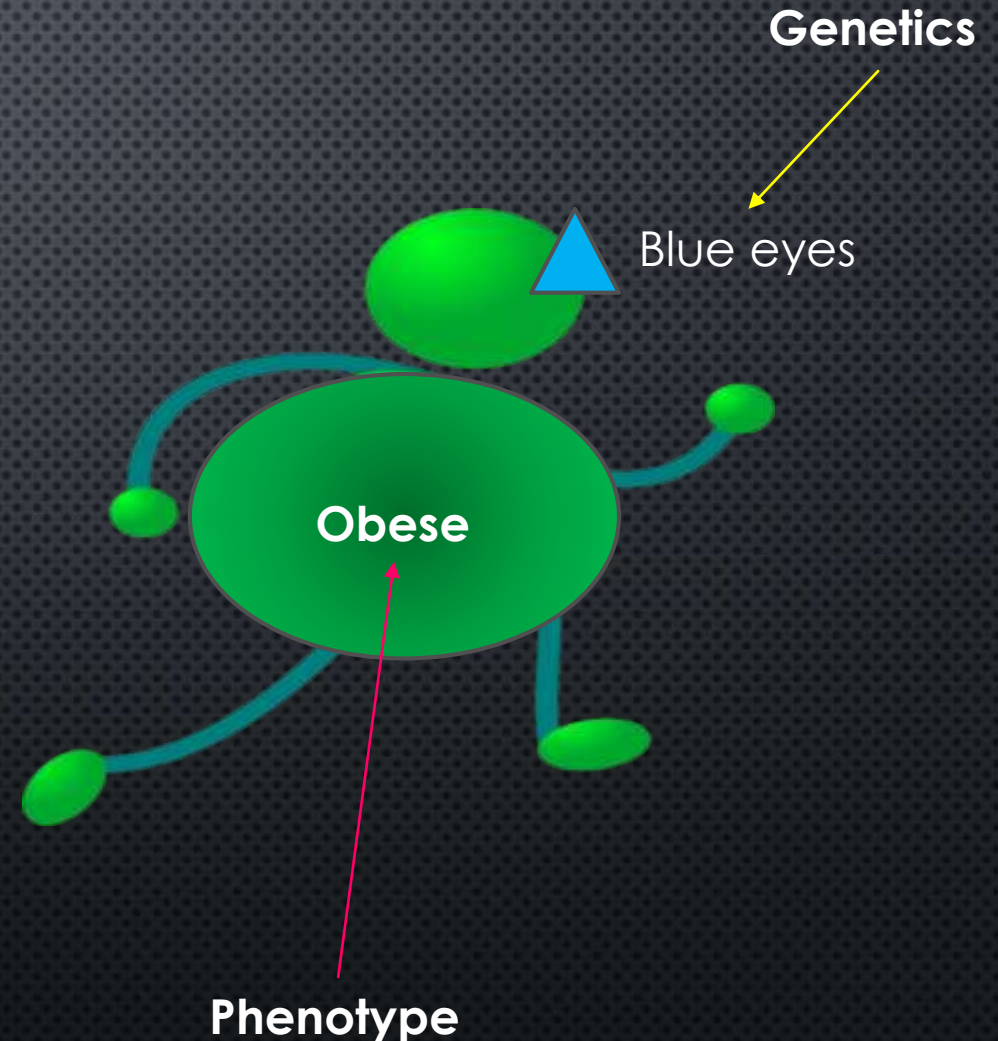


WILL GENOTYPE OR PHENOTYPE DRIVE CARDIAC INTERVENTIONS: PRECISION MEDICINE

**Professor Robert Chilton
University of Texas Health Science Center
San Antonio, Texas
Director of Cath Lab
Director clinical proteomics center**

HELPFUL DEFINITIONS

- 'GENOTYPE' IS USUALLY USED WHEN TALKING ABOUT THE GENETICS OF A PARTICULAR TRAIT (LIKE *EYE COLOR*)
- "PHENOTYPE" USED WHEN TALKING ABOUT *PHYSICAL* OR *BIOCHEMICAL* CHARACTERISTICS OF AN INDIVIDUAL



SUMMARY

- GENETIC INFORMATION IS GOOD ENOUGH TO ATTRIBUTE CAUSE TO INDIVIDUALBUT
- PHENOTYPE ... USUALLY CONVEYS MOST INFORMATION CLINICALLY

Electrophysiology

PHENOTYPE



Atrial fibrillation (AF) is the most common sustained cardiac arrhythmia in humans

1 in 10 patients over age 80

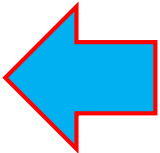
Prevalence in the United States projected to roughly double by the year 2050 to an estimated 6-12 million

GENOTYPE
4q25 locus

INCREASED RISK OF ATRIAL FIBRILLATION ON CHROMOSOME 4Q25: 2 LOCATIONS

Table 2 | Association by age at diagnosis in Iceland and by AF sub-phenotype in the United States

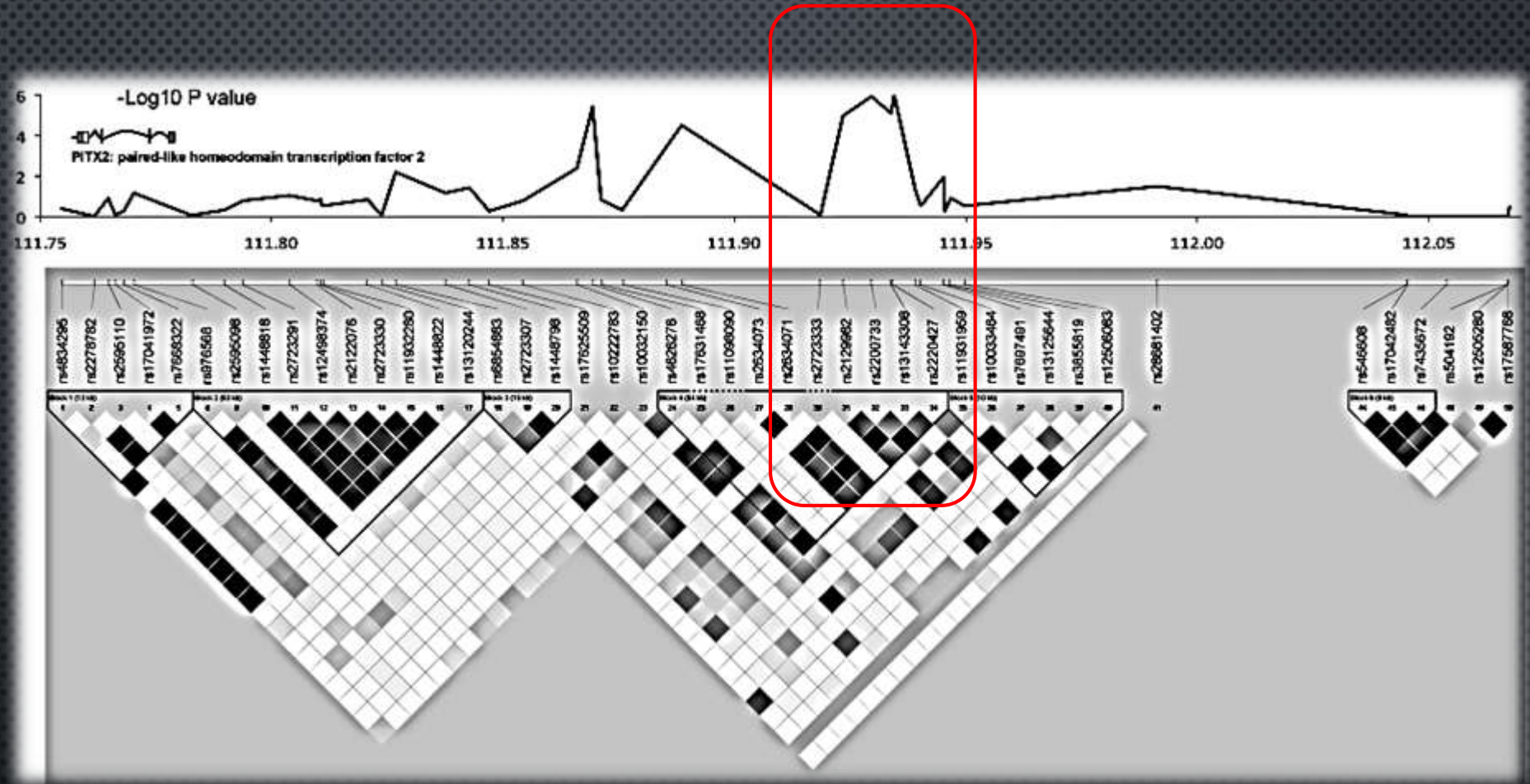
Sample (cases/controls)	Male (%)	Age (yr)	OR (95% CI)		P
			rs2200733*	rs10033464*†	
Iceland‡					
Diagnosis at age ≤60 yr (510/17,714)	77.8	50.7 ± 8.4	2.12 (1.77–2.54)	1.69 (1.34–2.12)	6.3 × 10 ^{−18}
Diagnosis at age 60–70 yr (654/17,714)	66.2	65.6 ± 2.9	1.88 (1.60–2.21)	1.44 (1.18–1.77)	6.7 × 10 ^{−15}
Diagnosis at age 70–80 yr (958/17,714)	58.9	75.0 ± 2.8	1.60 (1.39–1.84)	1.23 (1.03–1.47)	7.5 × 10 ^{−11}
Diagnosis at age >80 yr (679/17,714)	47.4	85.6 ± 4.2	1.20 (1.01–1.43)	1.31 (1.08–1.60)	0.0044
United States					
Lone AF (251/804)	81.7	46.1 ± 11.5	2.32 (1.80–2.99)	1.68 (1.19–2.37)	1.2 × 10 ^{−10}
AF + hypertension (67/804)	74.6	54.5 ± 10.2	2.23 (1.43–3.48)	1.66 (0.90–3.04)	0.0010
Other AF (318/804)	52.8	75.2 ± 11.3	1.44 (1.12–1.84)	0.97 (0.69–1.37)	0.015



