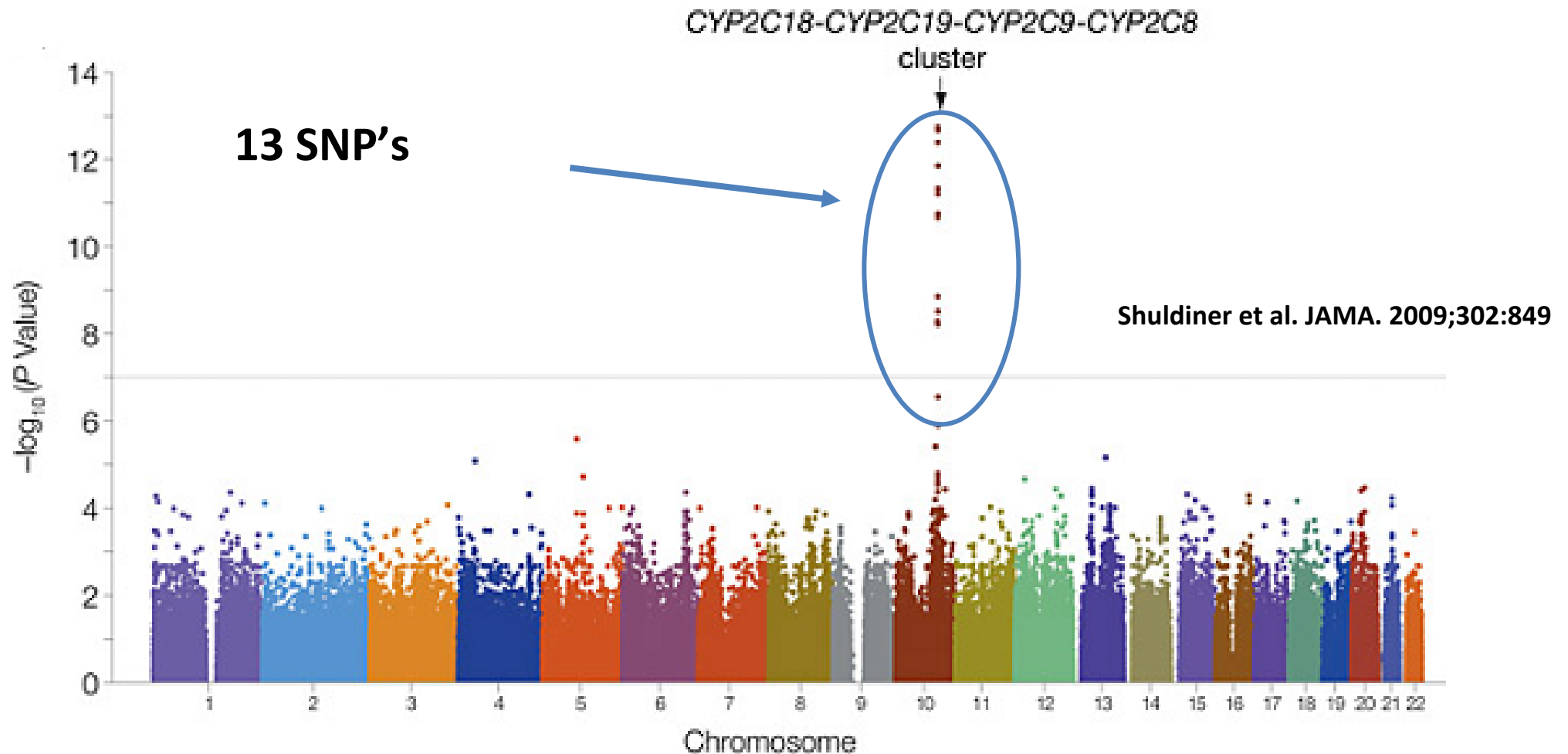


***Genetic Risk of
Clopidogrel Response Variability:
Cilostazol as a possible solution***

***Kyung Woo Park, MD, PhD
Cardiovascular Center,
Seoul National University Hospital***



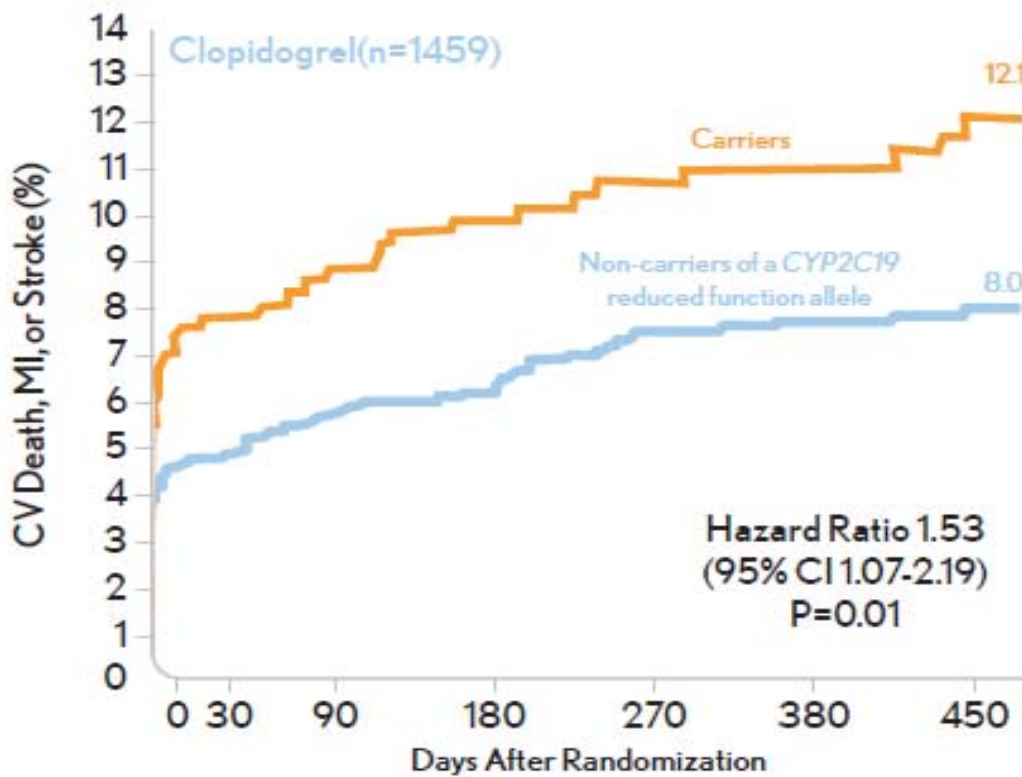
Decreased Response to Clopidogrel: GWAS Data



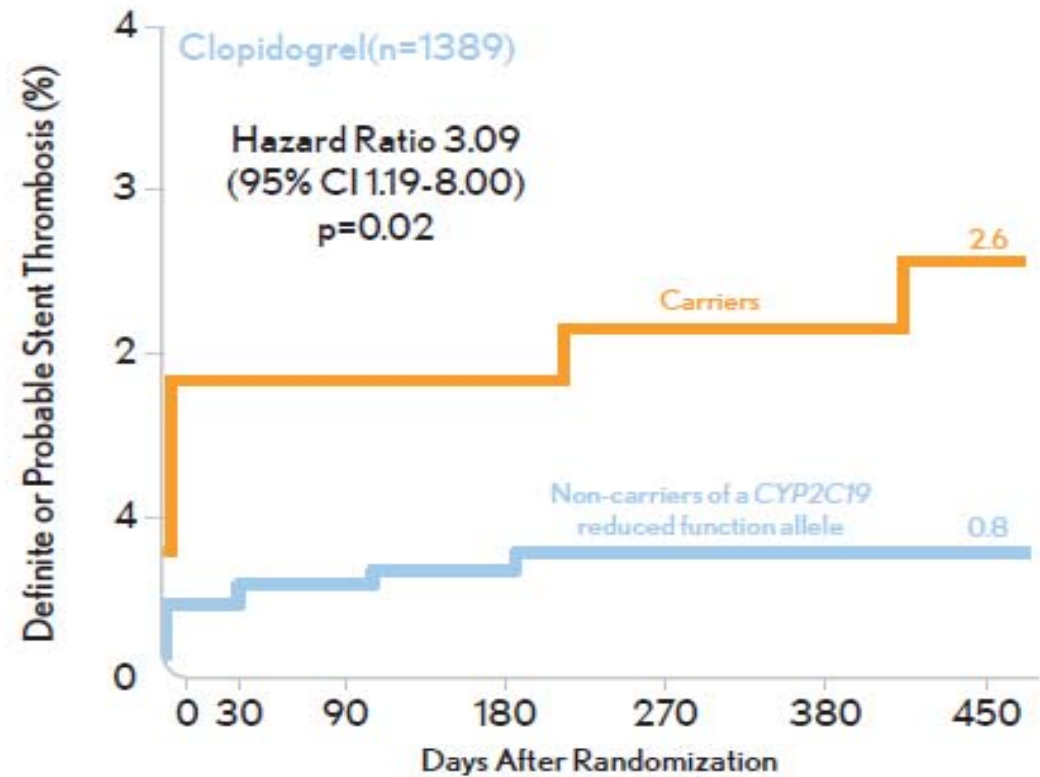
- 1) Clopidogrel response was highly heritable.
- 2) Cluster of 13 SNP's strongly associated with clopidogrel response ($p < 10^{-7}$). (locus on 10q24)
- 3) CYP2C19*2 accounted for most or all of the 10q24 association signal (~12% of response variability)
- 4) Majority of variation in clopidogrel response remains unexplained.

CYP2C19 LOF allele as a genetic risk of Ischemic Events

Pre-specified subgroup analyses of TRITON TIMI 38



Number at Risk:	0	30	90	180	270	380	450
Carrier	395	364	360	348	306	270	181
Non-Carrier	1064	1009	999	980	870	755	542



Number at Risk:	0	30	90	180	270	380	450
Carrier	375	368	366	359	316	279	186
Non-Carrier	1014	1004	1001	989	885	765	547

Ethnicity as a risk of HOPR?

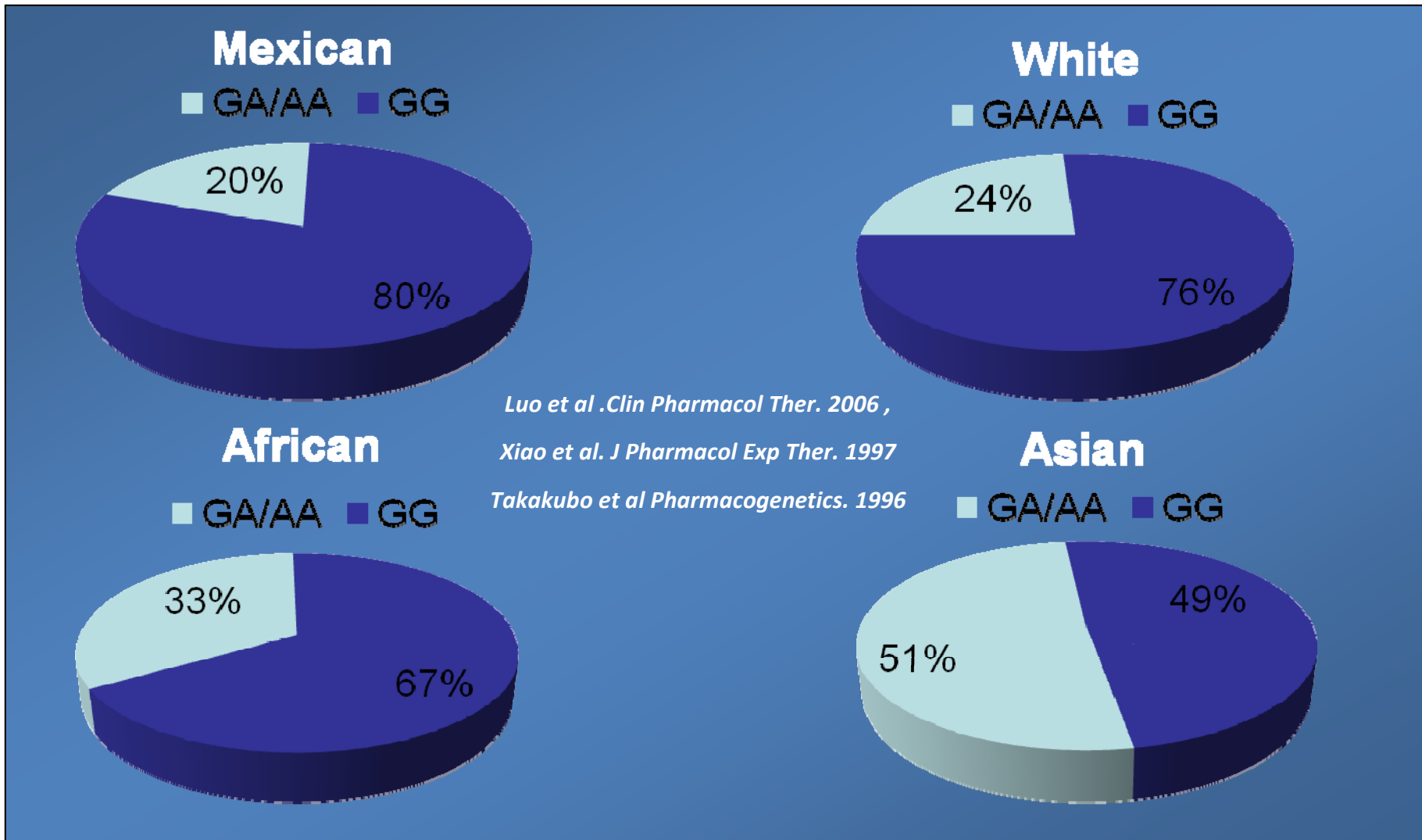
Characteristic	Mean Residual Platelet Reactivity (PRU)		P Value
	Characteristic present	Characteristic absent	
Age > 75 yrs	214 ± 77	201 ± 79	0.161
Men	200 ± 77	220 ± 82	0.041
Non-Caucasian ethnicity	229 ± 79	202 ± 78	0.047
Diabetes mellitus	220 ± 73	196 ± 80	0.005
⋮	⋮	⋮	⋮

