

Pharmacogenomics

Treating the Individual Asthma Patient

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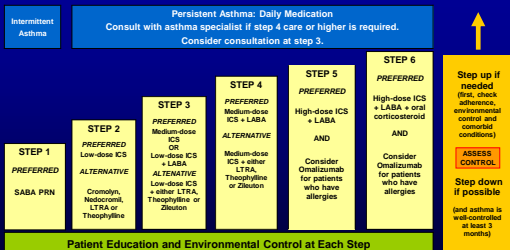
Too much of a good thing?

- 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
- Not working and rarely leaving the house

- On exam: Morbidly obese with scattered wheezes
- FEV1 70% predicted
- 7% eosinophils
- Sinus CT mucosal thickening
- pH probe mild reflux

- 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

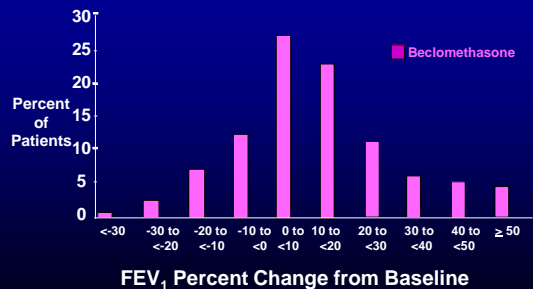


Quick-Relief Medication for All Patients:
 • SABA as needed for symptoms. Intensity of treatment depends on severity of symptoms: up to 3 treatments at 20-minute intervals as needed. Short course of systemic oral corticosteroids may be needed.
 • Caution: Increasing of β -agonist or use >2x/week for symptoms control indicates inadequate control and the need to step up treatment.

Partners' Asthma Center

NHLBI, National Asthma Education and Prevention Program, Expert Panel Report 3, page 517. Available at: <http://www.nhlbi.nih.gov/guidelines/index.htm>, Accessed 2.8.07.

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



Malmstrom et al. Ann Int Med 130: 487-95, 1999

OUTLINE

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

What Polymorphisms Do We Look For?

- SNPs- Single Nucleotide Polymorphisms eg Guanine to Cytosine
- Insertions-additional gene sequences
 - CNVs- Copy Number Variants
 - Change in the number of repeats of a sequence
- Deletions-removal of a nucleotide or nucleotides
- Genomic/Non-Genomic
 - Genomic- in areas of the gene that are know to code for or regulate a gene
- Macro-level – Race

