

Pharmacogenomics

Personalizing Asthma and Allergy Therapy

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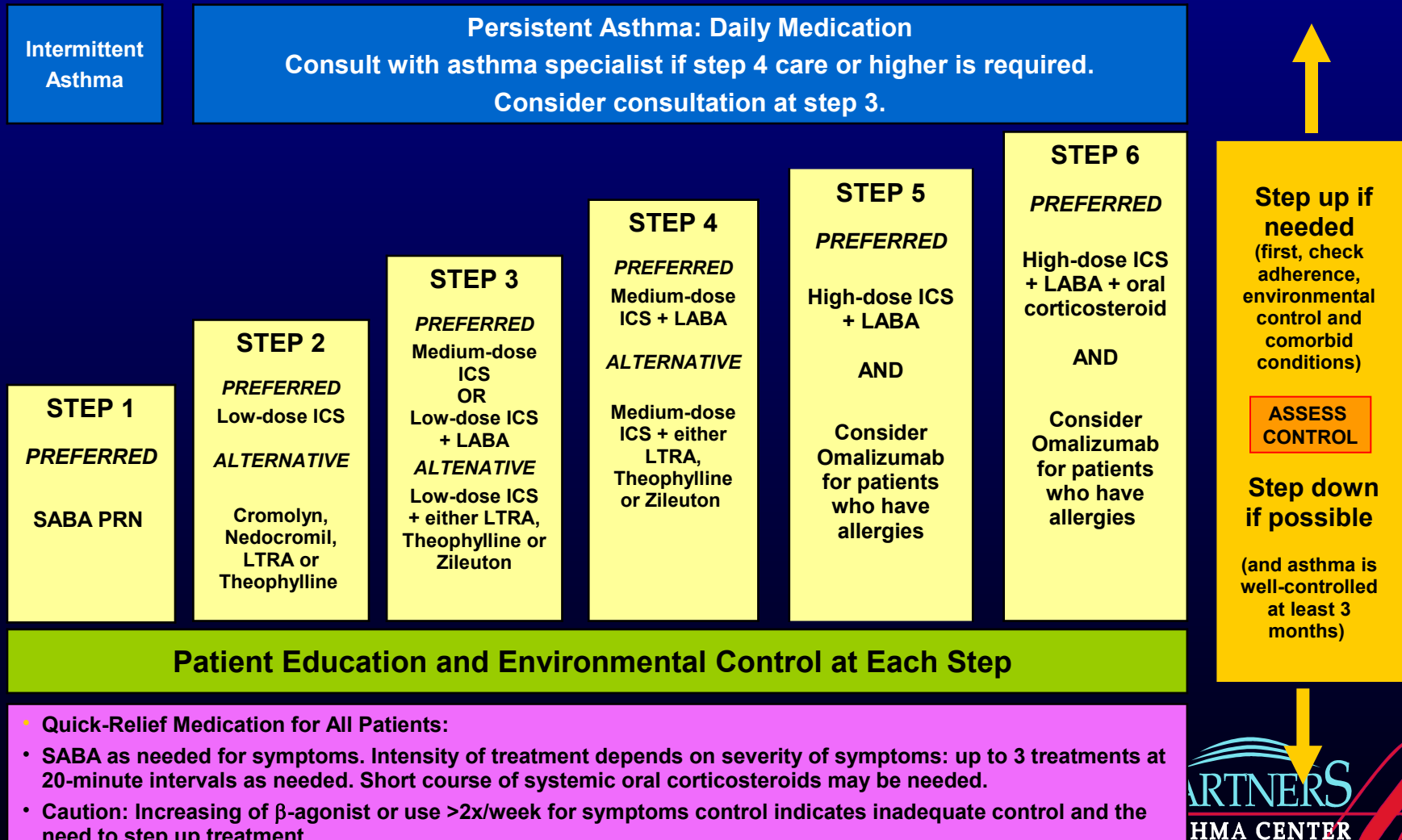
Partners' Asthma Center

Too much of a good thing?

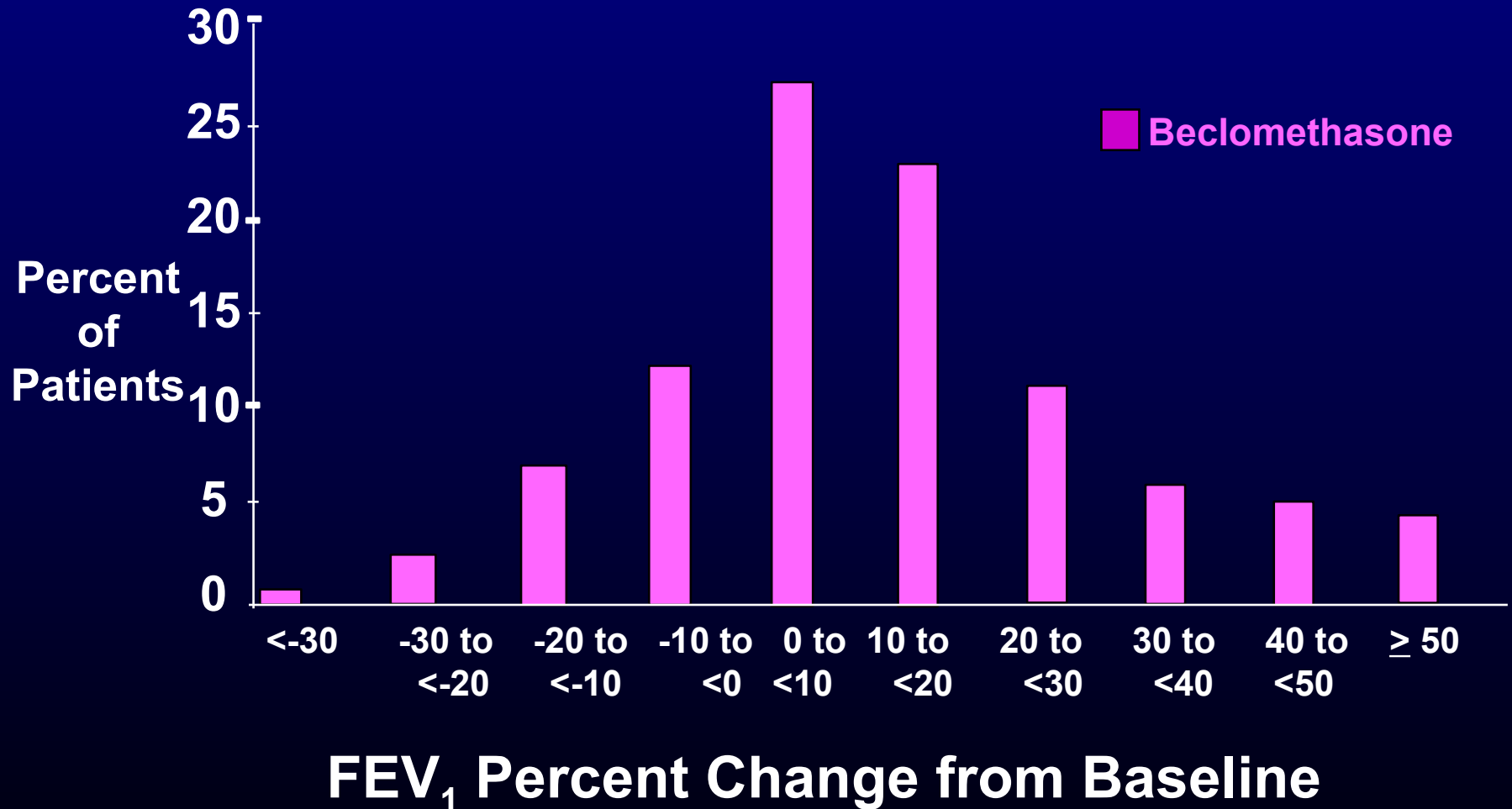
- 27 yo male with severe asthma with multiple hospitalizations
- Unable to reduce prednisone below 15 mg/d
- Uses nebulized and MDI beta-agonists 10-12 times a day
- Unable to tolerate inhaled corticosteroids because they make him wheeze
- Morbidly obese with scattered wheezes
- FEV1 70% predicted
- 7% eosinophils

