarker review	12
Chapter 4	13
4.1 Protocol for systematic reviews and meta-analyses	13
4.2 Standard data extraction form for systematic reviews	13
4.3 Full list of studies included in systematic reviews and meta-analyses	24
4.4 Quality assessment of studies included in systematic reviews and meta-analyses	24
4.4.1 HLA-B*15:02	24
4.4.2 HLA-A*31:01	24
4.5 Calculation of allele frequencies for HLA-B*15:02 and HLA-A*31:01	24
4.6 Simulation code	24
Table 8: Comparison of papers included in previous meta-analyses and our meta-analysis	24
Chapter 5	26
Table 9: Previous systematic reviews of discrete choice experiments	26
Table 10: Search terms used by previous systemic reviews to locate discrete choice experiments	

Table 11: Data extraction sheet prepared for systematic review of discrete choice experiments	30
Table 12: Data extracted from each paper in the systematic review of discrete choice experiments (1)	32
Table 13: Data extracted from each paper in the systematic review of discrete choice experiments (2)	35
Chapter 6	37
6.1 Survey of healthcare professionals	37
6.2 Survey of patients	55
6.3 Final discrete choice experiment, as shown to participants (abacavir example)	66
Chapter 7	124
7.1 Stata code for DCE analysis	
7.2 Data matrices for each DCE	125
7.3 Bootstrapped beta-coefficients for each DCE	125
7.4 Utility modelling for each DCE	125
Appendix References	125

NA

# Chapter 2

Table 1: PharmGKB 'Very Important Pharmacogenes' as of 2020

Gene	Gene product	Drug(s)	Overview of guideline (s)

ABCB1	P-glycoprotein (P-gp)	Many, including: anti-depressants, anti-virals, chemotherapeutics, opioids, steroids,	Encodes a transporter protein, which effluxes many substrates. This leads to a number of drug-drug interactions (e.g. rifampin and contraceptives) that can reduce efficacy. Also responsible for drug resistance in some cancers (1).
ABCG2	BCRP transporter protein	Many, including: anti-virals, chemotherapeutics, anti-fungals	Encodes a transporter protein, similar to P-gp. Leads to a number of drug-drug interactions (e.g. pitavastatin and cyclosporin), and can reduce efficacy. Responsible for drug resistance in some cancers (2).
ACE	ACE enzyme	ACE inhibitors, statins	Contradictory reports on the effect of <i>ACE</i> variants on ACE inhibitor response, but most focus on just one variant so further variant testing required (3).
ADRB1	Beta-1-adrenergic receptor	Anti-hypertensives, beta-blockers	The beta-1-adrenergic receptor mediates heart rate and contractility. Variants affect the efficacy of drugs treating hypertension, coronary artery disease, and heart failure (4, 5).
ADRB2	Beta-2-adrenergic receptor	Anti-hypertensives, beta-blockers	This receptor is expressed in cardiac myocytes and in bronchial and vascular smooth muscle cells. Variants affect the efficacy of drugs treating hypertension, congestive heart failure, and asthma (6).
CACNA1S	Dihydropyridine receptor (DHPR) [alpha subunit]	Volatile anaesthetics	DHPR is a voltage-gated calcium channel in skeletal muscle. Variants are linked to malignant hyperthemia susceptibility and hypokalaemic periodic paralysis (7).
CFTR	CFTR protein	Ivacaftor	Variants in <i>CFTR</i> cause cystic fibrosis. Over 1800 variants have been reported. Ivacaftor targets several specific <i>CFTR</i> variants, responsible for 4-5% of cystic fibrosis cases (8).
COMT	Catechol-O- methyltransferase	Many, including: anti-psychotics, drugs for management of Parkinson's disease, opioids,	Variants in <i>COMT</i> are associated with requiring higher doses of drugs in schizophrenia and higher doses of morphine in pain (9).
CYP2A6	CYP2A6 enzyme  Many, including: anti-virals, chemotherapeutics, nicotine, steroids		Encodes an enzyme affects the metabolism of nicotine (polymorphisms are linked to smoking behaviours). Also has a key role in the metabolism of many other drugs (10).

CYP2B6	CYP2B6 enzyme	Many, including: anti-depressants, anti-virals, chemotherapeutics	Encodes an enzyme responsible for metabolism of 4% of the top 200 drugs. Variants are associated with increased risk of ADRs from cyclophosphamide, efavirenz, and bupropion (11).				
CYP2C19	CYP2C19 enzyme	Many, including: anti-depressants, anti-platelet drugs, proton pump inhibitors	Encodes a liver enzyme with polymorphisms leading to reduced or absent enzyme activity. Loss-of-function mutations are associated with lower efficacy of several drugs (12).				
CYP2C8	CYP2C8 enzyme	Many, including: anti-diabetics, chemotherapeutics, opioids, statins	Encodes a liver enzyme that metabolises several large compounds. Has many polymorphisms associated with variability in drug response (13).				
CYP2C9	CYP2C9 enzyme	Many, including: anticoagulants, chemotherapeutics, NSAIDs, statins	Encodes a liver enzyme responsible for metabolic clearance of 15-20% of drugs. This leads to drug-drug interactions, reduced efficacy, or increased risk of ADRs, depending on the variant and the drug (14).				
CYP2D6	CYP2D6 enzyme	Many, including: anti-depressants, anti-hypertensives, chemotherapeutics, opioids	Encodes a liver enzyme involved in metabolism of up to 25% of commonly-used drugs. Leads to drug-drug interactions, reduced efficacy, or increased risk of ADRs, depending on the variant and the drug (15).				
CYP3A4	CYP3A4 enzyme	Many, including: anti-platelet drugs, chemotherapeutics, antibiotics	Encodes an enzyme that is responsible for metabolism of 40-50% of drugs in use. Different polymorphisms responsible for variability in response to many drugs (16).				
CYP3A5	CYP3A5 enzyme	Many, including: anti-virals, benzodiazepines, chemotherapeutics	Encodes an enzyme responsible for metabolism of many common drugs. Different polymorphisms responsible for variability in response to many drugs (17).				
CYP4F2	CYP4F2 enzyme	Vitamins E and K, anti-parasitic drugs	Encodes an enzyme that metabolises many endogenous compounds, affecting the dosing of warfarin (18).				
DPYD	DPYD enzyme Fluoropyrimidines		Polymorphisms that lead to decreased DPYD activity increase the risk of toxicity from standard doses of fluoropyrimidine drugs (e.g. 5-fluorouracil, capecitabine) (19).				

DRD2	Dopamine receptor D2	Anti-parkinsonian medications, anti-psychotics	Many drugs for Parkinson's disease use this receptor. More research has been done on the link between variants and response to anti-psychotic drugs (20).
F5	Factor V coagulation factor	Oral contraceptives	There is a well-established link between the Factor V Leiden polymorphism and VTE. Users of oral contraceptives with this polymorphism have a higher risk of VTE than wild-type users (21).
G6PD	G6PD enzyme	Anti-diabetes drugs, anti-malarials, chemotherapeutics	G6PD deficiency was one of the first mechanisms found to be linked to variable drug response. G6PD-deficient individuals are at higher risk of adverse drug reactions in response to several triggers, including some drugs (22).
GSTP1	GSTP1 isoform of GST enzyme	Chemotherapeutics, particularly platinum agents	The rs1695 polymorphism is associated with reduced enzyme activity, causing drug resistance and toxicity (23).
HLA-B	HLA-B cell surface molecule	Abacavir, allopurinol, carbamazepine, flucloxacillin, phenytoin	Strongly associated with several ADRs, particularly SJS/TEN in response to carbamazepine and phenytoin. Testing is strongly recommended by several agencies (24).
MTHFR	MTHFR enzyme	Chemotherapeutics and anti- rheumatic drugs	Increased risk of methotrexate toxicity with rs1801131 polymorphism. Also linked to survival in 5-fluorouracil-treated patients (25).
MT-RNR1	MT-RNR1 ribosomal RNA in mitochondria	Aminoglycoside antibiotics	The 1555A>G variation is strongly linked to hearing loss following aminoglycoside antibiotic use – 100% of aminoglycoside patients with the variant develop hearing loss (26).
NAT2	NAT2 enzyme	Many, including: antibiotics, anti- inflammatories, vasodilators	Variations in <i>NAT2</i> are linked to drug-induced hepatotoxicity with anti-TB drugs and ADRs with hydralazine treatment. Variations are also linked to patients requiring higher doses of sulfamethoxazole (27).
NUDT15 [MTH2]	NUDT15 enzyme	Thiopurine drugs	One variant (rs25108385) linked to toxicity from azathioprine and mercaptopurine (28).

RYR1	RYR1 calcium channel	Inhalational anaesthetics and depolarising muscle relaxants	'Dozens' of variants in <i>RYR1</i> increase the risk of malignant hyperthermia in response to anaesthesia (29).		
SLC19A1	RFC1 transporter protein	Folic acid, vitamin B12, methotrexate	The GG genotype increases the risk of spina bifida in infants.  Other variants are associated with an increased risk of methotrexate toxicities (30).		
SLCO1B1	OATP1B1	Many, including: ACE inhibitors, antibiotics, chemotherapeutics, statins	OATP1B1 is an active transport protein, responsible for mediating drug hepatic clearance. Some phenotypes result in impaired hepatic access, reducing drug efficacy (31).		
TPMT	ТРМТ	Thiopurine drugs (e.g. 6-mercaptopurine, azathioprine)	Encodes the TPMT enzyme that metabolises thiopurine drugs. Higher TPMT activity reduces efficacy of these drugs, lower activity increases the risk of ADRs (32).		
TYMS	TYMS enzyme	5-fluorouracil, methotrexate	Over-expression is linked to fluorouracil resistance in cancers. Other variants are associated with better responses to chemotherapy (33).		
UGT1A1	UGT1A1 enzyme	Irinotecan	The *28 and *6 alleles are associated with irinotecan toxicities, particularly neutropenia (34).		
VKORC1	VKORC1 enzyme	Warfarin	Encodes a key enzyme in the vitamin K cycle. Vitamin K is a key component of coagulation factor proteins. Warfarin inhibits VKORC1, leading to a reduction in coagulation factor proteins. High and low dose variants of <i>VKORC1</i> have been found (35).		

Table 1 - Very important pharmacogenes as designated by PharmGKB (<u>www.pharmgkb.org</u>) as of October 2020 (36, 37). ADR = adverse drug reaction. NSAIDs = non-steroidal anti-inflammatory drugs. SJS/TEN = Stevens-Johnson syndrome/toxic epidermal necrolysis. TB = tuberculosis. VTE = venous thromboembolism.

Table 2: Full data extraction of trials included in Chapter 3 biomarker review

Registration	<u>Trial</u> name	Start year	Year of results publicati on	References source	Trial design	<u>Biomarker</u>	Biomarker application	Drug of interest	Sample size (n randomised)	<u>Age</u>	<u>Sex (%</u> <u>F/M)</u>	<u>Race</u>	Location	
ISRCTN3074830	TARGET	2005	2011	2005 protocol	Biomarker strategy design (without	TPMT Prevention	strategy design	Prevention Azathiopr	333	Mean 43.2 (non- genotyped)	50.6/49.4 (non- genotyped)	White/South Asian/Black/ mixed or other (non- genotyped)	- UK	
8	(38, 39)	2003	2011	authors (39)	biomarker assessment in control arm)	11 1011	of ADRs	ine	ine	Mean 41.0 (genotyped )	50.3/49.7 (genotyped	White/South Asian/Black (genotyped)		
NCT01119300	EU- PACT	2011	2013	2009 protocol paper	Biomarker strategy design (without	CYP2C9*2 Improving		Warfarin	455	Mean 66.9 (control)	42.1/57.9 (control)	White/Black/ Asian (control)	UK	
146161115555	(40)	2011	2010	10.2217/pgs. 09.125	biomarker assessment in control arm)	VKORC1	efficacy	vvanami	400	Mean 67.8 (genotyped )	35.8/64.2 (genotyped )	White/Black/ Asian (genotyped)	Sweden	
	SHIVA			2014 protocol	Enrichment	Hormone receptors	ptors Targeted	Targeted chemoth erapy agents	· ·		Median 63 (control)	72/28 (control)		_
NCT01771458	(41)	2012	2015	supplementar y (41)	design	PI3K/AKT/ mTOR RAF/MEK	therapies		erapy 195	Median 61 (genotyped	61/39 (genotyped	Not reported	France	
NCT01894230	GIST (42)	2013	2018	2016 rationale paper	Biomarker strategy design (with biomarker	SLCO1B1* 5	Improving adherence	Any statin	159	Mean 62.5 (control)	65.8/34.2 (control)	White/Black/ other	USA	

				10.2217/pgs- 2016-0065	assessment in control arm)					Mean 62.7 (genotyped	49.4/50.6 (genotyped	White/Black/ other	
NCT02664350	n/a (43)	2016	Results not yet publishe d	2018 protocol paper 10.1016/ j.cct.2018.03. 001	Biomarker strategy design (without biomarker assessment in control arm)	CYP2D6	Quality of life	Opioids	200 (forecast)	Not available	Not available	Not available	USA

Table 2: Full data extraction of the 5 randomised controlled trials included in Chapter 3 biomarker review.

# Table 3: Full list of evidence cited by TARGET trial in biomarker review

No.	Type of reference	<u>Authors</u>	<u>Year</u>	DOI
1	Observational – cohort	Dubinsky et al (44)	2000	10.1016/S0016-5085(00)70140-5
2	Observational – cohort	Weinshilboum and Sladek(45)	1980	n/a
3	Observational – case control	Lennard et al(46)	1989	10.1038/clpt.1989.119
4	Observational – cohort	McLeod et al(47)	1994	10.1038/clpt.1994.4
5	Observational – cohort	Yates et al(48)	1997	10.7326/0003-4819-126-8-199704150-00003
6	Guidelines	British Society of Rheumatology [not found]	2000	n/a
7	Observational – of assay use	Holme et al(49)	2002	10.1093/qjmed/95.7.439
8	Observational – cohort	Bloomfeld & Onken (50)	2003	10.1046/j.1365-2036.2003.01392.x
9	Observational – cohort	McLeod et al (51)	1999	10.1046/j.1365-2141.1999.01416.x
10	Observational – cohort	Black et al (52)	1998	10.7326/0003-4819-129-9-199811010-00007
11	Observational – cohort	Pandya et al (53)	2002	10.1016/S0041-1345(02)02963-9
12	Expert opinion	Seidman(54)	2003	n/a

13	Observational – cohort	Murphy & Atherton (55)	2002	10.1046/j.1365-2133.2002.04922.x
14	Systematic review	Phillips et al	2001	10.1001/jama.286.18.2270
15	Case study	Tavadia et al(56)	2000	10.1067/mjd.2000.103980
16	Cost-effectiveness study	Marra et al (57)	2002	n/a
17	Qualitative work	Tan et al (58)	1997	10.1046/j.1365-2133.1997.d01-1198.x

Table 3 - evidence cited for biomarker inclusion by the TARGET randomised controlled trial. This is based on the 2005 protocol for TARGET (39)

## Table 4: Full list of evidence cited by EU-PACT trial in biomarker review

No.	Type of reference	<u>Authors</u>	Year	DOI
1	Editorial	Rosendaal (59)	1996	10.1056/NEJM199608223350810
2	Observational – retrospective cohort ◆	James et al (60)	1992	n/a
3	Observational – cohort	Penning-van Beest et al (61)	2001	n/a
4	Observational – case control	Hylek et al (62)	1996	10.1056/NEJM199608223350802
5	Editorial	Pirmohamed (63)	2006	10.1111/j.1365-2125.2006.02806.x
6	Observational – case control	Penning-van Beest et al (64)	2002	10.1016/S0895-4356(01)00485-1
7	Observational – cohort	Carlquist et al (65)	2006	10.1007/s11239-006-9030-7
8	Observational – cohort	Schalekamp et al (66)	2007	10.1007/s00228-007-0268-6
9	Observational – case control	Schalekamp et al (67)	2008	10.1001/archinternmed.2007.32
10	Observational – cohort	Gage et al (68)	2004	10.1160/TH03-06-0379
11	Observational – cohort (healthy volunteers)	Bodin et al (69)	2005	10.1182/blood-2005-01-0341
12	Observational – cohort	Wadelius et al (70)	2009	10.1182/blood-2008-04-149070
13	Observational – cohort	Schalekamp et al (71)	2006	10.1016/j.clpt.2006.04.006
14	Observational – cohort	Schalekamp et al (72)	2007	10.1038/sj.clpt.6100036
15	Observational – case control	Reitsma et al (73)	2005	10.1371/journal.pmed.0020312
16	Observational – cohort	Wadelius et al (74)	2005	10.1038/sj.tpj.6500313
17	Observational – cohort	D'Andrea et al (75)	2005	10.1182/blood-2004-06-2111
18	Observational – cohort (GWAS)	Takeuchi et al (76)	2009	10.1371/journal.pgen.1000433

19	Observational – cohort	Caldwell et al (77)	2008	10.1182/blood-2007-11-122010
20	Observational – cohort	Schelleman et al (78)	2008	10.1038/clpt.2008.101
21	Observational – cohort	Klein et al (79)	2009	10.1056/NEJMoa0809329.
22	Observational – cohort	Perini et al (80)	2008	10.1038/clpt.2008.166
23	Observational – cohort	Sconce et al (81)	2005	10.1182/blood-2005-03-1108
24	Observational – cohort	Tham et al (82)	2006	10.1016/j.clpt.2006.06.009
25	Observational – cohort	Gage et al (83)	2008	10.1038/clpt.2008.10
26	Cost-effectiveness analysis	Eckman et al (84)	2009	10.7326/0003-4819-150-2-200901200-00005
27	Cost-effectiveness analysis	Schalekamp et al (85)	2006	10.1016/j.clpt.2006.03.008
28	Literature review	Hughes and Pirmohamed (86)	2007	10.2165/00019053-200725110-00001

Table 4- evidence cited for biomarker inclusion by the EU-PACT randomised controlled trial. Based on 2009 protocol for EU-PACT (40).

