Growing National Concern About Adverse Drug Reactions



digmostra

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- IOM, Committee of Health Care in America Report: "To Err is Human: Building a Safer Health System", Decmber 1, 1999.
- GAO, Report to Congressional Requestors. "Adverse Drug Events", January, 2000.
- FDA/CDER Spontaneous Reporting System (SRS) contains adverse drug reaction reports. Expanded Adverse Events Reporting System (AERS) under development.

Introduction



Diagnostics

- Human Genome Project is accelerating genetic studies
- Human Genome Project has been accelerated
- Surrogate markers are expected to improve treatment efficacy
- Adverse reaction markers critical to understand toxicity
- Need for high information content diagnostic assays to address unmet clinical needs is increasing
- Examples of high information content assays include variant analysis, resequencing and expression (mRNA) profiling

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Challenges for Genetic and Cancer Diagnostic Testing



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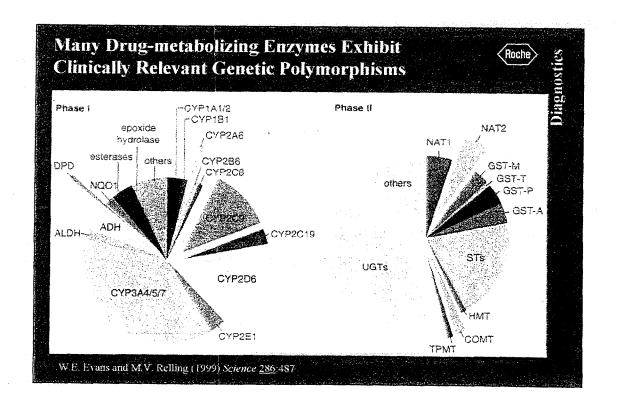
Scientific: Correlation of disease phenotypes and treatment efficacy with pathogens and genes need to be established

Technological: Diagnostic assay formats and platforms that permit reliable, cost-effective testing with significant predictive value with high throughput and automation need to be developed

Clinical: Indications and defined utility need to be demonstrated

Regulatory: Timely approval of medicines and assays

Psychosocial: Acceptance of genetic testing





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CYP2D6 Substrates

Antiarrythmics

Propafenone, Encainide

Beta-blockers

Timolol, Metoprolol

Antidepressants

Desimipramine, Clomipramine, Imipramine, Amytriptyline, Fluoxetine

Neuroleptics

Haloperidol, Thioridazine. Risperidone, Venlafaxine

Miscellaneous

Codeine, Debrisoquine, Phenformin, Indoramin

CYP2C19 substrates

Omeprazole
Amytriptyline (in part)
Imipramine
Certain barbiturates
Chlorproguanil
Proguanil
Citalopram
Diazepam
Mephenytoin
Phenytoin
Propranolol
Cyclophosphamide

Impact of Human CYP450 Polymorphisms on Drug Treatment in Poor Metabolizers

Polymorphic enzyme	Decreased clearance	Adverse effects	Reduced prodrug activation
CYP2C9	S-Warfarin Phenytoin Losartan	Bleeding Ataxia	Losartan
*	Tolbutamide NSAIDs	Hypoglycaemia Gl-bleeding (?)	
CYP2C19	Omeprazole Diazepam	Sedation	Proguanil
CYP2D6	Tricyclic antidepressants	Cardiotoxicity	Tramadol Codeine Ethylmorphine
	Haloperidol Anti-amhythmic drugs Perphenazine	Parkinsonism (?) Arrhythmias	
	Perhexiline SSRIs Zuclopenthixol S-Mianserin Tolteradine	Neuropathy Nausea	

Abbreviations: NSAIDs, nonsteroidal anti-inflammatory drugs; SSRIs, selective serotonin reuptake inhibitors.



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M. Ingelman-Sundberg, et al TiPS 20: 342, Aug, 1999

GeneChip CYP450 Microarray: A combination of controls assures validity For every block, "Column 11" control probes 1) Hybridization Positive control probe 2) Hybridization Negative control probe 3) Exon sense strand control probe 4) Exon antisense strand control probe

Issues Regarding Regulation of High Density Genetic Variation Diagnostics



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- What regulatory path is most appropriate for these new technologies?
 - CBER classification of HIV Resistance assays
 - Class II (Special Controls)
 - · Guidance Document
 - Regulation of Genetic Tests
 - · Classification based on Intended Use?



Diagnostic

- Possible Intended Uses, Impact on Regulatory Requirements
 - Identification or measurement of a physical or biochemical parameter
 - Well-characterized mutations, phenotype predictions
 - Identification of the clinical use of the measurement
 - Treatment decisions based on antiviral resistance, drug metabolic profiles
 - Identification of or implication of an effect on clinical outcome
 - · Treatment outcome

http://www.fda.gov/edrh/modact/genspec.html

Analytical Performance of Human Genotyping Assays: Validation Samples



mostice

- Establish reference panels of genomic DNA samples representative of common allelic variants, genotyped by independent method(s)
- When possible, obtain one or more genomic DNA samples representative of rare allelic variants (<1% allelic frequency)
- When samples carrying rare variants are available only in extremely limited quantity, clone by PCR to generate renewable resource
- When samples carrying rare allelic variants unavailable, generate relevant mutations in plasmids