- Switched to high dose ipratropium bromide (anticholinergic) and montelukast (leukotriene antagonist)
- Beta-agonists tapered
- 3 months later
 - Able to tolerate ICS
 - Off oral prednisone
- No hospitalizations in last 8 years

Stepwise Approach for Managing Asthma in Patients ≥ 12 Years of Age



45% of Patients Do Not Have an FEV₁ Response to ICS



GOAL

Bring you up to date on techniques that are allowing us to specify particular medications for individual patients

- Define pharmacogenomics and techniques
- Review developments in pharmacogenomics as they relate to use of beta agonists, leukotriene modifiers, and corticosteroids
- Review how we will use this information for treatment decisions in the future

PHARMACOGENOMICS

- Study of how genetic differences influence the variability in patients' responses (therapeutic and adverse) to drugs
 - 50-60% of genetic variability associates with variability in therapeutic responses

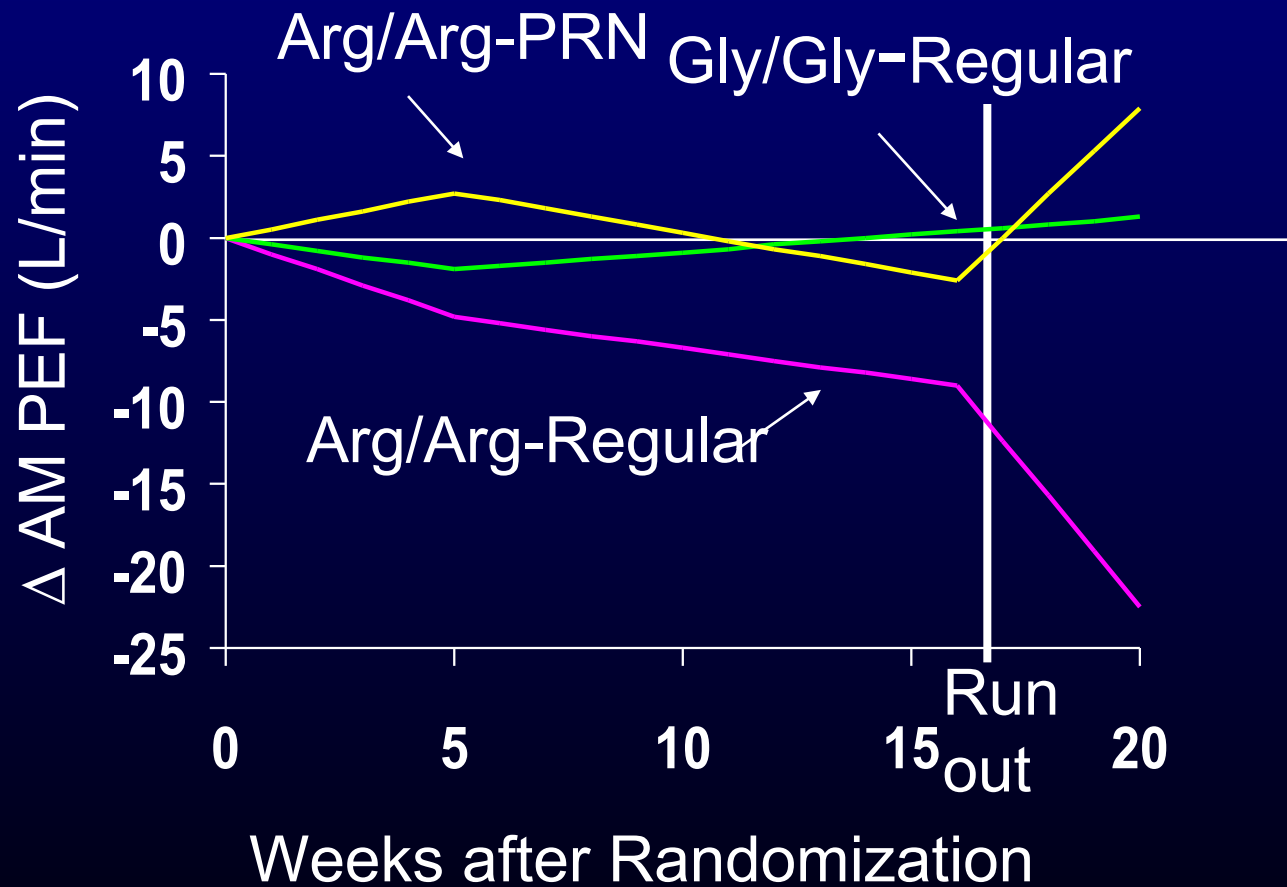
How Do We Look?

- Candidate Gene and Pathway Approaches
- GWAS
- Expression profiling in cells of responsive and non-responsive individuals