## Table 5: Full list of evidence cited by SHIVA trial in biomarker review

No.	Type of reference	Authors	<u>Year</u>	DOI
1	Randomised controlled trial	Slamon et al (87)	2001	10.1056/NEJM200103153441101
2	Observational – cohort	Lièvre et al (88)	2008	10.1200/JCO.2007.12.5906
3	Case study	Joensuu et al (89)	2001	10.1056/NEJM200104053441404
4	Literature review ◆	DiMasi & Grabowski (90)	2007	10.1200/JCO.2006.09.0803
5	Literature review ◆	Von Hoff (91)	1998	n/a
6	Randomised controlled trial	Thatcher et al (92)	2005	10.1016/S0140-6736(05)67625-8
7	Randomised controlled trial	Mok et al (93)	2009	10.1056/NEJMoa0810699
8	Observational – cohort	Von Hoff et al(94)	2010	10.1200/JCO.2009.26.5983

9	Editorial	Doroshow (95) [comment on Von Hoff 2010]	2010	10.1200/JCO.2010.31.1472
10	Randomised controlled trial	Kim et al (96)	2011	10.1158/2159-8274.CD-10-0010

Table 5 - evidence cited for biomarker inclusion by the SHIVA randomised controlled trial. Based on 2014 protocol for SHIVA (41).

## Table 6: Full list of evidence cited by GIST trial in biomarker review

<u>No.</u>	Type of reference	Author(s)	<u>Year</u>	<u>DOI</u>
1	Epidemiology – American Heart Association	Mozaffarian et al(97)	2015	10.1161/CIR.000000000000157
2	Editorial	Greenland and Lauer(98)	2015	10.1001/jama.2015.7434
3	Meta-analysis (of IPD)	Cholesterol Treatment Triallists' Collaborators(99)	2012	10.1016/S0140-6736(12)60367-5
4	Meta-analysis (of IPD)	Cholesterol Treatment Triallists' Collaborators(100)	2015	10.1016/S0140-6736(14)61368-4
5	Cochrane review	Taylor et al(101)	2013	10.1002/14651858.CD004816.pub5
6	Guidelines - American College of Cardiology/American Heart Association Task Force	Stone et al(102)	2013	10.1016/j.jacc.2013.11.002
7	Observational – cohort	Pencina et al(103)	2014	10.1056/NEJMoa1315665
8	Literature review	Hirsh et al(104)	2015	10.1016/j.jacc.2015.05.030
9	Observational – cohort	Bermingham et al(105)	2011	10.1016/j.clinthera.2011.07.007
10	Observational – cohort	Ho et al(106)	2008	10.1016/j.ahj.2007.12.011
11	Observational – cohort	Vodonos et al(107)	2015	10.1016/j.ejim.2015.02.014
12	Observational – cohort	Franklin et al(108)	2015	10.1002/pds.3787
13	Systematic review	De Vera et al(109)	2014	10.1111/bcp.12339
14	Literature review	Osterburg and Blaschke(110)	2005	10.1056/NEJMra050100
15	Qualitative study	Fung et al(111)	2010	n/a
16	Qualitative study	Cohen et al(112)	2012	10.1016/j.jacl.2012.03.003
17	Literature review	Ong et al(113)	2012	10.2217/pgs.12.2
18	Guidelines – American College of Cardiology/American Heart Association/National Heart, Lung and Blood Institute	Pasternak et al(114)	2002	10.1016/S0735-1097(02)02030-2
19	Expert opinion	Thompson et al(115)	2006	10.1016/j.amjcard.2005.12.013
20	Literature review/expert opinion	Alfirevic et al(116)	2014	10.1038/clpt.2014.121

21	Literature review	Patel et al(117)	2015	10.1016/j.atherosclerosis.2015.03.025
22	Guidelines – European Atherosclerosis	Stroes et al(118)	2015	10.1093/eurheartj/ehv043
	Society			
23	Observational – case control	Link et al(119)	2008	10.1056/NEJMoa0801936
24	Meta-analysis	Hou et al(120)	2015	10.1097/MD.00000000001268
25	Observational – cohort	Pasanen et al(121)	2006	10.1007/s00228-006-0123-1
26	In vitro work	Kimoto et al(122)	2012	10.1021/mp300379q
27	Randomised controlled trial	Voora et al(123)	2009	10.1016/j.jacc.2009.04.053
28	Guidelines – Clinical Pharmacogenetics	Wilke et al(124)	2012	10.1038/clpt.2012.57
	Implementation Consortium			
29	Observational – cohort	Birmingham et al(125)	2015	10.1007/s00228-014-1801-z
30	Observational – cohort/meta-analysis	de Keyser et al(126)	2014	10.1097/FPC.000000000000018
31	Sub-study of larger randomised	Danik et al(127)	2013	10.1016/j.ahj.2013.01.025
	controlled trial			
32	Sub-study of larger randomised	Martin et al(128)	2011	10.1111/j.1365-2125.2011.04090.x
	controlled trial			
33	Literature review	Niemi et al(129)	2011	10.1124/pr.110.002857
34	Guidelines – Clinical Pharmacogenetics	Ramsey et al(130)	2014	10.1038/clpt.2014.125
	Implementation Consortium			
35	Literature review	Voora and Ginsburg(131)	2012	10.1016/j.jacc.2012.01.067
36	Observational – cohort	Donnelly et al(132)	2011	10.1038/clpt.2010.255
37	Observational – cohort (pilot study)	Li et al(133)	2014	10.3390/jpm4020147

Table 6 - evidence cited for biomarker inclusion by the GIST randomised controlled trial. Based on 2016 protocol for GIST (42).

# <u>Table 7: Full list of evidence cited by Precision Medicine Guided Treatment for Cancer Pain trial in biomarker review</u>

No.	Type of reference	Author(s)	Year	DOI
1	Guidelines – National Cancer Institute	PDQ Supportive and Palliative Care	2002	n/a
		Editorial Board (134)		
2	Guidelines – National Comprehensive	National Comprehensive Cancer	2017	n/a
	Cancer Network	Network (135)		
3	Randomised controlled trial	Temel et al (136)	2010	10.1056/NEJMoa1000678
4	Guidelines – European Association for	Caraceni et al (137)	2012	10.1016/S1470-2045(12)70040-2
	Palliative Care			
5	Expert panel	Fine et al (138)	2009	10.1016/j.jpainsymman.2009.06.002
6	Observational – cohort	Zhao et al (139)	2014	10.1200/JCO.2013.50.6071

7	Randomised controlled trial (in twins)	Angst et al (140)	2012	10.1016/j.pain.2012.02.022
8	Literature review	Fillingim et al (141)	2008	10.1111/j.1601-0825.2008.01458.x
9	Observational – cohort	Gan et al (142)	2007	10.1007/BF03256239
10	Case study	Susce et al (143)	2006	10.1016/j.pnpbp.2006.03.018
11	Guidelines – Clinical Pharmacogenetics	Crews et al (144)	2014	10.1038/clpt.2013.254
	Implementation Consortium			
12	Observational – cohort	Baber et al (145)	2015	10.1038/tpj.2015.3
13	Case study	Ciszkowski et al (146)	2009	10.1056/NEJMc0904266
14	Case study	Gasche et al (147)	2004	10.1056/NEJMoa041888
15	Randomised controlled trial	Eckhardt et al (148)	1998	10.1016/S0304-3959(98)00021-9
16	Randomised controlled trial	Lötsch et al (149)	2009	10.1016/j.pain.2009.03.023
17	Observational – cohort	Andreassen et al (150)	2012	10.1007/s00228-011-1093-5
18	Randomised controlled trial	Samer et al (151)	2010	10.1111/j.1476-5381.2010.00673.x
19	Randomised controlled trial	Zwisler et al (152)	2009	10.1111/j.1742-7843.2009.00378.x
20	Observational – cohort	Zwisler et al (153)	2010	10.1111/j.1399-6576.2009.02104.x

Table 7 - evidence cited for biomarker inclusion by the Precision Medicine Guided Treatment for Cancer Pain trial. Based on 2018 publication (43).

#### 4.1 Protocol for systematic reviews and meta-analyses

See: citation (154)

Danielle Johnson, Andrea Jorgensen. The influence of HLA-B\*15:02 and HLA-A\*31:01 on the risk of developing adverse skin reactions to carbamazepine: protocol for two systematic reviews. PROSPERO 2019 CRD42019161000 Available from: https://www.crd.vork.ac.uk/prospero/display\_record.php?ID=CRD42019161000

#### 4.2 Standard data extraction form for systematic reviews

DATA EXTRACTION FORM: SYSTEMATIC REVIEW OF PGx STUDIES OF HLA-B\*15:02 AND CARBAMAZEPINE-INDUCED HYPERSENSITIVITY REACTIONS

PDF ID:	Lead Author:	Publication Date:
1. General	Notes of Interest	
	al comments of interest here	
par any gonoro		
*******	*******	******
2. Study De	<u>esign</u>	
) Design:	RCT prospective c	cohort retrospective cohort case-control
ase-control + hea	althy subjects	other(describe)

b) Sample size (if case-control study state numbers separately): healthy	total	cases	controls
c) Is justification/calculation given for sample size ?	yes no		
d) Is a priori power to detect effect sizes of varying degrees quote	d? yes n	0	
**************************	*****		
3. <u>Participants</u>			
a) Ethnic groups included: White Black African Indian Bangladeshi Pakistani African American Other(describe)	Black Caribbean Chinese	Black other	Japanese
b) What is inclusion/exclusion criteria ? Describe.			
Inclusion			
Cases:			
Controls:			
Exclusion			

c) What reactions were included?	SJS	TEN	SJS/TEN	DRESS	MPE	other
(describe)						

#### d) Patient characteristics (please continue on additional sheet if not enough space)

Characteristic	Units	Overall	Subgroup Cases (n=	Subgroup Controls (n=	Subgroup Healthy volunteers (n=	Other Subgroup (n=
Age (mean+/- SD; range) /(median; IQR) <sup>1</sup>						
No. males n (%)						
Ethnicity: White n (%)						
Black African n (%)						
Black Caribbean n (%)						
Black other n (%)						

Indian n (%)					
Indian ii (70)					
Bangladeshi n (%)					
Pakistani n (%)					
(11)					
OL: (0/)					
Chinese n (%)					
Japanese n (%)					
African American n (%)					
Amenican in (70)					
Other					
Indication for carbamazepine:					
n (%)					
11 (70)					
Epilepsy/seizures					
Dovobiotrio					
Psychiatric					
Neuralgia					
Neuropathic pain					
Neuropatrile pairi					
Autism					
Other					
011					
Other					
	1	l .	1	l .	

Other			
Other			

#### Notes

- - 4. Genotyping
- a) What variants were investigated? (please continue on additional sheet if not enough space)

Variant	Total genotyped	Number with the variant	Number without the variant	Comment <sup>1</sup>

1. Add any relevant comments here e.	g. if the row corr	espond	s to a spe	ecific ethnic	subgroup within the study)
b) Are results given for all? yes no(ex	kplain)				
c) Were genotyping personnel blinded to or	utcome/case-cor	ntrol sta	tus?	yes	not mentioned
d) i. Was test for HWE undertaken at each S	SNP?		,	yes	not mentioned
ii. If yes, what test was used ? Chi-square	Fisher's Exact	other	(describe	e) not state	ed n/a
iii. What p-value cut-off was used ?iv. Are results of testing provided ? yes	no n/a	n/a			
v. If yes, how many SNPs were found to o	deviate ?		n/a		
vi. If yes and more than one deviates, are	reasons for devi	iation ex	plored ?		
yes	not mentioned	n/a			
vii. Are deviating SNPs excluded from ana e) i. Is method of genotyping described?	lyses ?	yes	no i	unclear	n/a
e) i. is memou or genoryping described ?		yes	110		

ii. If yes, describe briefly		
f) i. Were genotype QC methods used	? yes	not mentioned
ii. If yes, what method was used ? re	esequencing all patients	resequencing random sample
resequencing extreme patients regenoty	yping all patients	regenotyping random sample
regenotyping extreme patients other (de	escribe) not state	ed n/a
iii. If yes, were results quoted ? yes		
iv) If so what was degree of agreeme	nt ?	n/a
g) Were genotype frequencies compar	red with previously pul	olished for same population ?
yes	not mentioned	
5. <u>Analysis</u>	**********	**********
a) If more than one ethnic group incl	<u>-</u>	s this adjusted for in analysis ?

b) i. Is extent of missing data stated ? yes	no						
ii. If yes, are reasons for missingness ex	cplored ?	yes	no	none a	re missing		n/a
iii. Are any checks undertaken for missin	ngness at	random	?	yes	n	ot me	entioned
iv. Is missing genotype data imputed ?	yes	no	unclea	r			
v. If so, how ? multiple imputation	other (des	cribe)				n/a	
vi. Are numbers contributing to each ana	alysis quo	ted ?	yes	no			
vii. If so, are numbers different to total sar	mple size	?	yes	no	n/a		
6. <u>Outcomes</u>							
a) Was justification given for choice	of outcom	es?		yes(de	scribe belo	w)	no
Justification for choice of outcomes							

b)	List any particular outcome(s) that appear(s) to be suppressed (describe)
	********

#### 7, Results

Outcome	Outcome	HLA-B*15:02 present	HLA-B*15:02 not present	p-value vs outcome present	Definition	Test unde rtake n	Additional Comments*
ADR	Present						
overall	Absent; controls						
	Absent; HV						
	Present						
SJS	Absent; controls						
	Absent; HV						
	Present						
TEN	Absent; controls						

	Absent; HV					
	Present					
SJS/TEN	Absent; controls					
	Absent; HV					
	Present					
DRESS	Absent; controls					
	Absent; HV					
	Present					
MPE	Absent; controls					
	Absent; HV					
Other	Present					
(describe)	Absent; controls					
	Absent; HV					
Other	Present					
(describe)	Absent; controls					
	Absent; HV					

<sup>\*</sup> e.g. particular ethnicities only, and whether the absent includes CBZ-tolerant patients or healthy patients. HV = healthy volunteers

#### 4.3 Full list of studies included in systematic reviews and meta-analyses

See https://www.dropbox.com/scl/fi/fozqjovdao5qr1da2wwhg/allpapers\_table.xlsx?dl=0&rlkey=nop19t2e0sukgme5ice8f0q88

#### 4.4 Quality assessment of studies included in systematic reviews and meta-analyses

#### 4.4.1 HLA-B\*15:02

See https://www.dropbox.com/scl/fi/5pvj5sp03gehx8kejet1y/Study-quality-1502.xlsx?dl=0&rlkey=dugxpbi1nvvqaqbolv83iwljw

#### 4.4.2 HLA-A\*31:01

See https://www.dropbox.com/scl/fi/r7c6qqvhr3a8vir9hvkh2/Study-quality-3101.xlsx?dl=0&rlkey=w4f5a6qz9dcahaxdcm7gke1gg

#### 4.5 Calculation of allele frequencies for HLA-B\*15:02 and HLA-A\*31:01

See

https://www.dropbox.com/scl/fi/0avm3a85rj3gg7x49eyti/allele\_freq\_1502\_3101\_20211109.xlsx?dl=0&rlkey=aa96zen4gbywrfyc89pgn9\_vnq

#### 4.6 Simulation code

See https://github.com/dkj201/simulation for full details.