Price MJ et al, Circulation 2009

- **Non-Caucasian ethnicity :**
 1. has higher residual platelet activity
 2. an independent predictor of high on-treatment plt reactivity
(OR: 3.05, 95% CI: 1.49 to 6.28, p=0.002)

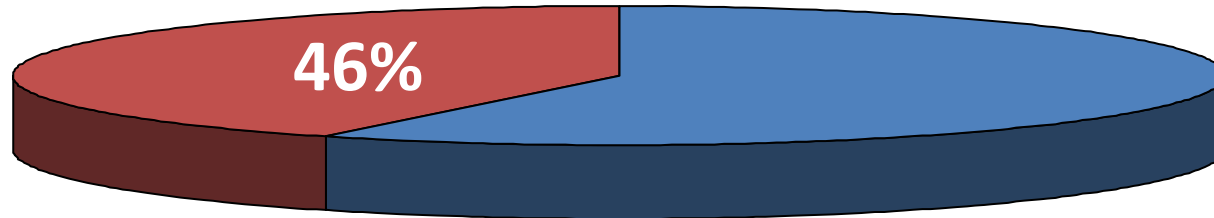


Different CYP2C19 *2 Allele Frequency



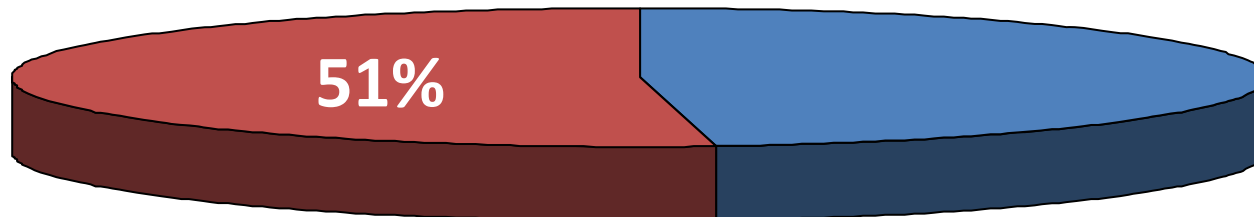
Different CYP2C19 LOF Frequency ***: according to Asian Ethnicity***

Japanese Population

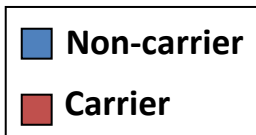


Sawada T et al. Circulation J 2011.

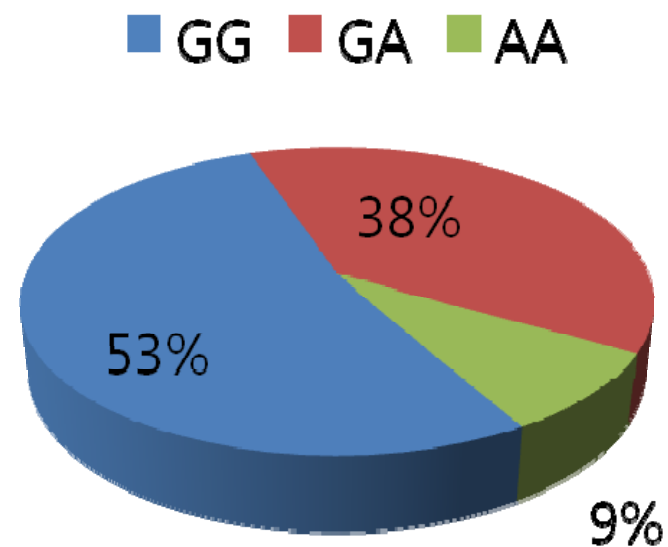
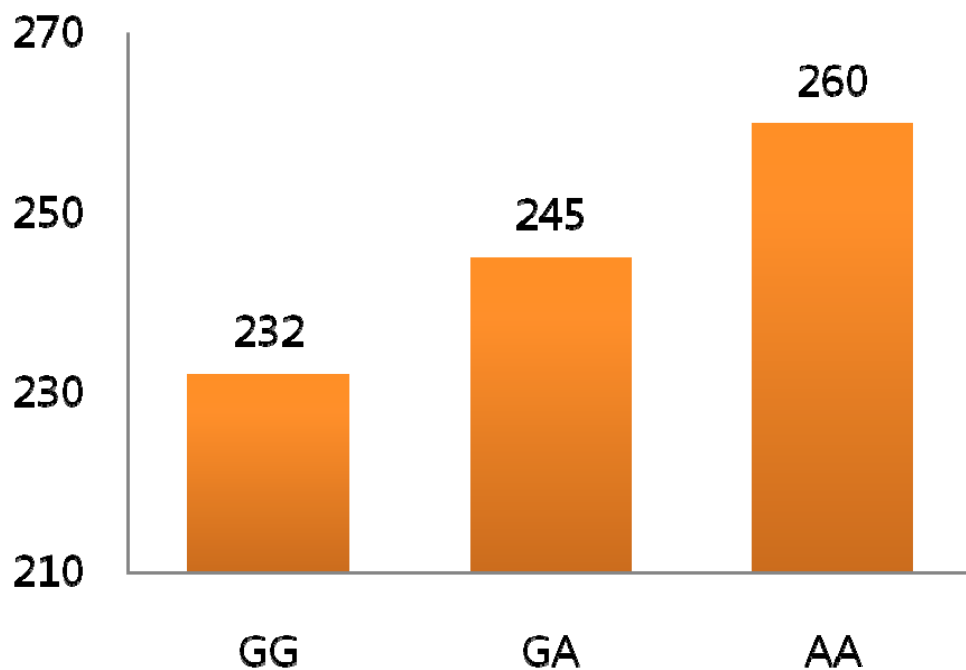
Chinese Population



Zhou Q et al. Pharmacogenomics J 2009

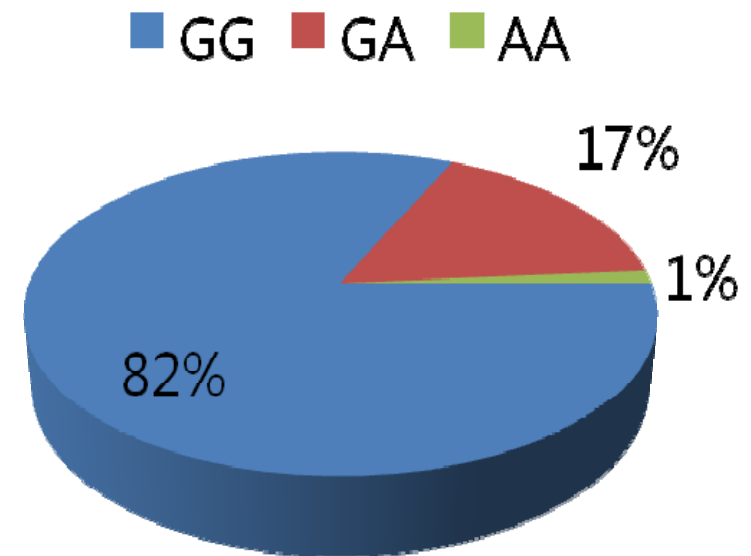
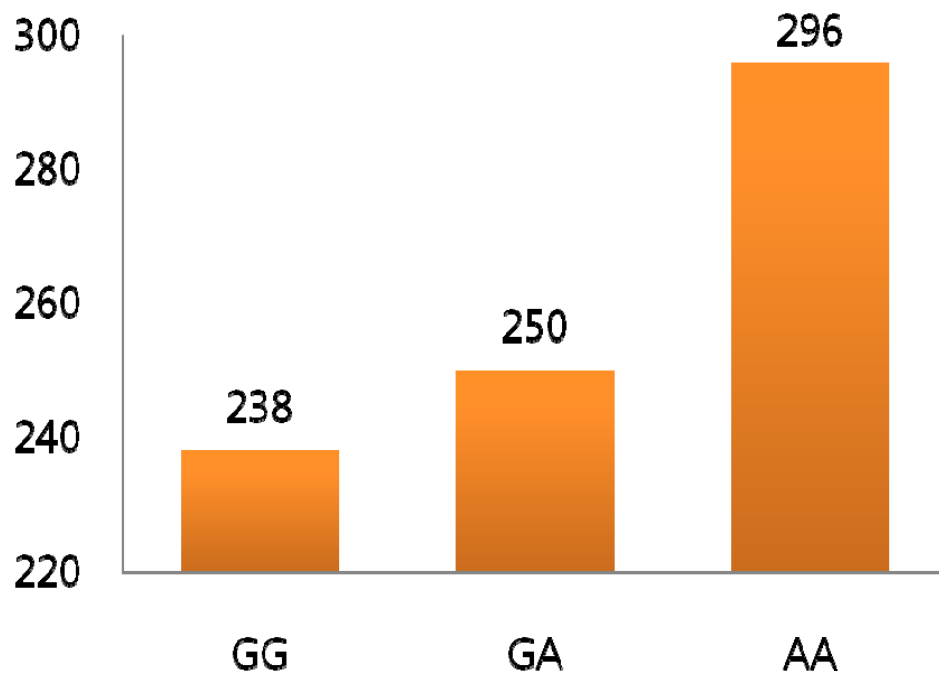