PHENOTYPE

MORE POST OP ATRIAL FIBRILLATION ASSOCIATED WITH RS2200733 POLYMORPHISM AT THE 4Q25 LOCUS

CABG Genomics
discovery cohort
(N=959)
postoperative AF
(n=289)



Circ Cardiovasc Genet. 2009;2:499-506

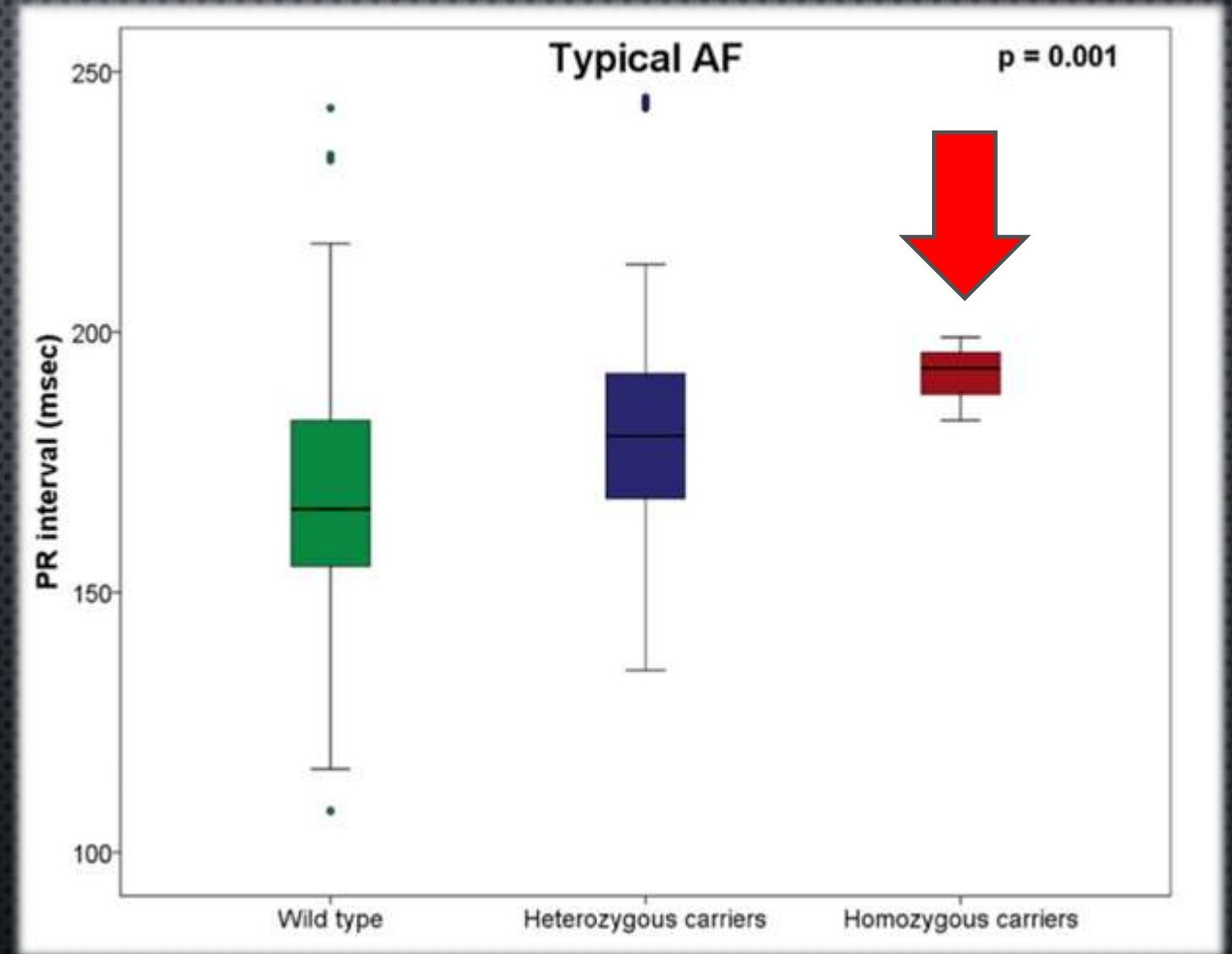
Single Nucleotide Polymorphism Database record ID number ("rs#"; e.g. rs206437)

HOMOZYGOUS CARRIERS HAVE INCREASED RISK FOR AFIB WITH INCREASED PR INTERVAL

rs2200733 polymorphism at the 4q25 locus was associated with **PR interval** duration in patients typical AF with no history of AF

N=1403 controls
N=269 typical Afib

GENOTYPE



Am J Cardiol. 2014 January 15; 113(2): 309–313

ELECTROCARDIOGRAPHIC PR INTERVAL DURATION (MILLISECONDS) BASED ON RS2200733 GENOTYPE

	Total	CC	CT	TT	P-value
Lone atrial fibrillation	168 [152–188]	162 [148–184]	178 [157–196]	176 [156–181]	0.038
Typical atrial fibrillation	171 [156–187]	166 [155–183]	180 [168–192]	196 [188–206]	0.001
No atrial fibrillation	156 [144–170]	156 [144–169]	156 [144–170]	170 [149–176]	0.4

Are Genetic Tests for Atherosclerosis Ready for Routine Clinical Use?