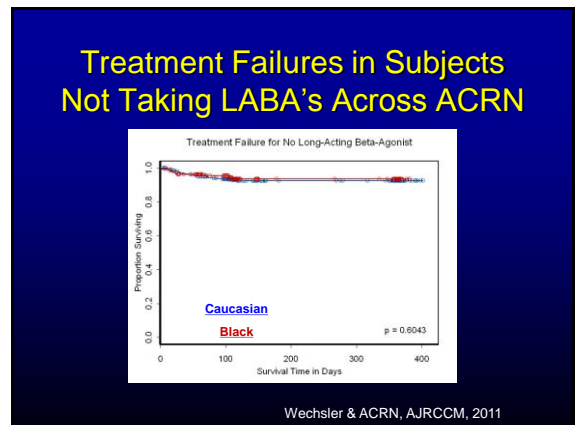
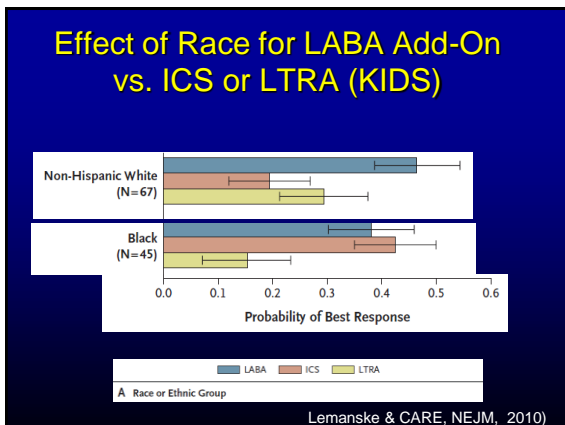
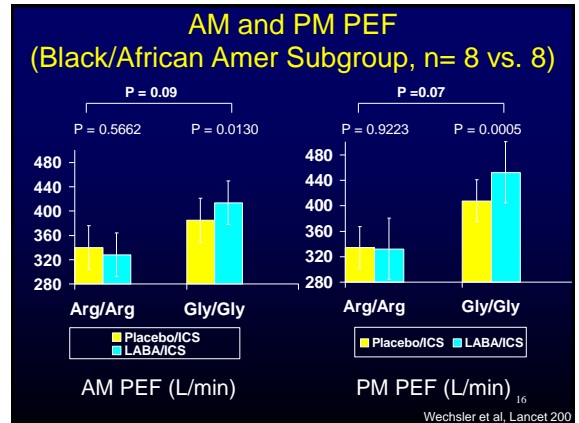
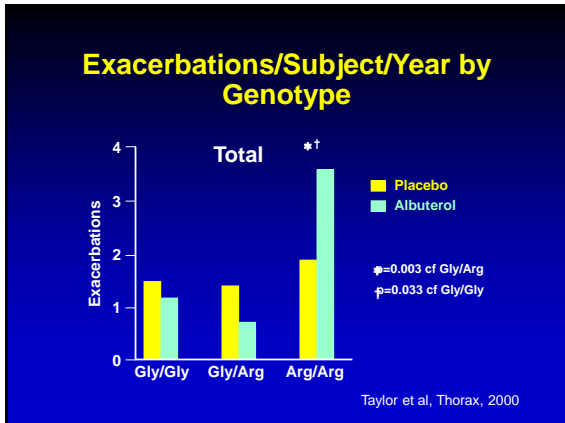
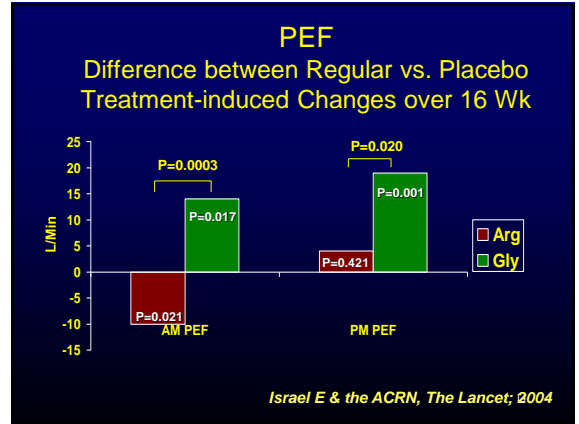
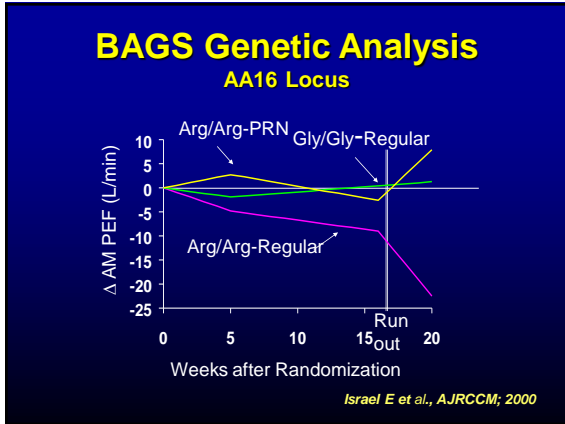
How Do We Look?

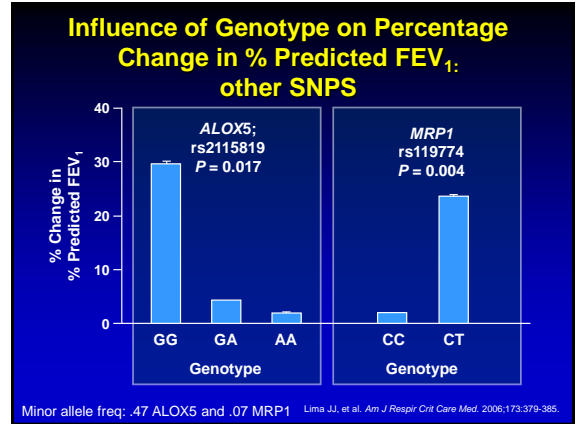
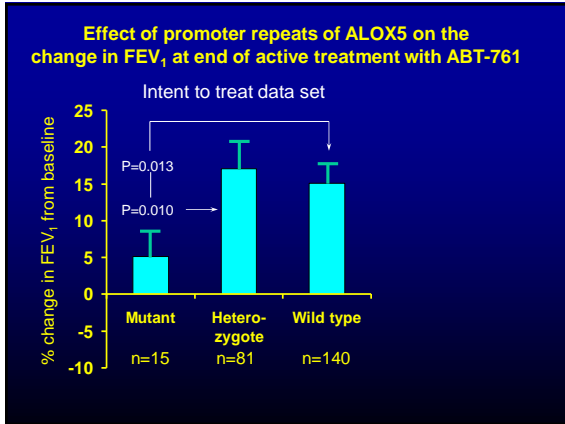
- Candidate Gene and Pathway Approaches
- GWAS
- Expression profiling in cells of responsive and non-responsive individuals

How Do We Confirm?

