Host Genes and Infectious Diseases

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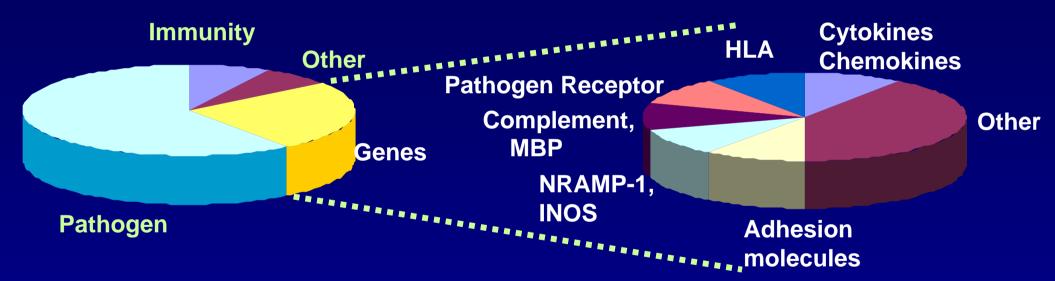
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CDC



A Model of Infectious Diseases

All Risk Factors

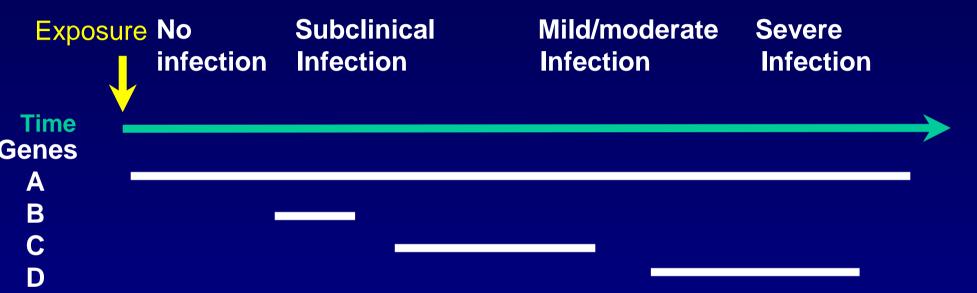
Genetic Risk Factors





Host Genes and Course of Infectious Diseases

Potential Timing of Impact of Host Genes





Identifying Host Genes Associated with Infectious Diseases

- Genetic approach
 - Candidate gene: twins, families, populations
 - Linkage: twins, families, populations

- Issues to consider
 - Race/ethnicity
 - Sample size
 - Laboratory tools
 - Analytic tools
 - Attributable risks
 - Ethical issues



Host Genes and Infectious Diseases

- Pathogenesis
- Resistance
- Severity
- Vaccine design/evaluation
- Drug design/response

- Malaria
- HIV
- TB

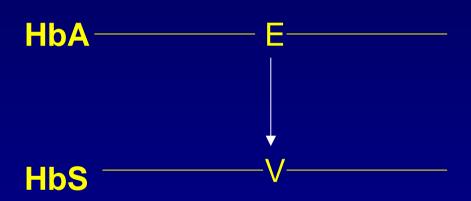
Host Genes and Malaria

- Reduced severity of disease
 - Hemoglobin/RBC proteins (HbS, HbC, α-thall, β-thall, Duffy, G6PD, Erythrocyte Band 3)
 - HLA alleles (B53, DR2)
 - INOS-954 promoter (C allele)
- Increased severity of disease
 - TNF α –308 promoter (A) homozygosity
 - ICAM-1 coding region (T) allele



Hemoglobin S and Malaria

Single nucleotide polymorphism in the β -hemoglobin gene (Chr11)



Under low O₂ tension HbS polymerizes and RBCs assume sickle shape and adhere easily to blood capillaries

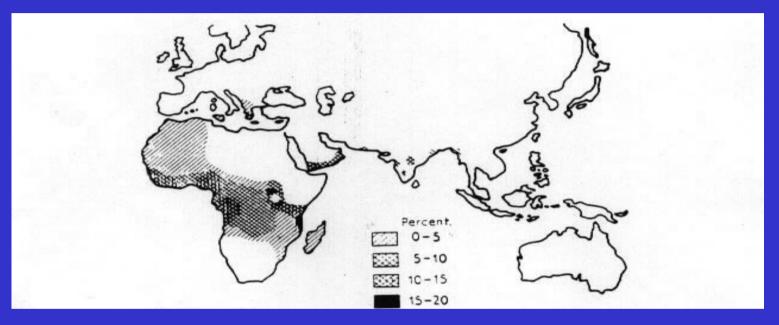


Sickled Red Blood Cell





Geographic Distribution of the Sickle Cell (HbS) Gene



Adapted from Harrison et al.