CYP2C19*2



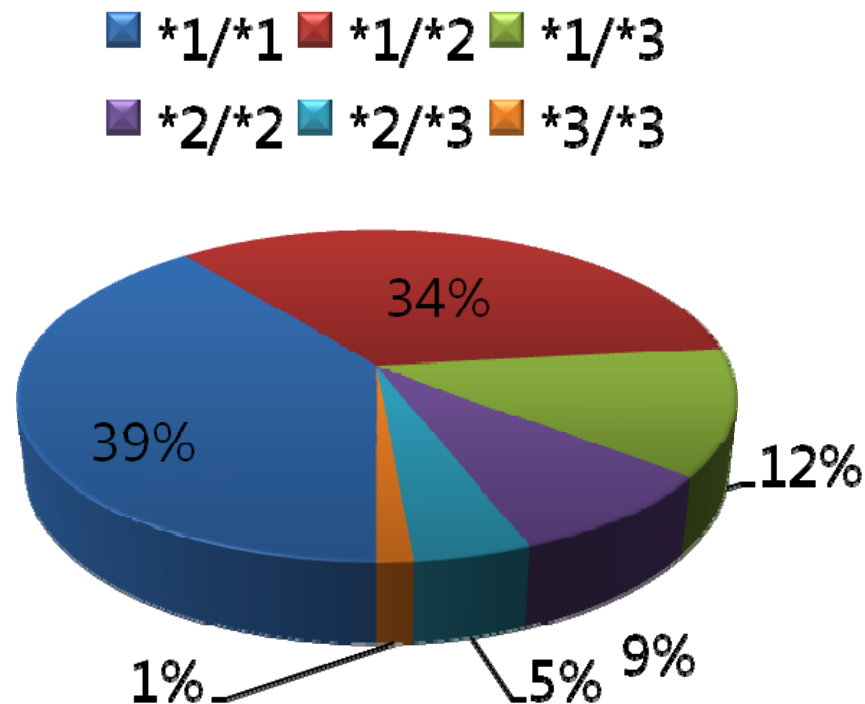
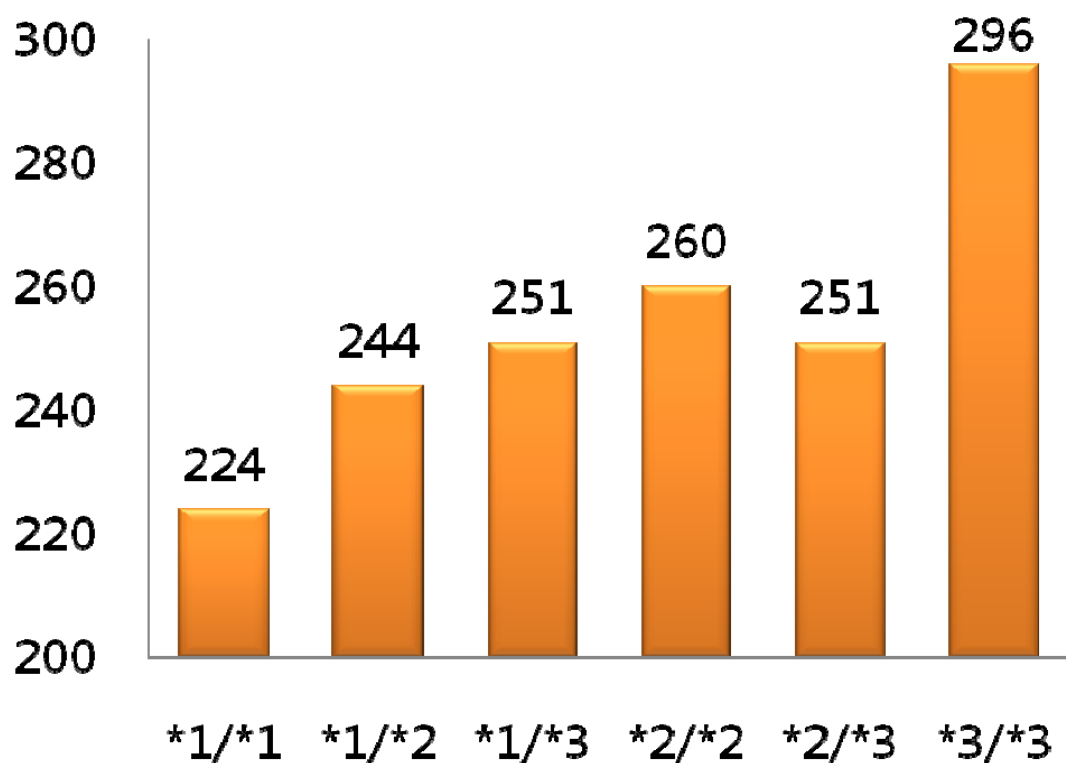
	GG	GA	AA	All	p-value
Freq	848	613	139	1600	
Expected	833.0	642.9	124.0		
PRU	232.5±82.6	245.7±79.4	260.3±72.0	240.0±81.0	P<0.001
Major G 0.72 (wild), Minor A 0.28 (mutant), $\chi^2 = 5.463$, p = 0.062)					

CYP2C19*3



	GG	GA	AA	All	p-value
Freq	1308	267	22	1597	
Expected	1301.1	280.7	15.1		
PRU	236.8.5±81.7	250.8±76.9	296.2±58.4	240.0±81.0	P<0.001
Major G 0.90 (wild), Minor A 0.10 (mutant), $\chi^2 = 3.813$, p = 0.050)					

CYP2C19 *2 & *3 combined : CROSS VERIFY cohort



	*1/*1	*1/*2	*1/*3	*2/*2	*2/*3	*3/*3	p-value
Freq	625	540	195	139	71	22	
PRU	224.7±82.3	244.8±81.1	251.4±80.7	260.3±72.0	251.0±64.9	296.2±58.4	<0.001

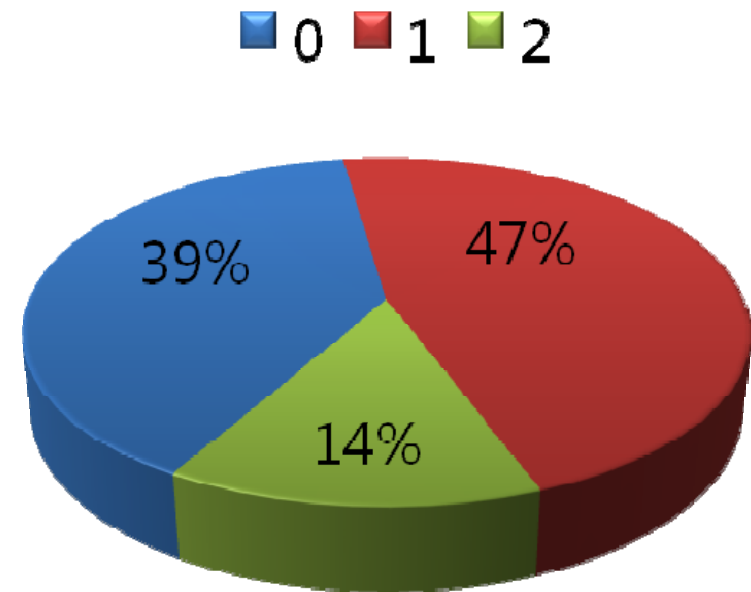
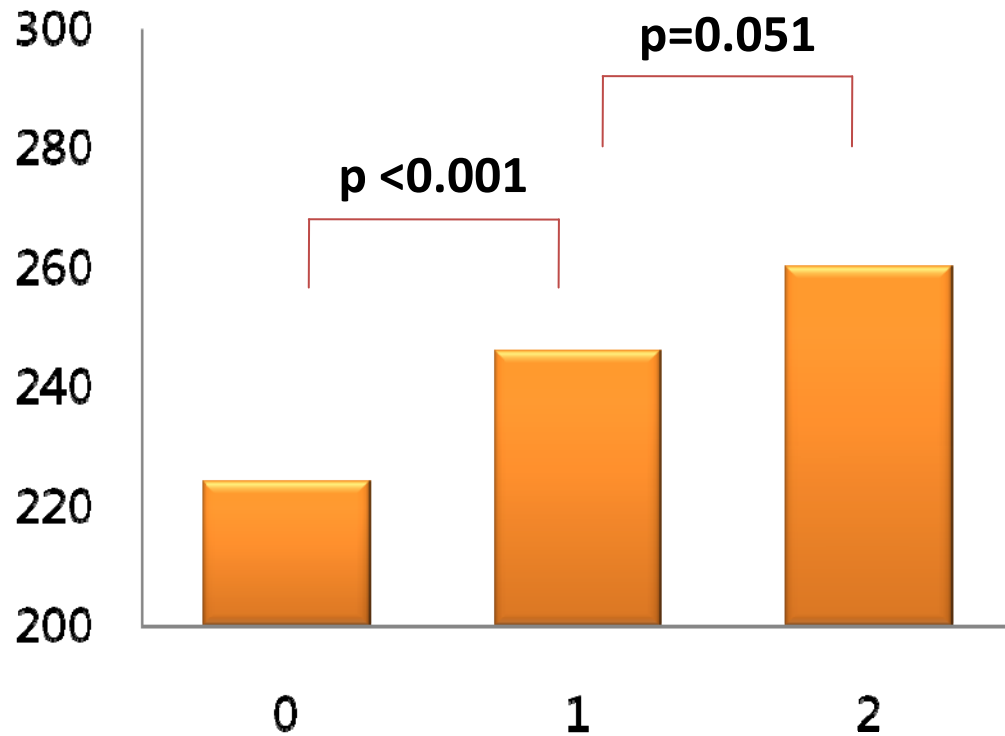
Unpublished data from the CROSS VERIFY cohort



Seoul National University Hospital Cardiovascular Center

Number of 'loss of fxn' alleles & PRU:

CROSS VERIFY cohort



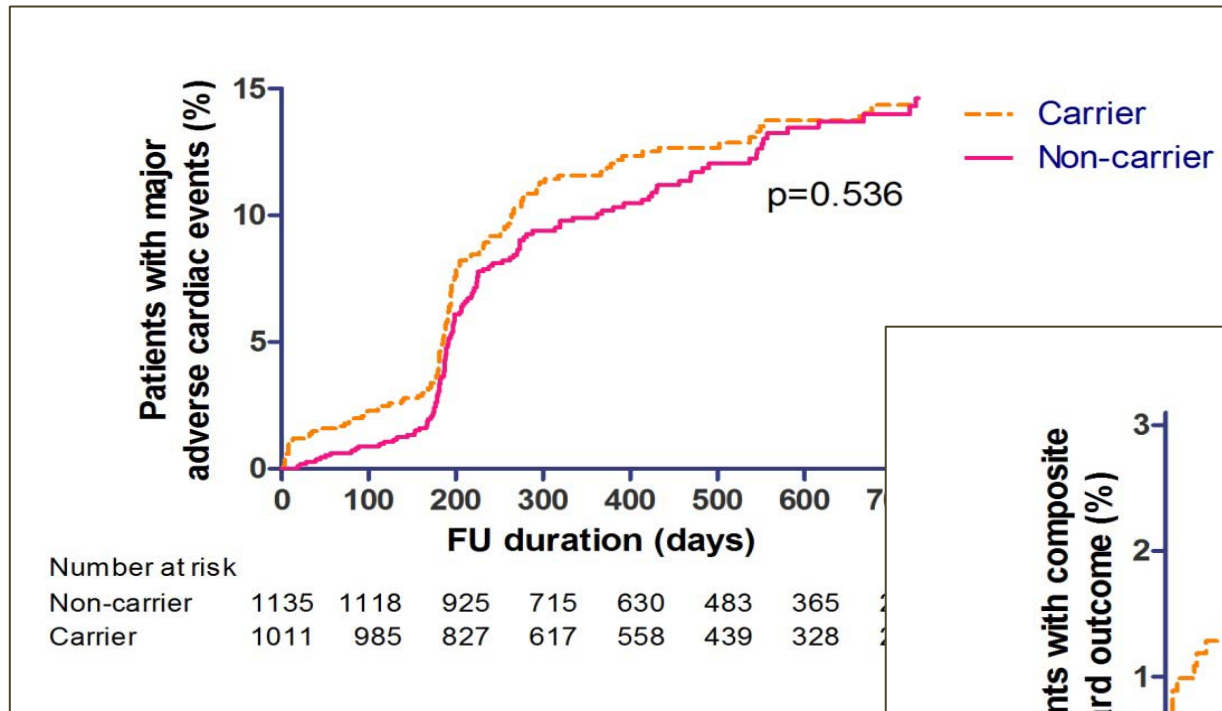
	Zero (*1/*1)	One (*1/*2, *1/*3)	Two (*2/*2, *2/*3, *3/*3)	p-value
Freq	625	735	232	
PRU	224.7±82.4	246.5±81.0	260.9±69.5	<0.001