- Prospective studies
 - Labor-intensive
 - Less subject to confounding
- Association studies
 - Require large numbers
 - Allow easy replication in multiple populations

Beta-Agonists



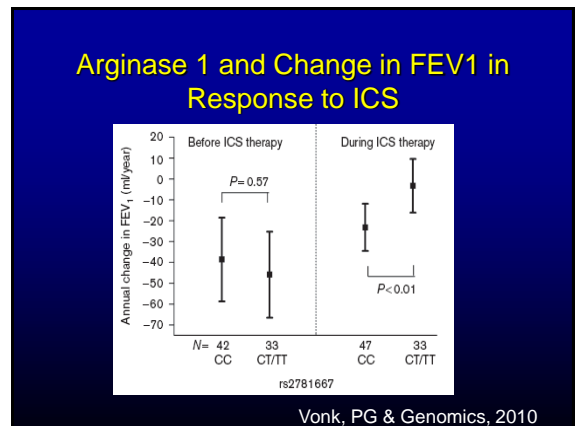
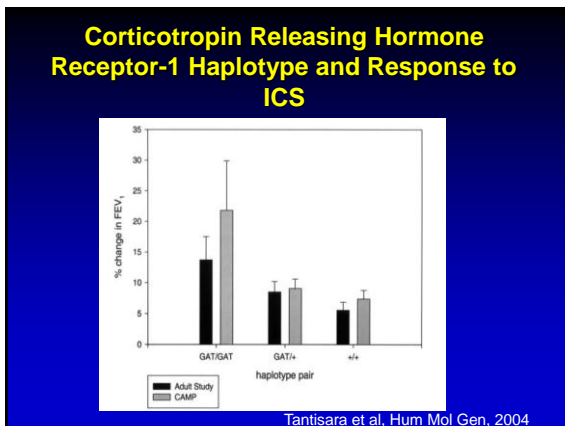


Influence of LT Pathway Polymorphisms on Asthma Exacerbation Risk

Gene (variant)	Genotype	Frequency (n)	Odds Ratio (95% CI)	P Value
ALOX5 (repeat variant)	5/5	0.64 (37)	1.0	0.045
	5/X	0.36 (21)	0.27 (0.08, 0.97)	
	AA	0.49 (28)	1.0	
LTA4H (rs2660845)	AG	0.44 (25)	4.0 (1.23, 12.99)	0.133
	GG	0.08 (5)	4.5 (0.63, 31.95)	
	AA	0.51 (30)	1.0	
LTC4S (rs730012)	AC	0.38 (22)	0.24 (0.07, 0.83)	0.106
	CC	0.11 (6)	0.16 (0.02, 1.49)	
	AA	0.51 (30)	1.0	

Lima JJ, et al. *Am J Respir Crit Care Med*. 2006;173:379-385.

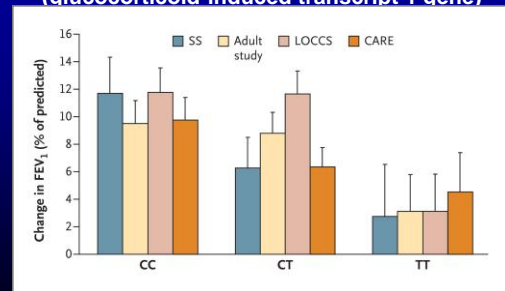
Corticosteroid Responses



Additional Associations with Corticosteroid Responses

- TBX21 - encodes the transcription factor T-bet (induction of Th1 and suppression of Th2)
 - associated with large improvement in airway hyperresponsiveness in response to corticosteroid therapy. (Tantisira PNAS 2004 and Ye J Clin Pharm Ther. 2009)
 - Only 5% frequency
- Adenyl Cyclase 9 (AC9) -activated by ADRB2
 - SNP associated with improved bronchodilator response with ICS (Tantisira, Hum Mol Genet. 2005 and Kim, J Clin Pharm Ther. 2011)

Changes in Lung Function and GLCC11 Polymorphism (glucocorticoid-induced transcript 1 gene)



Tantisira KG et al. N Engl J Med 2011

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