Nina P. Paynter, Paul M Ridker and Daniel I. Chasman

<http://dx.doi.org/10.1161/CIRCRESAHA.115.306360> Circulation Research. 2016;118:607-619

Originally published February 18, 2016

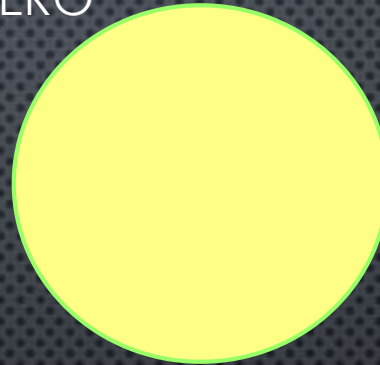
Familial hypercholesterolemia (FH) -inherited disorder high concentration of LDL

Xanthomas, deposits of cholesterol in peripheral tissues, and ↑↑ atherosclerosis from cholesterol deposition in the arterial wall



Mutations in the LDL-receptor gene (LDLR)
Asp461Asn mutation in LDLR gene

HETERO



2 times higher

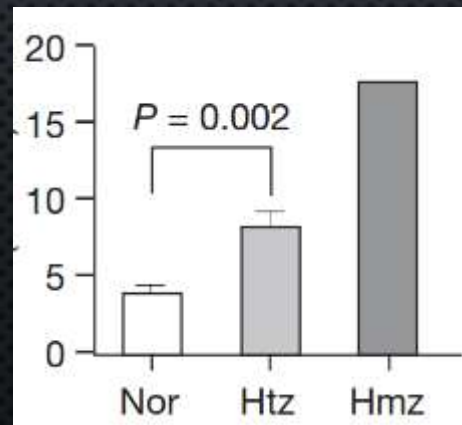
4 times higher



HOMO

Khachadurian in 1964 in Lebanese FH pedigrees...PHENOTYPE

Plasma Cholesterol mmol/L

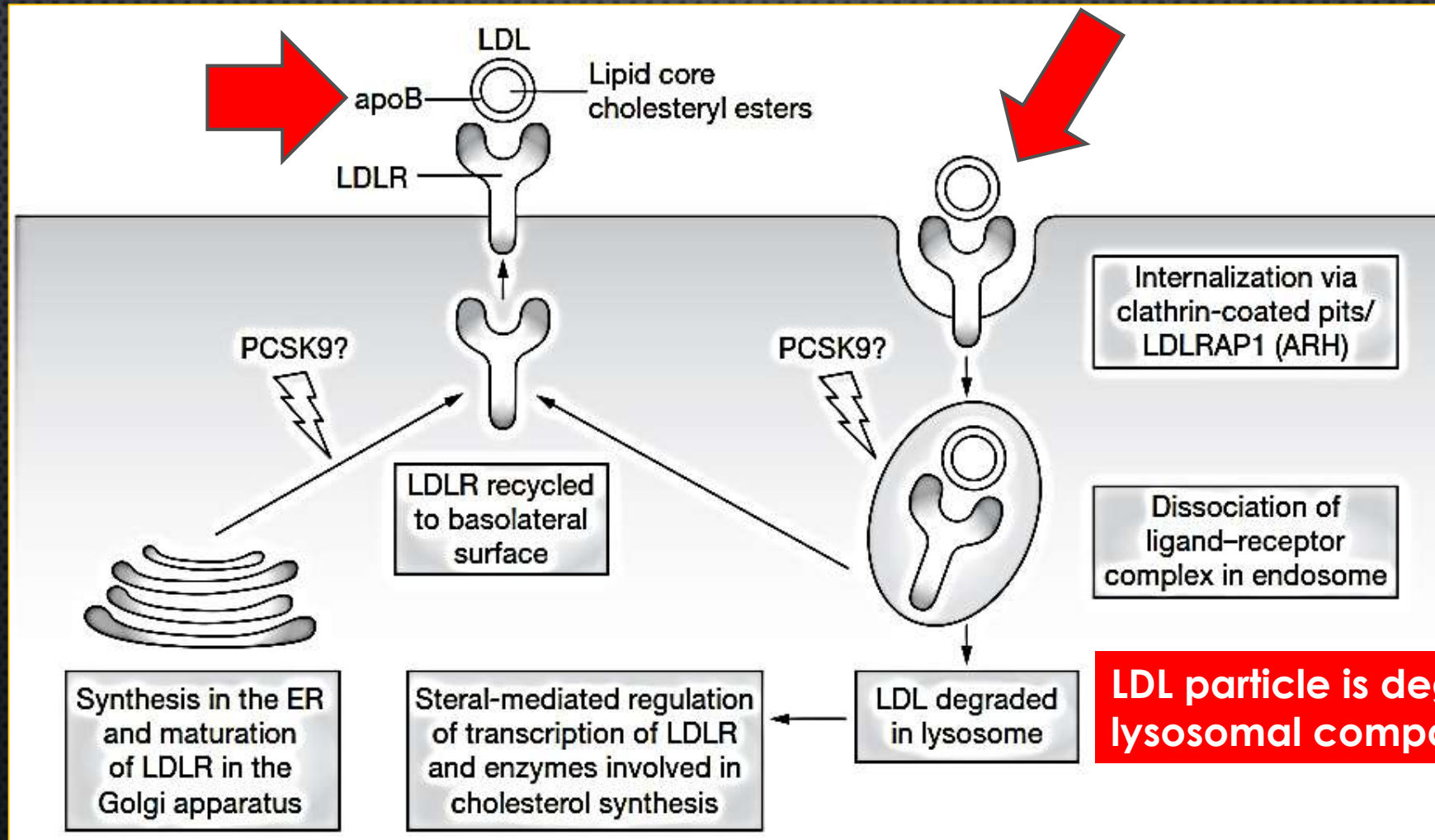


Normal

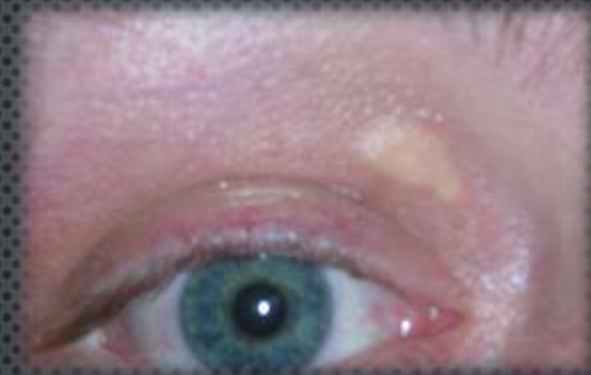


Specific for apolipoprotein B in LDL

LDL receptor--cell-surface glycoprotein -
synthesized as an immature protein -
and processed in the Golgi apparatus--
mature form that is transported to the cell
surface