HbS gene distribution: 15-40% in Sub Saharan Africa

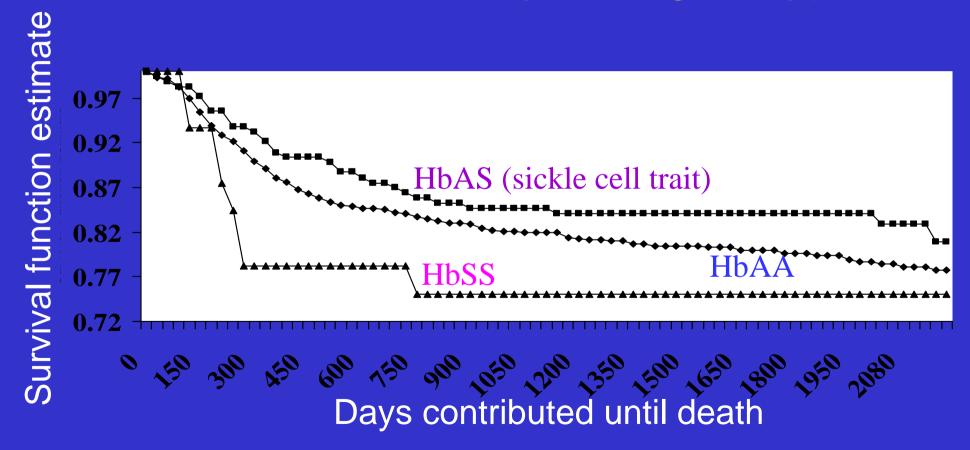


Hemoglobin S and Malaria

- Protective effect of heterozygosity against malaria anemia and severe malaria known for decades
- Mechanism of protection not clear
- Timing of protection unknown



CDC Study, Kisumu, Kenya Survival of Children by HbS genotype





Survival Advantage of HbS is Before the Onset of Immunity to Malaria

Comparison group	Relat. risk	95% CI	p- value
HbAS vs HbAA during 2 - 16 months of age	0.45	0.2-0.8	.0001
HB AS vs HbAA in first 2 months or > 16 months of age	1.2	0.7-2.1	0.5



Summary - Malaria

- CDC Study:
 - Sickle cell trait confers highly significant protection against overall mortality
 - ➤ The time window of protection is between 2 to 16 months of age
- Malaria and Host Genetics
 - > Complex, multifactorial, multigenic
 - > Implications for pathogenesis, vaccine design

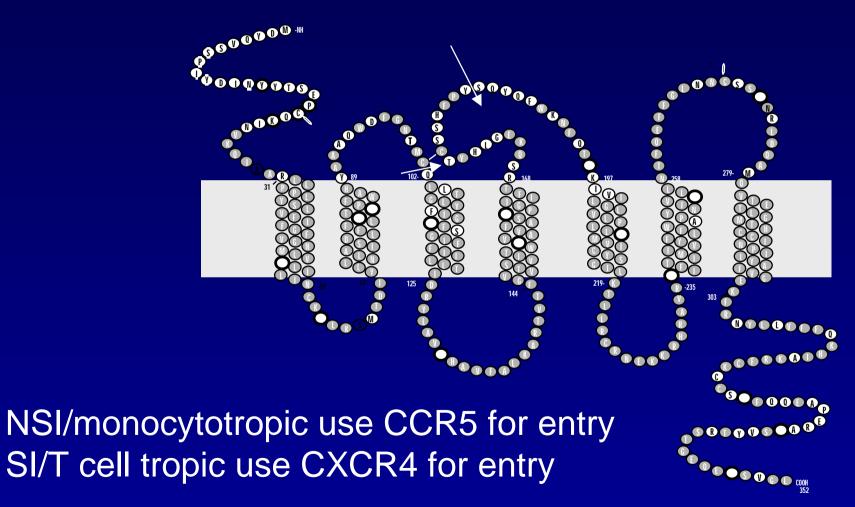


HIV

- Host Genes Impact Multiple Aspects of Disease
 - Resistance: HLA, CCR5
 - Disease progression: HLA, CCR, SDF, MBP, Vit DR, IL-10, TNF
 - Vaccine responses: HLA
 - Drug responses: HLA, MDR1