Beta-Agonists

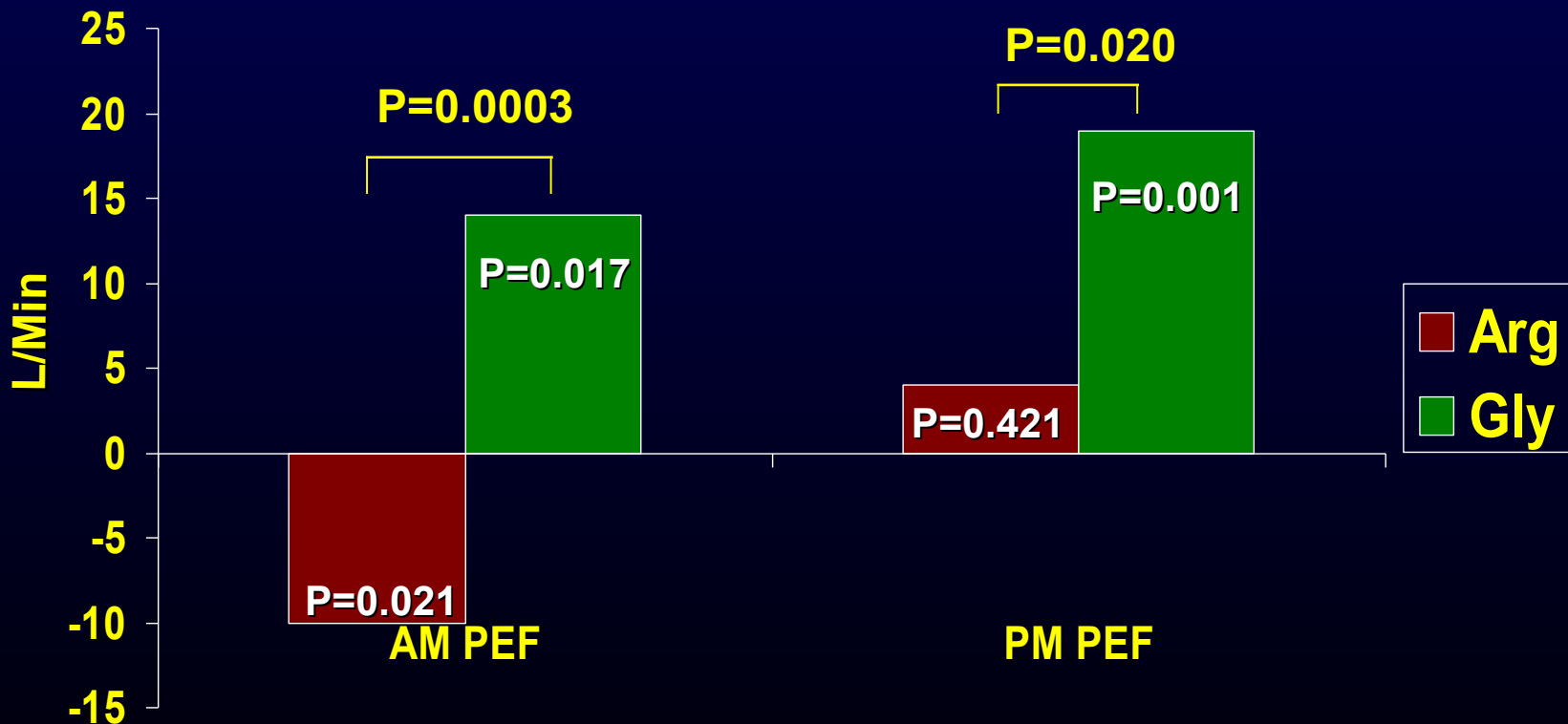
BAGS Genetic Analysis

AA16 Locus

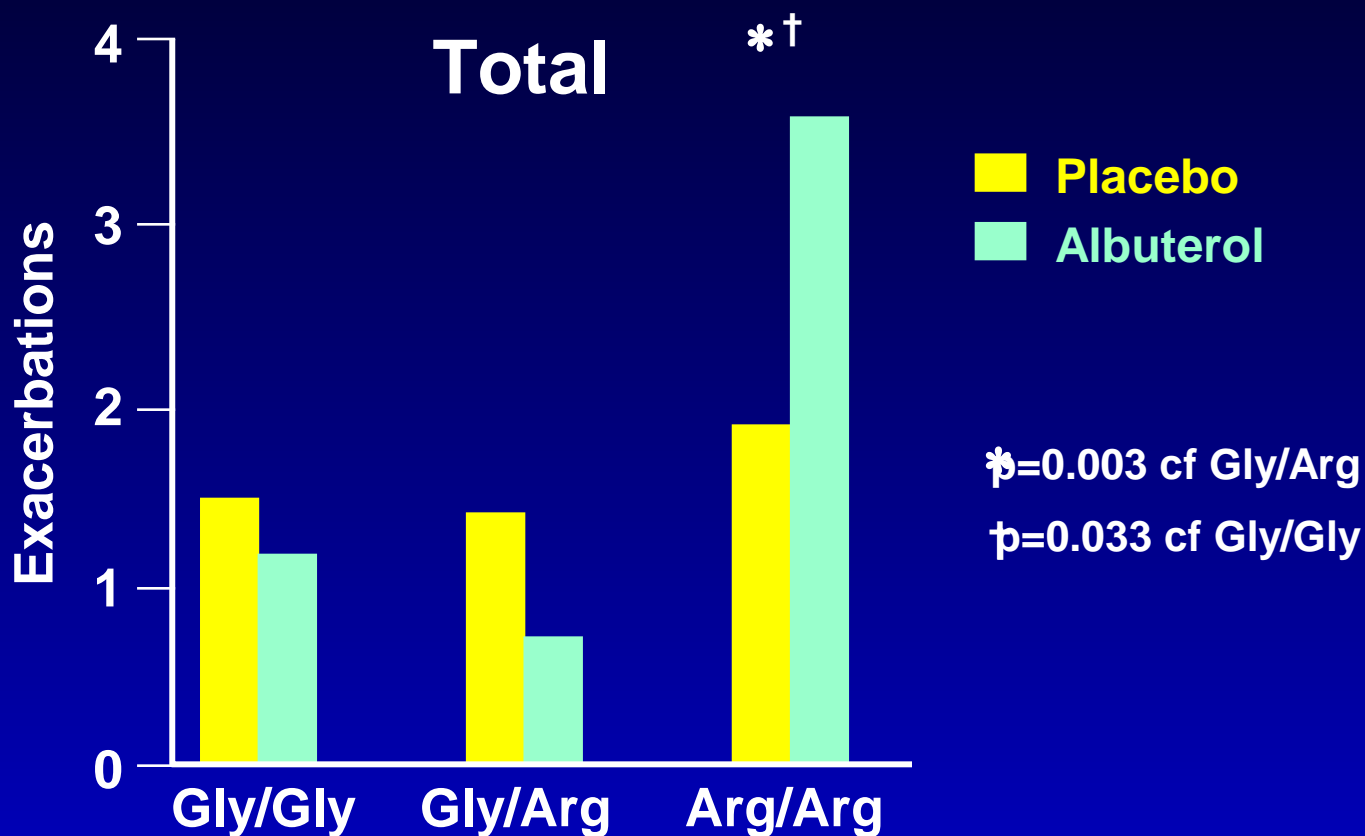


PEF

Difference between Regular vs. Placebo
Treatment-induced Changes over 16 Wk