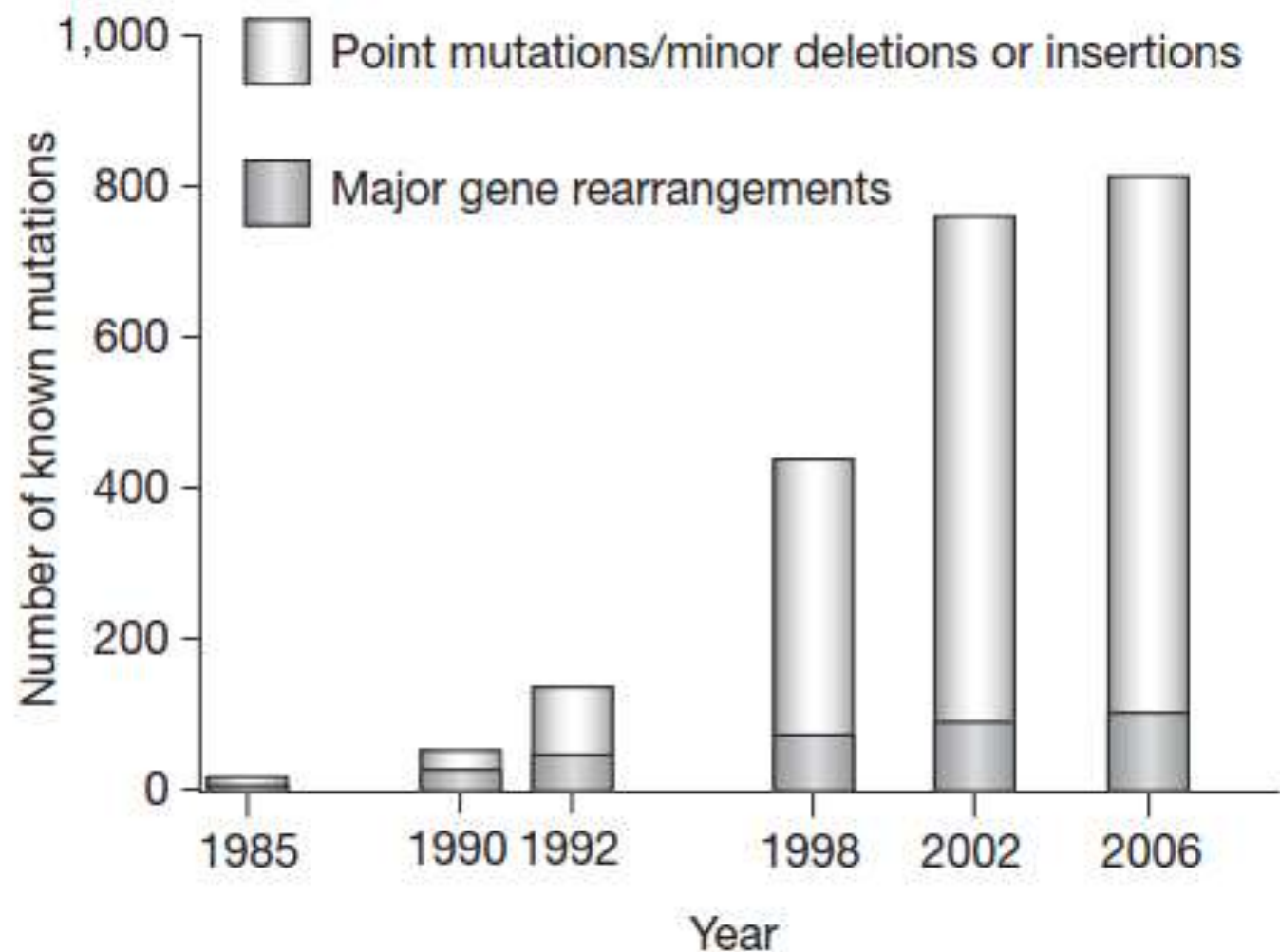
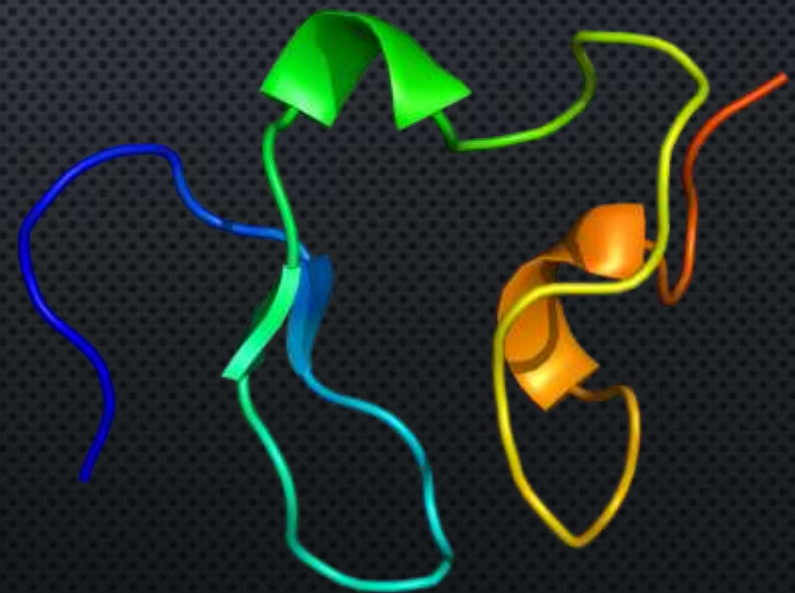


PHENOTYPE



Xanthelasma above right eye

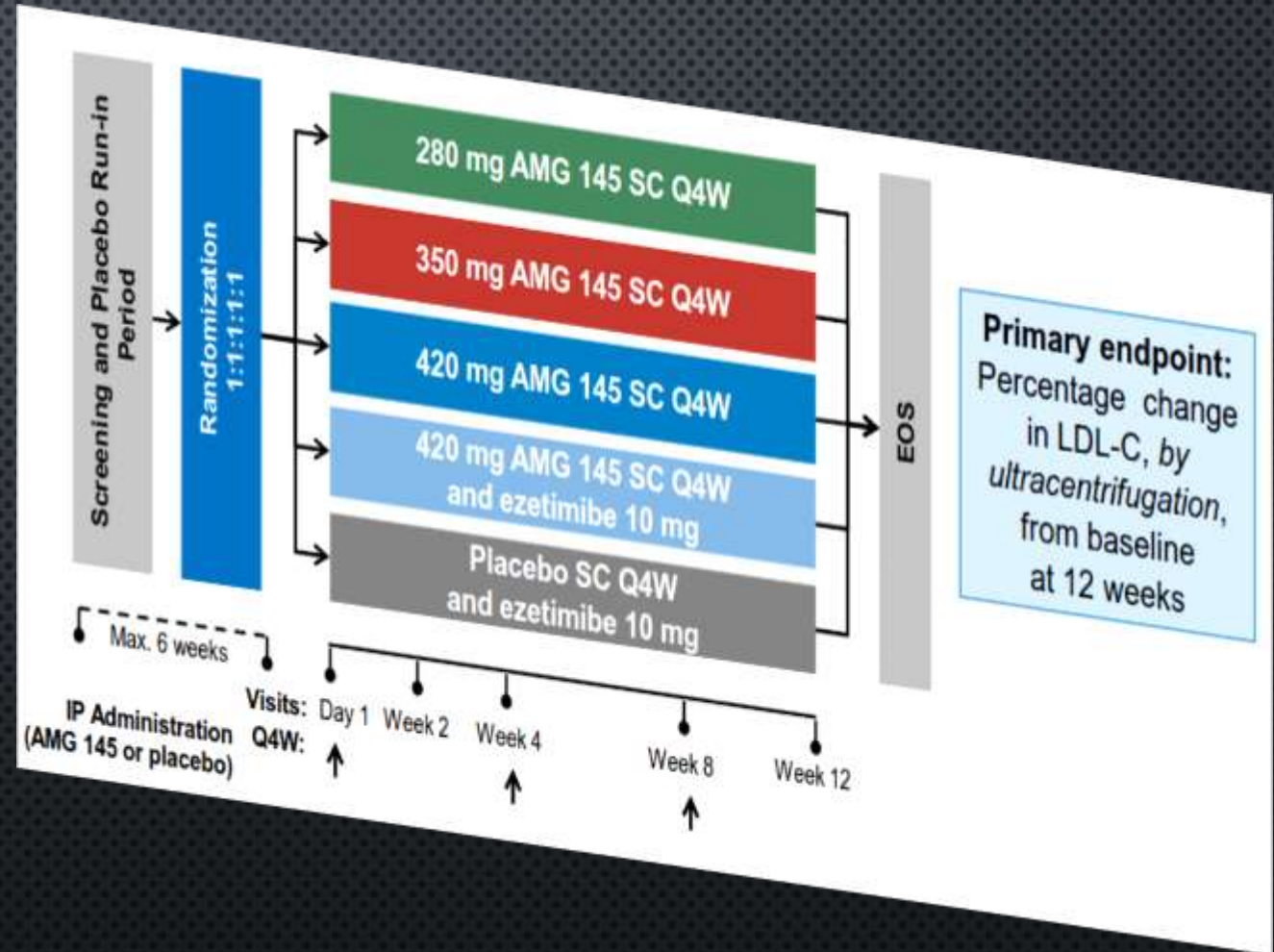
LDL receptor protein



Asp461Asn mutation in LDLR gene-GENOTYPE

GOAL ACHIEVEMENT AFTER UTILIZING AN ANTI-PCSK9 ANTIBODY IN STATIN INTOLERANT SUBJECTS (GAUSS)

- 10% TO 20% OF PATIENTS CANNOT TOLERATE STATINS
- PLASMA PROPROTEIN CONVERTASE SUBTILISIN/KEXIN TYPE 9 (PCSK9) PLAYS A PIVOTAL ROLE IN CELLULAR CHOLESTEROL HOMEOSTASIS, BY BINDING TO, AND MEDIATING THE RECYCLING OF LDL RECEPTORS
- AMG 145 -HUMAN MONOCLONAL ANTIBODY BINDS TO PCSK9 BLOCKS ITS INTERACTION WITH LDL-Rs, INCREASING THEIR RECYCLING AND REMOVAL OF LDL-C
- GLOBAL, RANDOMIZED, DOUBLE-BLIND, CONTROLLED STUDY