Table 8: Comparison of papers included in previous meta-analyses and our meta-analysis.

Previous meta-analysis	Included paper [by ancestry subgroup]	Included in current MA?	Reasons
	Hung et al 2006 [Han Chinese]	Υ	NA
Yip et al 2012 (155)	Wu et al 2010 [Han Chinese]	Υ	NA
	Liao et al 2010 [Han Chinese]	N	Previously excluded as is meeting abstract only
	Zhang et al 2011 [Han Chinese]	Y	NA

	Wang et al 2011 [Han Chinese]	Y	NA
	Locharernkul et al 2008 [Thai]	N	SJS and TEN reported separately.
	Tassaneeyakul et al 2010 [Thai]	Y	NA
	Kulkantrakorn et al 2012 [Thai]	N	Data is included within Tassaneeyakul 2010
	Then et al 2011 [Malaysian]	N	SJS only, not SJS/TEN
	Alfirevic et al 2006 [White] *	N	No HLA-B*15:02 positive patients
	Hung et al 2006 [Han Chinese]	Y	NA
	Liao et al 2010 [Han Chinese]	N	Previously excluded as is meeting abstract only
	Wu et al 2010 [Han Chinese]	Υ	NA
	Zhang et al 2011 [Han Chinese]	Y	NA
Tangamornsuksan et al 2013 (156)	Shi et al 2012 [Han Chinese]	Y	NA
	Locharernkul et al 2008 [Thai]	N	SJS and TEN reported separately
	Tassaneeyakul et al 2010 [Thai]	Y	NA
	Kim et al 2011 [Korean]	N	SJS only, not SJS/TEN
	Then et al 2011 [Malaysian]	N	SJS only, not SJS/TEN
	Niihara et al 2012 [Japanese] *	N	No HLA-B*15:02 positive patients
	Cheung et al 2013 [Han Chinese]	Y	NA
Grover et al 2014 (157)	Hung et al 2006 [Han Chinese]	Υ	NA
	Shi et al 2012 [Han Chinese]	Y	NA
	Wang et al 2011 [Han Chinese]	Υ	NA
	Wu et al 2010 [Han Chinese]	Y	NA
	Zhang et al 2011 [Han Chinese]	Υ	NA

Kim et al 2011 [Korean]	N	SJS only, not SJS/TEN
Then et al 2011 [Malaysian]	N	SJS only, not SJS/TEN
Kulkantrakorn et al 2012 [Thai]	N	Data is included within Tassaneeyakul 2010
Locharernkul et al 2008 [Thai]	N	SJS and TEN reported separately.
Tassaneeyakul et al 2010 [Thai]	Υ	NA

Table 8 - Comparison of papers included in previous meta-analyses and our meta-analysis.

### Table 9: Previous systematic reviews of discrete choice experiments

Reference	Summary	Aim	No. of included DCEs	Date range included	Populations	Key findings
Ryan and Gerard (2003) (158)	Systematic review of DCEs in health economics	To identify current practice in DCEs in health economics. Also evaluated study quality	34	1990- 2000	Patients Community Health insurance consumer	No studies judged to be of 'strong' design quality
de Bekker- Grob, <i>et al.</i> (2010) (159)	Updated systematic review of DCEs in health economics	To update the previous review (158) with current practice in DCEs in health economics	114	2001- 2008	Not reported	Large increase in use of DCEs in health economics, with changing methodologies
Clark, <i>et al.</i> (2014) (160)	A further updated systematic review of DCEs in health economics	To update the previous review (159) with current practice in DCEs in health economics	179	2009- 2012	Not reported	Further increase in use of DCEs, particularly in countries other than the UK

Soekhai, <i>et al.</i> (2019) (161)	A large systematic review of health-related DCEs	To provide an overview of DCEs in health economics, and their applications and methods	301	2013 - 2017	Patients Healthcare workers General public	Reflects on more modern DCE methods and design, but problems with poor reporting continue
Guerra, <i>et</i> <i>al.</i> (2019) (162)	A small systematic review of patient preferences for breast cancer treatment	To qualitatively synthesis information from DCEs about breast cancer treatment in patients	5	- (May) 2019	Patients	Patients most highly value attributes related to side-effects
Trapero- Bertran, <i>et</i> <i>al.</i> (2019) (163)	A systematic review of DCEs relating to priority setting in health care provision	To identify attributes for designing a DCE, to develop and validate a framework for decision making on health technologies	72	2008 - 2015	Patients Policy makers Providers General public	Better quality DCEs in heath care provision are needed, especially in areas other than oncology
Bien, <i>et al.</i> (2017) (164)	A systematic review of patient preferences for cancer treatment	To focus on DCEs in cancer treatment and assess the significance of different attribute types	28	2010- (April) 2016	Patients General population Healthcare professionals	Attributes related to side- effects were most often significant
Harrison, <i>et al.</i> (2017) (165)	Systematic review of DCEs comparing patients' and healthcare professionals' views	Review DCEs that that elicited opinions from both patients and healthcare professionals and examine concordance	38	1995- (July) 2015	Patients General public Parents/caregivers Healthcare professionals	Patients and healthcare professionals value attributes differently and more DCEs should incorporate this analysis
Vass, <i>et al.</i> (2017) (166)	Systematic review of qualitative methods in DCE practice	Explore the use of qualitative methods in healthcare related DCEs and explore their perceived usefulness with authors	254	2001- (June) 2012	Not reported	Only a minority of DCEs used 'extensive' qualitative work, although authors agree on its usefulness
Harrison, <i>et al.</i> (2014) (167)	Systematic review of DCEs that included a risk attribute	To highlight the use of risk in DCEs, and recommend ways to improve risk communication	117	1995- (April) 2013	Patients Healthcare professionals Public Parents Decision makers	Recommendation that risks should be placed in context and methods used to communicate risk should be validated

Table 9 - A summary of previous systematic reviews of discrete choice experiments. The highlighted rows show 4 linked reviews, each using the same methods to update on the field of health related DCEs. DCEs = discrete choice experiments

Table 10: Search terms used by previous systemic reviews to locate discrete choice experiments

	Ryan & Gerard (2003) (158)	de Bekker- Grob, <i>et</i> <i>al.</i> (2010) (159)	Clark, <i>et</i> <i>al.</i> (2014) (160)	Soekhai, et al. (2019) (161)	Guerra, et al. (2019) (162)	Trapero- Bertran, et al. (2019) (163)	Bien, <i>et</i> <i>al.</i> (2017) (168)	Harrison, et al. (2017) (165)	Vass, et al. (2017) (166)	Harrison, et al. (2014) (167)
Choice behaviour					<b>✓</b>					✓
Choice Behaviour (MeSH term)					✓					
Choice experiment					✓					
Choice model					✓					
Conjoint	✓		✓				✓			
Conjoint analysis	✓	✓	✓	✓	✓	✓	✓		✓	
Conjoint- analysis					✓					
Conjoint analysis/ measurement/ study/ choice								✓		<b>✓</b>
Conjoint choice experiment			<b>√</b>	<b>✓</b>			<b>√</b>			
Conjoint choice experiment(s)		<b>✓</b>				✓			<b>✓</b>	
Conjoint choice experiments	<b>✓</b>			<b>✓</b>						
Conjoint measurement	✓	✓	✓	✓		✓	✓		✓	

Conjoint studies	✓	<b>✓</b>	<b>✓</b>	✓		<b>✓</b>	<b>✓</b>		✓	
DCE			✓							
Discrete choice					✓			✓		✓
Discrete-choice					<b>✓</b>					
Discrete choice conjoint experiment			<b>√</b>	<b>✓</b>			<b>✓</b>			
Discrete choice conjoint experiments				<b>√</b>						
Discrete choice experiment			<b>√</b>	<b>√</b>			1			
Discrete choice experiment(s)		<b>✓</b>				<b>✓</b>			<b>✓</b>	
Discrete choice experiments										
Discrete choice model(I)ing		<b>✓</b>	<b>✓</b>			<b>✓</b>	<b>✓</b>		<b>✓</b>	
Discrete choice modeling				✓						
Discrete choice modelling	✓			✓						

					ı	1				
Discrete-choice										
Functional measurement	✓	✓	✓	✓		✓			✓	
Paired comparison					<b>✓</b>			<b>✓</b>		✓
Paired comparisons	✓	✓	✓	✓		<b>✓</b>			✓	
Pairwise choice					✓			<b>✓</b>		✓
Pairwise choices	✓	✓	✓	✓		✓			✓	
Part-worth utilities	✓	✓	✓	✓		<b>✓</b>			✓	
Part-worth utility								<b>✓</b>		✓
Patient preference					✓		✓			
Patient Preference (MeSH term)					✓					
Preference					✓					
Stated preference	✓	✓	<b>√</b>	✓	✓	<b>✓</b>	✓	✓	✓	✓

Table 10 - Grid showing search terms used by previous systematic reviews to locate discrete choice experiments.

## Table 11: Data extraction sheet prepared for systematic review of discrete choice experiments

Title	
Authors	
DOI	
Link	

Year of publication	
Country of origin	
Pharmacogenetic phenotype	
Disease	
Biomarker(s)	
Sample size	
Population type	
Population age	
Population race or ethnicity	
Population gender (% female)	
Response rate	
Number of choice tasks seen by each participant	
Number of possible choice tasks	
Survey method	
Number of attributes	
Attribute domains	
Design plan	
Design software	
Method(s) used to create choice sets	
Estimation procedures	

Validity checks	
Qualitative work	
Learning from this DCE	

Table c11 - Standard data extraction prepared for systematic review of discrete choice experiments. DCE = discrete choice experiment. DOI = digital object identifier.

## Table 12: Data extracted from each paper in the systematic review of discrete choice experiments (1)

First author / reference	Year	Country	PGx	Disease	Biomarker(s)	Sample size	Pop.	Age	Race/ ethnicity	Gender %F	Response rate	Learning
Ballinger (169)	2017	USA	Risk of ADRs (peripheral neuropathy and congestive heart failure)	Breast cancer	HER2 (negative only)	417	Patients	35% under 50 65% 50 or over	88% Caucasian 5% African American 4% Hispanic 3% Other	Not reported	Not reported	Previous experience of an ADR affects preferences
Boeri (170)	2018	UK	Risk of ADRs (10kg weight gain), hyper- responsiveness genotype	Schizophrenia	n/a	67	HCPs	Not reported	Not reported	41%	95.7%	HCPs are open to PGx but this varies depending on experience
Chan (171)	2013	Singaporo	Risk of ADRs (major bleeding	CV disease	CYP2C9	197	General public	Mean: 52.5	100% Chinese	72.7%	83.5%	Presentation of warfarin and
Chan (171) 20	2013	113 Singapore		CV disease	VKORC1	191	Patients	Mean 57.4	100% Chinese	26.9%	53.8%	ADRs to general public

Dong (172)	2016	Singapore	Risk of ADRs (SJS), cost of genetic test, change in cost of medications depending on genetic test result	Gout	HLA-B*58:01	189	Patients	Mean 57.1	61% Chinese 27% Indian 10% Malay 2% Other	34%	Not reported	Uptake as a useful output of a DCE
Herbild (173)	2009	Denmark	Time spent with dosage adjustments due to lack of effect or ADRs (not specified), likelihood of improvements from genetic test	Depression	CYP2D6	323	General public	Not reported	Not reported	53%	46%	Participants that don't normally pay for healthcare can consider cost in decision making
			If test predicts	Breast cancer	Oncotype DX	150	Patients	Mean 54.5	81% White 10% African- American 6% Hispanic/Latino 3% Asian/Asian American	Breast 100%		
Issa (174)	2013	USA	risk of recurrence, likelihood of benefit from drugs, and risk of ADRs (not specified)	Colorectal cancer	KRAS UGT1A1	150	Patients	Mean 42 (colorecta	74% White 10% African- American 9% Hispanic/Latino 3% Asian/Asian American 3% American- Indian/Alaska Native 1% Hawaiian Native/Pacific Islander	Colorectal 46%	42.2%	Detailed report of focus group methodology for attribute and level selection
Liede (175)	2017	Int	Risk of ADR (teeth and jaw problems, uterine cancer), type of ADR (effect on fertility, effect on female hormones)	Breast cancer	BRCA1 BRCA2	622	Patients*	Mean 41	Not reported	100%	53.5%	External validity checks are useful, where possible
Marshall (176)	2016	Canada	Risk of ADR [categorical] (temporary side-effects, permanent side-effects), likelihood of	Breast cancer	HER2	1004	General public	Mean 49	84% White 6% Chinese 2% Indigenous 1% Black 5% Other 2% Not answered	100%	Not reported	High levels of evidence assured if a test is insurance covered

			benefit from chemotherapy									
Marshall (177)	2017	USA	Risk of ADR (that makes you unable to do everyday activities or take care of yourself)	None	Whole genome sequencing	410	General public	Not reported	Not reported	Not reported	47.0%	Pictograms to represent risks
			Risk of ADR (nausea, hair				General public group A ‡	Mean 48.2	Not reported	48.5%	65%	
Najafzadeh (178)	2013	Canada	loss, skin rash, fatigue), severity of ADR (mild,	Cancer	None – hypothetical test	1096	General public group B	Mean 47.6	Not reported	50.6%	69%	Participants can tolerate complex information
			moderate, severe)				Patients	Mean 58.2	Not reported	58.3%	64%	
Payne	2014	UK	Risk of ADR (azathioprine-	Autoimmune	TPMT	297	Patients	Mean 45.8	Not reported	56%	50%	Ethical considerations of
(179)	2011	UK	associated neutropenia)	disease	IPMI	297	HCPs	Not reported	Not reported	Not reported	34%	including a cost attribute
Powell	2015	UK	Risk of ADR (mild skin rash, memory	Enilopoy	HLA-A*31:01	165	HCPs	Not reported	Not reported	Not reported	Not reported	Utility model for predicting uptake
(180)	2015	UK	problems, SJS), likelihood of benefit	Epilepsy	11LA-A 31:01	100	Patients	Median 38	90.2% White 3.7% Black 1.2% Asian 2.4% Mixed/multiple	66%	Not reported	based on results of DCE

Smith (181)	2014	USA	Risk of ADR (peripheral neuropathy, severe diarrhoea), type of ADR (moderate or severe), duration of ADR (1 year, just during treatment)	Breast cancer	None – hypothetical test	641	Patients	Most responde nts 50-59	90.6% Caucasian No other reported	99.7%	Not reported	Participants are able to comprehend and manage decisions based on genetic biomarkers
-------------	------	-----	---	---------------	--------------------------------	-----	----------	-------------------------------	--	-------	--------------	--

Table 12 - Data extracted from each paper in DCE systematic review. Does not include some fields extracted – DOI, title, full author list. \* Women with BRCA1 or BRCA2 mutations but unaffected by breast or ovarian cancer. ‡ The general public was split into 2 groups and each was presented with a unique scenario. Results were reported separately ADRs = adverse drug reactions. CV = cardiovascular. HCPs = healthcare professionals. PGx = pharmacogenetics. Pop. = population. SJS = Stevens Johnson syndrome.

Table 13: Data extracted from each paper in the systematic review of discrete choice experiments (2)

First author	Year	No. of choice tasks	No. of possible choice tasks	Survey method	No. of attributes	Attribute domains	Design plan	Design software	Method(s) used to create choice sets	Estimation procedures	Validity checks	Qualitative work
Ballinger (169)	2017	12	Not reported	Online	4	Health status Risk	Not reported	Not reported	Pragmatically chosen	Hierarchical Bayesian	Theoretic	Not reported
Boeri (170)	2018	26	26	Face to face	4	Health status Risk Time Other	Fractional factorial	NGene	D-efficiency	Random parameters logit	Not reported	Expert opinion
Chan (171)	2013	8	24	In clinic Online	4	Health care Money Risk Other	Not reported	Sawtooth CBC/Web	Not reported	Hierarchical Bayesian	Sen's expansion/ contraction	Pilot testing
Dong (172)	2016	9	32	Face to face	5	Money Risk Other	Main and interaction effects	SAS	D-efficiency	Latent class logit	Non- satiation Transitivity	Patient interviews Cognitive interviews
Herbild (173)	2009	8	32	Online	4	Health care Money Risk Time	Fractional factorial	SAS	D-efficiency	Conditional logistic regression	Theoretic	Expert opinion Focus groups [published]

Issa (174)	2013	20	Not reported	Online	5	Health care Health status Money Risk Other	Not reported	Sawtooth (custom)	Random pairing	Not reported	Sen's expansion/ contraction	Focus groups [published]
Liede (175)	2017	4	36	Online	7	Health status Risk Time Other	Fractional factorial	SAS	D-efficiency	Random parameters logit	External	Expert opinion Patient interviews
Marshall (176)	2016	12	500	Online	5	Health care Risk Other	Main and interaction effects	Sawtooth CBC/Web	D-efficiency	Hierarchical Bayesian	Sen's expansion/ contraction Theoretic	Expert opinion Focus groups Pilot testing [published]
Marshall (177)	2017	6	Not reported	Online	3	Health care Money Risk	Main effects	SAS	Not reported	Random parameters logit	Theoretic	Expert opinion Interviews
Najafzadeh (178)	2013	16	160	Online	7	Health status Money Risk Time Other	Fractional factorial	Sawtooth CBC/Web	D-efficiency	Conditional logit	Non- satiation Theoretic	Expert opinion Pilot testing
Payne (179)	2011	16	16	Post	5	Risk Time Other	Fractional factorial	Not reported	Unclear (Street and Burgess methods)	Random effects probit	Non- satiation Sen's expansion/ contraction Theoretic	Expert opinion Focus groups Interviews Pilot testing [published]
Powell (180)	2015	16 (HCPs) 8 (patients)	Not reported	Online	6 (HCPs) 5 (patients)	Money Risk Time Other	Fractional factorial	Not reported	Pairing with constant comparator (HCPs) Not reported (patients)	Random effects logit	Not reported (HCPs) Non- satiation (patients)	Interviews Focus groups
Smith (181)	2014	14	Not reported	Online	4	Health care Health status Risk	Not reported	Not reported	Not reported	Not reported	Sen's expansion/ contraction	Pilot testing

Table 13 - Further data extracted for each paper in DCE systematic review.

