International SAE Consortium, Ltd

An International Industrial Biomedical Consortium Researching the Genetic Basis of Drug Related Serious Adverse Events

Brookings / FDA event on Sentinel Initiative

January 11, 2009



SAEC's Mission

"The SAEC will identify and validate DNAvariants useful in predicting the risk of drug induced serious adverse events."





Key iSAEC Phase 1 Objectives

- Coordinate <u>international</u> network[s] for obtaining well phenotyped <u>cases</u> and <u>controls</u> for SAE PGx research [SSR and DILI]
- Using GWAS, explore cross drug/within drug DNA-variants useful in predicting the risk of SSR and DILI
- Create a <u>publicly available</u> "knowledge base" of cross drug safety PGx markers for predicting key SAEs
- Manage IP relating to PGx markers useful in predicting SAEs to ensure broad and open access
- Develop a <u>cross-disciplinary forums</u> to address clinical and scientific issues related to PGx of SAEs [e.g. Phenotype Standardization Program]
- Support the execution of the FDA Critical Path and related international safety PGx regulatory efforts





SAEC Web Site

http://www.saeconsortium.org



The Serieus Adverse Event Consortium (SAEC) is a nonprofit organization comprised of leading pharmacentical comparies, and academic institutions with scientific and situategic input from the U.S. Food and Drug Administration (FDA). The mission of the SAEC is to help identify and widdate DRA-walants useful in predicting the thild of ungerelated versions adverse events (CAE).

Patients respond differently to medicines and all medicines can have side effects in some people. The SAEC's work is based on the hypothesis that these differences have a genetic basis, and its research thatless will examine the impact genes can have individuals respond to medicines. The SAEC's initial studies are focused on identifying the genetic markers associated with drug-related fiver taskicly (the leading cause of acute time failure) and Steven Johnsons Syndrome SLS, asserts from disk mecrolis). All research results are made available publicly within 12 months of the completion of the study group's genotyping. The SAEC has developed a data gorial to provide the research community with free and unencumbered access to study data. Results obtained in initial studies can thus be reevaluated by researchers who can determine their validity as predictive markers.

Examples of severe adverse drug reactions

A number of severe adverse drug reactions are known. These include the conditions which are the initial focus of the Consortium's efforts:

- Serious Skin Rashes: Stevens-Johnson Syndrom (SJS) and Toxic Epidermal Necrosis (TEN) - related, rare, severe, mucocutaneous bistering disorders that are associated with over 200 medicines.
- <u>Drug Induced Liver Injury</u> (DILI) Hepatotoxicity caused by more than 30 different drugs in more than seven different classes, including NSAIDS, various antibiotics, analgesics and anti-discussion.



SAE Consortium SUIIDAY, FEBRUARY 24, 2008					
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Letter from the Chairman

November 2007

Dear Colleagues:

Welcome to the website of the International Serious Adverse Event Consortium [SAEC]. We're delighted you chose to spend a few minutes with us.

We launched the SAEC in August of 2007, as an industrial biomedical consortium, focused on identifying and validating DNAvariants useful in predicting the risk of drug induced, rare serious adverse events [SAEs]. Drug-induced, rare SAEs can be a





Phase 1 SSR Discovery Project - June` 09





Pharmacos Sourced SAE Cohorts



Sourcing SAE cases via Providers & EMRs Phase 1 Feasibility Project

- 2009-10 Feasibility Projects
- Focus: Using EMR and associated research systems to determine the feasibility of yielding high quality SAE cases.
- SAE targets/3/collaboration [of joint interest]
 - Cerner → Hepatotoxicity, TdP/PQT, and SSR
 - HMORN → Hepatotoxicity, EWG, and SSR
 - VA → Hepatotoxicity and Rhabdo/ Myopathy
- Standardized phenotype translated into EMR data ontology – feasibility of (retrospective) case detection in CDW
- Determine how many potential cases and key clinical data gaps



SAE Consortium

HMORN Sourced SAE Cohorts (Pilot)



SAE PSP Project Phase 1 - EMR ID of SAE Cases

- ADR research challenges → lack of phenotype standards and "coding" into EMR data ontologies
- PSP 1 Focus: Immunologic Related SAEs (SSR, DILI & AHHS) and TdP/PQT
- 2009-10 Project → Munir Pirmohamed, Chairman; Julian Arbuckle, Project Consultant
- EWG Steering Committee → Munir Pirmohamed (Liverpool), Ann Daly (Newcastle), Paul Watkins (UNC), Guru Aithal (Nottingham), Dan Roden (Nashville) & Dr. Elijah Behr (London)
- Organizing committee → Arthur Holden (SAEC), Michael Dunn (WT), Munir Pirmohamed (Clinical Chairman), and ShaAvhree Buckman (FDA)
- Expert Working Committees (3) → AHHS/SSR, DILI, and TdP/PQT
- Q4`09 Member CRF inputs solicited; representative involvement in 2010 consensus conference
- Pooled funding → SAEC project management and working group meetings, WT 2010 consensus conference, and FDA scientific writing consultants for proceedings write-up



Key Points from SAEC's Phase 1

- 1. Genomic technology and the scope of the SAEC's collaborative networks are helping to enable basic genomic research on drug induced SAEs ... a significant public health and drug industry/FDA challenge ... faster and more efficiently.
- 2. Robust statistical inference is possible from relatively small SAE cohorts [e.g. DILI research] and larger, well constructed control cohorts
- DNA variants may be useful in predicting and mitigating SAE risk for some drugs in some patients → certainly supportive of safety PGx studies as part of clinical development or post market surveillance studies
- 4. There are important genetic effects at root of many SAEs, but many of these effects will <u>likely</u> be drug and ethnicity specific, and vary in terms of their predictability and clinical utility
- 5. At the same time, we are finding genetic risk alleles [e.g. HLA B*5701] that predispose certain individuals to drug induced SAEs [e.g. SSR, Acute hypersensitivity reaction, or DILI] across multiple drugs.

SAE Consortium

SAEC Evolutionary Perspective



EMR based drug safety PGx science

- "Standardization" of phenotype
- "Functional/ compatible" ontology/ies
- Drug and ethnic diversity
- High quality phenotyping \rightarrow real time
- Adequate case & control numbers (e.g. 50 cases X drug X ethnicity/3X control)
- Supporting genomics skills and infrastructure
- Committed partners, with "results orientation"



The SAEC Has Established an Ideal Collaborative Framework for SAE Research 1

- ✓ Global orientation to deal with ethnic diversity
- ✓ Driving standardized phenotyping across SAE/ADRs
- ☑ Strong research design skills via Scientific Management Committee
- ✓ Strong network development skills → adequate size cohorts & optimal controls [*from academia, providers and companies*]
- ✓ Strong genomics "core"
- Excellent Data Analysis & Coordinating Center [@ Columbia] with state-of-the-art analysis pipeline and public website for data release
- ✓ Efficient IP pipeline to ensure all discovered markers are placed unencumbered into the public domain
- ✓ Strong, focused management
- \blacksquare Strong integration \rightarrow industry, academia, providers & regulators



1. Giacomini et at, *Nature*, Volume 446,4/07.