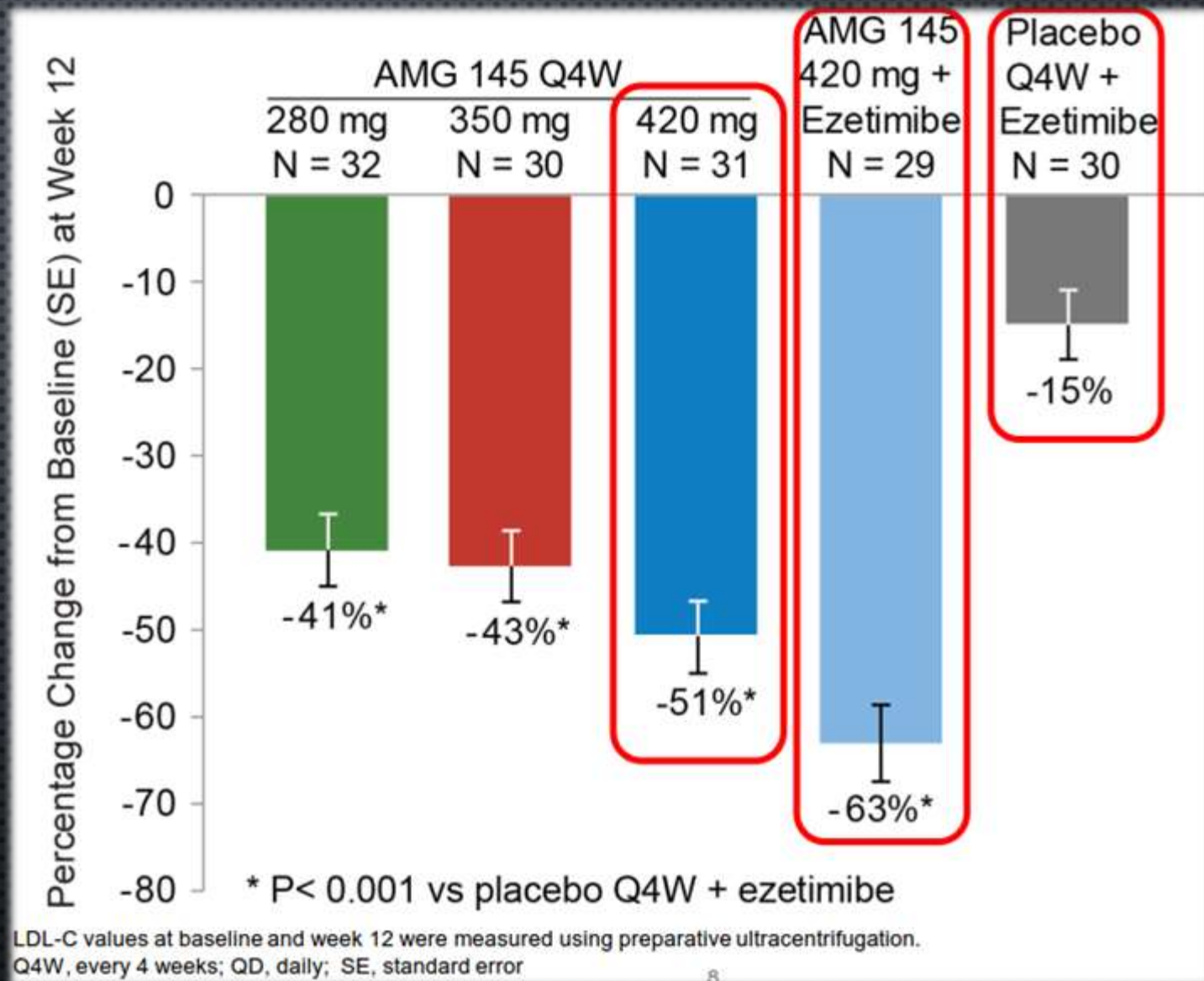
GAUSS TRIAL-BASELINE

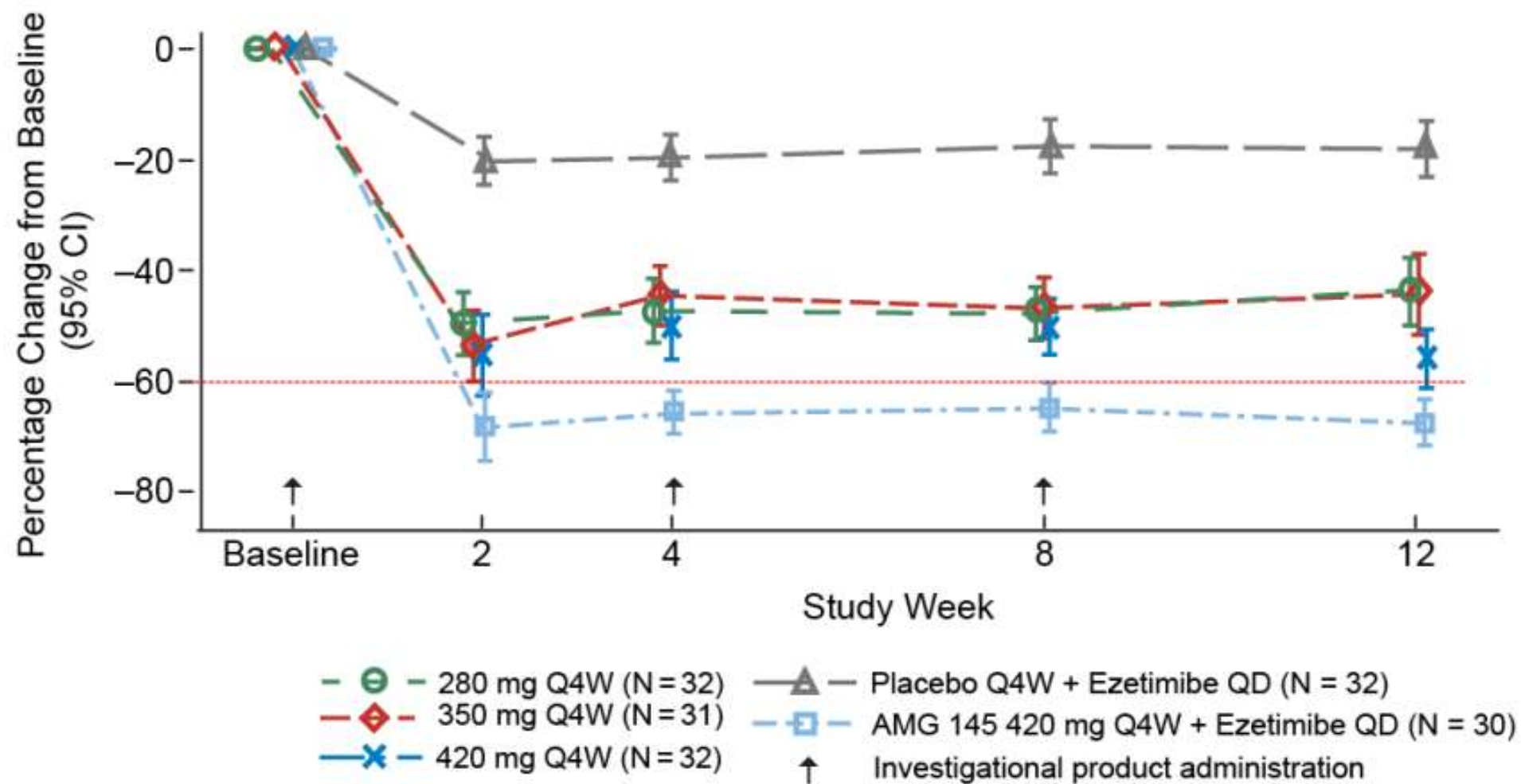
Characteristic	AMG 145 Q4W			AMG 145 420 mg + Ezetimibe N = 30	Placebo Q4W + Ezetimibe N = 32
	280 mg N = 32	350 mg N = 31	420 mg N = 32		
Sex, female, n (%)	18 (56)	21 (68)	20 (63)	23 (77)	18 (56)
Age, years, mean (SD)	62 (10)	62 (9)	60 (9)	62 (7)	62 (7)
LDL-C, mg/dL , mean (SD)*	195 (48)	190 (48)	204 (60)	194 (60)	183 (36)
Free PCSK9, ng/mL, mean (SD)	383 (98)	396 (129)	372 (87)	379 (111)	390 (91)