# Chapter 6

6.1 Survey of healthcare professionals

## Questionnaire for healthcare professionals

## Welcome

Thank you for participating in our survey. Your feedback is important for designing a questionnaire that will be useful to patients, academics, and anyone with an interest in pharmacogenetics.

This survey has been approved by the University of Liverpool Health and Life Sciences Research Ethics Committee (ref. 4736).

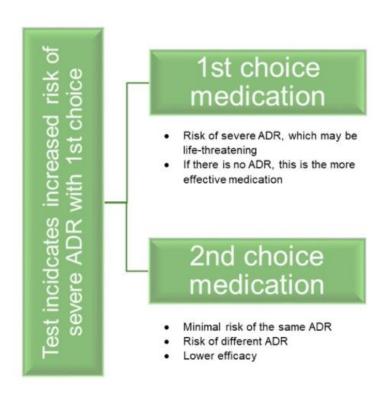
# Background information 🖸

Genetic testing can be used to predict and prevent adverse drug reactions (ADRs). This is termed pharmacogenetics and is one of the promises of personalised medicine. The <u>FDA</u> defines pharmacogenetics as 'variations in DNA sequence as related to drug response'.

Although there are some examples of this in use in the clinic (see below), pharmacogenetics is still a comparatively new field with many unknowns. We are therefore designing a survey to quantify the general public's views on genetic testing to prevent ADRs.

The scenario we will test is one where a patient has a genetic test indicating they are at increased risk of an ADR with a first-choice medication. We want to interrogate in which scenarios participants would choose a second-choice, potentially less effective medication, and when they would choose to risk the ADR with the first-choice medication.





You have been invited to take this survey since you are a professional interested in personalised medicine. The results of this survey will be used alongside opinions from patients and the general public, to design a national survey.



## **Examples**

## Abacavir, hypersensitivity syndrome in HIV, and HLA-B\*57:01

Abacavir is an antiretroviral drug for HIV treatment. A hypersensitivity syndrome initially presenting as fever, rash, nausea and vomiting, potentially leading to severe hypotension and death is strongly associated with the HLA-B\*57:01 allele, and can be avoided by withholding abacavir from patients testing positive for <u>HLA-B\*57:01</u>.

The British National Formulary, European Medicines Agency and British HIV Association recommend testing every patient before commencing <u>abacavir</u>.

## Carbamazepine, Stevens-Johnson syndrome in epilepsy and HLA-B\*15:02

Carbamazepine is a tricyclic anticonvulsant used to treat epileptic seizures, trigeminal neuralgia, and some psychiatric disorders. It is linked to the rare but extremely serious <u>Stevens-Johnson</u> <u>syndrome/toxic epidermal necrolysis (SJS/TEN) reactions</u>.

HLA-B\*15:02 is <u>strongly associated</u> with SJS/TEN in patients receiving carbamazepine. The <u>British National Formulary</u> specifies that individuals of Han Chinese or Thai origin are tested for the HLA-B\*15:02 allele, and to avoid carbamazepine unless there is no alternative.



# **Survey**

Below is a list of characteristics that might be considered important when deciding whether to order a pharmacogenetic test. Please select the top characteristic in each group that you think is the most important in making the decision to order a genetic test. Please then write a little to explain your choice.

* Tes	t characteristics 🖸
$\circ$	Time to result
$\bigcirc$	Cost of test
	Level of evidence for testing (e.g. one or more randomised controlled trials, compared to a test with only a genome-wide association study [GWAS] behind it)
$\bigcirc$	Coverage of the test (test can predict either severe ADRs only, or severe and mild ADRs)
$\bigcirc$	PPV (probability of experiencing the ADR if a positive result on the pharmacogenetic test – 'true positive')
	NPV (probability of not experiencing the ADR if a negative result on the pharmacogenetic test - 'true negative')
$\bigcirc$	If the test included in BNF
$\bigcirc$	Other (please specify)
Reas	on(s) for selection:   Output  Description:

* Medication choices 🖸
Efficacy/effectiveness of first- and second-choice medications
Risk of severe ADRs with first- and second-choice medications
Risk of mild ADRs with first- and second-choice medications
Ost/cost-effectiveness of first- and second-choice medications
Other (please specify)
Reason(s) for selection:

* Test information 🖸
O Information on specific gene polymorphism(s)
A panel of several pharmacogenes that may yield useful results to inform future prescribing decisions
○ Whole genome sequencing
What does the test result mean (easily understandable interpretation of the test result)
Other (please specify)
Reason(s) for selection:

* Practicalities 🖸
O How sample is collected (saliva, blood, etc)
○ Who is involved in ordering, interpreting and explaining results to patients
O Privacy of test results (restricted to doctor-patient, use for research, use by insurance companies)
Other (please specify)
Reason(s) for selection:

Please add any further cha please write down anything	_		_			
4						
Please could you write a lit	tle to help us	understa	ınd the reas	ons for ye	our choice	es. [Optional] 🔽
		Prev	Next			

# **Survey**

Below is a list of characteristics that might be considered important when deciding whether to order a pharmacogenetic test. Please select the top characteristic in each group that you think is the most important in making the decision to order a genetic test. Please then write a little to explain your choice.

* Te	est characteristics 모
0	Time to result
$\bigcirc$	Cost of test
0	Level of evidence for testing (e.g. one or more randomised controlled trials, compared to a test with only a genome-wide association study [GWAS] behind it)
$\bigcirc$	Coverage of the test (test can predict either severe ADRs only, or severe and mild ADRs)
$\bigcirc$	PPV (probability of experiencing the ADR if a positive result on the pharmacogenetic test – 'true positive')
0	NPV (probability of not experiencing the ADR if a negative result on the pharmacogenetic test – 'true negative')
$\circ$	If the test included in BNF
$\bigcirc$	Other (please specify)
Rea	ason(s) for selection: 🖸

* Medication choices 🔽
Efficacy/effectiveness of first- and second-choice medications
Risk of severe ADRs with first- and second-choice medications
Risk of mild ADRs with first- and second-choice medications
Ost/cost-effectiveness of first- and second-choice medications
Other (please specify)
Reason(s) for selection:

Test information 🔽
Information on specific gene polymorphism(s)
A panel of several pharmacogenes that may yield useful results to inform future prescribing decisions
Whole genome sequencing
What does the test result mean (easily understandable interpretation of the test result)
Other (please specify)
eason(s) for selection: $ abla$

* Practicalities 🗩
O How sample is collected (saliva, blood, etc)
O Who is involved in ordering, interpreting and explaining results to patients
O Privacy of test results (restricted to doctor-patient, use for research, use by insurance companies)
Other (please specify)
Reason(s) for selection:

Please add any furthe please write down an		-	-		_	
4						
Please could you writ	e a little to he	elp us understa	ınd the reas	ons for yo	ur choices. [	Optional] 🔽
				<u> </u>		
		Prev	Next			

# **Other Information**

Your group (tick all that apply) 🔽	
☐ GP	
Hospital doctor	
Other healthcare professional	
Academic	
Other (please specify)	
f hospital doctor, what is your speciality? 🔽	
f other healthcare professional, what is your speciality? 🔽	

* Have you ever ordered a genetic test for a pharmacogenetic purpose?
Pharmacogenetic referring to 'variations in DNA sequence as related to drug response'.
Examples might include -
<ul> <li>Abacavir, hypersensitivity syndrome in HIV, and HLA-B*57:01</li> <li>Carbamazepine, Stevens-Johnson syndrome in epilepsy and HLA-B*15:02</li> </ul>
$\Diamond$
○ Yes
○ No
○ Unsure
* Have you ever used the results of genetic testing (ordered by yourself, other medical staff, or direct-to-consumer testing) to inform prescribing or treatment of a patient?
○ Yes
○ No
○ Unsure
Prev Done Control of the Control of

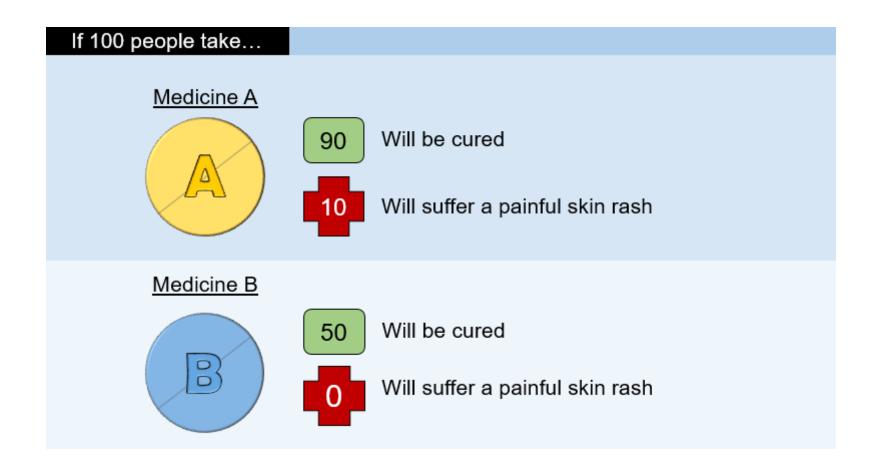
## 6.2 Survey of patients

# Introduction to this survey

Genetic testing can be used to predict the risk of side-effects of some medicines.

For example, some medicines come with a risk of a painful and potentially serious skin rash as a side-effect. In the example below, if **100 people** with an illness take **medicine A**, **90 will be cured**. However, **10 of them** will suffer the potentially serious skin rash as a side-effect.

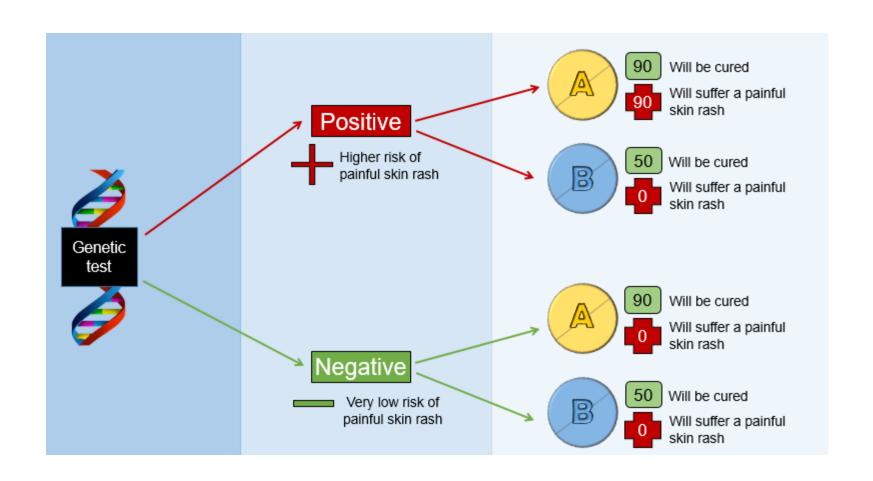
Medicine B is less effective (**only 50 out of 100 people will be cured**), but has **no risk** of this side-effect.



You might already know which medicine you would choose if you were in this position.

However, by using genetic testing, we may be able to tell which people are more likely to suffer the painful skin rash. In this case, if someone has a negative genetic test result, we can say they have a **very low risk** of the painful skin rash if they take medicine A. They might then choose to take medicine A, to have a better chance of a cure.

But if they test positive, they would be at **higher risk** of the painful skin rash if they take medicine A. They could still choose to take it, hoping for a cure. Or, they could choose medicine B instead, where there is **no risk** of skin rash - but also less chance of a cure.



However, this is still a very new field and there are lots of things we don't know.

We wish to find out people's opinions about genetic testing to predict the risk of drug side effects, and what helps people decide if they want to have a genetic test.



- \* 1. Firstly, to your knowledge, have you had any genetic tests? lacktriangle
- Yes
- O No
- O Don't know
- O Prefer not to answer

* 2. Imagine you have an illness where you can take medicine A or medicine B. The pictures above can help with this. There is a risk of a serious side-effect with medicine A, but it is <b>more effective</b> (works better) than B. There is <b>no risk</b> of the serious side-effect with medicine B. A genetic test can help you and your doctor predict the risk of you getting the serious side-effect with medicine A.
Below is a list of things that might be considered important about a genetic test. Please choose the <b>top</b>
5 things you think are most important. $\square$
Risk of developing a severe side-effect without the genetic test
Risk of developing a mild side-effect without the genetic test
How well medicine A works
How well medicine B works
Accuracy of the genetic test (no test is 100% accurate)
How much evidence is there to show that the genetic test works in predicting risk of the serious side- effect (for example, it is used regularly by doctors, or maybe it has only been used in clinical trials. You could even be one of the first to try it)
☐ Time to wait for the result
Level of information you receive from the test (you could have all your genes analysed ('gene panel test'), or just the ones relevant for medicine A)
Who sees the test results
If your doctor recommends you receive the genetic test

If most other people in your situation choose to take the genetic test
Who delivers the test results to you
Cost of the test to you personally
Cost of the test to the NHS
Cost of the medicine A to the NHS
Cost of the medicine B to the NHS
Severity of the disease being treated by these medications
How sample is collected (blood, saliva, biopsy, etc)
3. Please add any further things you think are important. There are no wrong answers, please write down anything that comes to mind.

	Which of these do you think is the <b>most important thing</b> to consider when deciding whether or not se the genetic test? 🔽
0	Severity of the disease being treated (it could be something that affects your life but is not life-threatening, or it could be something more serious)
0	How much evidence is there to show that the genetic test works in predicting risk of the serious side- effect (for example, it could be used regularly by doctors, or maybe it has only been used in clinical trials. You could even be one of the first to try it)
$\bigcirc$	How accurate the genetic test is (no test is 100% accurate)
$\bigcirc$	Risk of getting the serious side-effect if you take medicine A
$\bigcirc$	How well medicine A and medicine B work
0	<b>Who you see for the genetic testing service</b> (for example, your GP, a hospital doctor, a nurse, a genetic counsellor)
$\bigcirc$	How the test is done (for example, a blood sample, a saliva sample, a biopsy)
0	<b>Privacy of your test results</b> (for example, only you and your doctor, or use for research also, or also being shared with an employer or life insurance company)
$\bigcirc$	Something else (please specify)

5. Please could you write a little to explain your choice.  $\ \ \ \ \ \$ Prev Next

We would now like to ask some questions about you and how you interpret risk.

It can be difficult to clearly communicate the risk (chance) of serious side-effects. Below are four different ways it has been done in the past, using an example of two risks for comparison.