Unpublished data from the CROSS VERIFY cohort

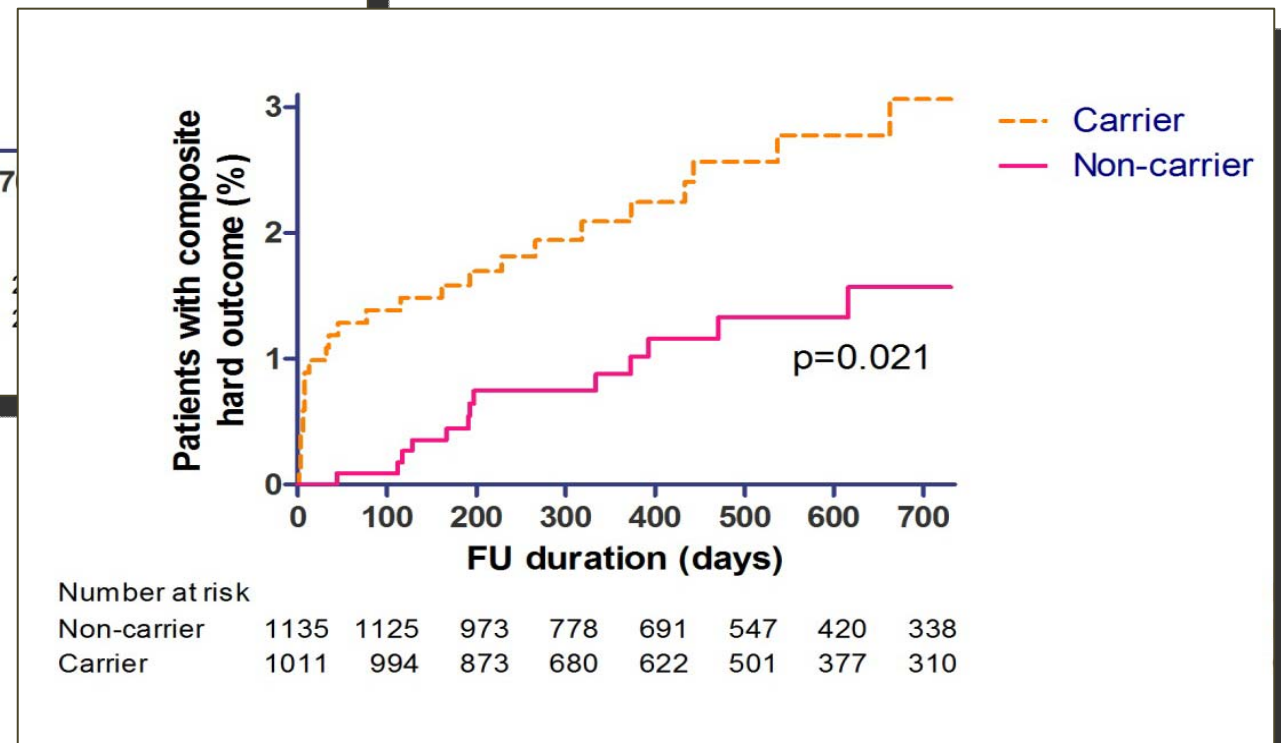


Seoul National University Hospital Cardiovascular Center

Relationship btw genetic risk and outcome in Koreans: Associated with only hard outcomes (from the SKY registry)



**Hard outcome
(CD, MI, and ST)**



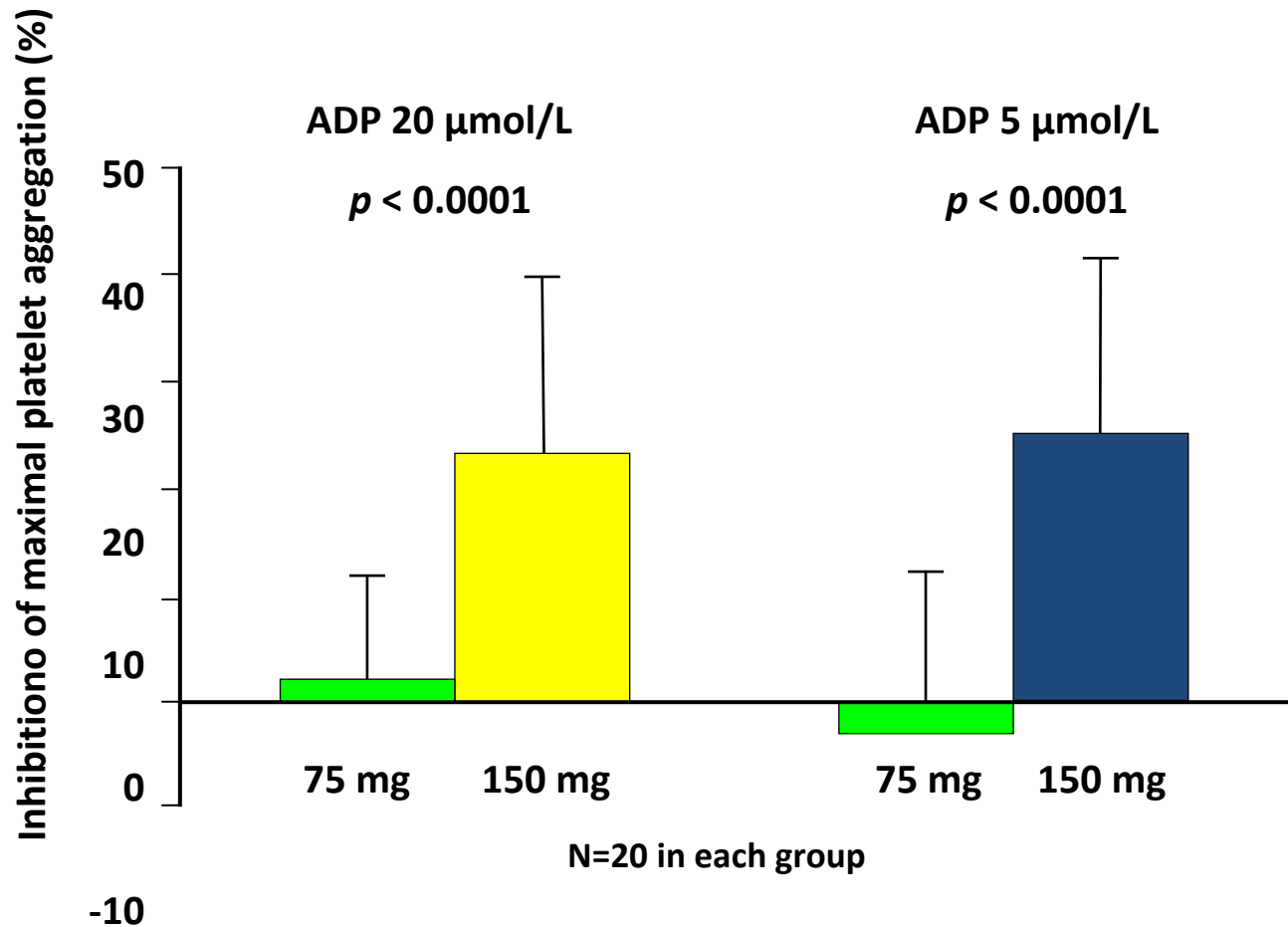
**All MACE
(including revascularization)**

Doubling the dose of plavix??



Increase dose of clopidogrel: *OPTIMUS*

Randomized comparison of a high clopidogrel maintenance dose in patients with diabetes mellitus and coronary artery disease





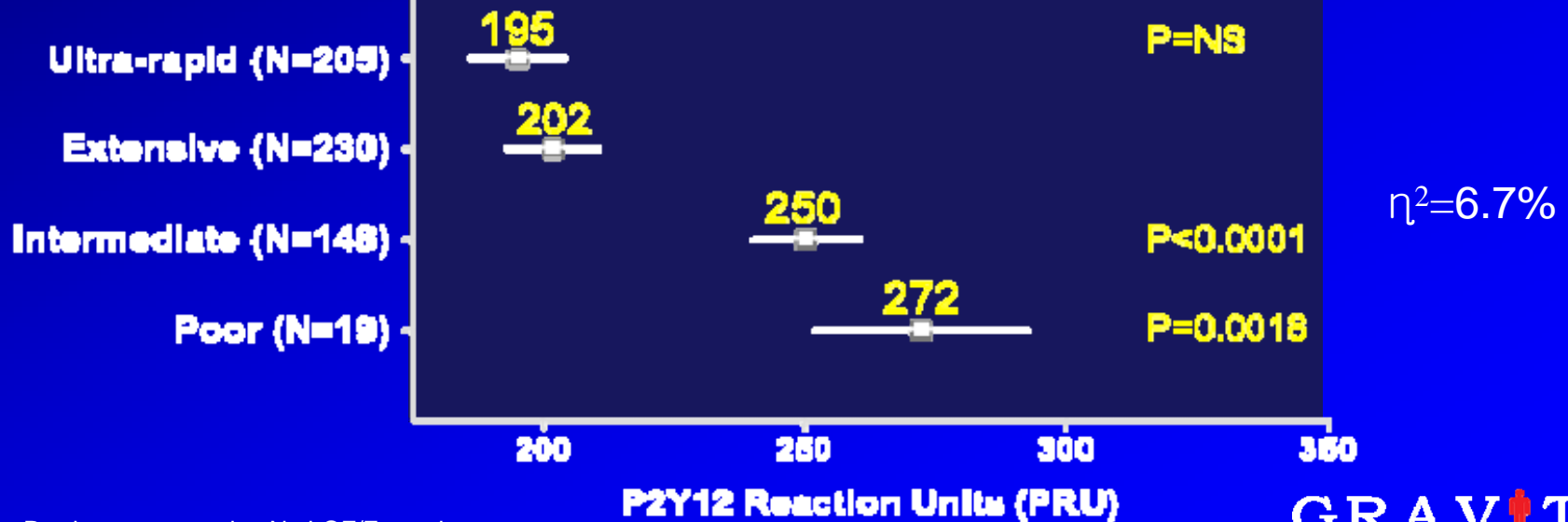
GENOTYPE INFORMATION & FUNCTIONAL TESTING

Platelet Reactivity on Clopidogrel Post-PCI Is Associated With *CYP2C19* Genotype & Phenotype

CYP2C19 genotype



Metabolic phenotype



P2Y12 Reaction Units (PRU)

Least squared means. P values compared to No LOF/Extensive.

η^2 : portion of variance explained by the genotype or phenotype in the multivariate generalized linear model

GRAVITAS

Courtesy of M. Price, Price M et al. ACC 2011



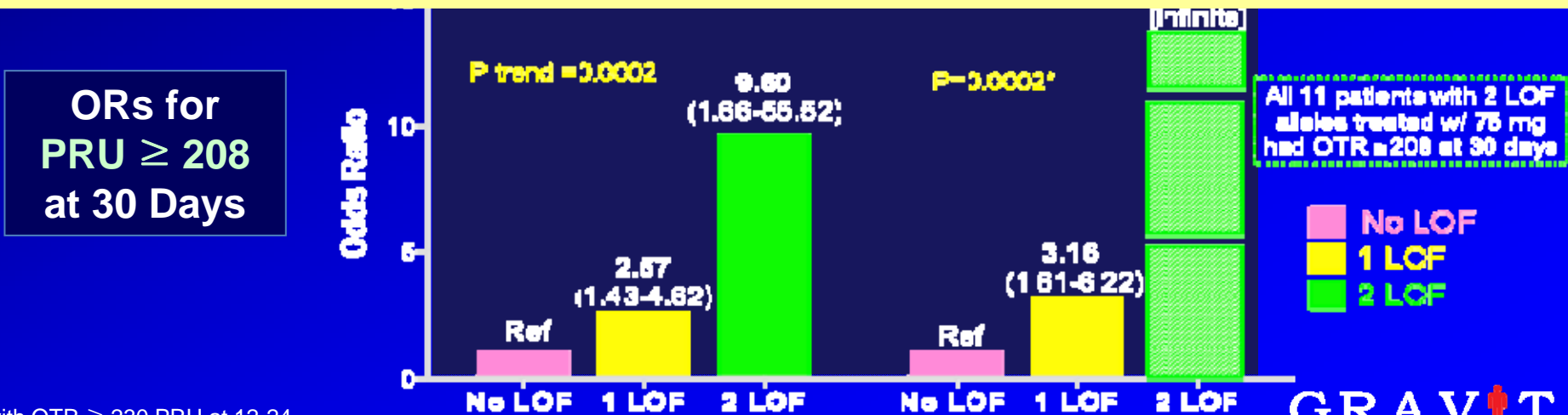
GENOTYPE INFORMATION & FUNCTIONAL TESTING

CYP2C19 LOF Allele Is Associated With Higher Risk of Persistently High OTR at 30 Days Regardless of Dose



ORs for PRU ≥ 230 at 30 Days

Therefore, doubling the dose of Clopidogrel cannot be a solution to overcome the genetic risk of clopidogrel response variability.