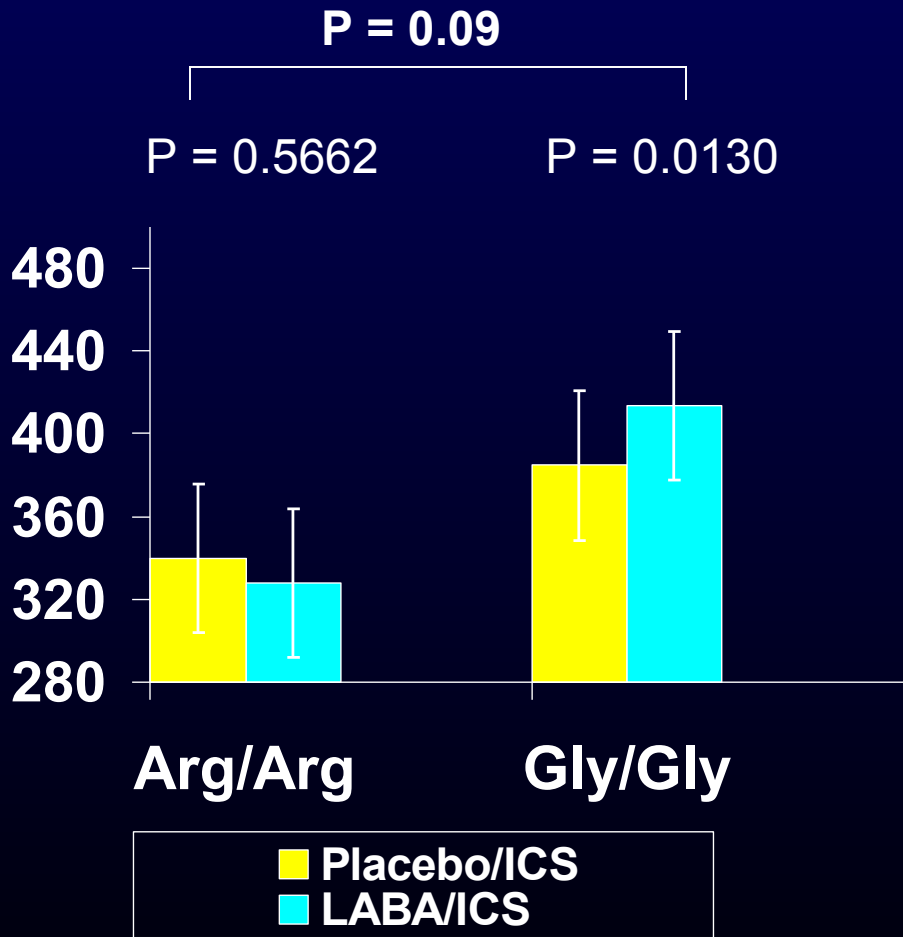


Exacerbations/Subject/Year by Genotype

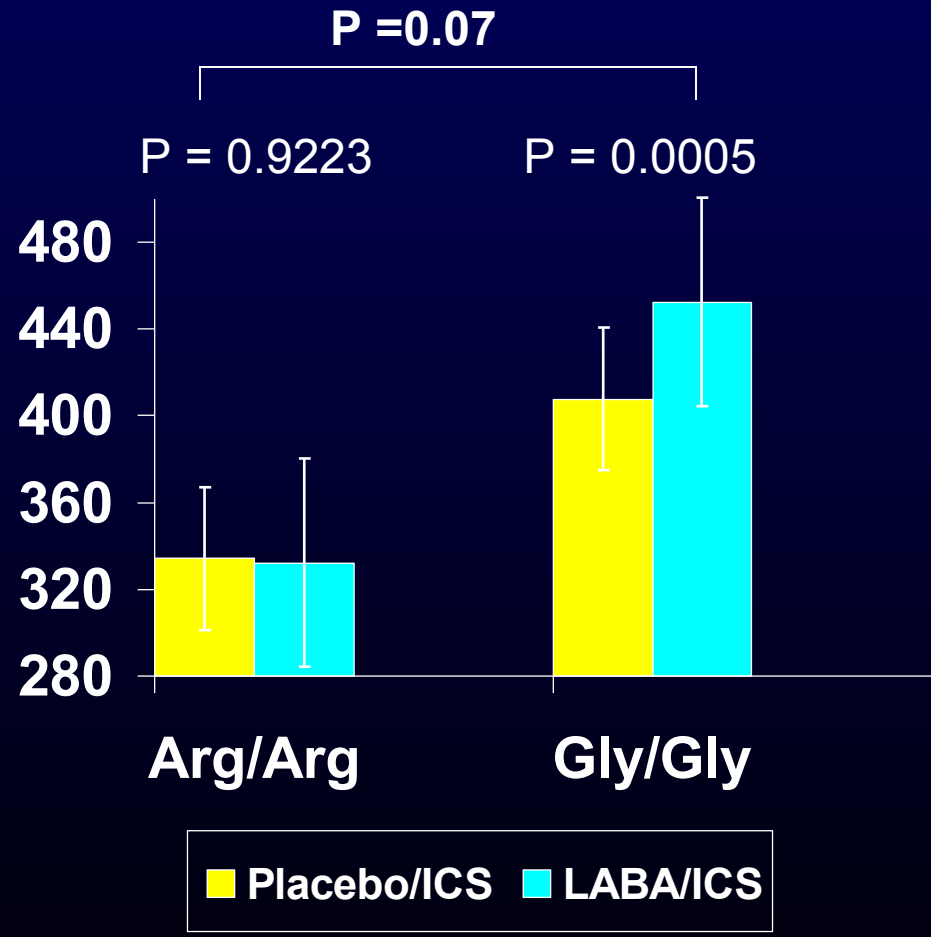


AM and PM PEF

(Black/African Amer Subgroup, n= 8 vs. 8)