2 out of 100 people experience the serious side-effect 45 out of 100 people experience the serious side-effect

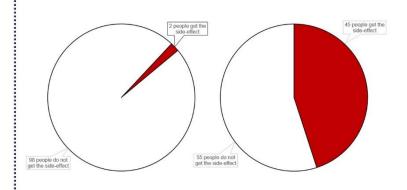
# 2 out of 100 people experience the serious side-effect 45 out of 100 people experience the serious side-effect

## Written

2 out of 100 people experience the serious side-effect

45 out of 100 people experience the serious side-effect

## **Boxes**



## Pictograms

## **Proportion pie charts**

* 6.	Which of these ways of communicating risk do you think is the clearest? 🔽
$\circ$	Written
$\circ$	Boxes
$\bigcirc$	Pictograms
$\bigcirc$	Proportion pie chart
$\circ$	None of these
$\circ$	Other (please specify)
* 7.	What is your age group? 🖸
	18-24
	25-34
	35-44
	45-54
	55 +
	Prefer not to answer



## 6.3 Final discrete choice experiment, as shown to participants (abacavir example)

See <a href="https://ctrc.liv.ac.uk/InDevelopment/DCE?p=test12345">https://ctrc.liv.ac.uk/InDevelopment/DCE?p=test12345</a>

Full surveys for all 8 DCEs are located here: <a href="https://www.dropbox.com/sh/yv97hf3g82pfdqv/AAD13JMDjgzvVXILUk2G20TEa?dl=0">https://www.dropbox.com/sh/yv97hf3g82pfdqv/AAD13JMDjgzvVXILUk2G20TEa?dl=0</a>





# **Explanation**

This is a project about the general public's opinion of using genetic testing to decide on the best treatment options for a patient. It is known as Public Opinions of PERsonalised medicine (POPPER).

The purpose of this survey is to examine whether people would choose to have a genetic test in different situations. There is some information at the start of the survey that you will need to read.

You will be asked to **imagine you have been diagnosed with a particular serious condition**, and need to be treated for it with a drug. It is known that the drug can cause side-effects in some people. You will then be asked if you would want a genetic test to help the doctor understand whether you are at risk of the side-effects, and therefore help them choose the most appropriate treatment approach for you in some different scenarios.

You might find it upsetting to imagine being diagnosed with a serious condition. You can exit the survey at any time, without giving a reason. If you have any particular concerns, you can contact one of the researchers or another organisation for support (details below)

We will not be collecting any personal information as part of this study. We will only be collecting your preferences. There are questions about your age group and gender at the end of the study, but these are **optional** and if you choose to complete them they cannot be used to identify you.

Your data will be stored on computer servers at the University of Liverpool for up to 5 years and aggregate results from the project may be published in scientific papers. The survey is being conducted through JISC, and you can view their privacy policy here.

# **Consent**

Participation in this study is completely voluntary. If you decide not to participate there will not be any negative consequences. Please be aware that if you decide to participate, you may stop participating at any time and you may decide not to answer any specific question.

Because the data you provide is anonymous, it will not be possible to remove your data from the survey once you submit it.

By participating you are indicating that you have read the description of the study, are over the age of 18, and that you agree to the terms as described.

By clicking 'I consent' you agree to participate in this research study.

I consent

I do not consent

## La Details of researchers

This research is being conducted by Danielle Johnson, a postgraduate research student at the University of Liverpool. The work is supervised by Professor Andrea Jorgensen, Professor of Biostatistics, at the University of Liverpool. If you have any queries, you can contact danielle.johnson@liverpool.ac.uk or ethics@liverpool.ac.uk

This research has been approved by the University of Liverpool Health and Life Sciences Research Ethics Committee (Human participants, tissues, and databases), reference 4736. This work was funded by the MRC HTMR Network of Hubs for Trials Methodology Research (MR/L004933/2). Danielle Johnson was awarded a PhD studentship (R19) funded by the MRC HTMR Network.

# Safeguarding

For mental health support: Samaritans: 116 123 (24/7) or email jo@samaritans.org

# 

For advice about cancer: Macmillan Cancer Support: 0808 808 00 00 (8am-8pm, 7 days a week) or online

For advice about HIV: Terrence Higgins Trust: 0808 802 1221 (10am-6pm weekdays, 10am-1pm weekends) or info@tht.org.uk

For advice about epilepsy: Epilepsy Society: 01494 601 400 (9am-4pm Monday to Friday, 9am-7:30pm Wednesdays) or helpline@epilepsysociety.org.uk

For advice about heart disease: British Heart Foundation: 0300 330 3311 (9am-5pm weekdays, 10am-4pm Saturdays) or hearthelpline@bhf.org.uk, or online chat

Upon consenting to take part, participants are randomised to one of eight DCEs. The example with abacavir is shown here.

0% complete

# Page 1: Introduction I

Imagine you have been diagnosed with HIV.

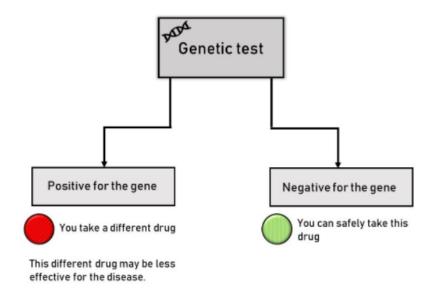
Your doctor advises you that the best treatment is a medicine called **ABACAVIR**. However, some people experience side-effects from this medicine. These can be **severe and potentially life-threatening**.

One of these is **hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

## Genetic testing

We know that people with a particular type of gene in their DNA are more likely to suffer from sideeffects, and **testing for this gene is now advised before prescribing abacavir in several countries.** 

The test can help doctors choose a different approach, to reduce your risk of having the side-effect. People with a particular type of this gene will be prescribed a different drug, but this may be less effective for treating HIV.



Given this information, and imagining that you are in this situation, you have two options to choose from:

- 1. You can choose to have the gene test. If you test positive, you would receive a different drug, which might be less effective for treating HIV.
- 2. You can choose not to be tested, and take abacavir regardless of your risk of side-effects.

In this survey we will ask you to choose one of 2 genetic tests (Test A or Test B), with different characteristics. You can also choose not to have a test (no test).

Both tests involve a single saliva swab taken from inside the mouth. The results are available to the doctor in 1-2 days.

Submit and continue >

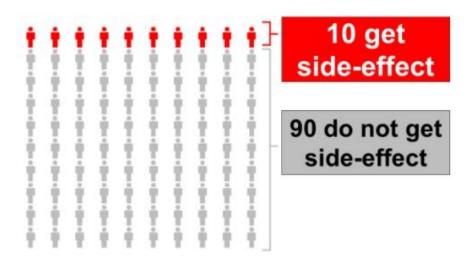
5% complete

# Page 2: Introduction II

Each of the tests presented to you (Test A, Test B, or no test) is described in this survey according to 5 different characteristics, which are as follows:

### 1. Chance of serious side-effect from this medication

This is how many people will experience the serious side-effect. For example, 1 in 10 means that if 100 people take the drug, 10 people would experience the side-effect. This will look like this:



If you choose the 'no test' option, you will not have this information on your genetic risk of this sideeffect.

#### 2. Cost of the test to the NHS

This is how much it costs the NHS to buy the test. If you choose the 'no test' option, this is £0.

### 3. Use of your data for further research by universities and other researchers

Whenever you have any medical treatment, you might be asked if you want to participate in research. For these genetic tests, this would mean allowing universities and researchers access to your test results. Your test results, together with many other people's test results, would be looked at to improve medical knowledge.

This can be done in two ways. Both of these would only be done with your expressed permission.

The first is to provide your test data, but without any further information about you. Researchers would not know your name or any identifiable information. They would not be able to contact you – your data would be anonymous. There would be no way to link your data back to you.

The second way asks you to give permission for researchers to request access to your medical record along with your genetic test results. This would only be for research that requires this level of detail. Allowing this gives the researchers an option to request access to much more information that might help move research along faster. They may also be able to contact you in the future if there are any clinical trials or studies that you could help by being a part of.

Patient: Steve Bennett

NHS number: 00000001

Date of birth: 01/02/75

Sex: M

Contact number: 07771112223

Ethnicity: White

Blood type: O positive

Disease: HIV

Previous medical conditions:

High blood pressure

Epilepsy Depression

Genetic test results:

Gene 2.....Positive
Gene 3.....Negative
Negative

If you choose the 'no test' option, none of your data will be used for research.

#### 4. Number of medicines the test can be used to inform

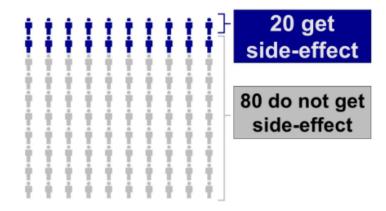
Your genetic test results tells your doctor what medicine or medicines are most suitable for you. Some genetic tests only look at one gene. With one of these single gene tests, the result is typically only used to guide the prescribing of a single medicine. However, a genetic panel test would provide enough information to cover 25 or 50 different medicines. Should you need to take any of these medicines in the future, your doctor could look up the results from the genetic panel test before prescribing.

If you choose the 'no test' option, you will not have this information.

#### 5. Risk of any serious side-effect from any medicine over the next 10 years

A serious side-effect is one that causes sufficient concern or harm that you need to go to hospital. The diagrams show your overall chance of having a serious side-effect from **any medicine you might take in the next 10 years**, excluding the one mentioned above.

This will also look like the pictogram you saw above.



If you choose the 'no test' option, you will not have this information.

### Page 3: Question 1

These characteristics are outlined in the questionnaire below, with some variations between the choices. In each case, choose whether you would prefer to **have test A, test B, or no test**.

Take care with these. Some choices may look very similar at first glance but they are all different.

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

1	Test A	Test B	No test
Chance of serious side-effect from this medicine	15 in 100  15 in 100  15 get side-effect  85 do not get side-effect	3 in 100  3 in 100  3 get side-effect  97 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, but no contact (anonymous)	Yes, and they can contact me (linked)	
Number of medicines the test can be used to inform	1	50	
Cost of the test to the NHS	£30	£30	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	5 in 100  5 get side-effect  95 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	

		Submit and continue >
O No test		
O Test B		
O Test A		
Which would you choose?	* Required	

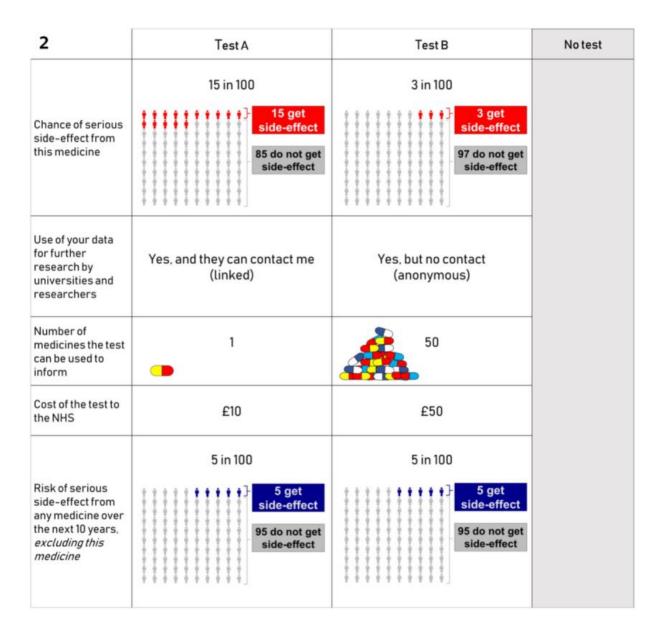
# Page 4: Question 2

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:



Which would you choose	? * Required	
○ Test A		
O Test B		
O No test		

### Page 5: Question 3

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

3	Test A	Test B	Notest
Chance of serious side-effect from this medicine	15 in 100  15 in 100  15 get side-effect  85 do not get side-effect	5 in 100  5 in 100  5 get side-effect  95 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, but no contact (anonymous)	No	
Number of medicines the test can be used to inform	25	25	
Cost of the test to the NHS	£10	£50	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	5 in 100  5 get side-effect  95 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	

Which would you choose? * Required	
<ul><li>○ Test A</li><li>○ Test B</li><li>○ No test</li></ul>	
	Submit and continue >

### Page 6: Question 4

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

4	Test A	Test B	No test
Chance of serious side-effect from this medicine	15 in 100  15 get side-effect  85 do not get side-effect	5 in 100  5 in 100  5 get side-effect  95 do not get side-effect	
Use of your data for further research by universities and researchers	No	Yes, and they can contact me (linked)	
Number of medicines the test can be used to inform	25	25	
Cost of the test to the NHS	£30	£30	
Risk of serious side-effect from any medicine over the next 10 years. excluding this medicine	2 in 100  2 get side-effect  98 do not get side-effect	20 in 100  20 get side-effect  80 do not get side-effect	

Which would you choose?	* Required
○ Test A	
O Test B	
O No test	

# Page 7: Question 5

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

5	Test A	Test B	Notest
Chance of serious side-effect from this medicine	5 in 100  5 get side-effect  95 do not get side-effect	15 in 100  15 get side-effect  85 do not get side-effect	
Use of your data for further research by universities and researchers	No	Yes, but no contact (anonymous)	
Number of medicines the test can be used to inform	1	25	
Cost of the test to the NHS	£10	£50	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	2 in 100  2 get side-effect  98 do not get side-effect	20 in 100  20 get side-effect  80 do not get side-effect	

Which would you choose?	* Required
O Test A	
O Test B	
O No test	

# Page 8: Question 6

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

6	Test A	Test B	Notest
Chance of serious side-effect from this medicine	15 in 100  15 get side-effect  85 do not get side-effect	3 in 100  3 get side-effect  97 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, and they can contact me (linked)	Yes, but no contact (anonymous)	
Number of medicines the test can be used to inform	50	1	
Cost of the test to the NHS	£30	£30	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	5 in 100  5 get side-effect  95 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	

Which would you choose?	* Required
<ul><li>○ Test A</li><li>○ Test B</li></ul>	
O No test	

# Page 9: Question 7

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

7	Test A	Test B	Notest
Chance of serious side-effect from this medicine	3 in 100  3 get side-effect  97 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, and they can contact me (linked)	No	
Number of medicines the test can be used to inform	1	50	
Cost of the test to the NHS	£30	£30	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	20 in 100  20 get side-effect  80 do not get side-effect	2 in 100  2 get side-effect  98 do not get side-effect	

Which would you choos	se? * Required		
<ul><li>Test A</li><li>Test B</li><li>No test</li></ul>			

# Page 10: Question 8

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

8	Test A	Test B	Notest
Chance of serious side-effect from this medicine	5 in 100  5 get side-effect  95 do not get side-effect	15 in 100  15 get side-effect  85 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, but no contact (anonymous)	Yes, and they can contact me (linked)	
Number of medicines the test can be used to inform	25	25	
Cost of the test to the NHS	£10	£50	
Risk of serious side-effect from any medicine over the next 10 years. excluding this medicine	20 in 100  20 get side-effect  80 do not get side-effect	2 in 100  2 get side-effect  98 do not get side-effect	

Which would you choose? * Required	
○ Test A ○ Test B	
O No test	

### Page 11: Question 9

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

9	Test A	Test B	Notest
Chance of serious side-effect from this medicine	5 in 100  5 get side-effect  95 do not get side-effect	3 in 100  3 get side-effect  97 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, and they can contact me (linked)	No	
Number of medicines the test can be used to inform	25	1	
Cost of the test to the NHS	£50	£10	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	20 in 100  20 get side-effect  80 do not get side-effect	2 in 100  2 get side-effect  98 do not get side-effect	

Which would you choose? * Required	
<ul><li>○ Test A</li><li>○ Test B</li></ul>	
O No test	
	Submit and continue >

### Page 12: Question 10

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

10	Test A	Test B	Note
Chance of serious side-effect from this medicine	3 in 100  3 get side-effect  97 do not get side-effect	15 in 100  15 get side-effect  85 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, and they can contact me (linked)	No	
Number of medicines the test can be used to inform	50	1	
Cost of the test to the NHS	£30	£30	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	2 in 100  2 get side-effect  98 do not get side-effect	20 in 100  20 get side-effect  80 do not get side-effect	

Which would you choose? * Required	
<ul><li>○ Test A</li><li>○ Test B</li></ul>	
O No test	
	Submit and continue >

# Page 13: Question 11

### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

11	Test A	Test B	No test
Chance of serious side-effect from this medicine	3 in 100  3 get side-effect  97 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, but no contact (anonymous)	Yes, and they can contact me (linked)	
Number of medicines the test can be used to inform	50	1	
Cost of the test to the NHS	£50	£10	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	5 in 100  5 get side-effect  95 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	

Which would you choose?	* Required
<ul><li>Test A</li><li>Test B</li><li>No test</li></ul>	

# Page 14: Question 12

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

12	Test A	Test B	Notest
Chance of serious side-effect from this medicine	5 in 100  5 get side-effect  95 do not get side-effect	3 in 100  3 get side-effect  97 do not get side-effect	
Use of your data for further research by universities and researchers	No	Yes, and they can contact me (linked)	
Number of medicines the test can be used to inform	50	1	
Cost of the test to the NHS	£50	£10	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	2 in 100  2 get side-effect  98 do not get side-effect	20 in 100  20 get side-effect  80 do not get side-effect	

Which would you choose? * Required	
○ Test A	
○ Test B	
O No test	
	Submit and continue >

# Page 15: Question 13

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

13	Test A	Test B	Notest
Chance of serious side-effect from this medicine	3 in 100  3 in 100  3 get side-effect  7 do not get side-effect	15 in 100  15 get side-effect  85 do not get side-effect	
Use of your data for further research by universities and researchers	No	Yes, but no contact (anonymous)	
Number of medicines the test can be used to inform	1	50	
Cost of the test to the NHS	£10	£50	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	20 in 100  20 get side-effect  80 do not get side-effect	2 in 100  2 get side-effect  98 do not get side-effect	

Which would you choose? * Required	
○ Test A ○ Test B	
O No test	

Submit and continue >

### Page 16: Question 14

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

14	Test A	Test B	No test
Chance of serious side-effect from this medicine	3 in 100  3 get side-effect  7 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	
Use of your data for further research by universities and researchers	No	Yes, but no contact (anonymous)	
Number of medicines the test can be used to inform	25	25	
Cost of the test to the NHS	£50	£10	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	5 in 100  5 get side-effect  95 do not get side-effect	5 in 100  5 get side-effect  95 do not get side-effect	

Which would you choose	? * Required
<ul><li>Test A</li><li>Test B</li><li>No test</li></ul>	

Submit and continue >

## Page 17: Question 15

#### Reminder

The serious side-effect from this medicine:

**hypersensitivity** (consisting of a painful/itchy rash, nausea and vomiting, fever, and anaphylaxis – breathing problems, wheezing, and losing consciousness).