ORs for PRU ≥ 208 at 30 Days

Patients with OTR ≥ 230 PRU at 12-24 hours after PCI. Adjusted ORs.



Courtesy of M. Price. Price M et al. ACC 2011

Adding cilostazol as a third agent??



Multicenter Randomized Trial Evaluating the Efficacy of Cilostazol on Ischemic Vascular Complications After Drug-Eluting Stent Implantation for Coronary Heart Disease

Results of the CILON-T (Influence of CILostazol-based triple antiplatelet therapy ON ischemic complication after drug-eluting stenT implantation) Trial

Jung-Won Suh, MD,*† Seung-Pyo Lee, MD,* Kyung-Woo Park, MD,* Hae-Young Lee, MD,* Hyun-Jae Kang, MD,* Bon-Kwon Koo, MD,* Young-Seok Cho, MD,† Tae-Jin Youn, MD,† In-Ho Chae, MD,† Dong-Ju Choi, MD,† Seung-Woon Rha, MD,‡ Jang-Ho Bae, MD,§ Taek-Geun Kwon, MD,§ Jang-Whan Bae, MD,|| Myeong-Chan Cho, MD,|| Hyo-Soo Kim, MD*
Seoul, Seongnam, Daejeon, and Cheongju, Korea

Objectives

We aimed to test whether cilostazol has beneficial effects in the real-world patients treated with intracoronary drug-eluting stents (DES).

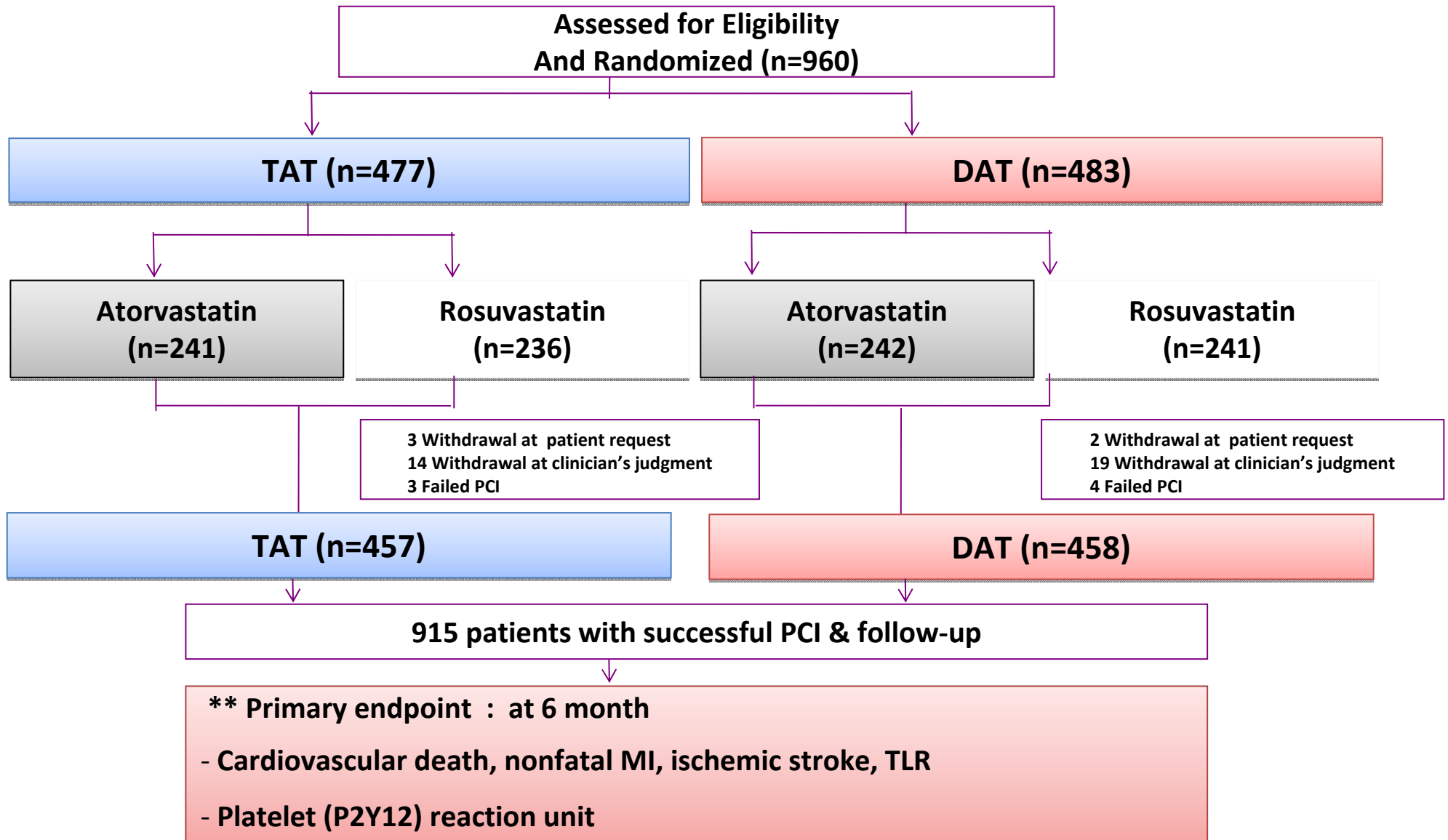
Background

The addition of cilostazol on the conventional dual antiplatelet therapy has been reported to reduce platelet reactivity and to improve clinical outcomes after percutaneous coronary intervention in previous studies.

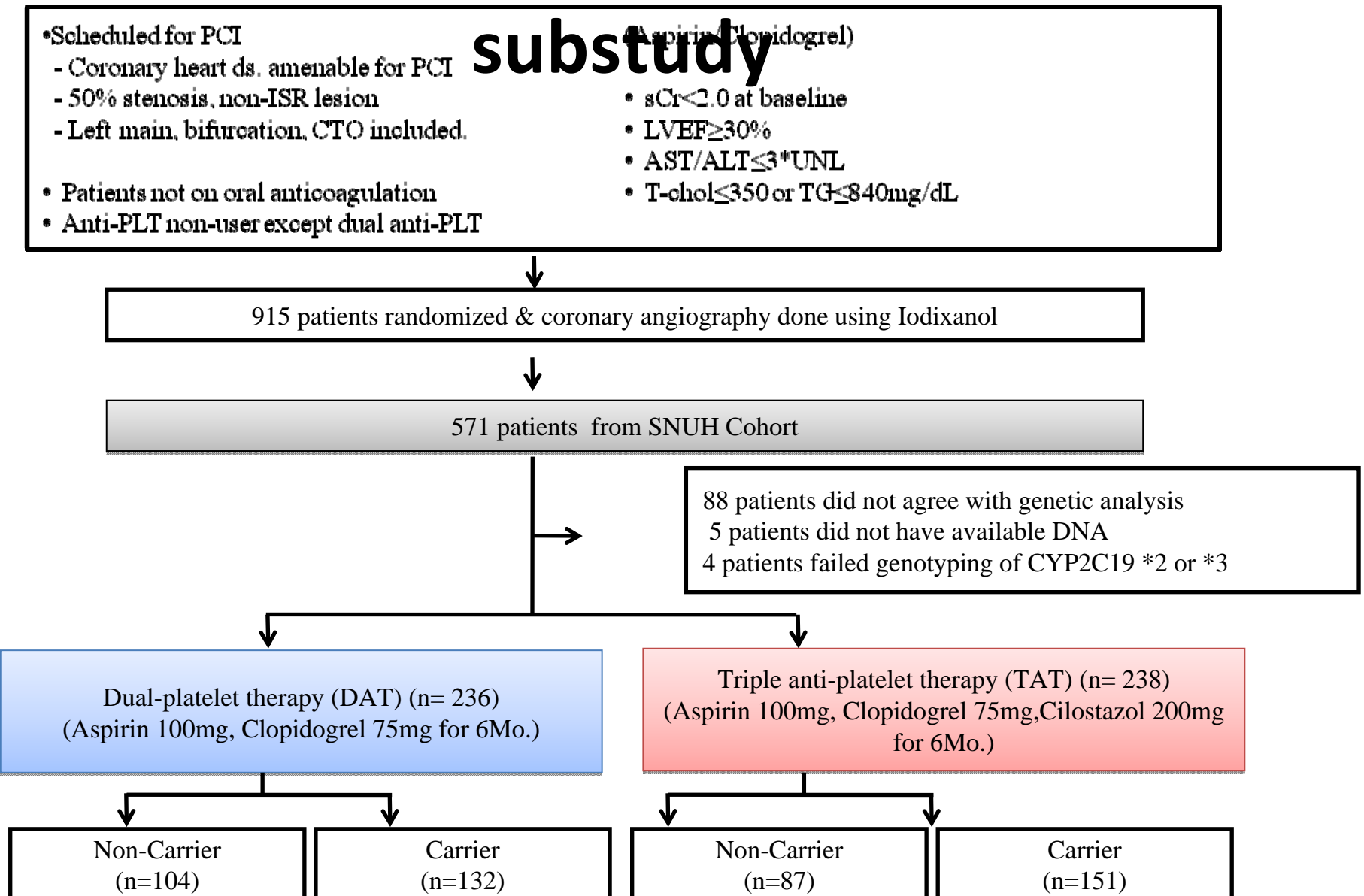
Methods

In a randomized multicenter trial, we enrolled 960 patients who received DES. They were randomized to receive either dual antiplatelet therapy (DAT) (aspirin and clopidogrel) or triple antiplatelet therapy (TAT) (aspirin, clopi-

CILON-T Trial Design



Add Cilostazol: CILON-T genetic



Genotyping & Analysis

- **Genotyping of 3 SNPs of CYP2C19**

- TaqMan™ Assay :

- CYP2C19*2 (P227P, rs4244285)
 - CYP2C19*3 (W212X, rs4986893)

- SNaPshot™ Multiplex Analysis

- CYP2C19*17(806C/T, rs12248560)

- **Statistical Analysis**

- Frequency

- Categorical : count (%)
 - Continuous : mean \pm SD

- Testing differences between groups

- Categorical : chi-square test or Fisher's exact test
 - Continuous : unpaired Student's test or 1-way ANOVA



Baseline Characteristics I

	DAT		TAT		p-value*
	Non-Carrier	Carrier	Non-Carrier	Carrier	
Demographic characteristics					
Age (yr)	63.25±8.34	62.83±8.30	62.82±9.14	64.11±9.02	0.601
Men	74 (71.2%)	101 (76.5%)	58 (65.9%)	97 (64.2%)	0.045
Body mass index (kg/m ²)	25.96±7.19	27.68±8.11	27.00±6.52	27.03±7.43	0.882
Smoker	22 (21.2%)	32 (24.2%)	16 (18.2%)	28 (18.5%)	0.251
Hypertension	73 (70.2%)	87 (65.9%)	59 (67.0%)	103 (68.2%)	0.917
Diabetes	25 (24.0%)	44 (33.3%)	30 (34.1%)	60 (39.7%)	0.054
Dyslipidemia	41 (39.4%)	51 (38.6%)	50 (56.8%)	68 (45.0%)	0.026
Chronic kidney disease	1 (1.0%)	1 (0.8%)	3 (3.4%)	3 (2.0%)	0.285
H/O PCI	14 (13.5%)	12 (9.1%)	8 (9.1%)	16 (10.6%)	0.695
H/O CABG	5 (4.8%)	5 (3.8%)	3 (3.4%)	4 (2.6%)	0.428
H/O Myocardial infarction	3 (2.9%)	8 (6.1%)	4 (4.5%)	9 (6.0%)	0.722
Congestive heart failure	0 (0%)	1 (0.8%)	0 (0%)	0 (0%)	0.494
Cerebrovascular accident	6 (5.8%)	7 (5.3%)	5 (5.7%)	7 (4.6%)	0.767
Peripheral artery disease	3 (2.9%)	3 (2.3%)	1 (1.1%)	3 (2.0%)	0.540