Chemokine Receptor (CCR5) used by HIV

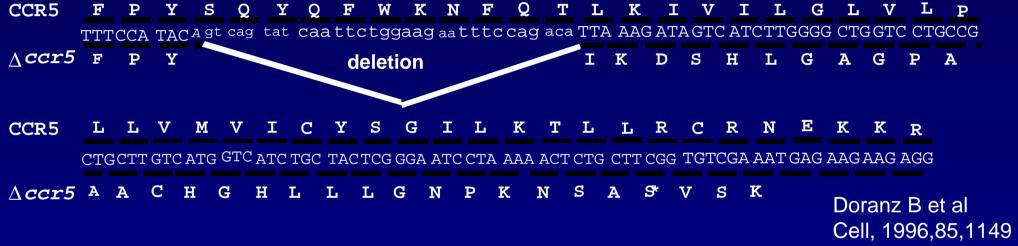


Cell Membrane

Rucker et al, Cell, 1996, 87:437



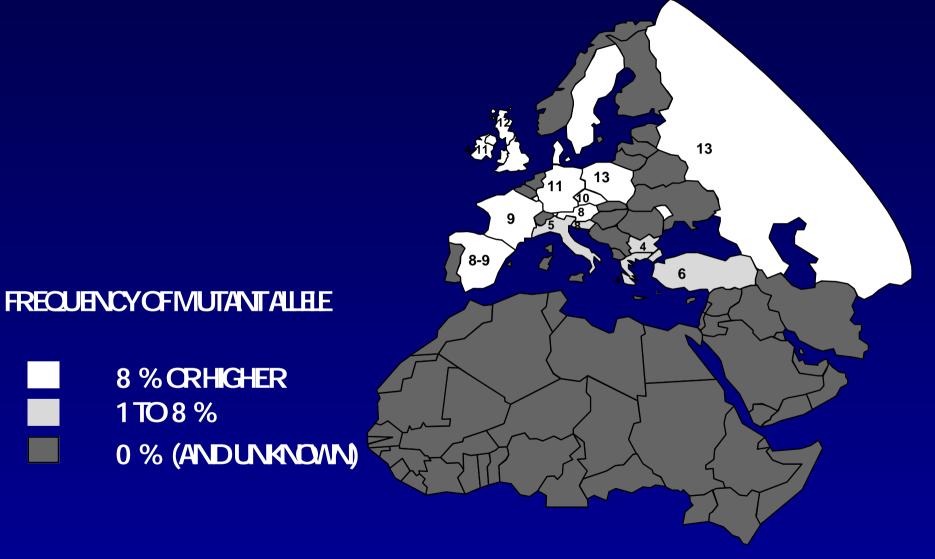
CCR5 and the 32 bp Deletion



CCR5 △ 32 Genotype

10% Caucasians are heterozygous, 1% homozygous Complete resistance to HIV infection (with NSI variants) Presence delays progression to AIDS Attractive therapeutic target

Geographic Variation in Δ 32 CCR5 Distribution

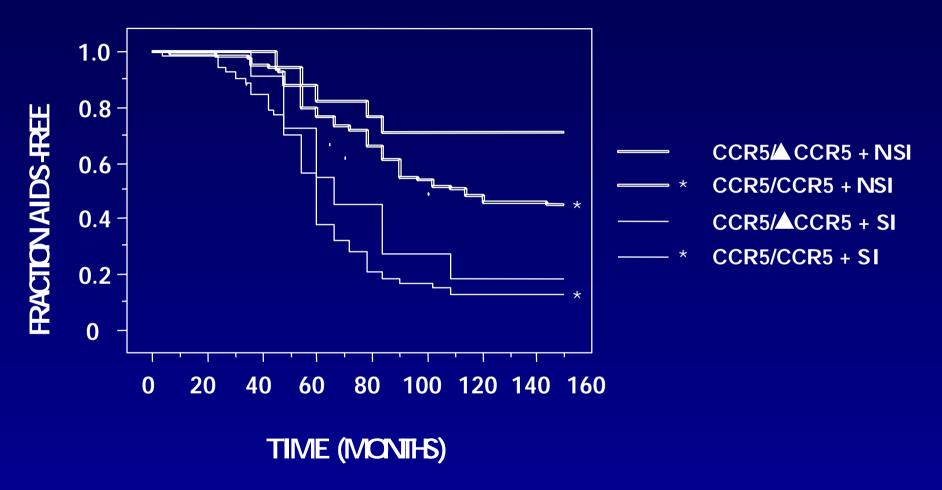




1TO8%



Effect of \triangle 32 CCR5 Genotype on HIV Disease Progression





HLA Genotype influences Multiple Aspects of HIV Infection and Disease

- Resistance to infection (HLA-A2 supertype, HLA-A11, HLA discordance)
- Rate of progression to AIDS
 - Delayed: HLA-B27, B57
 - Accelerated: HLA-A1,B8,DR3
- Vaccine responses
 - Enhanced (canarypox HIV vaccine: HLA-B27)
- Drug reactivity (abacavir: HLA-B57)



HLA type and Responses to a Candidate HIV Vaccine

Vaccine Protein	HLA-B27 +	HLA-B27-	OR
Gag	64	15	10.3
Env	36	11	4.6
Any	40	11	5.4

% CTL responses observed at 6 weeks post vaccination with ALVAC-HIV Canarypox vaccines. Kaslow, R et al, JV, 2001, 75; 8681