The serious side-effect from any medicine you might take in the next 10 years:

15	Test A	Test B	Notest
Chance of serious side-effect from this medicine	5 in 100  5 in 100  5 get side-effect  95 do not get side-effect	15 in 100  15 get side-effect  85 do not get side-effect	
Use of your data for further research by universities and researchers	Yes, but no contact (anonymous)	No	
Number of medicines the test can be used to inform	1	25	
Cost of the test to the NHS	£30	£10	
Risk of serious side-effect from any medicine over the next 10 years, excluding this medicine	2 in 100  2 in 100  2 get side-effect  98 do not get side-effect	20 in 100  20 get side-effect  80 do not get side-effect	

vvnich would you choose? * Required	
<ul><li>Test A</li><li>Test B</li><li>No test</li></ul>	

Submit and continue >

94% complete		
Page 18: Final questions	5	
What is your age group?		
<ul> <li>18 - 24</li> <li>25 - 34</li> <li>35 - 44</li> <li>45 - 54</li> <li>55 - 64</li> <li>65+</li> </ul>		
What is your gender?		
O Female O Male O Another		

O Prefer not to answer

Have you ever had a genetic test?
<ul><li>Yes</li><li>No</li><li>Don't know</li><li>Prefer not to answer</li></ul>
Have you suffered from the illness described in this survey?
<ul><li>Yes</li><li>No</li><li>Don't know</li><li>Prefer not to answer</li></ul>
How difficult did you find this survey to complete, on a scale of 1 (not difficult at all) to 10 (almost impossible)?
Please select >
Do you have any comments on this survey? (please do not include any identifiable information here)
Finish 🗸

# Final page

### Thank you for completing this survey.

Thank you for completing this survey. Your data has been stored anonymously, and it is not possible to identify you or to remove it from the servers. However, if you have any concerns, you can contact

danielle.johnson@liverpool.ac.uk or

ethics@liverpool.ac.uk

If you have any particular concerns about HIV you may find it helpful to contact <u>Terrence Higgins Trust</u>: 0808 802 1221 (10am-6pm weekdays, 10am-1pm weekends) or <u>info@tht.org.uk</u>

You can also contact the researchers directly.

For 24/7 mental health support, you can contact <u>Samaritans</u>: 116 123 (24/7) or email <u>jo@samaritans.org</u>

## Click here to submit your responses

### Chapter 7

### 7.1 Stata code for DCE analysis

```
**This is correct for ABACAVIR
ssc install outreg2
clear
cd "M:\Stata"
capture log close
log using abacavirwtp.log, replace text
set more off
*import, using first row as variable names
import excel using datamatrix abacavir.xlsx , firstrow
*check first 2 rows look right
list in 1/2
*run the model
xtlogit pref asc no adr today 1 adr today 2 privacy yes1 privacy yes2 medsno cost future adr 1 future adr 2, re i
(personid)
*export results to excel
outreg2 using abacavirresults, excel
**save here as abacavir boot.dta**
*using output of the previous abacavir run (abacavir boot.dta)
use "M:\Stata\abacavir boot.dta"
bootstrap, saving("M:\Stata\abacavir boot results mean.dta") reps(1000) seed(123) : xtlogit pref asc no adr today 1
adr today 2 privacy yes1 privacy yes2 medsno cost future adr 1 future adr 2, re i (personid)
generate pref_adr_today_0 = -1*(pref_b_adr_today_1+pref_b_adr_today_2)
generate pref privacy no = -1*(privacy yes1+privacy yes2)
generate pref future adr 0 = -1*(pref b future adr 1+pref b future adr 2)
```

export excel using "M:\Stata\abacavir\_boot\_results.xls", firstrow(variables)
log close

### 7.2 Data matrices for each DCE

See <a href="https://www.dropbox.com/sh/gp40k9go4qgrmnr/AAB7ZXZqM3ImSmA3YUbyqIZva?dl=0">https://www.dropbox.com/sh/gp40k9go4qgrmnr/AAB7ZXZqM3ImSmA3YUbyqIZva?dl=0</a>

### 7.3 Bootstrapped beta-coefficients for each DCE

See https://www.dropbox.com/sh/oygat71gu9ch0d3/AACPdb\_2y3ujNsAcscJ60kQHa?dl=0

### 7.4 Utility modelling for each DCE

See https://www.dropbox.com/sh/69eka4kmyx8e680/AADgWAMGzpPYkPHAcw8UJUq\_a?dl=0

### **Appendix References**

- 1. Hodges LM, Markova SM, Chinn LW, Gow JM, Kroetz DL, Klein TE, et al. Very important pharmacogene summary: ABCB1 (MDR1, Pglycoprotein). Pharmacogenetics and genomics. 2011;21(3):152-61.
- 2. Fohner AE, Brackman DJ, Giacomini KM, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for ABCG2. Pharmacogenetics and genomics. 2017;27(11):420-7.
- 3. Thorn CF, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for angiotensin-converting enzyme. Pharmacogenetics and genomics. 2010;20(2):143-6.
- PharmGKB. Very Important Pharmacogene: ADRB1 2020 [Available from: <a href="https://www.pharmgkb.org/vip/PA166170369">https://www.pharmgkb.org/vip/PA166170369</a>.
- 5. Sandilands AJ, O'Shaughnessy KM. The functional significance of genetic variation within the beta-adrenoceptor. British journal of clinical pharmacology. 2005;60(3):235-43.
- 6. Litonjua AA, Gong L, Duan QL, Shin J, Moore MJ, Weiss ST, et al. Very important pharmacogene summary ADRB2. Pharmacogenetics and genomics. 2010;20(1):64-9.
- 7. Sangkuhl K, Dirksen RT, Alvarellos ML, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for CACNA1S. Pharmacogenetics and genomics. 2020;30(2).
- 8. McDonagh EM, Clancy JP, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for CFTR. Pharmacogenetics and genomics. 2015;25(3):149-56.
- 9. PharmGKB. Very Important Pharmacogene: COMT 2020 [Available from: https://www.pharmgkb.org/vip/PA166165409.
- 10. McDonagh EM, Wassenaar C, David SP, Tyndale RF, Altman RB, Whirl-Carrillo M, et al. PharmGKB summary: very important pharmacogene information for cytochrome P-450, family 2, subfamily A, polypeptide 6. Pharmacogenetics and genomics. 2012;22(9):695-708.

- 11. Thorn CF, Lamba JK, Lamba V, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for CYP2B6. Pharmacogenetics and genomics. 2010;20(8):520-3.
- 12. Scott SA, Sangkuhl K, Shuldiner AR, Hulot JS, Thorn CF, Altman RB, et al. PharmGKB summary: very important pharmacogene information for cytochrome P450, family 2, subfamily C, polypeptide 19. Pharmacogenetics and genomics. 2012;22(2):159-65.
- 13. Aquilante CL, Niemi M, Gong L, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for cytochrome P450, family 2, subfamily C, polypeptide 8. Pharmacogenetics and genomics. 2013;23(12):721-8.
- 14. Van Booven D, Marsh S, McLeod H, Carrillo MW, Sangkuhl K, Klein TE, et al. Cytochrome P450 2C9-CYP2C9. Pharmacogenetics and genomics. 2010;20(4):277-81.
- 15. Owen RP, Sangkuhl K, Klein TE, Altman RB. Cytochrome P450 2D6. Pharmacogenetics and genomics. 2009;19(7):559-62.
- 16. PharmGKB. Very Important Pharmacogene: CYP3A4 2020 [Available from: <a href="https://www.pharmgkb.org/vip/PA166169915">https://www.pharmgkb.org/vip/PA166169915</a>.
- 17. Lamba J, Hebert JM, Schuetz EG, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for CYP3A5. Pharmacogenetics and genomics. 2012;22(7):555-8.
- 18. Alvarellos ML, Sangkuhl K, Daneshjou R, Whirl-Carrillo M, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for CYP4F2. Pharmacogenetics and genomics. 2015;25(1):41-7.
- 19. PharmGKB. Very Important Pharmacogene: DPYD 2020 [Available from: <a href="https://www.pharmgkb.org/vip/PA166169426">https://www.pharmgkb.org/vip/PA166169426</a>.
- 20. Mi H, Thomas PD, Ring HZ, Jiang R, Sangkuhl K, Klein TE, et al. PharmGKB summary: dopamine receptor D2. Pharmacogenetics and genomics. 2011;21(6):350-6.
- 21. PharmGKB. Very Important Pharmacogene: F5 2020 [Available from: https://www.pharmgkb.org/vip/PA166169437.
- 22. McDonagh EM, Thorn CF, Bautista JM, Youngster I, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for G6PD. Pharmacogenetics and genomics. 2012;22(3):219-28.
- 23. PharmGKB. Very Important Pharmacogene: GSTP1 2020 [Available from: https://www.pharmgkb.org/vip/PA166169438.
- 24. Barbarino JM, Kroetz DL, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for human leukocyte antigen B. Pharmacogenetics and genomics. 2015;25(4):205-21.
- 25. PharmGKB. Very Important Pharmacogene: MTHFR 2020 [Available from: https://www.pharmgkb.org/vip/PA166169429.
- 26. Barbarino JM, McGregor TL, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for MT-RNR1. Pharmacogenetics and genomics. 2016;26(12):558-67.
- 27. McDonagh EM, Boukouvala S, Aklillu E, Hein DW, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for N-acetyltransferase 2. Pharmacogenetics and genomics. 2014;24(8):409-25.
- 28. PharmGKB. Very Important Pharmacogene: NUDT15 2020 [Available from: <a href="https://www.pharmgkb.org/vip/PA166178335">https://www.pharmgkb.org/vip/PA166178335</a>.
- 29. Alvarellos ML, Krauss RM, Wilke RA, Altman RB, Klein TE. PharmGKB summary: very important pharmacogene information for RYR1. Pharmacogenetics and genomics. 2016;26(3):138-44.
- 30. Yee SW, Gong L, Badagnani I, Giacomini KM, Klein TE, Altman RB. SLC19A1 pharmacogenomics summary. Pharmacogenetics and genomics. 2010;20(11):708-15.
- 31. Oshiro C, Mangravite L, Klein T, Altman R. PharmGKB very important pharmacogene: SLCO1B1. Pharmacogenetics and genomics. 2010;20(3):211-6.
- Wang L, Pelleymounter L, Weinshilboum R, Johnson JA, Hebert JM, Altman RB, et al. Very important pharmacogene summary: thiopurine S-methyltransferase. Pharmacogenetics and genomics. 2010;20(6):401-5.
- 33. PharmGKB. Very Important Pharmacogene: TYMS 2020 [Available from: <a href="https://www.pharmgkb.org/vip/PA166165418">https://www.pharmgkb.org/vip/PA166165418</a>.

- 34. Barbarino JM, Haidar CE, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for UGT1A1. Pharmacogenetics and genomics. 2014;24(3):177-83.
- 35. Owen RP, Gong L, Sagreiya H, Klein TE, Altman RB. VKORC1 pharmacogenomics summary. Pharmacogenetics and genomics. 2010;20(10):642-4.
- 36. PharmGKB. VIPs: Very Important Pharmacogenes 2020 [Available from: https://www.pharmgkb.org/vips.
- 37. Hippman C, Nislow C. Pharmacogenomic Testing: Clinical Evidence and Implementation Challenges. J Pers Med. 2019;9(3).
- 38. Newman WG, Payne K, Tricker K, Roberts SA, Fargher E, Pushpakom S, et al. A pragmatic randomized controlled trial of thiopurine methyltransferase genotyping prior to azathioprine treatment: the TARGET study. Pharmacogenomics. 2011;12(6):815-26.
- 39. Ollier B, Newman B, Payne K, Poulton K, Andrews J, Elliott R, et al. The TARGET Study TPMT: AZA Response to Genotyping and Enzyme Testing Protocol: A prospective randomised controlled trial of thiopurine methyltransferase (TPMT) genotyping in the management of patients, prior to commencement of AZA treatment. In: University of Manchester, editor. 2005.
- 40. van Schie RM, Wadelius MI, Kamali F, Daly AK, Manolopoulos VG, de Boer A, et al. Genotype-guided dosing of coumarin derivatives: the European pharmacogenetics of anticoagulant therapy (EU-PACT) trial design. Pharmacogenomics. 2009;10(10):1687-95.
- 41. Le Tourneau C, Delord JP, Goncalves A, Gavoille C, Dubot C, Isambert N, et al. Molecularly targeted therapy based on tumour molecular profiling versus conventional therapy for advanced cancer (SHIVA): a multicentre, open-label, proof-of-concept, randomised, controlled phase 2 trial. Lancet Oncol. 2015;16(13):1324-34.
- 42. Singh K, Peyser B, Trujillo G, Milazzo N, Savard D, Haga SB, et al. Rationale and design of the SLCO1B1 genotype guided statin therapy trial. Pharmacogenomics. 2016;17(17):1873-80.
- 43. Mosley SA, Hicks JK, Portman DG, Donovan KA, Gopalan P, Schmit J, et al. Design and rational for the precision medicine guided treatment for cancer pain pragmatic clinical trial. Contemp Clin Trials. 2018;68:7-13.
- 44. Dubinsky MC, Lamothe S, Yang HY, Targan SR, Sinnett D, Theoret Y, et al. Pharmacogenomics and metabolite measurement for 6-mercaptopurine therapy in inflammatory bowel disease. Gastroenterology. 2000;118(4):705-13.
- 45. Weinshilboum RM, Sladek SL. Mercaptopurine pharmacogenetics: monogenic inheritance of erythrocyte thiopurine methyltransferase activity. American journal of human genetics. 1980;32(5):651-62.
- 46. Lennard L, Van Loon JA, Weinshilboum RM. Pharmacogenetics of acute azathioprine toxicity: relationship to thiopurine methyltransferase genetic polymorphism. Clinical pharmacology and therapeutics. 1989;46(2):149-54.
- 47. McLeod HL, Lin JS, Scott EP, Pui CH, Evans WE. Thiopurine methyltransferase activity in American white subjects and black subjects. Clinical pharmacology and therapeutics. 1994;55(1):15-20.
- 48. Yates CR, Krynetski EY, Loennechen T, Fessing MY, Tai HL, Pui CH, et al. Molecular diagnosis of thiopurine S-methyltransferase deficiency: genetic basis for azathioprine and mercaptopurine intolerance. Annals of internal medicine. 1997;126(8):608-14.
- 49. Holme SA, Duley JA, Sanderson J, Routledge PA, Anstey AV. Erythrocyte thiopurine methyl transferase assessment prior to azathioprine use in the UK. QJM: monthly journal of the Association of Physicians. 2002;95(7):439-44.
- 50. Bloomfeld RS, Onken JE. Mercaptopurine metabolite results in clinical gastroenterology practice. Alimentary pharmacology & therapeutics. 2003;17(1):69-73.
- 51. McLeod HL, Coulthard S, Thomas AE, Pritchard SC, King DJ, Richards SM, et al. Analysis of thiopurine methyltransferase variant alleles in childhood acute lymphoblastic leukaemia. British journal of haematology. 1999;105(3):696-700.
- 52. Black AJ, McLeod HL, Capell HA, Powrie RH, Matowe LK, Pritchard SC, et al. Thiopurine methyltransferase genotype predicts therapy-limiting severe toxicity from azathioprine. Annals of internal medicine. 1998;129(9):716-8.

