

# iPLEX<sup>®</sup> ADME CYP2C9 / VKORC1 Panel<sup>1</sup> v1.0

Available through Assays by Sequenom

Investigate and confirm pharmacogenetic biomarkers in CYP2C9 and VKORC1 genes associated with drug metabolism with high accuracy and rapid results.

The iPLEX ADME CYP2C9 / VKORC1 Panel is a set of 45 pre-designed and pre-validated SNP assays for use in the screening of polymorphisms in CYP2C9 (36 SNPs) and VKORC1 (9 SNPs) genes. Screening for CYP2C9 and VKORC1 variants allows clinical researchers develop dosing protocols and surveillance techniques toward model drugs and experimental biomarkers.

- Analyze 45 mutations in 2 pharmacogenetically relevant genes
- Obtain biologically relevant data with CYP2C9 / VKORC1 panel covering most of the known haplotypes for both genes
- Use as little as 30 ng of DNA per sample

## Contact:

**Sequenom Inc.** USA: 1 877 4GENOME  
3595 John Hopkins Ct EU: (+49) 40-899676-0  
San Diego, CA 92121 AP: (+61) 7 3845 3691



Gene	Alleles			
<b>CYP2C9</b>	CYP2C9*1A	CYP2C9*9	CYP2C9*22	
	CYP2C9*1B	CYP2C9*10	CYP2C9*23	
	CYP2C9*1C	CYP2C9*11A	CYP2C9*24	
	CYP2C9*1D	CYP2C9*11B	CYP2C9*25	
	CYP2C9*2A	CYP2C9*12	CYP2C9*26	
	CYP2C9*2B	CYP2C9*13	CYP2C9*27	
	CYP2C9*2C	CYP2C9*14	CYP2C9*28	
	CYP2C9*3A	CYP2C9*15	CYP2C9*29	
	CYP2C9*3B	CYP2C9*16	CYP2C9*30	
	CYP2C9*4	CYP2C9*17	CYP2C9*31	
	CYP2C9*5	CYP2C9*18	CYP2C9*32	
	CYP2C9*6	CYP2C9*19	CYP2C9*33	
	CYP2C9*7	CYP2C9*20	CYP2C9*34	
	CYP2C9*8	CYP2C9*21	CYP2C9*35	
	<b>Gene</b>	<b>Alleles</b>		
	<b>VKORC1</b>	*1	BHT4	
*2/H1		H2/H5		
*2A		H4		
*2B		H6		
*3		H7A		
*3F;BHT3		H7B		
*4		H8		
*7RE		H9		
BHT2RE				

<sup>1</sup> The iPLEX ADME CYP2C9 / VKORC1 Panel is For Research Use only. Not for use in diagnostic procedures.

SEQUENOM<sup>®</sup>

## Analysis Method

Screening with iPLEX ADME CYP2C9/VKORC1 panel is performed by PCR amplification and primer extension using the iPLEX® ADME CYP2C9/VKORC1 reagents. The panel consists of 3 multiplexed wells that are run on each sample using 10ng of input DNA per well. Mutations are detected after the sample run.

## Equipment and Software Required

- MassARRAY® System<sup>1</sup>
- MassARRAY NanoDispenser<sup>1</sup>
- Typer Software<sup>1</sup> version 4.0.20

## Ordering Information:

2 x 384 Format

10 x 384 Format

10 x 96 Format

iPLEX ADME CYP2C9/VKORC1 PCR Primers<sup>1</sup>

iPLEX ADME CYP2C9/VKORC1 Extend Primers<sup>1</sup>

PCR Reagent Sets

iPLEX Pro Extension Reagent Set<sup>1</sup>

To place an order, please contact the order desk at [orderdesk@sequenom.com](mailto:orderdesk@sequenom.com), or call (858) 202-9215.