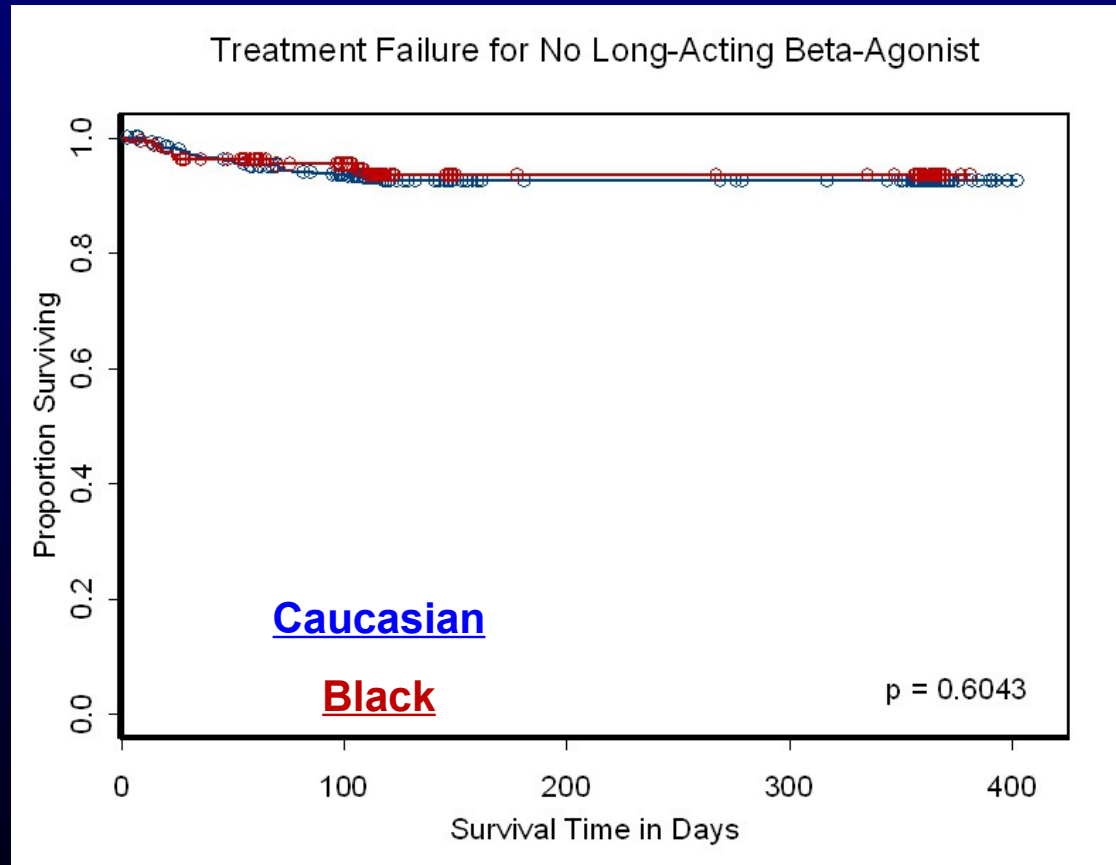


AM PEF (L/min)



PM PEF (L/min)

Treatment Failures in Subjects Not Taking LABA's Across ACRN



Treatment Failures in Subjects Taking LABA +ICS

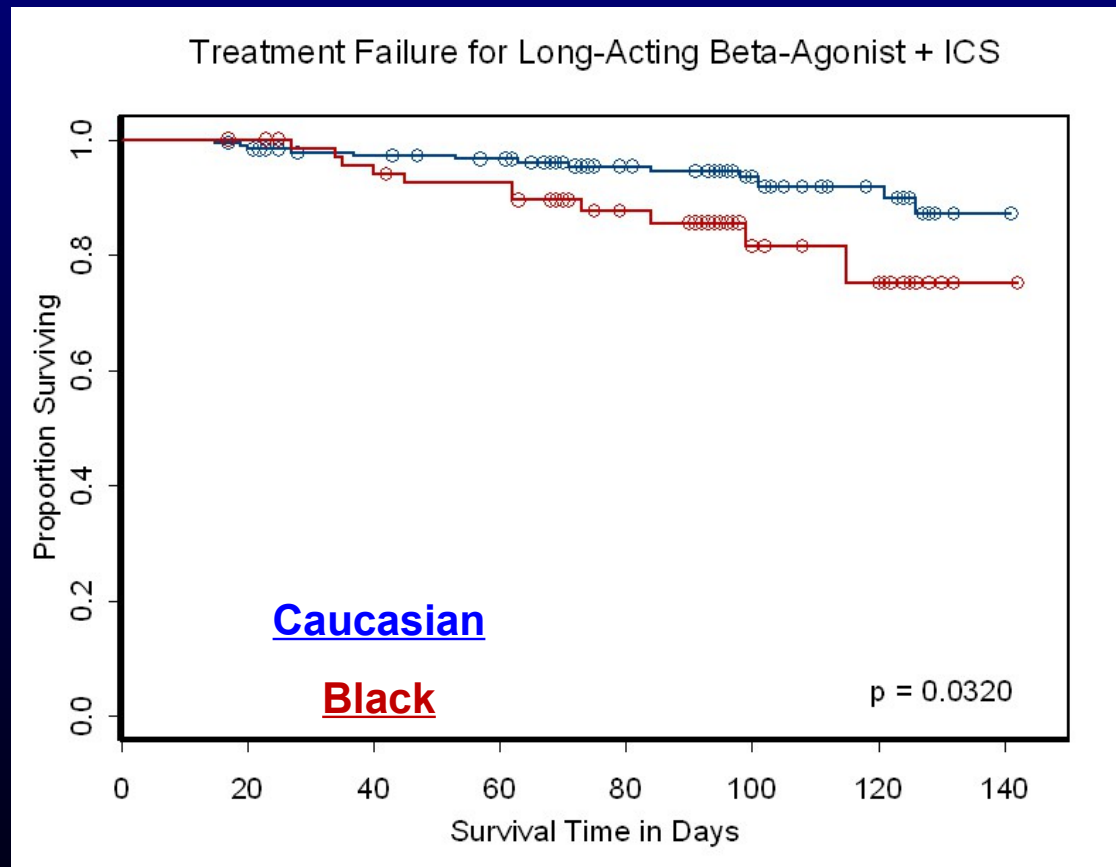


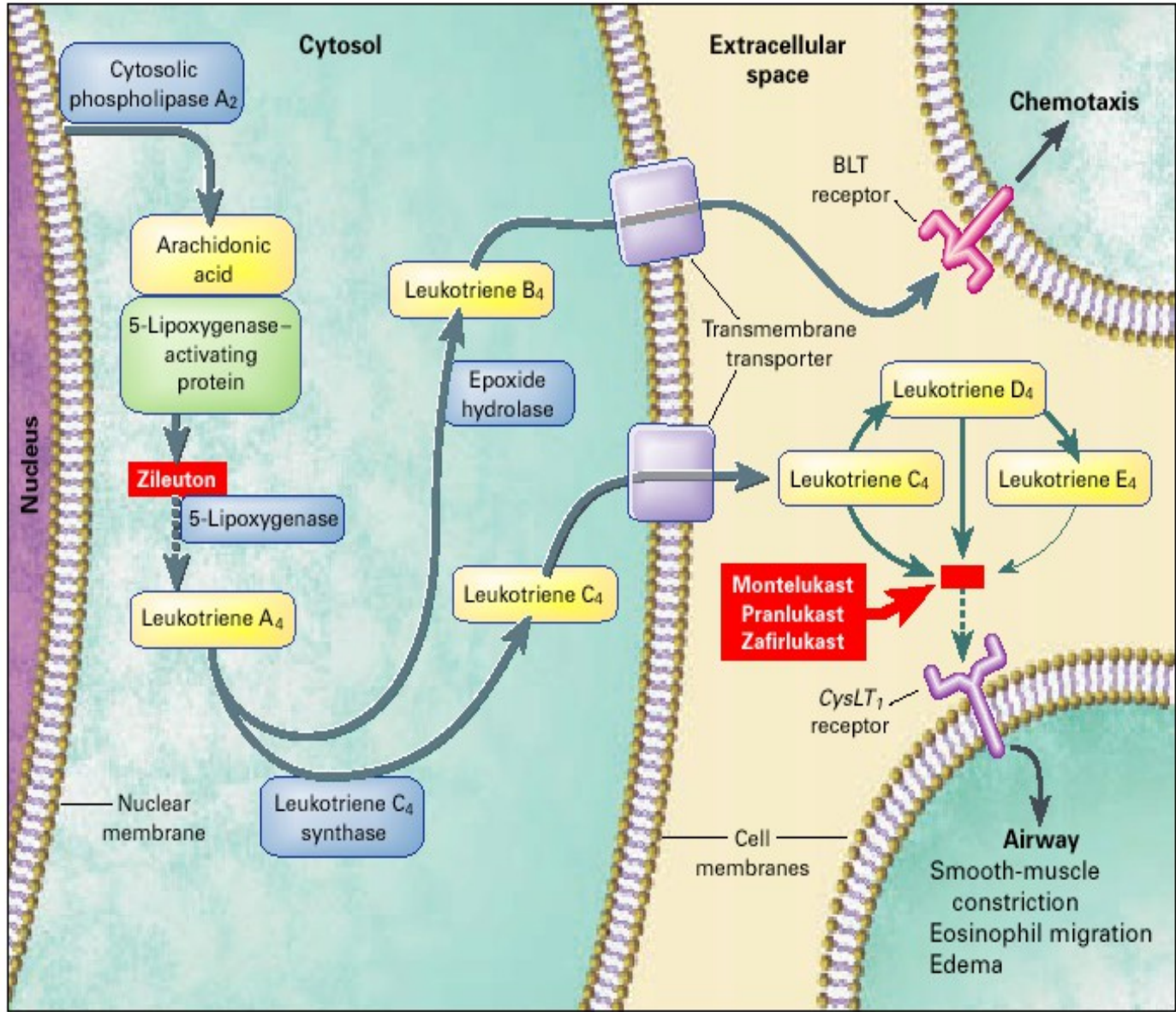
Table 1. LABA added to ICS are not as effective in Blacks: Effect of LABA Added on to ICS in Blacks vs. mixed populations

