

**Using the VerifyNow P2Y12 Assay to Optimize Dual Antiplatelet
Therapy for Patients with Acute Stroke or Transient Ischemic
Attack: Associations between P2Y12 Reaction Units and
CYP2C19 Polymorphisms**



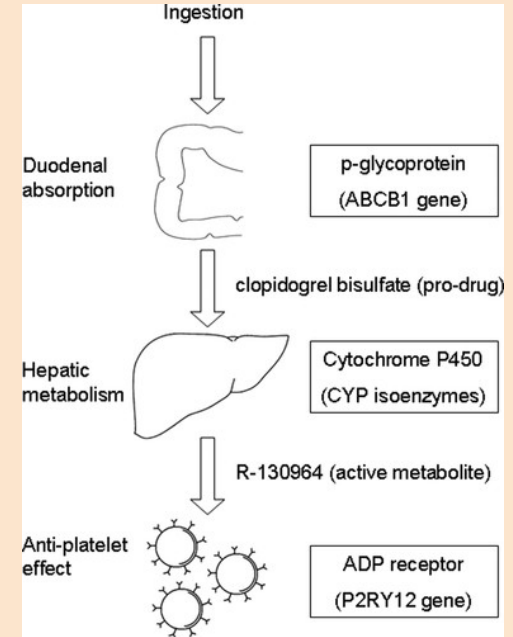
**THE QUEEN'S
MEDICAL CENTER**

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Background

- Certain patients with acute ischemic stroke (IS) or transient ischemic attack (TIA) are treated with dual antiplatelet therapy (DAPT) to reduce risk for recurrent stroke
 - CHANCE and POINT trials
- DAPT in these trials consisted of clopidogrel and aspirin
 - Clopidogrel requires hepatic transformation to active metabolite via the *CYP2C19* enzyme



Anderson et al., 2010, *Stroke*



Background

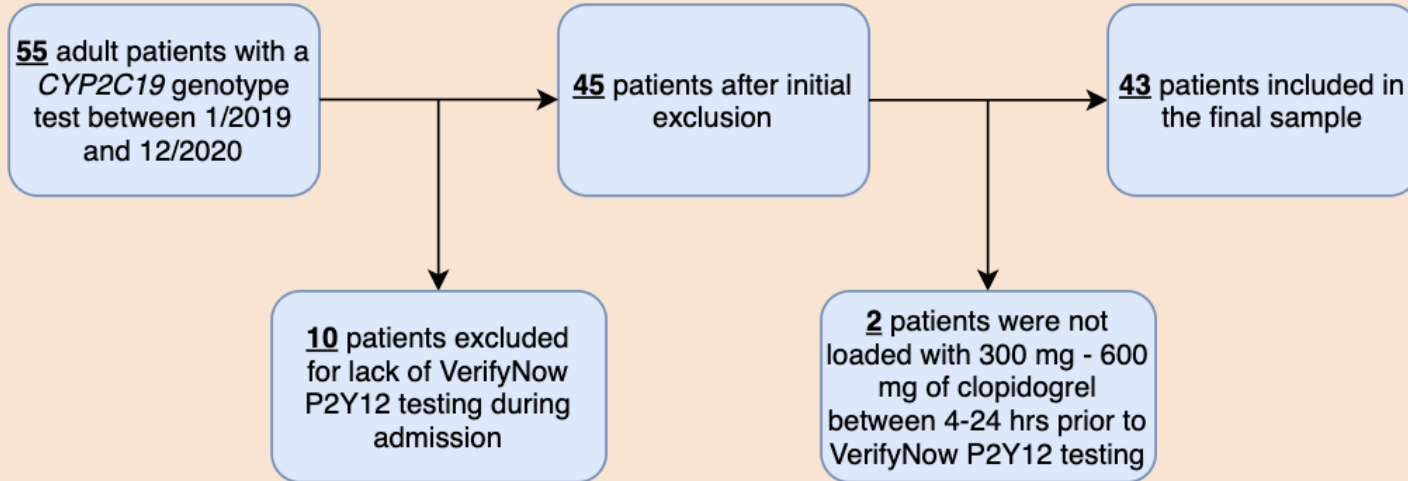
- Polymorphisms exist for the *CYP2C19* gene
 - Some are loss-of-function (LoF), some are gain-of-function
 - LoF alleles are associated with lowered clopidogrel metabolism
- VerifyNow P2Y12 assay measures platelet reactivity units (PRUs) in patients loaded with clopidogrel
 - High PRU (>180) suggest poor platelet inhibition from clopidogrel



Study Objectives

- 1) What are the sensitivity and specificity of a high post-load PRU for *CYP2C19* LoF genotype?
- 2) Are there comorbidities, concurrent medications, or racial/ethnic variables that may influence the association between high post-load PRU and *CYP2C19* LoF genotype?