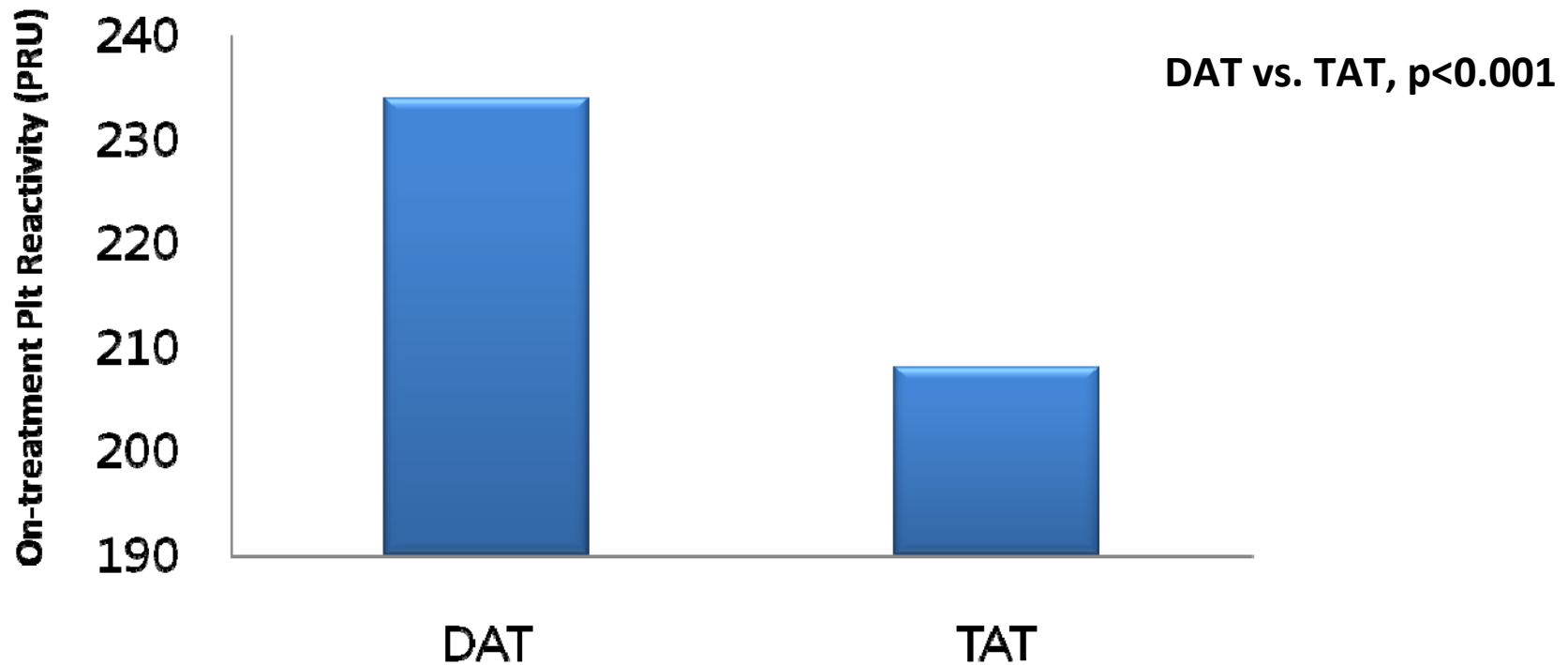


Baseline Characteristics II

	DAT		TAT		p-value*
	Non-Carrier	Carrier	Non-Carrier	Carrier	
Laboratory finding					
Hematocrit	39.9±4.42	40.04±6.15	40.20±4.67	39.87±4.84	0.829
Creatinine (mg/dL)	1.06±0.18	1.17±0.32	1.09±0.23	1.07±0.22	0.572
Cholesterol (mg/dL)	174.43±46.25	165.86±36.53	173.36±37.05	174.18±38.20	0.222
- Triglyceride	153.05±94.34	134.51±76.87	151.15±102.02	144.91±73.20	0.597
- HDL-Cholesterol	42.92±10.84	42.11±10.38	44.00±11.68	42.72±10.53	0.481
- LDL-Cholesterol	106.63±39.34	103.00±34.36	101.32±32.59	105.81±34.58	0.968
CRP (mg/L)	3.70±4.92	3.41±5.82	54.9±450.78	7.64±34.67	0.243
Concomitant Medication					
ACEi	25 (24.3%)	20 (15.4%)	16 (18.2%)	28 (18.7%)	0.860
ARB	14 (13.6%)	20 (15.4%)	11 (12.5%)	20 (13.3%)	0.498
Beta-blocker	41 (39.8%)	56 (43.1%)	39 (44.3%)	68 (45.3%)	0.487
Calcium channel blocker	25 (24.3%)	34 (26.2%)	26 (29.5%)	47 (31.3%)	0.231
Statin	36 (35%)	39 (30%)	35 (39.8%)	58 (38.6%)	0.199
Proton pump inhibitor	0 (0%)	1 (0.8%)	0 (0%)	3 (2%)	0.624



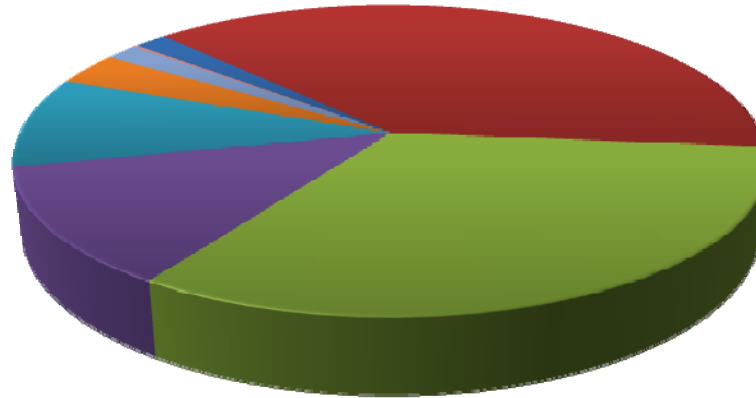
TAT has higher antiplatelet activity



PRU	n	mean PRU
All patients	474	221.71 ± 86.09
DAT	236 (49.8%)	234.15 ± 78.85
TAT	238 (50.2%)	208.86 ± 90.32