	<u>Blacks</u> Bailey 08	<u>Blacks</u> Spector 11	<u>Blacks</u> Brown 12	Mixed Shapiro 09	Mixed Ind 03	Mixed Kavuru 00
N	239	311	742	84	160	92
Duration	12 mo	3 mo	12 mo	3 mo	6 mo	3 mo
ICS dose (mcg/day)	200	640	640	500	500	200
Δ FEV1 L (Δ %)	0.11 (3%)		0.09 (4%)	0.23 (10%)	-	0.23 (10%)
Δ AM PEF L/min	16	18	10	38	25	35
Δ Rescue puffs/d	0.2 (ns)	.6		1.4	-	1.5
Δ % Sx free days	1.9 (ns)		1.9 (ns)	18.4	21	15.4

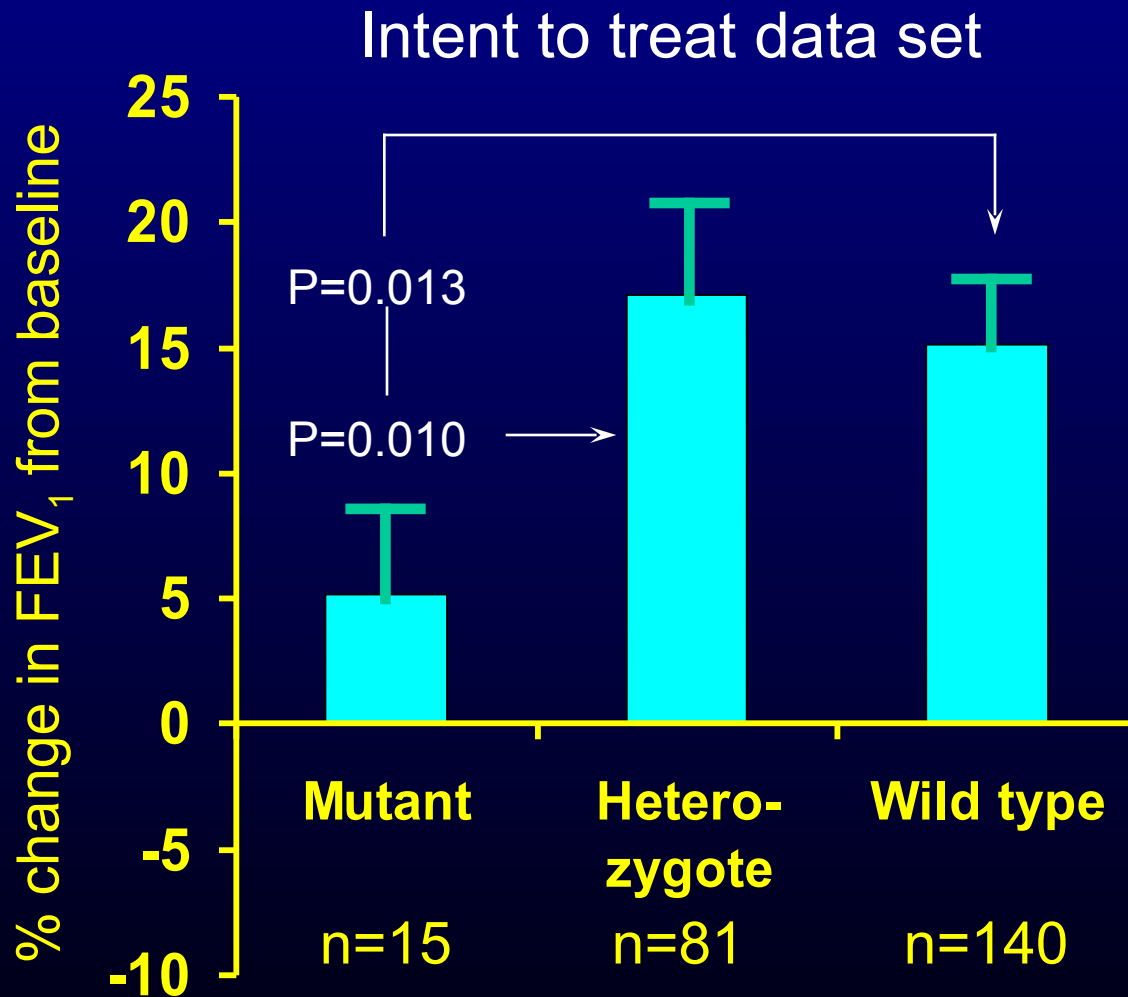
Conclusions

- Arg16Arg polymorphism identifies patients who may not do as well with regular beta-agonists
 - It may identify Blacks who do less well with LABAs
- Blacks may not realize as much benefit from LABAs as Caucasians