STATIN INTOLLERANCE

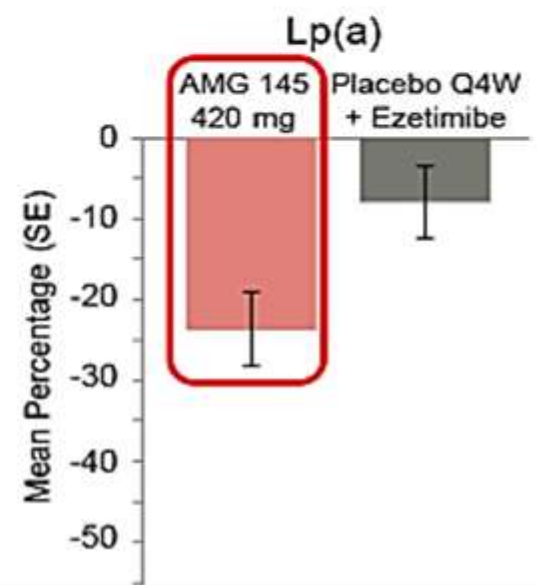
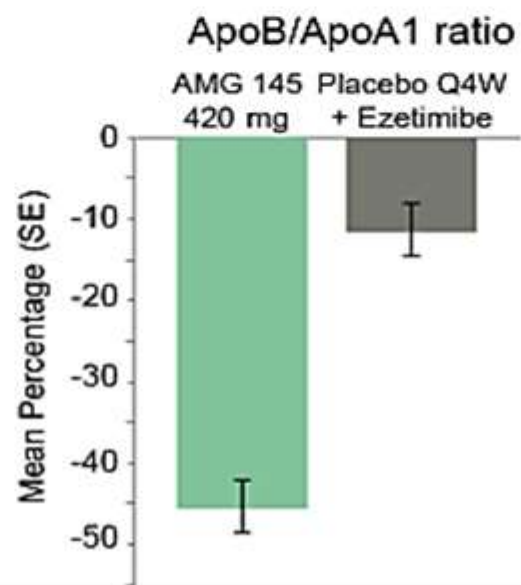
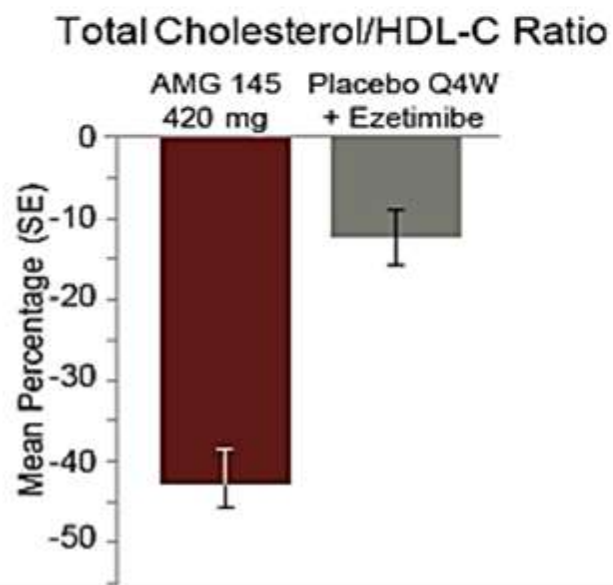
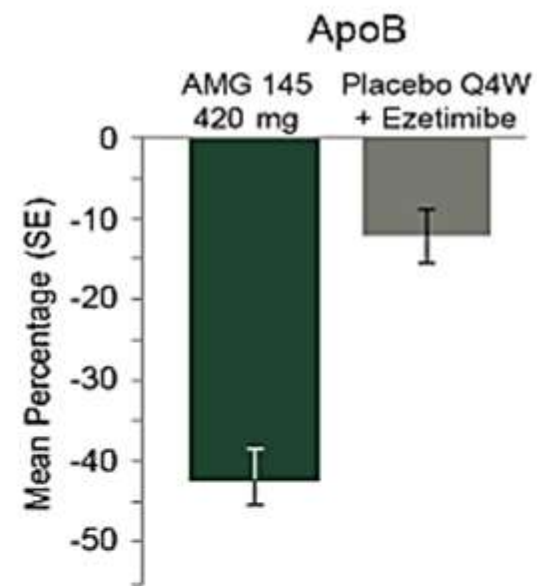
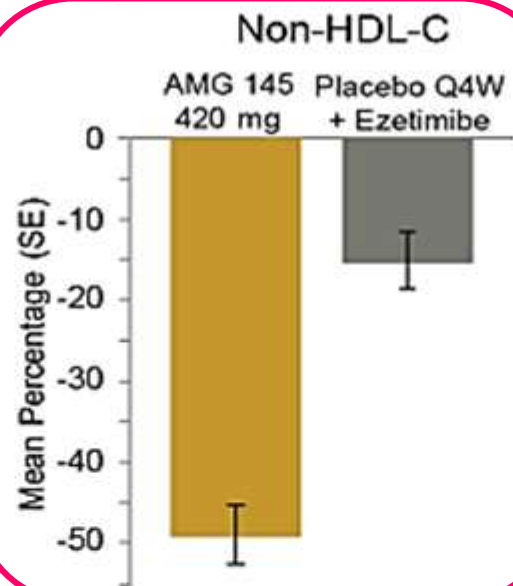
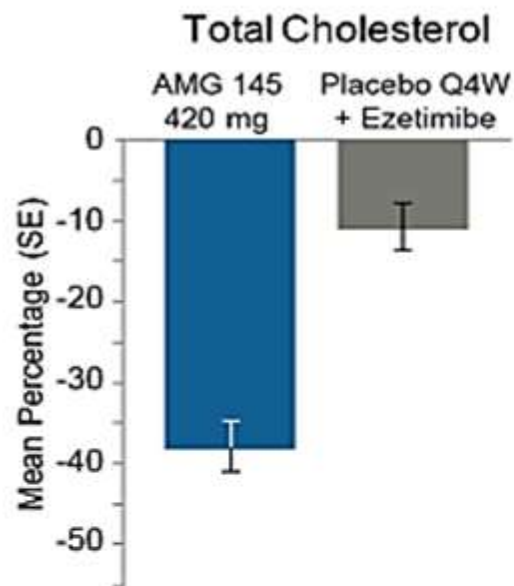
Characteristic	AMG 145 Q4W			AMG 145 420 mg + Ezetimibe N = 30	Placebo Q4W + Ezetimibe N = 32
	280 mg N = 32	350 mg N = 31	420 mg N = 32		
Statins failed (muscle-related events)					
≥ 1, n (%)	32 (100)	31 (100)	32 (100)	30 (100)	32 (100)
≥ 2, n (%)	28 (53)	24 (77)	23 (72)	21 (70)	25 (78)
≥ 3, n (%)	11 (34)	11 (35)	12 (38)	6 (20)	11 (34)
Worst statin-related events, any statin					
Myalgia, n (%)	31 (97)	30 (97)	29 (91)	29 (97)	29 (91)
Myositis, n (%)	3 (9)	3 (10)	2 (6)	2 (7)	4 (13)
Rhabdomyolysis, n (%)	0 (0.0)	0 (0.0)	1 (3)	0 (0)	0 (0)

PRIMARY ENDPOINT





Percentage Change from Baseline



Adverse Events, Patient Incidence, n (%)	AMG 145			AMG 145 420 mg + Ezetimibe 10 mg N = 30	Placebo Q4W + Ezetimibe N = 32
	280 mg N = 32	350 mg N = 31	420 mg N = 32		
Treatment-emergent AEs	22 (68.8)	15 (48.4)	18 (56.3)	20 (66.7)	19 (59.4)
Serious AEs*	2 (6.3)	1 (3.2)	1 (3.1)	0 (0.0)	0 (0.0)
Deaths	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Treatment-related AEs	8 (25.0)	3 (9.7)	6 (18.8)	5 (16.7)	7 (21.9)
Muscle-related AEs					
Myalgia	5 (15.6)	1 (3.2)	1 (3.1)	6 (20.0)	1 (3.1)
Muscle fatigue	2 (6.3)	0 (0.0)	0 (0.0)	0 (0.0)	1 (3.1)
Muscle spasms	1 (3.1)	2 (6.5)	0 (0.0)	0 (0.0)	3 (9.4)
AEs leading to discontinuation	0 (0.0)	1 (3.2)	1 (3.1)	1 (3.3)	2 (6.3)
Other most commonly reported AEs					
Nasopharyngitis	2 (6.3)	2 (6.5)	1 (3.1)	3 (10.0)	5 (15.6)
Nausea	2 (6.3)	1 (3.2)	1 (3.1)	0 (0.0)	1 (3.1)
Fatigue	4 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)	2 (6.3)