- 53. Pandya B, Thomson W, Poulton K, Bruce I, Payne D, Qasim F. Azathioprine toxicity and thiopurine methyltransferase genotype in renal transplant patients. Transplant Proc. 2002;34(5):1642-5.
- 54. Seidman EG. Clinical use and practical application of TPMT enzyme and 6-mercaptopurine metabolite monitoring in IBD. Reviews in gastroenterological disorders. 2003;3 Suppl 1:S30-8.
- 55. Murphy LA, Atherton D. A retrospective evaluation of azathioprine in severe childhood atopic eczema, using thiopurine methyltransferase levels to exclude patients at high risk of myelosuppression. The British journal of dermatology. 2002;147(2):308-15.
- 56. Tavadia SM, Mydlarski PR, Reis MD, Mittmann N, Pinkerton PH, Shear N, et al. Screening for azathioprine toxicity: a pharmacoeconomic analysis based on a target case. Journal of the American Academy of Dermatology. 2000;42(4):628-32.
- 57. Marra CA, Esdaile JM, Anis AH. Practical pharmacogenetics: the cost effectiveness of screening for thiopurine s-methyltransferase polymorphisms in patients with rheumatological conditions treated with azathioprine. The Journal of rheumatology. 2002;29(12):2507-12.
- 58. Tan BB, Lear JT, Gawkrodger DJ, English JS. Azathioprine in dermatology: a survey of current practice in the U.K. The British journal of dermatology. 1997;136(3):351-5.
- 59. Rosendaal FR. The Scylla and Charybdis of oral anticoagulant treatment. The New England journal of medicine. 1996;335(8):587-9.
- 60. James AH, Britt RP, Raskino CL, Thompson SG. Factors affecting the maintenance dose of warfarin. J Clin Pathol. 1992;45(8):704-6.
- 61. Penning-van Beest FJ, van Meegen E, Rosendaal FR, Stricker BH. Characteristics of anticoagulant therapy and comorbidity related to overanticoagulation. Thromb Haemost. 2001;86(2):569-74.
- 62. Hylek EM, Skates SJ, Sheehan MA, Singer DE. An analysis of the lowest effective intensity of prophylactic anticoagulation for patients with nonrheumatic atrial fibrillation. The New England journal of medicine. 1996;335(8):540-6.
- 63. Pirmohamed M. Warfarin: almost 60 years old and still causing problems. British journal of clinical pharmacology. 2006;62(5):509-11.
- 64. Penning-van Beest FJ, Geleijnse JM, van Meegen E, Vermeer C, Rosendaal FR, Stricker BH. Lifestyle and diet as risk factors for overanticoagulation. Journal of clinical epidemiology. 2002;55(4):411-7.
- 65. Carlquist JF, Horne BD, Muhlestein JB, Lappe DL, Whiting BM, Kolek MJ, et al. Genotypes of the cytochrome p450 isoform, CYP2C9, and the vitamin K epoxide reductase complex subunit 1 conjointly determine stable warfarin dose: a prospective study. Journal of thrombosis and thrombolysis. 2006;22(3):191-7.
- 66. Schalekamp T, van Geest-Daalderop JH, Kramer MH, van Holten-Verzantvoort AT, de Boer A. Coumarin anticoagulants and cotrimoxazole: avoid the combination rather than manage the interaction. Eur J Clin Pharmacol. 2007;63(4):335-43.
- 67. Schalekamp T, Klungel OH, Souverein PC, de Boer A. Increased bleeding risk with concurrent use of selective serotonin reuptake inhibitors and coumarins. Archives of internal medicine. 2008;168(2):180-5.
- 68. Gage BF, Eby C, Milligan PE, Banet GA, Duncan JR, McLeod HL. Use of pharmacogenetics and clinical factors to predict the maintenance dose of warfarin. Thromb Haemost. 2004;91(1):87-94.
- 69. Bodin L, Verstuyft C, Tregouet DA, Robert A, Dubert L, Funck-Brentano C, et al. Cytochrome P450 2C9 (CYP2C9) and vitamin K epoxide reductase (VKORC1) genotypes as determinants of acenocoumarol sensitivity. Blood. 2005;106(1):135-40.
- 70. Wadelius M, Chen LY, Lindh JD, Eriksson N, Ghori MJ, Bumpstead S, et al. The largest prospective warfarin-treated cohort supports genetic forecasting. Blood. 2009;113(4):784-92.
- 71. Schalekamp T, Brasse BP, Roijers JF, Chahid Y, van Geest-Daalderop JH, de Vries-Goldschmeding H, et al. VKORC1 and CYP2C9 genotypes and acenocoumarol anticoagulation status: interaction between both genotypes affects overanticoagulation. Clinical pharmacology and therapeutics. 2006;80(1):13-22.

- 72. Schalekamp T, Brasse BP, Roijers JF, van Meegen E, van der Meer FJ, van Wijk EM, et al. VKORC1 and CYP2C9 genotypes and phenprocoumon anticoagulation status: interaction between both genotypes affects dose requirement. Clinical pharmacology and therapeutics. 2007;81(2):185-93.
- 73. Reitsma PH, van der Heijden JF, Groot AP, Rosendaal FR, Buller HR. A C1173T dimorphism in the VKORC1 gene determines coumarin sensitivity and bleeding risk. PLoS Med. 2005;2(10):e312.
- 74. Wadelius M, Chen LY, Downes K, Ghori J, Hunt S, Eriksson N, et al. Common VKORC1 and GGCX polymorphisms associated with warfarin dose. Pharmacogenomics J. 2005;5(4):262-70.
- 75. D'Andrea G, D'Ambrosio RL, Di Perna P, Chetta M, Santacroce R, Brancaccio V, et al. A polymorphism in the VKORC1 gene is associated with an interindividual variability in the dose-anticoagulant effect of warfarin. Blood. 2005;105(2):645-9.
- 76. Takeuchi F, McGinnis R, Bourgeois S, Barnes C, Eriksson N, Soranzo N, et al. A genome-wide association study confirms VKORC1, CYP2C9, and CYP4F2 as principal genetic determinants of warfarin dose. PLoS genetics. 2009;5(3):e1000433.
- 77. Caldwell MD, Awad T, Johnson JA, Gage BF, Falkowski M, Gardina P, et al. CYP4F2 genetic variant alters required warfarin dose. Blood. 2008;111(8):4106-12.
- 78. Schelleman H, Chen J, Chen Z, Christie J, Newcomb CW, Brensinger CM, et al. Dosing algorithms to predict warfarin maintenance dose in Caucasians and African Americans. Clinical pharmacology and therapeutics. 2008;84(3):332-9.
- 79. International Warfarin Pharmacogenetics C, Klein TE, Altman RB, Eriksson N, Gage BF, Kimmel SE, et al. Estimation of the warfarin dose with clinical and pharmacogenetic data. The New England journal of medicine. 2009;360(8):753-64.
- 80. Perini JA, Struchiner CJ, Silva-Assuncao E, Santana IS, Rangel F, Ojopi EB, et al. Pharmacogenetics of warfarin: development of a dosing algorithm for brazilian patients. Clinical pharmacology and therapeutics. 2008;84(6):722-8.
- 81. Sconce EA, Khan TI, Wynne HA, Avery P, Monkhouse L, King BP, et al. The impact of CYP2C9 and VKORC1 genetic polymorphism and patient characteristics upon warfarin dose requirements: proposal for a new dosing regimen. Blood. 2005;106(7):2329-33.
- Tham LS, Goh BC, Nafziger A, Guo JY, Wang LZ, Soong R, et al. A warfarin-dosing model in Asians that uses single-nucleotide polymorphisms in vitamin K epoxide reductase complex and cytochrome P450 2C9. Clinical pharmacology and therapeutics. 2006;80(4):346-55.
- 83. Gage BF, Eby C, Johnson JA, Deych E, Rieder MJ, Ridker PM, et al. Use of pharmacogenetic and clinical factors to predict the therapeutic dose of warfarin. Clinical pharmacology and therapeutics. 2008;84(3):326-31.
- 84. Eckman MH, Rosand J, Greenberg SM, Gage BF. Cost-effectiveness of using pharmacogenetic information in warfarin dosing for patients with nonvalvular atrial fibrillation. Annals of internal medicine. 2009;150(2):73-83.
- 85. Schalekamp T, Boink GJ, Visser LE, Stricker BH, de Boer A, Klungel OH. CYP2C9 genotyping in acenocoumarol treatment: is it a cost-effective addition to international normalized ratio monitoring? Clinical pharmacology and therapeutics. 2006;79(6):511-20.
- 86. Hughes DA, Pirmohamed M. Warfarin pharmacogenetics: economic considerations. Pharmacoeconomics. 2007;25(11):899-902.
- 87. Slamon DJ, Leyland-Jones B, Shak S, Fuchs H, Paton V, Bajamonde A, et al. Use of chemotherapy plus a monoclonal antibody against HER2 for metastatic breast cancer that overexpresses HER2. The New England journal of medicine. 2001;344(11):783-92.
- 88. Lievre A, Bachet JB, Boige V, Cayre A, Le Corre D, Buc E, et al. KRAS mutations as an independent prognostic factor in patients with advanced colorectal cancer treated with cetuximab. Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2008;26(3):374-9.
- 89. Joensuu H, Roberts PJ, Sarlomo-Rikala M, Andersson LC, Tervahartiala P, Tuveson D, et al. Effect of the tyrosine kinase inhibitor STI571 in a patient with a metastatic gastrointestinal stromal tumor. The New England journal of medicine. 2001;344(14):1052-6.
- 90. DiMasi JA, Grabowski HG. Economics of new oncology drug development. Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2007;25(2):209-16.

- 91. Von Hoff DD. There are no bad anticancer agents, only bad clinical trial designs--twenty-first Richard and Hinda Rosenthal Foundation Award Lecture. Clinical Cancer Research. 1998;4(5):1079-86.
- 92. Thatcher N, Chang A, Parikh P, Rodrigues Pereira J, Ciuleanu T, von Pawel J, et al. Gefitinib plus best supportive care in previously treated patients with refractory advanced non-small-cell lung cancer: results from a randomised, placebo-controlled, multicentre study (Iressa Survival Evaluation in Lung Cancer). Lancet. 2005;366(9496):1527-37.
- 93. Mok TS, Wu YL, Thongprasert S, Yang CH, Chu DT, Saijo N, et al. Gefitinib or carboplatin-paclitaxel in pulmonary adenocarcinoma. The New England journal of medicine. 2009;361(10):947-57.
- 94. Von Hoff DD, Stephenson JJ, Jr., Rosen P, Loesch DM, Borad MJ, Anthony S, et al. Pilot study using molecular profiling of patients' tumors to find potential targets and select treatments for their refractory cancers. Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2010;28(33):4877-83.
- 95. Doroshow JH. Selecting systemic cancer therapy one patient at a time: is there a role for molecular profiling of individual patients with advanced solid tumors? Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2010;28(33):4869-71.
- 96. Kim ES, Herbst RS, Wistuba, II, Lee JJ, Blumenschein GR, Jr., Tsao A, et al. The BATTLE trial: personalizing therapy for lung cancer. Cancer discovery. 2011;1(1):44-53.
- 97. Mozaffarian D, Benjamin EJ, Go AS, Arnett DK, Blaha MJ, Cushman M, et al. Executive Summary: Heart Disease and Stroke Statistics—2015 Update. Circulation. 2015;131(4):434-41.
- 98. Greenland P, Lauer MS. Cholesterol Lowering in 2015: Still Answering Questions About How and in Whom. JAMA. 2015;314(2):127-8.
- 99. Cholesterol Treatment Trialists Collaborators. The effects of lowering LDL cholesterol with statin therapy in people at low risk of vascular disease: meta-analysis of individual data from 27 randomised trials. The Lancet. 2012;380(9841):581-90.
- 100. Cholesterol Treatment Trialists Collaborators. Efficacy and safety of LDL-lowering therapy among men and women: meta-analysis of individual data from 174 000 participants in 27 randomised trials. The Lancet. 2015;385(9976):1397-405.
- 101. Taylor F, Huffman MD, Macedo AF, Moore TH, Burke M, Davey Smith G, et al. Statins for the primary prevention of cardiovascular disease. The Cochrane database of systematic reviews. 2013(1):CD004816.
- 102. Stone NJ, Robinson JG, Lichtenstein AH, Bairey Merz CN, Blum CB, Eckel RH, et al. 2013 ACC/AHA guideline on the treatment of blood cholesterol to reduce atherosclerotic cardiovascular risk in adults: a report of the American College of Cardiology/American Heart Association Task Force on Practice Guidelines. J Am Coll Cardiol. 2014;63(25 Pt B):2889-934.
- 103. Pencina MJ, Navar-Boggan AM, D'Agostino RB, Sr., Williams K, Neely B, Sniderman AD, et al. Application of new cholesterol guidelines to a population-based sample. The New England journal of medicine. 2014;370(15):1422-31.
- 104. Hirsh BJ, Smilowitz NR, Rosenson RS, Fuster V, Sperling LS. Utilization of and Adherence to Guideline-Recommended Lipid-Lowering Therapy After Acute Coronary Syndrome: Opportunities for Improvement. J Am Coll Cardiol. 2015;66(2):184-92.
- 105. Bermingham M, Hayden J, Dawkins I, Miwa S, Gibson D, McDonald K, et al. Prospective analysis of LDL-C goal achievement and self-reported medication adherence among statin users in primary care. Clin Ther. 2011;33(9):1180-9.
- 106. Ho PM, Magid DJ, Shetterly SM, Olson KL, Maddox TM, Peterson PN, et al. Medication nonadherence is associated with a broad range of adverse outcomes in patients with coronary artery disease. Am Heart J. 2008;155(4):772-9.
- 107. Vodonos A, Ostapenko I, Toledano R, Henkin Y, Zahger D, Wolak T, et al. Statin adherence and LDL cholesterol levels. Should we assess adherence prior to statin upgrade? European journal of internal medicine. 2015;26(4):268-72.
- 108. Franklin JM, Krumme AA, Tong AY, Shrank WH, Matlin OS, Brennan TA, et al. Association between trajectories of statin adherence and subsequent cardiovascular events. Pharmacoepidemiol Drug Saf. 2015;24(10):1105-13.