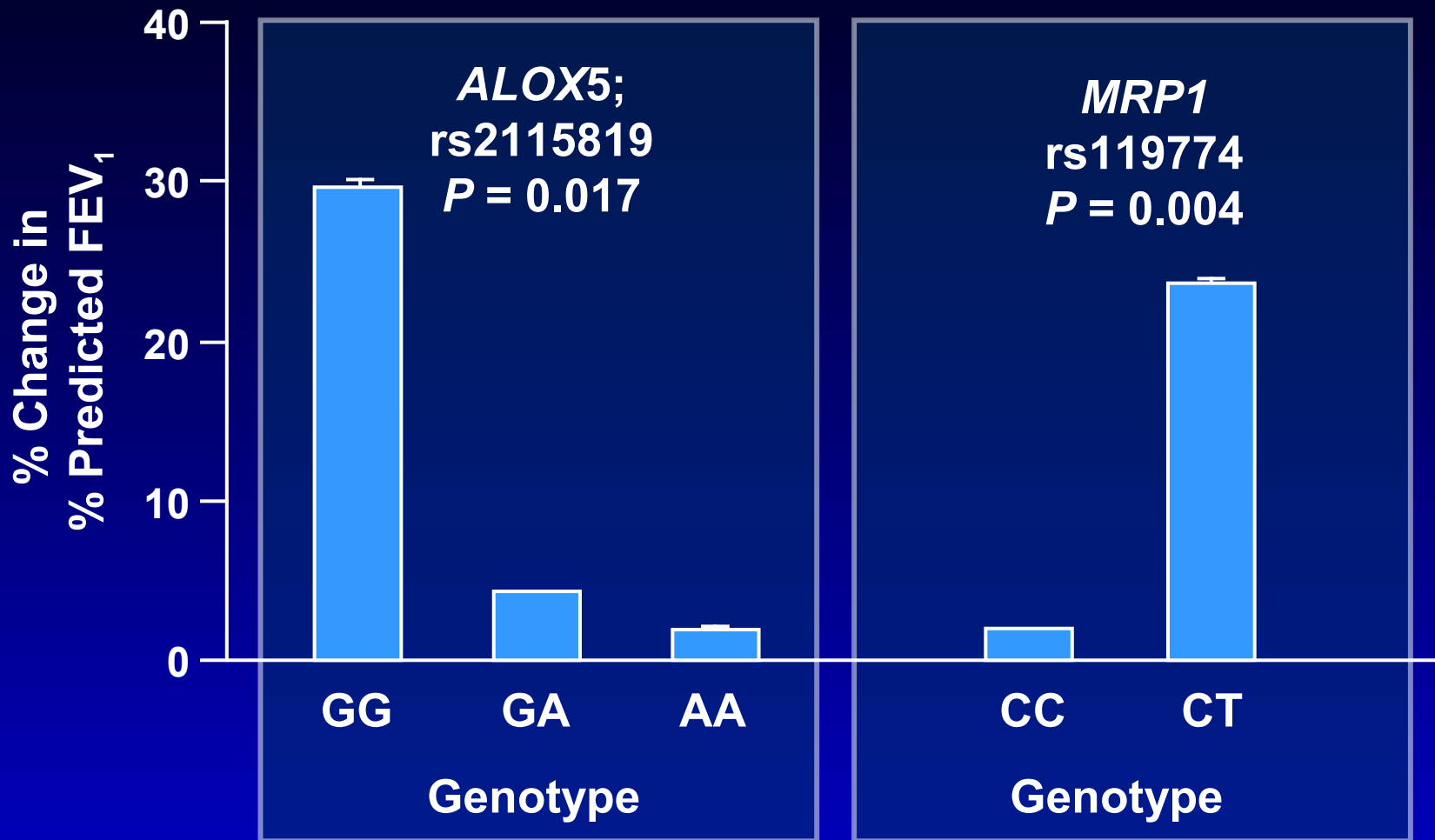
Leukotriene Modifiers



Effect of promoter repeats of ALOX5 on the change in FEV₁ at end of active treatment with ABT-761