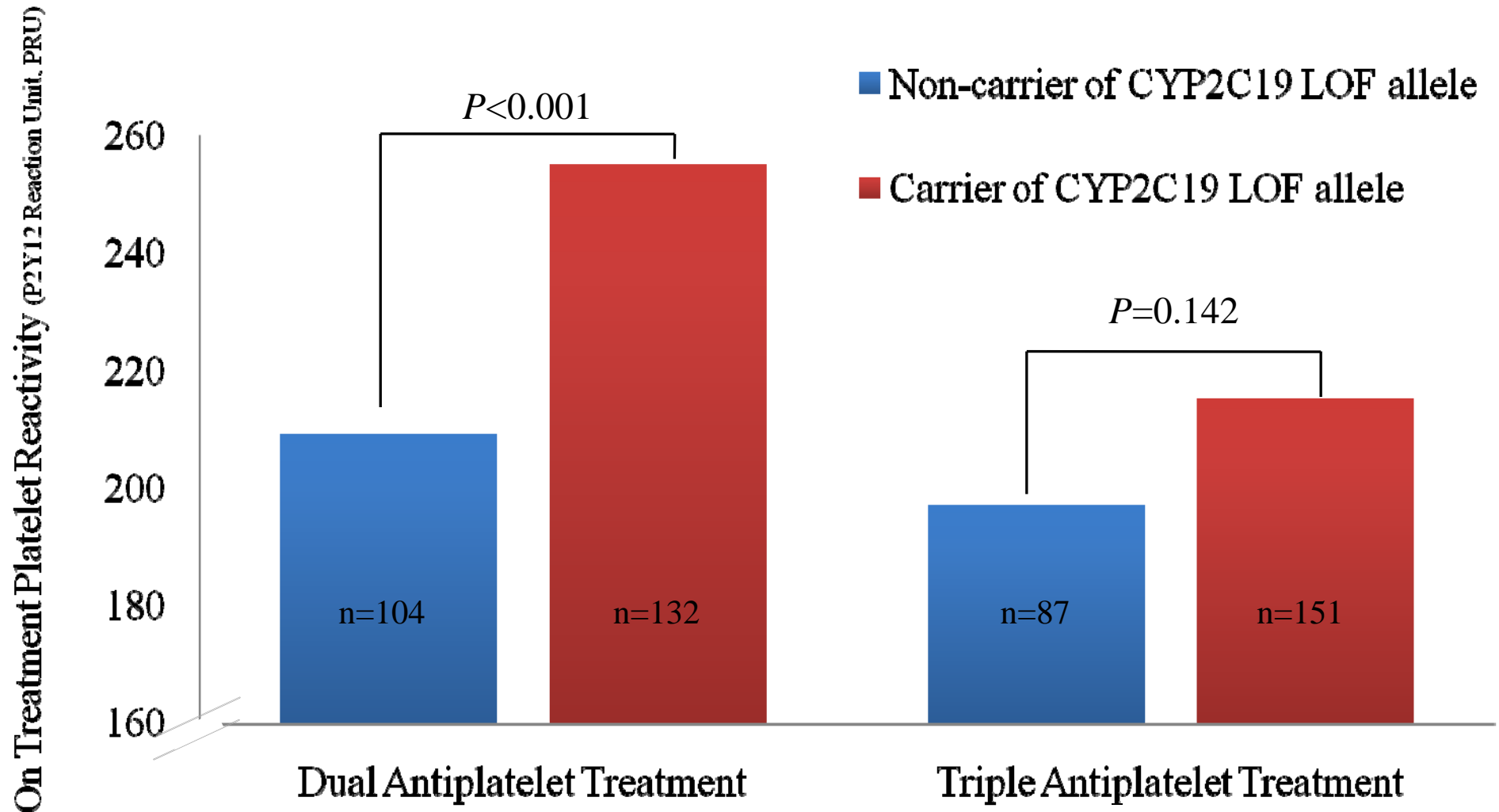
Genotype Distribution



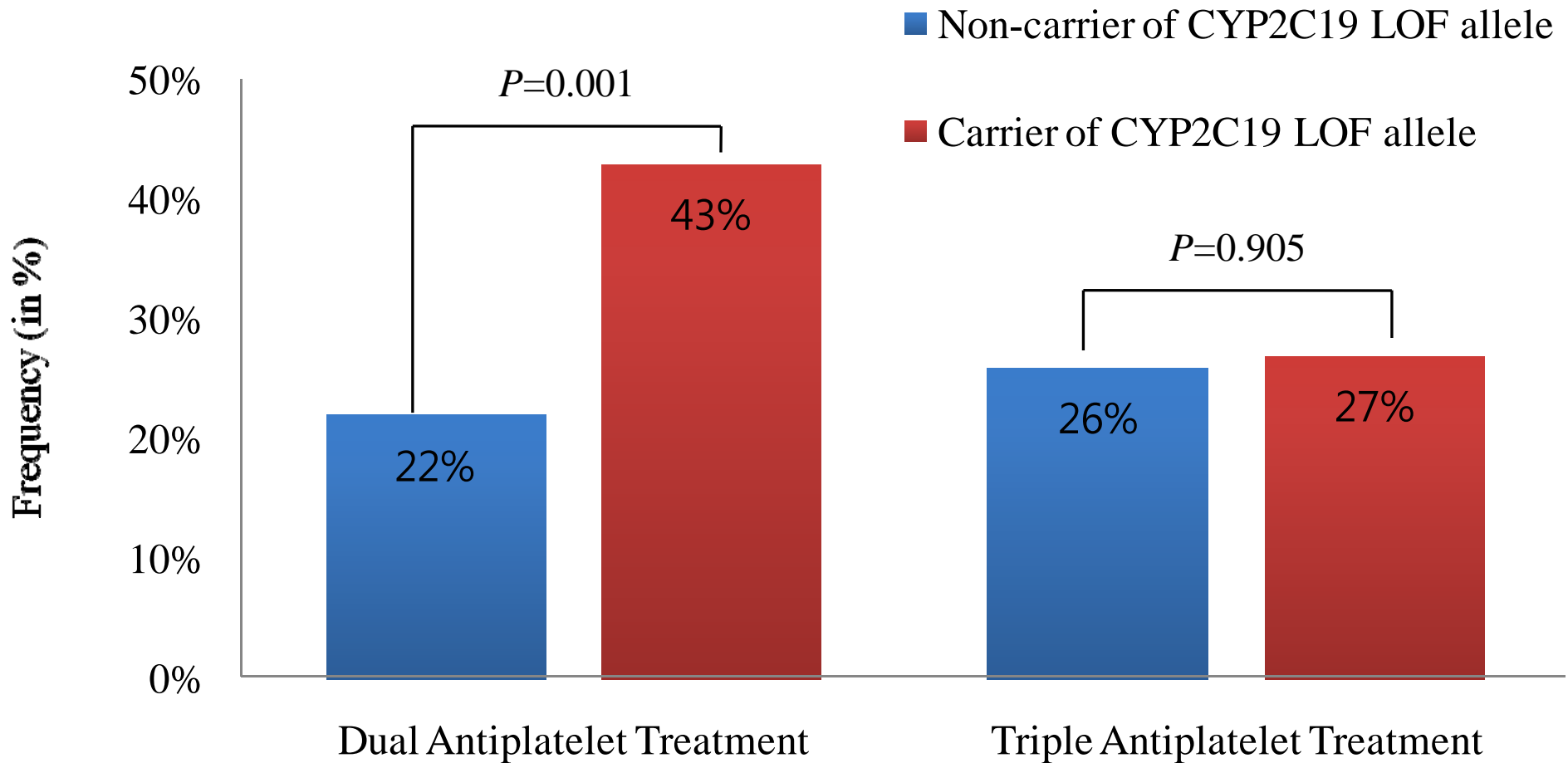
LOF allele (-)		LOF allele (+)	
*1/*17	8 (1.7%)	*1/*2	156 (32.9%)
*1/*1	183 (38.6)	*1/*3	58 (12.2%)
		*2/*2	44 (9.3%)
		*2/*3	16 (3.4%)
		*3/*3	8 (1.7%)
		*2/*17	1 (0.2%)
Total	191 (40.3%)		283 (59.7%)



CYP2C19 LOF associated with increased mean OPR in DAT, but not in TAT group

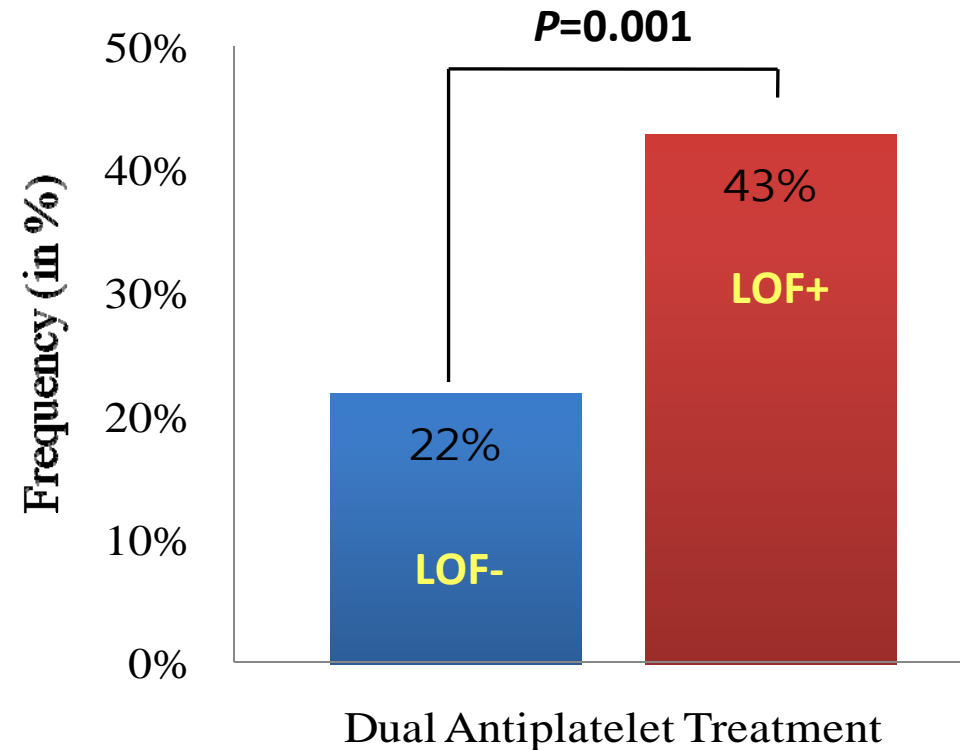
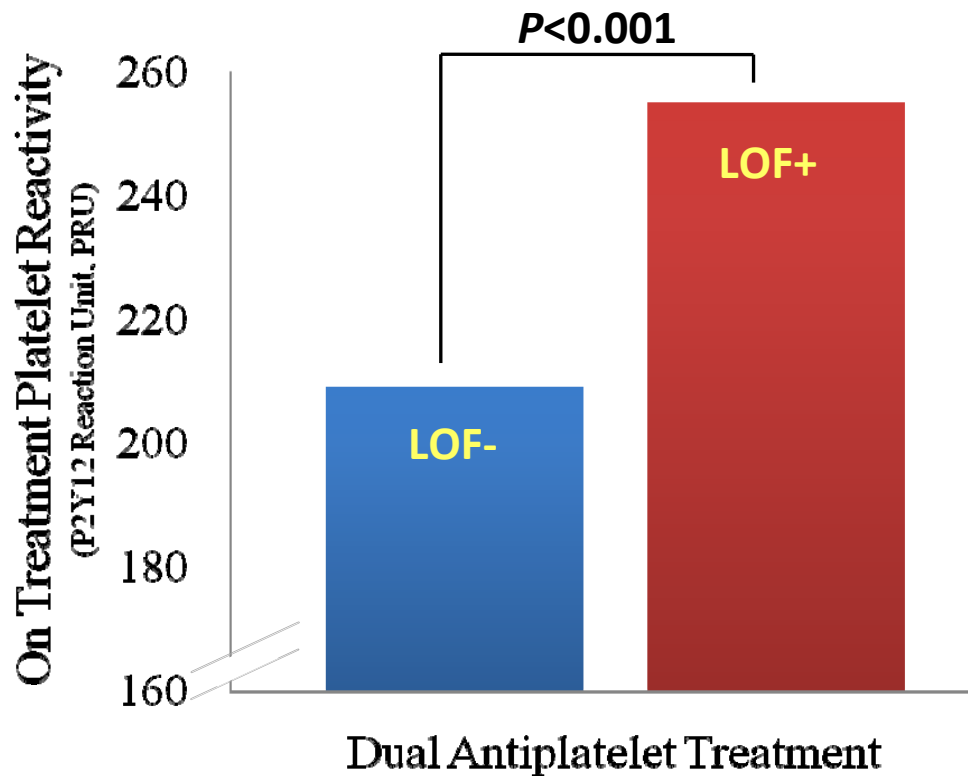