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ASSAYS BY SEQUENOM

## CYP2C9 Allele Nomenclature

Allele	Nucleotide changes	Amino acid change	dbSNP IDs
<b>CYP2C9*1A</b>	None		
<b>CYP2C9*1B</b>	-2665_-2664delTG; -1188T>C		rs71486745; rs4918758
<b>CYP2C9*1C</b>	-1188T>C		rs4918758
<b>CYP2C9*1D</b>	-2665_-2664delTG		rs71486745
<b>CYP2C9*2A</b>	-1188T>C; 430C>T	R144C	rs4918758; rs1799853
<b>CYP2C9*2B</b>	-2665_-2664delTG; -1188T>C; 430C>T	R144C	rs71486745; rs4918758; rs1799853
<b>CYP2C9*2C</b>	430C>T	R144C	rs1799853
<b>CYP2C9*3A</b>	1075A>C	I359L	rs1057910
<b>CYP2C9*3B</b>	-1188T>C; 1075A>C	I359L	rs4918758; rs1057910
<b>CYP2C9*4</b>	1076T>C	I359T	rs56165452
<b>CYP2C9*5</b>	1080C>G	D360E	rs28371686
<b>CYP2C9*6</b>	818delA	273Frame shift	rs9332131
<b>CYP2C9*7</b>	55C>A	L19I	rs67807361
<b>CYP2C9*8</b>	449G>A	R150H	rs7900194
<b>CYP2C9*9</b>	752A>G	H251R	rs2256871
<b>CYP2C9*10</b>	815A>G	E272G	rs9332130
<b>CYP2C9*11A</b>	1003C>T	R335W	rs28371685
<b>CYP2C9*11B</b>	-2665_-2664delTG; -1188T>C; 1003C>T	R335W	rs71486745; rs4918758; rs28371685
<b>CYP2C9*12</b>	1465C>T	P489S	rs9332239
<b>CYP2C9*13</b>	269T>C	L90P	rs72558187
<b>CYP2C9*14</b>	374G>A	R125H	rs73994288
<b>CYP2C9*15</b>	485C>A (linkage with -1188T>C can not be excluded)	S162X	rs72558190
<b>CYP2C9*16</b>	-1188T>C; 895A>G	T299A	rs4918758; rs72558192
<b>CYP2C9*17</b>	1144C>T	P382S	C09C1144T
<b>CYP2C9*18</b>	-1188T>C; 1075A>C; 1190A>C; 1425A>T	I359L; D397A	rs4918758; rs1057910; rs72558193; rs1057911
<b>CYP2C9*19</b>	-1188T>C; 1362G>C	Q454H	rs4918758; C09G1362C
<b>CYP2C9*20</b>	-1188T>C; 208G>C	G70R	rs4918758; C09G208C
<b>CYP2C9*21</b>	89C>T	P30L	C09C89T
<b>CYP2C9*22</b>	121A>G	N41D	C09A121T
<b>CYP2C9*23</b>	226G>A	V76M	C09G226A
<b>CYP2C9*24</b>	1060G>A Existence of the CYP2C9*2 polymorphism 430C>T on the same allele can not be excluded	E354K	C09G1060A
<b>CYP2C9*25</b>	353_362delAGAAATGGAA	118Frameshift	rs72558188
<b>CYP2C9*26</b>	-1188T>C; 389C>G	T130R	rs4918758; rs5031019
<b>CYP2C9*27</b>	-2665_-2664delTG; -1188T>C; 449G>T	R150L	rs71486745; rs4918758; rs7900194
<b>CYP2C9*28</b>	641A>T	Q214L	C09A641T
<b>CYP2C9*29</b>	835C>A	P279T	C09C835A
<b>CYP2C9*30</b>	1429G>A	A477T	C09G1429A
<b>CYP2C9*31</b>	980T>C	I327T	rs57505750
<b>CYP2C9*32</b>	1468G>T	V490F	C09G1468T
<b>CYP2C9*33</b>	395G>A	R132Q	rs72558184
<b>CYP2C9*34</b>	1004G>A	R335Q	C09G1004A
<b>CYP2C9*35</b>	374G>T; 430C>T	R125L; R144C	rs73994288; rs1799853

## VKORC1 Allele Nomenclature

Allele	Nucleotide changes	Amino acid change	dbSNP IDs
<b>*1</b>	None		None
<b>*2/H1</b>	7566C>T; 11639G>A; 6484C>T		rs2359612; rs9923231; rs9934438
<b>*2A</b>	7566C>T; 6853G>C; 11639G>A; 6484C>T		rs2359612; rs8050894; rs9923231; rs9934438
<b>*2B</b>	5808T>G; 6484C>T; 6853G>C; 7566C>T; 11639G>A		rs2884737; rs9934438; rs8050894; rs2359612; rs9923231
<b>*3</b>	9041G>A		rs7294
<b>*3F;BHT3</b>	8773C>T; 9041G>A	P83L	rs7200749; rs7294
<b>*4</b>	6009C>T		rs17708472
<b>*7RE</b>	7566C>T; 11639G>A		rs2359612; rs9923231
<b>BHT4RE</b>	6009C>T; 6853G>C		rs17708472; rs8050894
<b>BHT4</b>	7566C>T		rs2359612
<b>H2/H5</b>	7566C>T; 5808T>G; 11639G>A; 6484C>T		rs2884737; rs2359612; rs9923231; rs9934438
<b>H4</b>	7566C>T; 6853G>C		rs2359612; rs8050894
<b>H6</b>	6853G>C		rs8050894
<b>H7A</b>	9041G>A; 6853G>C; 8773C>T	P83L	rs7294; rs8050894; rs7200749
<b>H7B</b>	9041G>A; 6853G>C		rs7294; rs8050894
<b>H8</b>	861C>A; 9041G>A; 6853G>C		rs17708472; rs7294; rs8050894
<b>H9</b>	6009C>T; 861C>A; 6853G>C		rs17708472; rs17880887; rs8050894

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