Influence of Genotype on Percentage Change in % Predicted FEV₁: other SNPS

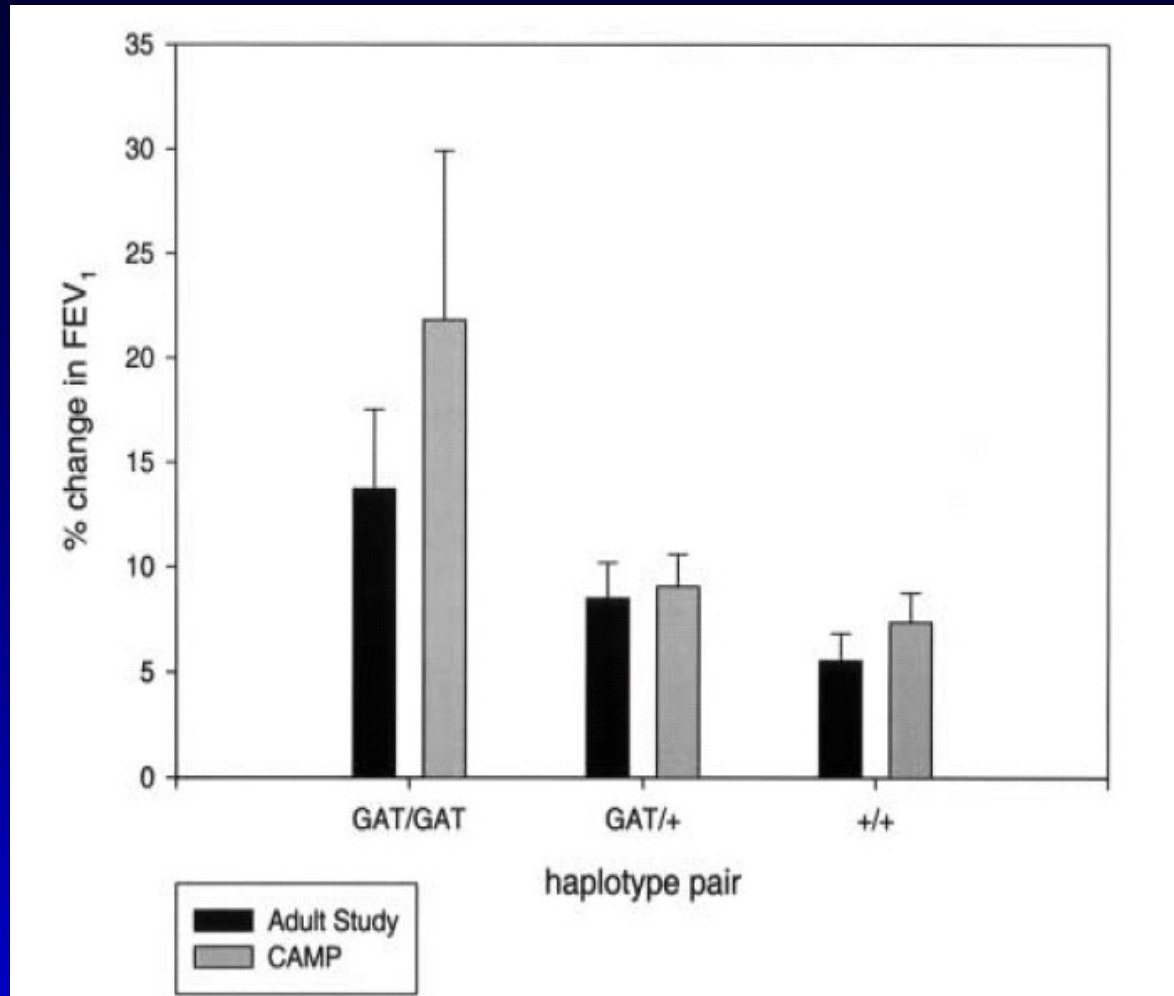


Influence of LT Pathway Polymorphisms on Asthma Exacerbation Risk

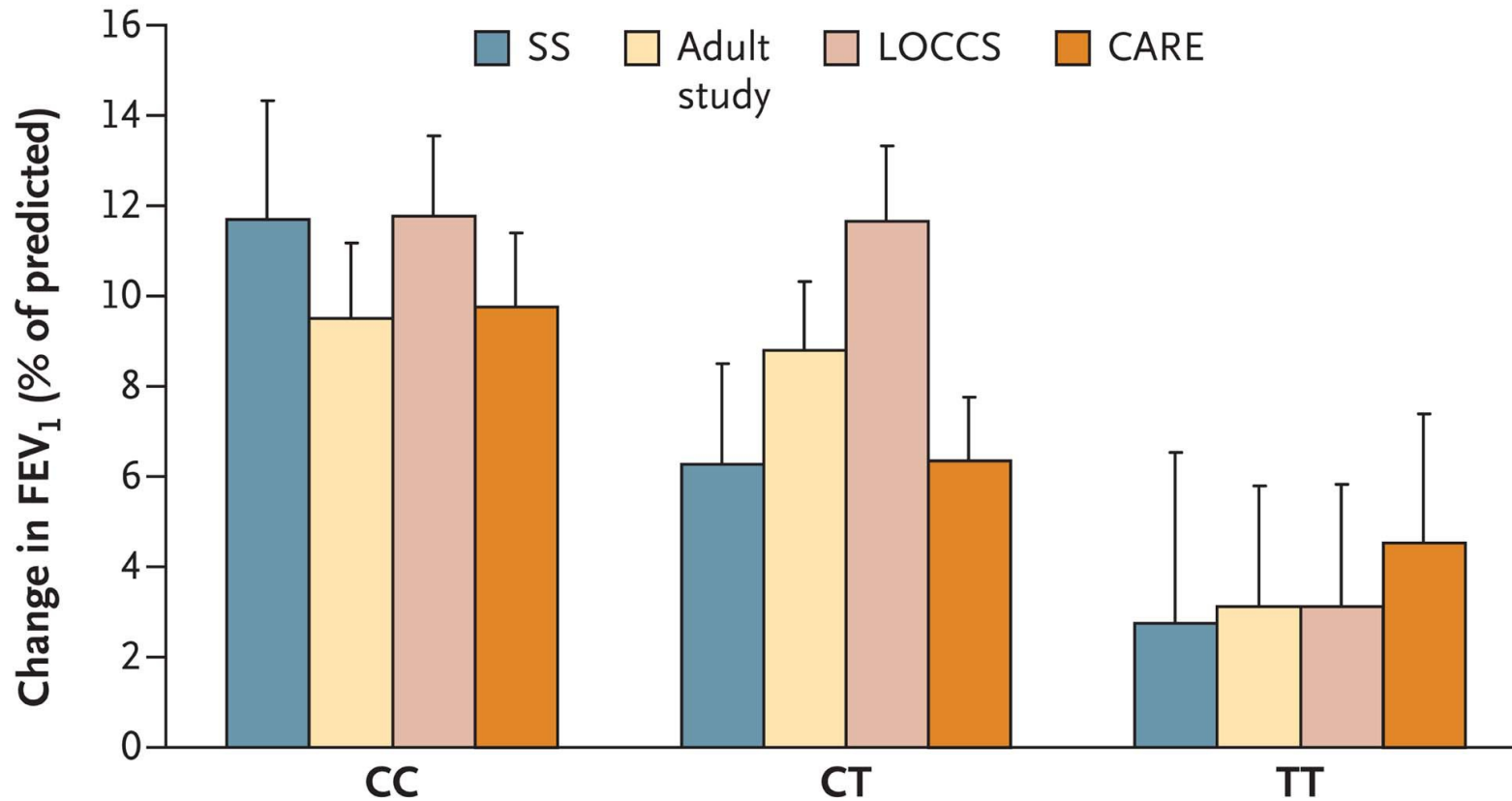
Gene (variant)	Genotype	Frequency (n)	Odds Ratio (95% CI)	P Value
<i>ALOX5</i> (repeat variant)	5/5	0.64 (37)	1.0	0.045
	5/X	0.36 (21)	0.27 (0.08, 0.97)	
<i>LTA4H</i> (rs2660845)	AA	0.49 (28)	1.0	0.021
	AG	0.44 (25)	4.0 (1.23, 12.99)	
	GG	0.08 (5)	4.5 (0.63, 31.95)	
<i>LTC4S</i> (rs730012)	AA	0.51 (30)	1.0	0.023
	AC	0.38 (22)	0.24 (0.07, 0.83)	
	CC	0.11 (6)	0.16 (0.02, 1.49)	

Corticosteroid Responses

Corticotropin Releasing Hormone Receptor-1 Haplotype and Response to ICS



Changes in Lung Function and GLCCI1 Polymorphism (glucocorticoid-induced transcript 1 gene)



Biologics

- No polymorphisms that associate with altered responses
- However there are polymorphisms that associate with the biology
 - FCER1 gene polymorphisms and IgE levels (Palmer, Clin Exp All 1999)
 - IL6R polymorphisms and circulating IL6R (Bleecker, JACI, 2012)

Additional Methods to Discover Candidate Genes

- In vitro expression profiling of cells from responsive and non-responsive populations
 - NFkB - transcription factor activating transcription of cytokines, chemokines, growth factors, cellular ligands, and adhesion molecules associated with asthma (Chapman Mol Cell Endocrinol. 1995)
 - FK506 binding protein 51 gene (FKBP51) in bronchial epithelium (Woodruff PNAS 2007)

Conclusions

- Genetic polymorphisms do associate with differential responses to asthma medications

Potential Applications

- Predict responders
- Predict those with increased tendency for adverse effects
- Individualize dosing
- Allow introduction of medications that have effects in a predictable population

What prevents us from using this information now?

- Repeatable cross-sectional and prospective studies in multiple populations that will put the associations on a firm foundation of data

WHY?

- Gene-gene interactions
 - May modify associations so that they may differ significantly among different populations and ethnic groups
- Gene-environment interactions
 - May result in differences among populations and ethnic groups

SUMMARY

- While we have not yet reached the level of specificity seen in cystic fibrosis, where a drug is targeted to one specific polymorphism, we are beginning to identify patterns of genetic change which will predict responses (or adverse effects) to asthma medications.
- Combining multiple studies in multiple populations with informatics for physicians will allow us to bring this information to the practice setting