- 109. De Vera MA, Bhole V, Burns LC, Lacaille D. Impact of statin adherence on cardiovascular disease and mortality outcomes: a systematic review. British journal of clinical pharmacology. 2014;78(4):684-98.
- 110. Osterberg L, Blaschke T. Adherence to medication. The New England journal of medicine. 2005;353(5):487-97.
- 111. Fung V, Sinclair F, Wang H, Dailey D, Hsu J, Shaber R. Patients' perspectives on nonadherence to statin therapy: a focus-group study. Perm J. 2010;14(1):4-10.
- 112. Cohen JD, Brinton EA, Ito MK, Jacobson TA. Understanding Statin Use in America and Gaps in Patient Education (USAGE): an internet-based survey of 10,138 current and former statin users. J Clin Lipidol. 2012;6(3):208-15.
- 113. Ong FS, Deignan JL, Kuo JZ, Bernstein KE, Rotter JI, Grody WW, et al. Clinical utility of pharmacogenetic biomarkers in cardiovascular therapeutics: a challenge for clinical implementation. Pharmacogenomics. 2012;13(4):465-75.
- 114. Pasternak RC, Smith SC, Bairey-Merz CN, Grundy SM, Cleeman JI, Lenfant C. ACC/AHA/NHLBI clinical advisory on the use and safety of statins. Journal of the American College of Cardiology. 2002;40(3):567-72.
- 115. Thompson PD, Clarkson PM, Rosenson RS, National Lipid Association Statin Safety Task Force Muscle Safety Expert P. An assessment of statin safety by muscle experts. Am J Cardiol. 2006;97(8A):69C-76C.
- 116. Alfirevic A, Neely D, Armitage J, Chinoy H, Cooper RG, Laaksonen R, et al. Phenotype standardization for statin-induced myotoxicity. Clinical pharmacology and therapeutics. 2014;96(4):470-6.
- 117. Patel J, Superko HR, Martin SS, Blumenthal RS, Christopher-Stine L. Genetic and immunologic susceptibility to statin-related myopathy. Atherosclerosis. 2015;240(1):260-71.
- 118. Stroes ES, Thompson PD, Corsini A, Vladutiu GD, Raal FJ, Ray KK, et al. Statin-associated muscle symptoms: impact on statin therapy-European Atherosclerosis Society Consensus Panel Statement on Assessment, Aetiology and Management. Eur Heart J. 2015;36(17):1012-22.
- 119. Group SC, Link E, Parish S, Armitage J, Bowman L, Heath S, et al. SLCO1B1 variants and statin-induced myopathy--a genomewide study. The New England journal of medicine. 2008;359(8):789-99.
- 120. Hou Q, Li S, Li L, Li Y, Sun X, Tian H. Association Between SLCO1B1 Gene T521C Polymorphism and Statin-Related Myopathy Risk: A Meta-Analysis of Case-Control Studies. Medicine. 2015;94(37):e1268.
- 121. Pasanen MK, Backman JT, Neuvonen PJ, Niemi M. Frequencies of single nucleotide polymorphisms and haplotypes of organic anion transporting polypeptide 1B1 SLCO1B1 gene in a Finnish population. Eur J Clin Pharmacol. 2006;62(6):409-15.
- 122. Kimoto E, Yoshida K, Balogh LM, Bi YA, Maeda K, El-Kattan A, et al. Characterization of organic anion transporting polypeptide (OATP) expression and its functional contribution to the uptake of substrates in human hepatocytes. Mol Pharm. 2012;9(12):3535-42.
- 123. Voora D, Shah SH, Spasojevic I, Ali S, Reed CR, Salisbury BA, et al. The SLCO1B1\*5 genetic variant is associated with statin-induced side effects. J Am Coll Cardiol. 2009;54(17):1609-16.
- Wilke RA, Ramsey LB, Johnson SG, Maxwell WD, McLeod HL, Voora D, et al. The clinical pharmacogenomics implementation consortium: CPIC guideline for SLCO1B1 and simvastatin-induced myopathy. Clinical pharmacology and therapeutics. 2012;92(1):112-7.
- 125. Birmingham BK, Bujac SR, Elsby R, Azumaya CT, Wei C, Chen Y, et al. Impact of ABCG2 and SLCO1B1 polymorphisms on pharmacokinetics of rosuvastatin, atorvastatin and simvastatin acid in Caucasian and Asian subjects: a class effect? Eur J Clin Pharmacol. 2015;71(3):341-55.
- de Keyser CE, Peters BJ, Becker ML, Visser LE, Uitterlinden AG, Klungel OH, et al. The SLCO1B1 c. 521T> C polymorphism is associated with dose decrease or switching during statin therapy in the Rotterdam Study. Pharmacogenetics and genomics. 2014;24(1):43-51.
- 127. Danik JS, Chasman DI, MacFadyen JG, Nyberg F, Barratt BJ, Ridker PM. Lack of association between SLCO1B1 polymorphisms and clinical myalgia following rosuvastatin therapy. Am Heart J. 2013;165(6):1008-14.

- 128. Martin NG, Li KW, Murray H, Putt W, Packard CJ, Humphries SE. The effects of a single nucleotide polymorphism in SLCO1B1 on the pharmacodynamics of pravastatin. British journal of clinical pharmacology. 2012;73(2):303-6.
- 129. Niemi M, Pasanen MK, Neuvonen PJ. Organic anion transporting polypeptide 1B1: a genetically polymorphic transporter of major importance for hepatic drug uptake. Pharmacol Rev. 2011;63(1):157-81.
- 130. Ramsey LB, Johnson SG, Caudle KE, Haidar CE, Voora D, Wilke RA, et al. The clinical pharmacogenetics implementation consortium guideline for SLCO1B1 and simvastatin-induced myopathy: 2014 update. Clinical pharmacology and therapeutics. 2014;96(4):423-8.
- 131. Voora D, Ginsburg GS. Clinical application of cardiovascular pharmacogenetics. J Am Coll Cardiol. 2012;60(1):9-20.
- 132. Donnelly LA, Doney AS, Tavendale R, Lang CC, Pearson ER, Colhoun HM, et al. Common nonsynonymous substitutions in SLCO1B1 predispose to statin intolerance in routinely treated individuals with type 2 diabetes: a go-DARTS study. Clinical pharmacology and therapeutics. 2011;89(2):210-6.
- 133. Li JH, Joy SV, Haga SB, Orlando LA, Kraus WE, Ginsburg GS, et al. Genetically guided statin therapy on statin perceptions, adherence, and cholesterol lowering: a pilot implementation study in primary care patients. J Pers Med. 2014;4(2):147-62.
- 134. P. D. Q. Supportive Palliative Care Editorial Board. Cancer Pain (PDQ(R)): Health Professional Version. PDQ Cancer Information Summaries. Bethesda (MD): National Cancer Institute (US); 2002.
- 135. National Comprehensive Cancer Network. NCCN clinical practice guidelines in oncology: Adult cancer pain. Version 2.2016 <a href="https://www.nccn.org/professionals/physician\_gls/default.aspx2016">https://www.nccn.org/professionals/physician\_gls/default.aspx2016</a> [
- 136. Temel JS, Greer JA, Muzikansky A, Gallagher ER, Admane S, Jackson VA, et al. Early palliative care for patients with metastatic non-small-cell lung cancer. The New England journal of medicine. 2010;363(8):733-42.
- 137. Caraceni A, Hanks G, Kaasa S, Bennett MI, Brunelli C, Cherny N, et al. Use of opioid analgesics in the treatment of cancer pain: evidence-based recommendations from the EAPC. Lancet Oncol. 2012;13(2):e58-68.
- 138. Fine PG, Portenoy RK, Ad Hoc Expert Panel on Evidence R, Guidelines for Opioid R. Establishing "best practices" for opioid rotation: conclusions of an expert panel. J Pain Symptom Manage. 2009;38(3):418-25.
- Thao F, Chang VT, Cleeland C, Cleary JF, Mitchell EP, Wagner LI, et al. Determinants of pain severity changes in ambulatory patients with cancer: an analysis from Eastern Cooperative Oncology Group trial E2Z02. Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2014;32(4):312-9.
- 140. Angst MS, Phillips NG, Drover DR, Tingle M, Ray A, Swan GE, et al. Pain sensitivity and opioid analgesia: a pharmacogenomic twin study. Pain. 2012;153(7):1397-409.
- 141. Fillingim RB, Wallace MR, Herbstman DM, Ribeiro-Dasilva M, Staud R. Genetic contributions to pain: a review of findings in humans. Oral Dis. 2008;14(8):673-82.
- 142. Gan SH, Ismail R, Wan Adnan WA, Zulmi W. Impact of CYP2D6 genetic polymorphism on tramadol pharmacokinetics and pharmacodynamics. Mol Diagn Ther. 2007;11(3):171-81.
- 143. Susce MT, Murray-Carmichael E, de Leon J. Response to hydrocodone, codeine and oxycodone in a CYP2D6 poor metabolizer. Prog Neuropsychopharmacol Biol Psychiatry. 2006;30(7):1356-8.
- 144. Crews KR, Gaedigk A, Dunnenberger HM, Leeder JS, Klein TE, Caudle KE, et al. Clinical Pharmacogenetics Implementation Consortium guidelines for cytochrome P450 2D6 genotype and codeine therapy: 2014 update. Clinical pharmacology and therapeutics. 2014;95(4):376-82.
- 145. Baber M, Chaudhry S, Kelly L, Ross C, Carleton B, Berger H, et al. The pharmacogenetics of codeine pain relief in the postpartum period. Pharmacogenomics J. 2015;15(5):430-5.
- 146. Ciszkowski C, Madadi P, Phillips MS, Lauwers AE, Koren G. Codeine, ultrarapid-metabolism genotype, and postoperative death. The New England journal of medicine. 2009;361(8):827-8.

- 147. Gasche Y, Daali Y, Fathi M, Chiappe A, Cottini S, Dayer P, et al. Codeine intoxication associated with ultrarapid CYP2D6 metabolism. The New England journal of medicine. 2004;351(27):2827-31.
- 148. Eckhardt K, Li S, Ammon S, Schanzle G, Mikus G, Eichelbaum M. Same incidence of adverse drug events after codeine administration irrespective of the genetically determined differences in morphine formation. Pain. 1998;76(1-2):27-33.
- Lotsch J, Rohrbacher M, Schmidt H, Doehring A, Brockmoller J, Geisslinger G. Can extremely low or high morphine formation from codeine be predicted prior to therapy initiation? Pain. 2009;144(1-2):119-24.
- 150. Andreassen TN, Eftedal I, Klepstad P, Davies A, Bjordal K, Lundstrom S, et al. Do CYP2D6 genotypes reflect oxycodone requirements for cancer patients treated for cancer pain? A cross-sectional multicentre study. Eur J Clin Pharmacol. 2012;68(1):55-64.
- 151. Samer CF, Daali Y, Wagner M, Hopfgartner G, Eap CB, Rebsamen MC, et al. The effects of CYP2D6 and CYP3A activities on the pharmacokinetics of immediate release oxycodone. Br J Pharmacol. 2010;160(4):907-18.
- Twisler ST, Enggaard TP, Noehr-Jensen L, Pedersen RS, Mikkelsen S, Nielsen F, et al. The hypoalgesic effect of oxycodone in human experimental pain models in relation to the CYP2D6 oxidation polymorphism. Basic Clin Pharmacol Toxicol. 2009;104(4):335-44.
- 153. Zwisler ST, Enggaard TP, Mikkelsen S, Brosen K, Sindrup SH. Impact of the CYP2D6 genotype on post-operative intravenous oxycodone analgesia. Acta Anaesthesiol Scand. 2010;54(2):232-40.
- 154. Johnson D, Jorgensen AL. The influence of HLA-B\*15:02 and HLA-A\*31:01 on the risk of developing adverse skin reactions to carbamazepine: protocol for two systematic reviews. CRD42019161000. 2019 [Available from: https://www.crd.york.ac.uk/PROSPERO/display\_record.php?RecordID=161000.
- 155. Yip VL, Marson AG, Jorgensen AL, Pirmohamed M, Alfirevic A. HLA genotype and carbamazepine-induced cutaneous adverse drug reactions: a systematic review. Clinical pharmacology and therapeutics. 2012;92(6):757-65.
- 156. Tangamornsuksan W, Chaiyakunapruk N, Somkrua R, Lohitnavy M, Tassaneeyakul W. Relationship between the HLA-B\*1502 allele and carbamazepine-induced Stevens-Johnson syndrome and toxic epidermal necrolysis: a systematic review and meta-analysis. JAMA Dermatol. 2013;149(9):1025-32.
- 157. Grover S, Kukreti R. HLA alleles and hypersensitivity to carbamazepine: an updated systematic review with meta-analysis. Pharmacogenetics and genomics. 2014;24(2):94-112.
- Ryan M, Gerard K. Using discrete choice experiments to value health care programmes: current practice and future research reflections. Applied health economics and health policy. 2003;2(1):55-64.
- 159. de Bekker-Grob EW, Ryan M, Gerard K. Discrete choice experiments in health economics: a review of the literature. Health Econ. 2012;21(2):145-72.
- 160. Clark MD, Determann D, Petrou S, Moro D, de Bekker-Grob EW. Discrete choice experiments in health economics: a review of the literature. Pharmacoeconomics. 2014;32(9):883-902.
- 161. Soekhai V, de Bekker-Grob EW, Ellis AR, Vass CM. Discrete Choice Experiments in Health Economics: Past, Present and Future. Pharmacoeconomics. 2019;37(2):201-26.
- 162. Guerra RL, Castaneda L, de Albuquerque RCR, Ferreira CBT, Correa FM, Fernandes RRA, et al. Patient Preferences for Breast Cancer Treatment Interventions: A Systematic Review of Discrete Choice Experiments. Patient. 2019;12(6):559-69.
- 163. Trapero-Bertran M, Rodriguez-Martin B, Lopez-Bastida J. What attributes should be included in a discrete choice experiment related to health technologies? A systematic literature review. PLoS One. 2019;14(7):e0219905.
- 164. Bien DR, Danner M, Vennedey V, Civello D, Evers SM, Hiligsmann M. Patients' Preferences for Outcome, Process and Cost Attributes in Cancer Treatment: A Systematic Review of Discrete Choice Experiments. The Patient Patient-Centered Outcomes Research. 2017;10(5):553-65.

- Harrison M, Milbers K, Hudson M, Bansback N. Do patients and health care providers have discordant preferences about which aspects of treatments matter most? Evidence from a systematic review of discrete choice experiments. BMJ open. 2017;7(5):e014719.
- 166. Vass C, Rigby D, Payne K. The Role of Qualitative Research Methods in Discrete Choice Experiments. Medical decision making: an international journal of the Society for Medical Decision Making. 2017;37(3):298-313.
- 167. Harrison M, Rigby D, Vass C, Flynn T, Louviere J, Payne K. Risk as an attribute in discrete choice experiments: a systematic review of the literature. Patient. 2014;7(2):151-70.
- 168. Bien DR, Danner M, Vennedey V, Civello D, Evers SM, Hiligsmann M. Patients' Preferences for Outcome, Process and Cost Attributes in Cancer Treatment: A Systematic Review of Discrete Choice Experiments. Patient. 2017;10(5):553-65.
- 169. Ballinger TJ, Kassem N, Shen F, Jiang G, Smith ML, Railey E, et al. Discerning the clinical relevance of biomarkers in early stage breast cancer. Breast Cancer Res Treat. 2017;164(1):89-97.
- 170. Boeri M, McMichael AJ, Kane JPM, O'Neill FA, Kee F. Physician-Specific Maximum Acceptable Risk in Personalized Medicine: Implications for Medical Decision Making. Medical decision making: an international journal of the Society for Medical Decision Making. 2018;38(5):593-600.
- 171. Chan SL, Wen Low JJ, Lim YW, Finkelstein E, Farooqui MA, Chia KS, et al. Willingness-to-pay and preferences for warfarin pharmacogenetic testing in Chinese warfarin patients and the Chinese general public. Per Med. 2013;10(2):127-37.
- 172. Dong D, Ozdemir S, Mong Bee Y, Toh SA, Bilger M, Finkelstein E. Measuring High-Risk Patients' Preferences for Pharmacogenetic Testing to Reduce Severe Adverse Drug Reaction: A Discrete Choice Experiment. Value in health: the journal of the International Society for Pharmacoeconomics and Outcomes Research. 2016;19(6):767-75.
- 173. Herbild L, Bech M, Gyrd-Hansen D. Estimating the Danish populations' preferences for pharmacogenetic testing using a discrete choice experiment. The case of treating depression. Value in health: the journal of the International Society for Pharmacoeconomics and Outcomes Research. 2009;12(4):560-7.
- 174. Issa AM, Tufail W, Atehortua N, McKeever J. A national study of breast and colorectal cancer patients' decision-making for novel personalized medicine genomic diagnostics. Per Med. 2013;10(3):245-56.
- 175. Liede A, Mansfield CA, Metcalfe KA, Price MA, Kathleen Cuningham Foundation Consortium for Research into Familial Breast C, Snyder C, et al. Preferences for breast cancer risk reduction among BRCA1/BRCA2 mutation carriers: a discrete-choice experiment. Breast Cancer Res Treat. 2017;165(2):433-44.
- 176. Marshall DA, Deal K, Bombard Y, Leighl N, MacDonald KV, Trudeau M. How do women trade-off benefits and risks in chemotherapy treatment decisions based on gene expression profiling for early-stage breast cancer? A discrete choice experiment. BMJ open. 2016;6(6):e010981.
- 177. Marshall DA, Gonzalez JM, MacDonald KV, Johnson FR. Estimating Preferences for Complex Health Technologies: Lessons Learned and Implications for Personalized Medicine. Value in health: the journal of the International Society for Pharmacoeconomics and Outcomes Research. 2017;20(1):32-9.
- 178. Najafzadeh M, Johnston KM, Peacock SJ, Connors JM, Marra MA, Lynd LD, et al. Genomic testing to determine drug response: measuring preferences of the public and patients using Discrete Choice Experiment (DCE). BMC health services research. 2013;13(1):454.
- 179. Payne K, Fargher EA, Roberts SA, Tricker K, Elliott RA, Ratcliffe J, et al. Valuing pharmacogenetic testing services: a comparison of patients' and health care professionals' preferences. Value in health: the journal of the International Society for Pharmacoeconomics and Outcomes Research. 2011;14(1):121-34.
- 180. Powell G, Holmes EA, Plumpton CO, Ring A, Baker GA, Jacoby A, et al. Pharmacogenetic testing prior to carbamazepine treatment of epilepsy: patients' and physicians' preferences for testing and service delivery. British journal of clinical pharmacology. 2015;80(5):1149-59.
- 181. Smith ML, White CB, Railey E, Sledge GW, Jr. Examining and predicting drug preferences of patients with metastatic breast cancer: using conjoint analysis to examine attributes of paclitaxel and capecitabine. Breast Cancer Res Treat. 2014;145(1):83-9.