Methods



Results

Table 1: Baseline Characteristics of Study Population, Stratified by Loss of Function Mutation in *CYP2C19*

	No LoF allele (n = 23)	At least one LoF allele (n = 32)	p-value
Age (mean [SD])	65.61 [11.42]	69.29 [11.14]	0.240
Male (%)	14 (60.9)	19 (61.3)	1.000
Race (%)			0.779
White	7 (30.4)	11 (35.5)	
Black	2 (8.7)	0 (0.0)	
Asian	10 (43.5)	14 (45.2)	
Native Hawaiian/ Pacific Islander	4 (17.4)	6 (19.4)	
P2Y12 Reaction Units (mean [SD])	139.45 [92.50]	193.83 [73.05]	0.037
Evidence of P2Y12 Inhibition (%)	14 (70.0)	8 (34.8)	0.046

Results

Table 2: Medical Characteristics of Study Population, Stratified by Platelet Receptor Blockade as Measured by P2Y12 Reaction Units

	No Platelet Receptor Blockade (n = 21)	Platelet Receptor Blockade (n = 22)	p-value
Age (mean [SD])	69.62 [9.51]	66.59 [12.05]	0.367
Male (%)	15 (71.4)	11 (50.0)	0.261
Diabetes Mellitus (%)			0.738
No	7 (30.4)	11 (35.5)	
Yes	2 (8.7)	0 (0.0)	
Pre-diabetes	10 (43.5)	14 (45.2)	
No data	4 (17.4)	6 (19.4)	
Methamphetamine Use (%)	0 (0.0)	2 (9.1)	0.490
No Cocaine Use (%)	21 (100.0)	22 (100.0)	N/A

Results

Table 2: Medical Characteristics of Study Population, Stratified by Platelet Receptor Blockade as Measured by P2Y12 Reaction Units (cont.)

	No Platelet Receptor Blockade (n = 21)	Platelet Receptor Blockade (n = 22)	p-value
Smoking Status (%)			0.332
Not a Smoker	9 (42.9)	14 (63.6)	
Current Smoker	3 (14.3)	3 (13.6)	
Former Smoker	9 (42.9)	5 (22.7)	
Excessive Alcohol Use (%)			0.135
No Excessive Use	20 (95.2)	19 (86.4)	
Binge Drinker	1 (4.8)	0 (0.0)	
Heavy Drinker	0 (0.0)	3 (13.6)	
Presence of LoF allele (%)	15 (71.4)	8 (36.4)	0.046



Results

Sensitivity and Specificity of VerifyNow P2Y12 Platelet Receptor Blockade (PRU < 180) for LoF Genotype

	Presence of LoF Allele	No LoF allele detected	Test Characteristic
No platelet inhibition (PRU > 180)	15	6	Sensitivity 65%
Platelet inhibition (PRU < 180)	8	14	Specificity 70%



Results

Sensitivity and Specificity of VerifyNow P2Y12 Platelet Receptor Blockade (PRU < 170) for LoF Genotype

	Presence of LoF Allele	No LoF allele detected	Test Characteristic
No platelet inhibition (PRU > 170)	17	6	Sensitivity 74%
Platelet inhibition (PRU < 170)	6	14	Specificity 70%



Results

Sensitivity and Specificity of VerifyNow P2Y12 Platelet Receptor Blockade (PRU < 150) for LoF Genotype

	Presence of LoF Allele	No LoF allele detected	Test Characteristic
No platelet inhibition (PRU > 150)	18	6	Sensitivity 78%
Platelet inhibition (PRU < 150)	5	14	Specificity 70%



Discussion

- In our study population, lack of platelet inhibition as defined by PRU > 180 has a sensitivity of 65% and specificity of 70% for *CYP2C19* LoF mutation
- Lowering the threshold for interpreting lack of platelet inhibition to PRU > 170 and 150 in our cohort increases the sensitivity to 74% and 78%, respectively, without affecting the specificity
- There was no significant difference in the race/ethnicity distributions between patients with and without LoF mutations
 - Previous studies have shown higher prevalence of LoF mutations in Asian populations, which may influence pre-test probability in clinical reasoning for some practices
 - In our study population, findings suggest that race/ethnicity are not reliable indicators of *CYP2C19* allele frequency
- Common medical comorbidities among patients undergoing testing do not associate with testing results



Study Strengths and Limitations

Strengths:

- Unique clinical protocol generated data to study test characteristics of the P2Y12 assay as a proxy for *CYP2C19* LoF genotype
- Patient population allowed inclusion of genetic information relevant to individuals of rarely studied ancestries

Limitations:

- Enriched sample precludes generalizability of findings
- Power limited by sample size



Conclusion

- 1) Among 43 patients who underwent *CYP2C19* genotyping at The Queen's Medical Center in 2019-2020, the VerifyNow P2Y12 assay was not a reliable proxy for *CYP2C19* loss of function mutation.
- 2) Larger studies are required to confirm test performance in a general patient population
- 3) Further studies will address the relationship between P2Y12 Reaction Units and clinical outcomes



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Citations

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