CYP2C19 LOF associated with HOPR in DAT, but not in TAT group

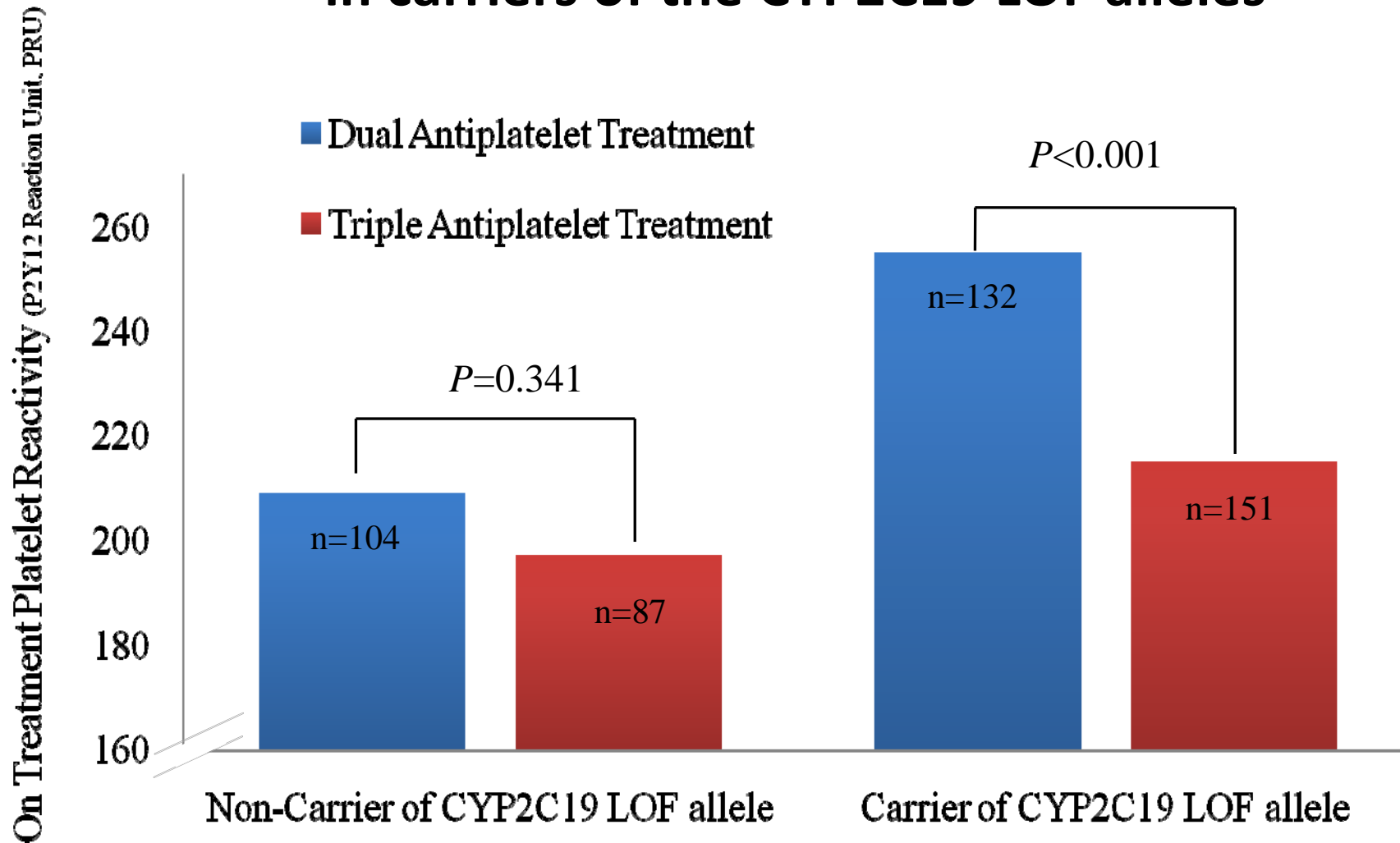


CYP2C19 LOF associated with HOPR in Koreans

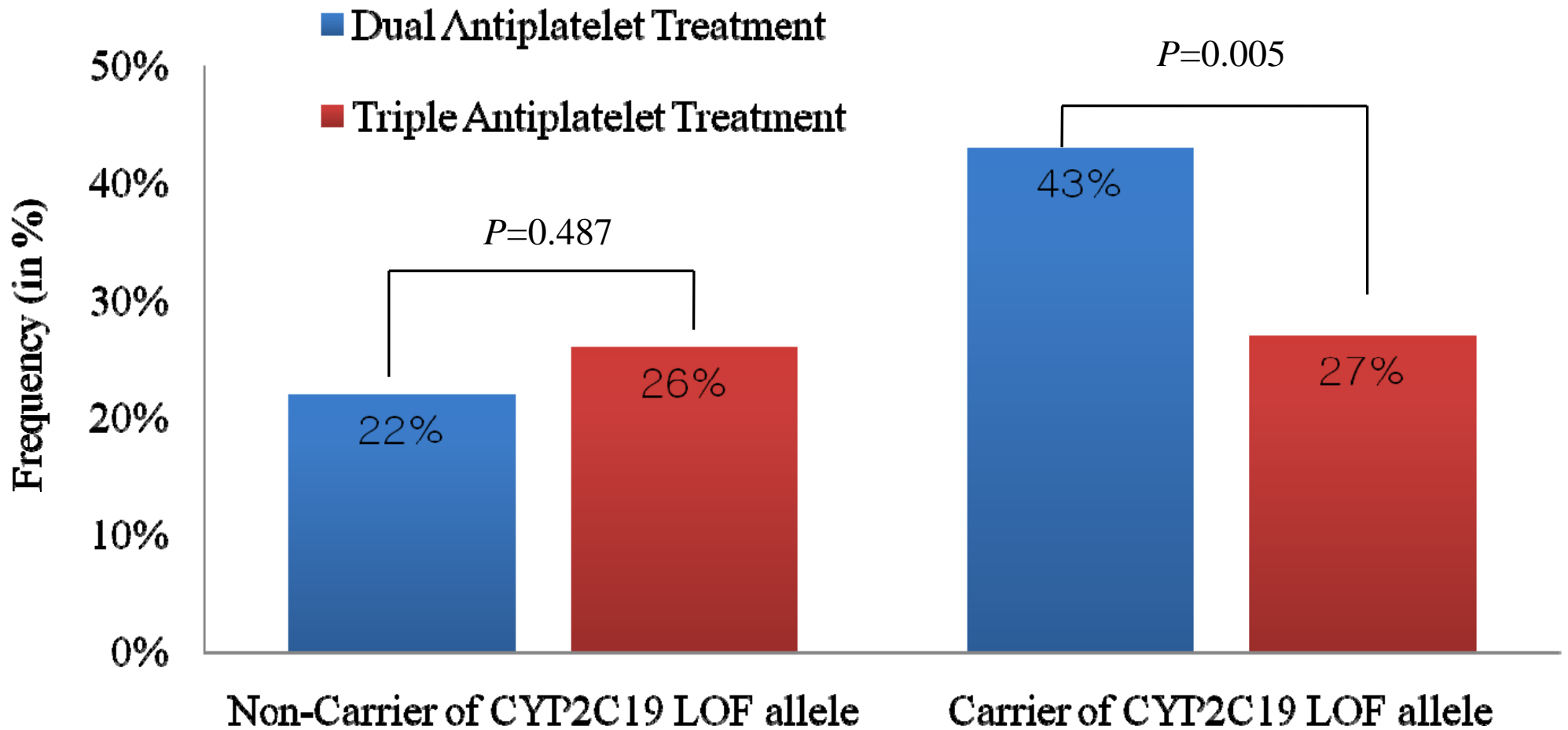
Genetic subanalysis of the CILON-T trial



TAT significantly reduces mean OPR in carriers of the CYP2C19 LOF alleles



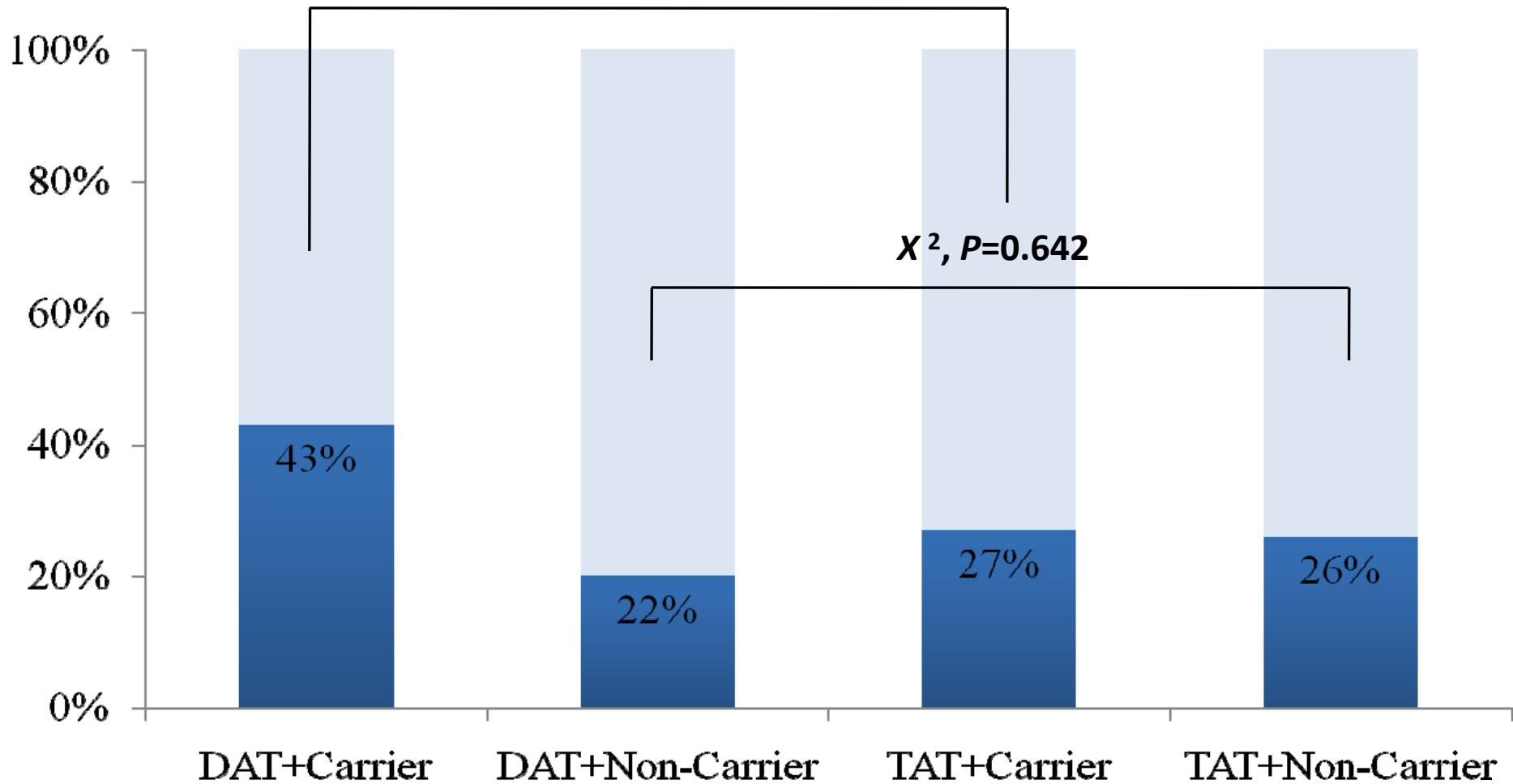
TAT significantly reduces the frequency of HOPR in carriers of the CYP2C19 LOF alleles



Frequency of HOPR

according to treatment and genotype
(using PRU>275 as definition of HOPR)

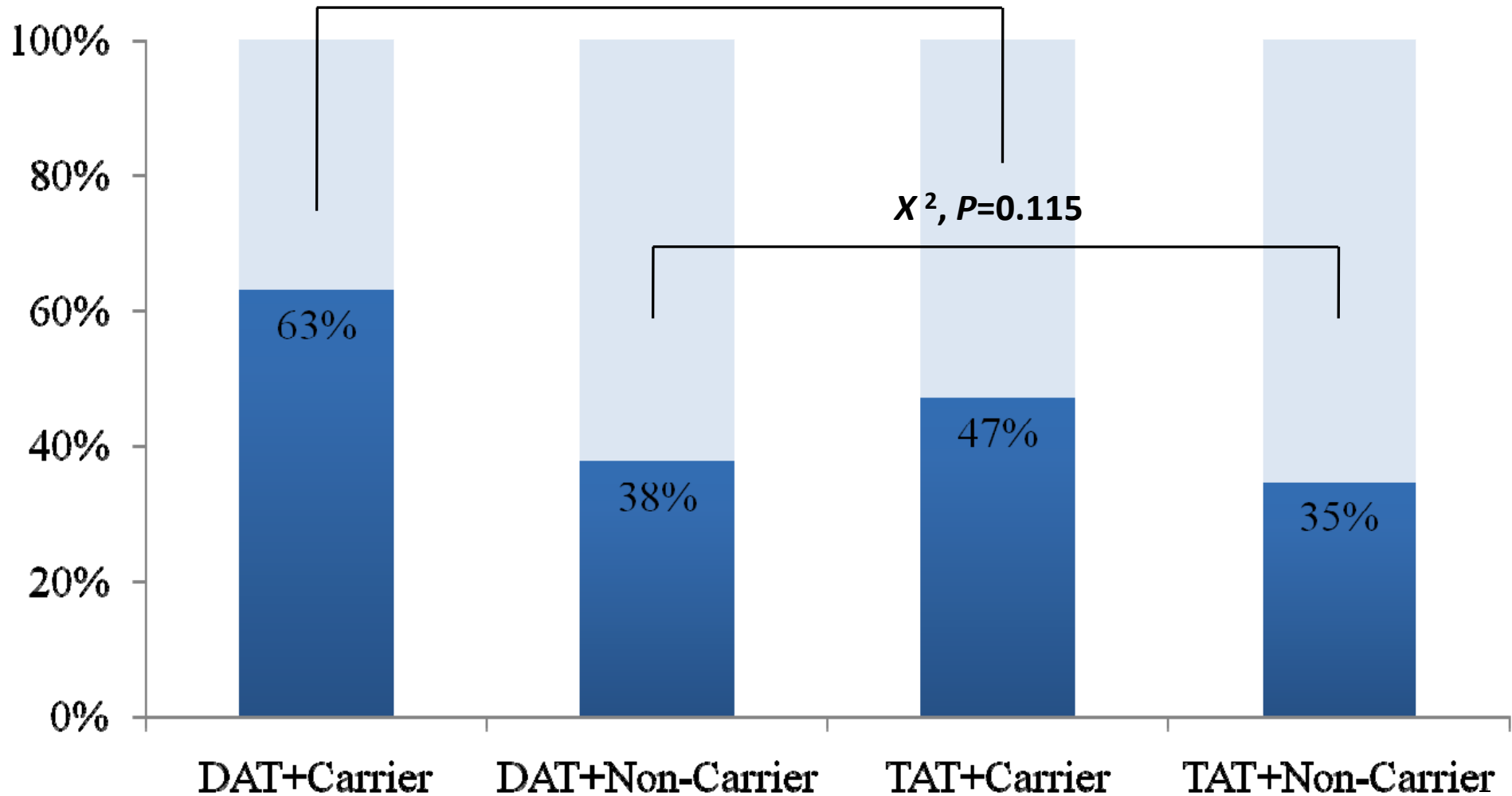
$\chi^2, P<0.001$



Frequency of HOPR

according to treatment and genotype
(using PRU>230 as definition of HOPR)

$\chi^2, P<0.001$



Independent predictors of HOPR

	OR	95% Confidence Interval		P-value
		Lower limit	Upper Limit	
Age in decade	1.34	1.03	1.73	0.029
Female gender	2.53	1.56	4.12	< 0.001
Antiplatelet regimen according to genotype				
DAT + Carrier	3.13	1.65	5.94	< 0.001
TAT + Non-Carrier	1.12	0.54	2.33	0.760
TAT + Carrier	1.25	0.66	2.36	0.491

Input variables: age (in decade), gender, cigarette smoking, diabetes, chronic kidney disease, antiplatelet treatment regimen according to CYP2C19 genotype



Summary and Conclusion

- 1. CYP2C19 LOF PM are associated with higher mean OPR and increased risk for HOPR, and with increased risk of hard outcome after PCI.***
- 2. The frequency of CYP2C19 LOF alleles are significantly higher in the Asian population.***
- 3. The addition of cilostazol resulted in lower mean OPR and reduction of risk for HOPR in patients with CYP2C19 LOF alleles.***