HLA Genotype Determines Hypersensitivity to Anti Retroviral Therapy

HLA type	Abacavir Hypersensitive	Abacavir Tolerant	OR
HLA-B57+	78	2	117
B57 Haplotype	72	0	822

% subjects in each group. Mallal S et al, Lancet, 2002, 359;9308 Abacavir = Ziagen (Trizivir when combined with AZT/3TC) Haplotype = HLA B57-DR7-DQ3

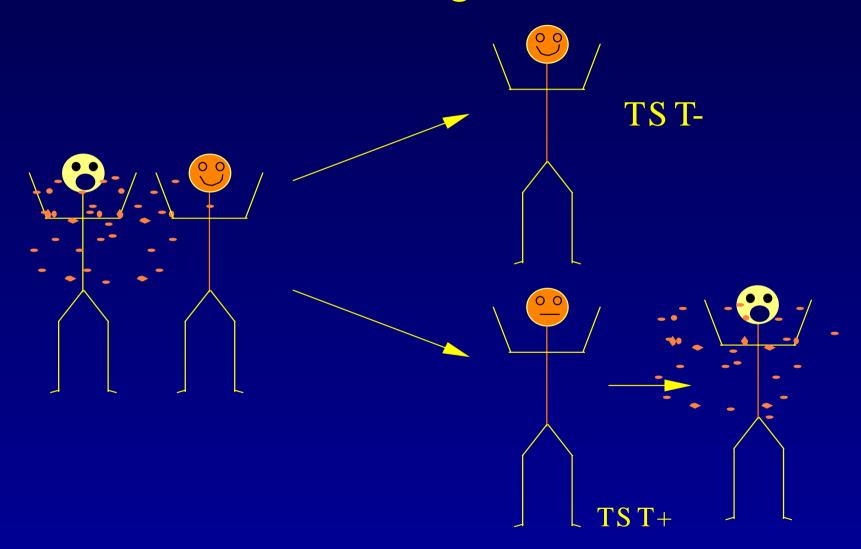


Host Genes and HIV

- Multigenic, complex, multiple aspects
- Implications for therapy and vaccine design

Host Genes and Mycobacterial Infections

TB Pathogenesis





Genes associated with TB

- Immune Genes
 - HLA class DR and DQ (6p21)
 - IL-1RA (2q14)
 - $IL-1\alpha$ (2q14)
 - MBL (10q11)

- Other Genes/loci
 - Vit DR (12q12)
 - NRAMP-1 (2q35)
 - X chromosome
 - Chromosome 15 locus

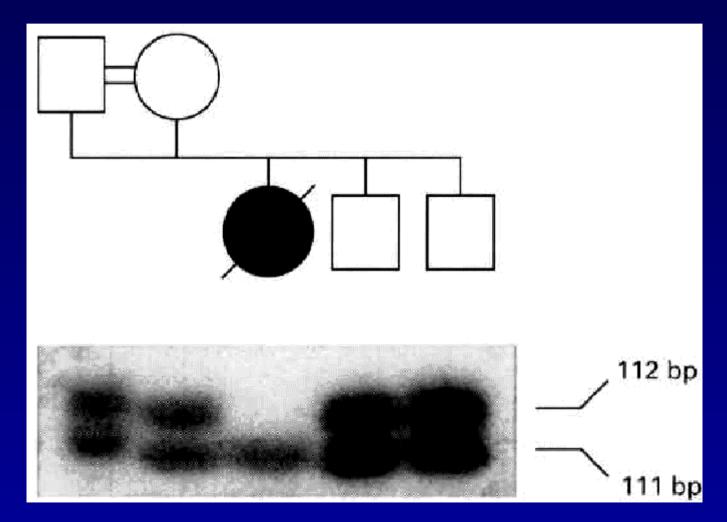


Importance of Family Studies

- IFN-γ R-1 (6q23)
 - Recurrent M. avium and other non-TB mycobacteria
 Newport et al, NEJM, 1996
 - Disseminated BCG infection Jouanguy et al, NEJM, 1996
- IL-12 Rβ (5q31)
 - Recurrent atypical mycobacterial infections
 De Jong et al, Science 1998



IFN-γ Receptor Deficiency in an Infant with Fatal BCG Infection





Future Directions Host Genes and Infectious Diseases in Public Health

Needs:

- define attributable risks
- define interaction of risk genotypes with other risk factors
- identify other genes

Long term

- identify populations at high risk for diseases
- Understand pathogenesis
- Develop new prevention or therapeutic strategies



The Future....

- New treatments based on genes associated with resistance/susceptibility
- Host genetic typing in the clinic?
- Host genetic typing in epidemics?
- Host genetic typing of populations for prevention or treatment?

CDC's genetics web site: www.cdc.gov/genetics