PHENOTYPE

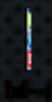
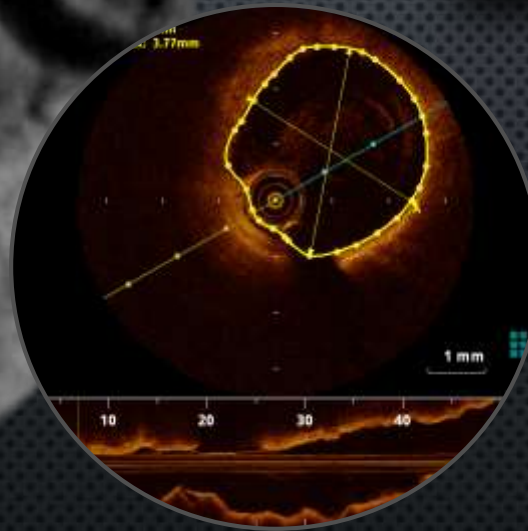
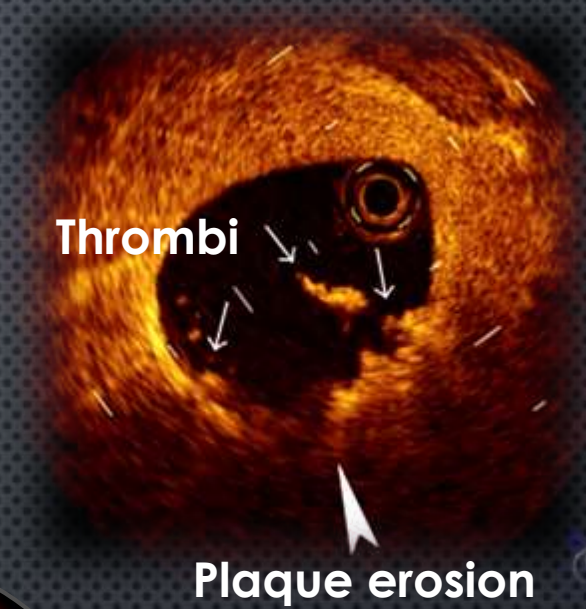
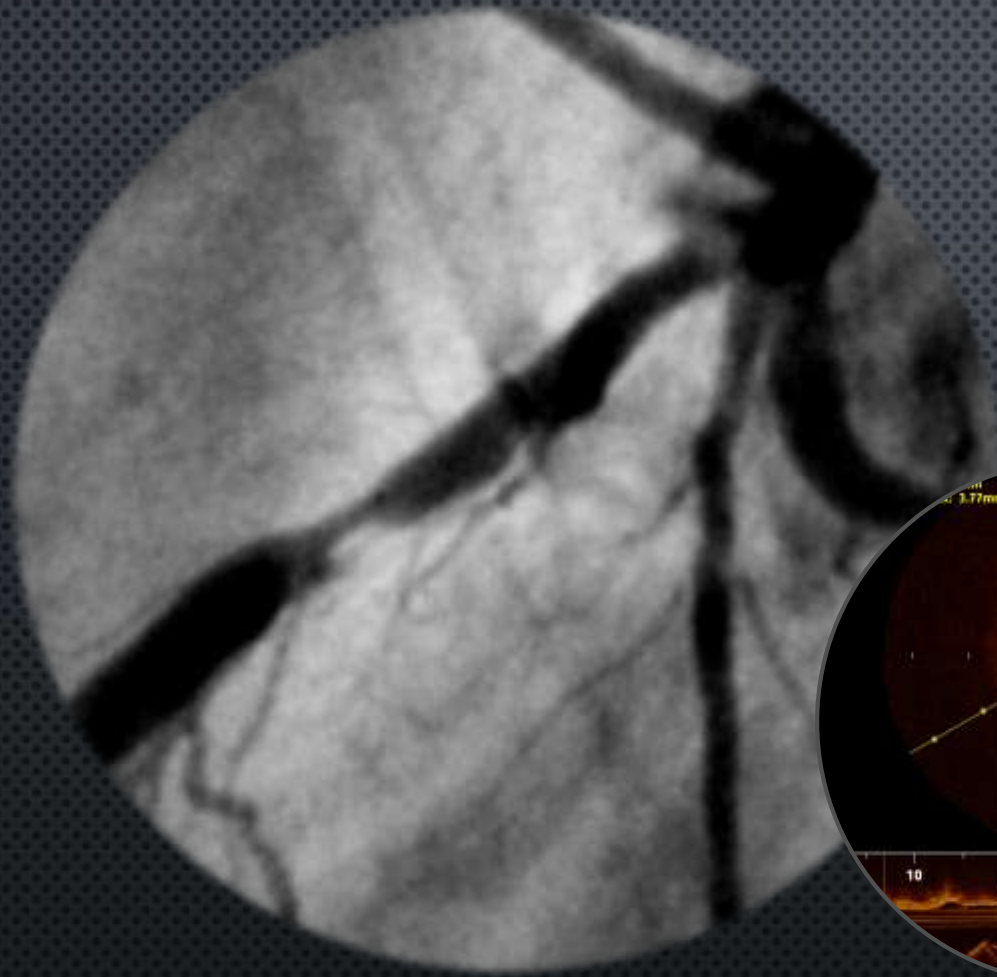


Mural thrombi



Acute coronary syndrome

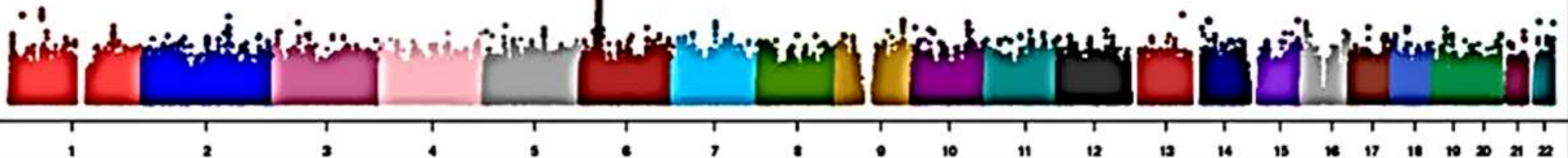
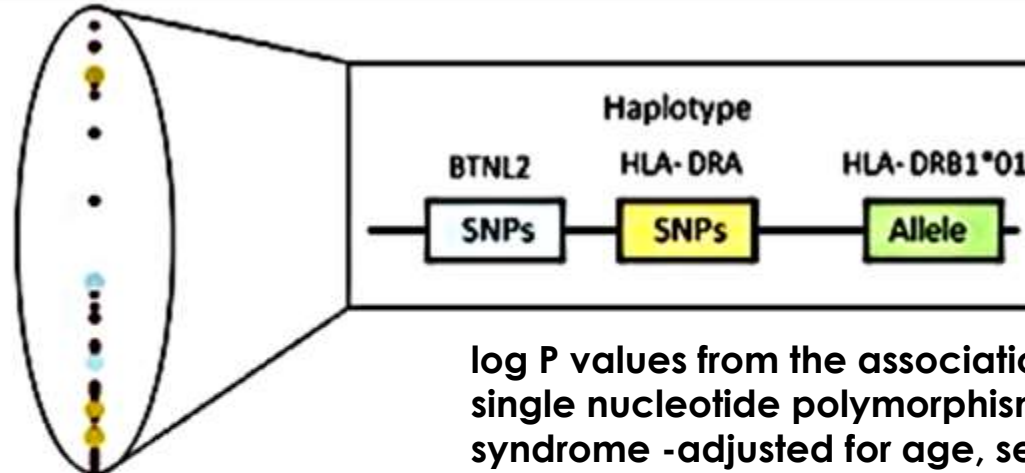
Vascular plaque rupture



