Sickle Cell Disease: A Look to the Future

David G. Nathan, M.D. San Diego, California 12/08/2011

Industrial Relationships

- Scientific Advisory Board: Aileron, Inc
- Consultant: NK Therapeutics, Inc







FIGURE 2 Photomicrograph of Hb SS crythrocytes stained for Hb F by the Kleihauer-Betke technique

The order of matter states are the states of the state of the local value of the states of the state of the states of the state

except that Mas CTO, approximately 30 #6/100 mi of	•	•		
ood, was used as an isotope.	Pa.	Inteversibly	Nonirreversibly	
RESULTS	tient	nickted ceils	sickled cells	· *
Clinical data and champetians. Bouting he	1	0.034 (±0.019)*	$0.068 (\pm 0.047)$	< 0.0
Canical add and observations. Routine ne-	2	$0.033 (\pm 0.013)$	0.071 (±0.036)	<0.0
natological data on the patients used in these	3	$0.034 (\pm 0.018)$	$0.091 (\pm 0.071)$	< 0.0

studies are shown in Table I. In no patient did	3	0.034 (土0.018) 0.026 (土0.019)	$0.091 (\pm 0.071)$ $0.058 (\pm 0.035)$	<0.01 <0.01
present or past symptomatology correlate with total Hb level or per cent Hb F. Scattergrams	Mean	0.032 (±0.004)	0.072 (±0.014)	<0.01

rotal Hb level or per cent Hb F. Scattergrams were constructed of Hb level vs. per cent Hb P, • Mean (±ss) of 50 cells.

1734 J. F. Bertles and P. F. A. Milner

matological data on the patients





FIGURE 2 Distribution of RBC on subconfluent culture plates stained with Giemsa after the fifth plate wash. Sickle RBC distribute themselves in rosette-like clusters around endothelial cells (A), whereas normal RBC are present in fewer numbers and are randomly distributed (B). $\times 375$.

Hebbel et al J. Clin. Invest. 65, 1980 157



Dana-Farber Cancer Institute, Boston *David Nathan *Donna Neuberg Medical College of Wisconsin, Milwaukee *Joshua Field Brigham and Women's, Boston Maureen Okam Children's Hospital, Boston Matthew Heeney, David Nathan Washington University, St. Louis **Elaine Majerus** Howard University, DC **Onyinye Onyekwere** Johns Hopkins University, Baltimore **Jeffrey Keefer**

*Phase I Core group La Jolla Institute of Allergy & Immunology *Joel Linden

David Nathan, Principal Investigator (clinical)
Joel Linden, Principal Investigator (basic science)
Donna Neuberg, Co-Investigator
Joshua Field, Co-Investigator

Hypoxia/Reoxygenation: A Mouse Model of ACS

NY1DD-Vehicle-H/R NY1DD-ATL146e-H/R



3 h hypoxia (8% O_2) 18h reperfusion (air) A2A agonist started 3 h after reperfusion

Human Blood







Hours since infusion start

Subject 9: NFkB during infusion



Mean fluorescence intensity

NF_KB+ Cells In Activated Gate: Dose Level 2



Hours

Proposed Future Studies

- Pilot Study of Regadenoson for the Treatment of Acute Chest Syndrome – Primary endpoint: Accrual
- Phase II Trial of Regadenoson for the Treatment of Acute Vaso-occlusive Episodes
 - Primary endpoint: Reduction in inflammatory biomarkers
- Anti-NKT cell monoclonal antibody

Acute and Chronic VOC •Pain and ACS leading to hospitalization are the tips of the iceberg.

 Thirty percent of adults with SCD report pain on a daily basis.
 Smith et al

2008



iNKT Cell Depletion by NKTT120 (Standard IgG1)



CONFIDENTIAL

DNA Sequence Polymorphism

- Common SNPs may be biologically trivial or evolutionarily constrained and thereby important in some way.
- Rare variants associated with mendelian disorders are much more apt to be important so full DNA sequencing should be useful (up to a point)
- But "guilt by association" often leads to a dangerous conclusion (SNPs around CRP show big effects in heart disease but obviously have no causal relationship)

Exceptions Prove the Rule

- BCI11A discovered in a common SNP search related to fetal hemoglobin and has a big effect size (ten percent)
- Three other SNPs related to fetal hemoglobin bring the total effect size to near fifty percent
- Alpha thalassemia is also clearly associated with decreased severity
- Yet published and unpublished GWAS that evaluate severity do not reveal HbF associated SNPs or SNPs that might relate to alpha globin synthesis. Faint association with TGF beta suggesting that inflammation might be important

WILL GENOME SEQUENCING REVEAL THERAPEUTIC LEADS IN SCD?

- "Unbiased" research=fishing expedition
- Hypothesis should precede not follow technology application in science
- "Guilt by association" is dangerous in medicine as well as law
- Poorly crafted GWAS are worse than no studies at all
- Best to compare lowest five percent and highest five percent for each severity category

SCD ATTACK POINTS

- Increase delay time with a small molecule without changing O2 affinity (cyanate which does change affinity and is neurotoxic) or a stapled peptide that would compete with Val 6
- Increase or introduce a hemoglobin (HbF or Hb Korle Bu) with hydrophobic binding to val 6 in HbS, prolongs delay time and inhibits stacking of polymers by forming hybrids (hydroxyurea and/or BCL11A inhibition)
- Hematopoietic stem cell transplant
- Gene therapy by excision and replacement by a non S allele
- Suppression of inflammation and ischemia reperfusion injury
- Prenatal diagnosis
- Control asthma