Presented
by
Frederico Goodsaid, Ph.D
at the
Nonclinical Studies Subcommittee
of the
Advisory Committee for Pharmaceutical Science

March 9, 2000

TaqMan Gene Expression Arrays for Accurate Quantification of **Toxicity Targets**



PE Biosystems

Test Procedure

- RNA isolation
- · cDNA generation (may be skipped with one-step RT-PCR)
- · Card loading and sealing
- Thermal cycling
- Data Analysis





Agenda

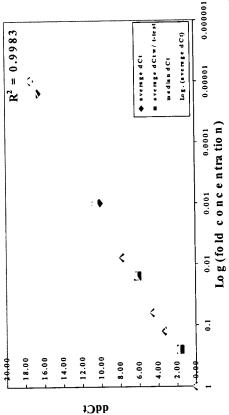
- Test Procedure
- Specifications

Validation Results

PE Biosystems



Dynamic Range in the Cytokine Card



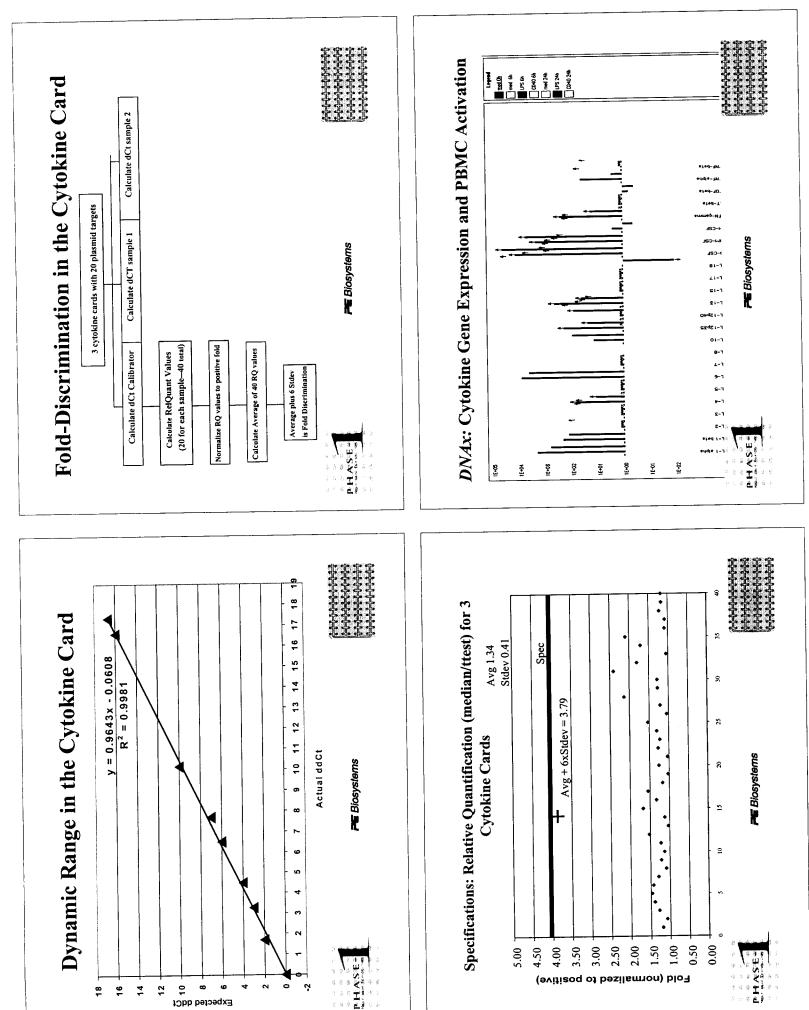








FE Biosystems



Expected ddCt

Source Precision Medicine: Cytokine Gene Expression and PBMC Activation \$ E+03 <u>E</u>+02 Ē ş

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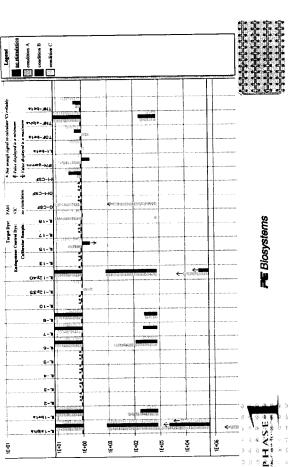
Phase 1 Molecular Toxicology: Cytokine Gene

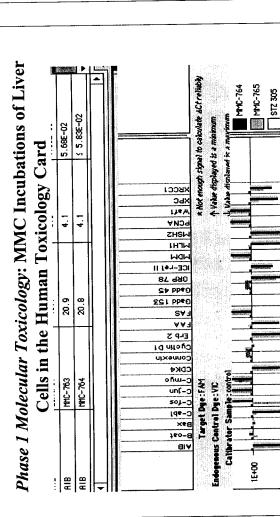
Expression in Liver Cell Cultures

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PE Biosystems

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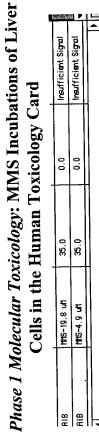
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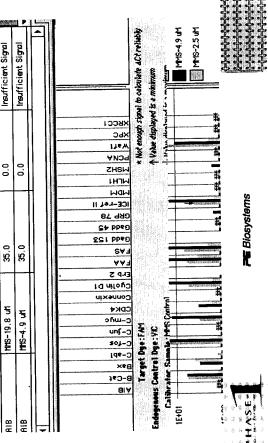
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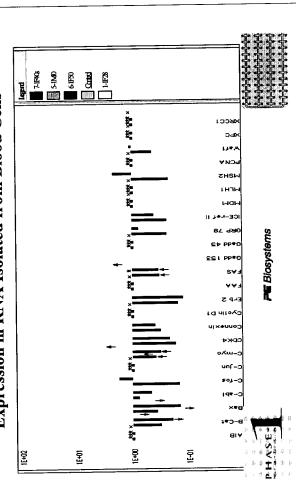
PE Biosystems

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Phase I Molecular Toxicology: Carnustine Incubations of Liver Cells in the Human Toxicology Card

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Phase I Molecular Toxicology: Toxicology Gene Expression in RNA Isolated from Blood Cells



Phase 1 Molecular Toxicology: Cytokine Gene Expression in RNA Isolated from Blood Cells