GENOTYPE MARKERS FOR INCREASED RISK FOR ACUTE CORONARY SYNDROME

Associations are highly significant on BTNL2 and HLA-DRA SNPs on chromosome 6p21.3 for acute coronary syndrome

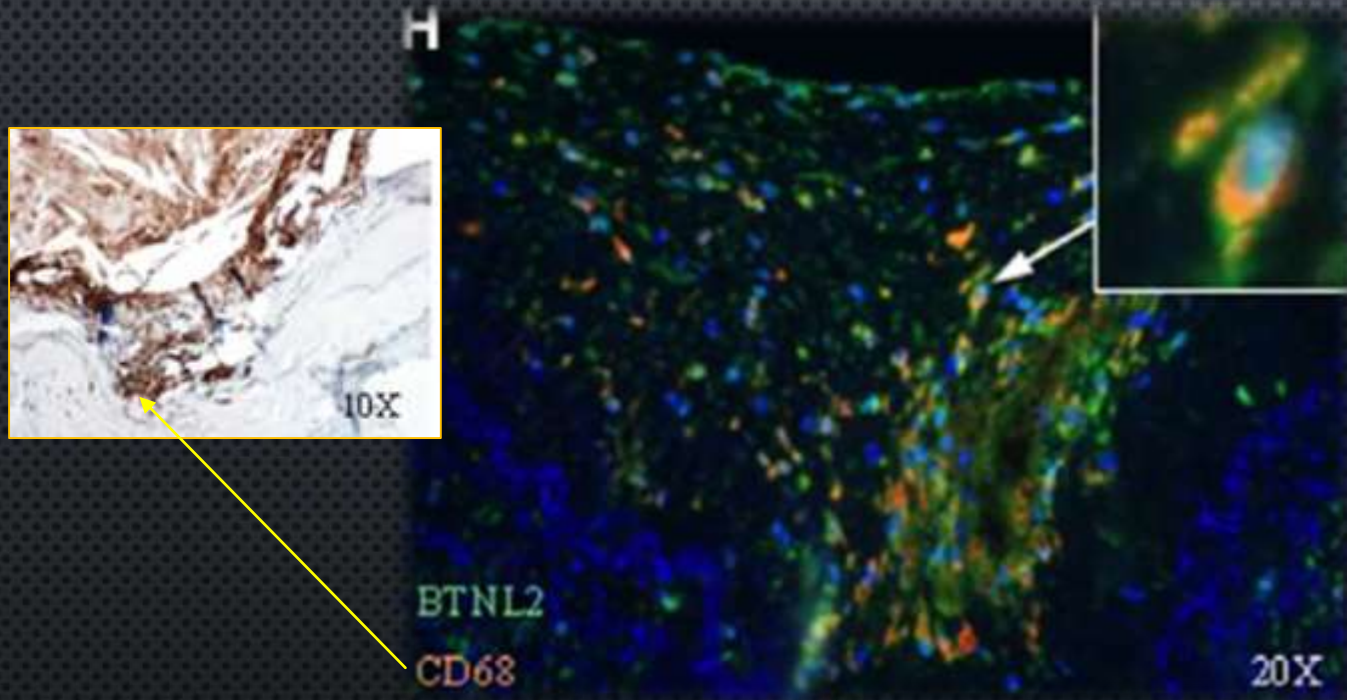
Chromosomes



Manhattan plots of the discovery population with acute coronary syndrome

Circ Cardiovasc Genet. 2016;9:55-63

ACS PLAQUES HAVE INCREASED MACROPHAGES AND BTNL2 TOGETHER




Immunofluorescence staining for colocalization of BTNL2 and CD68 in coronary artery macrophages



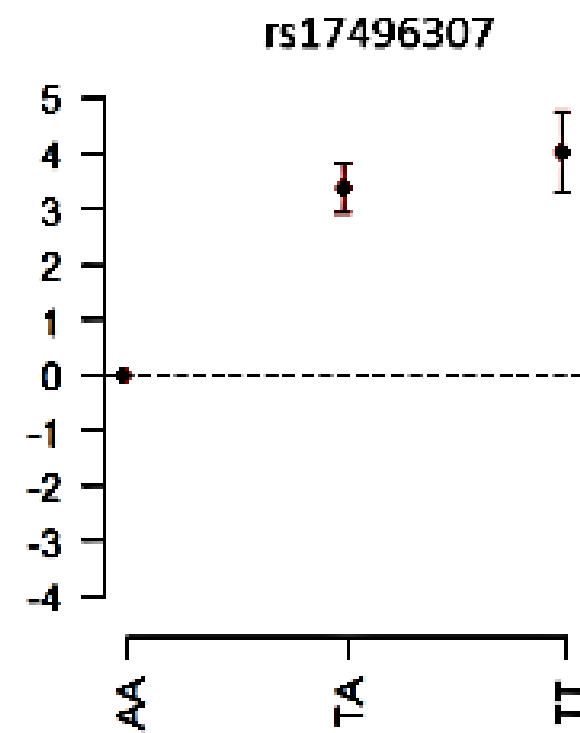
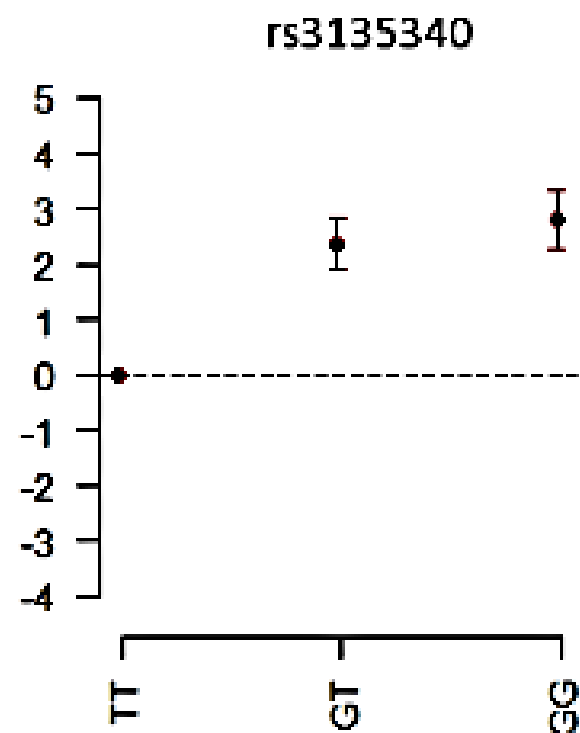
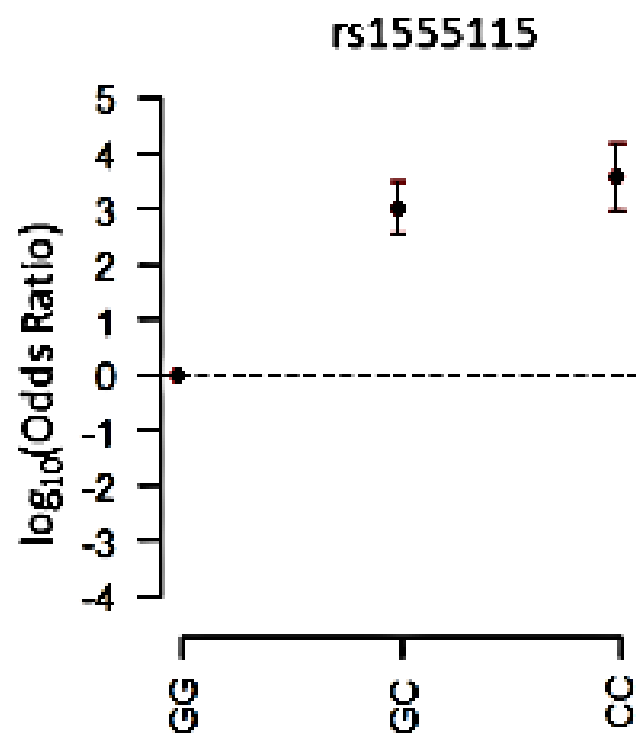
Increased immune reaction
BTNL2 inhibits FOXP3+T cell proliferation

BTNL2 is in green and CD68 in red,
and they merge in yellow

Circ Cardiovasc Genet. 2016;9:55-63



HLA-DRB1*01 Positive Individuals



SUMMARY: EXERCISE AND STAY SKINNY

Thank you