



Ribonomics, Inc.

Bill Phelps, R&D Director

Genome Tri-Conference 2002: Santa Clara, CA

February 24, 2002

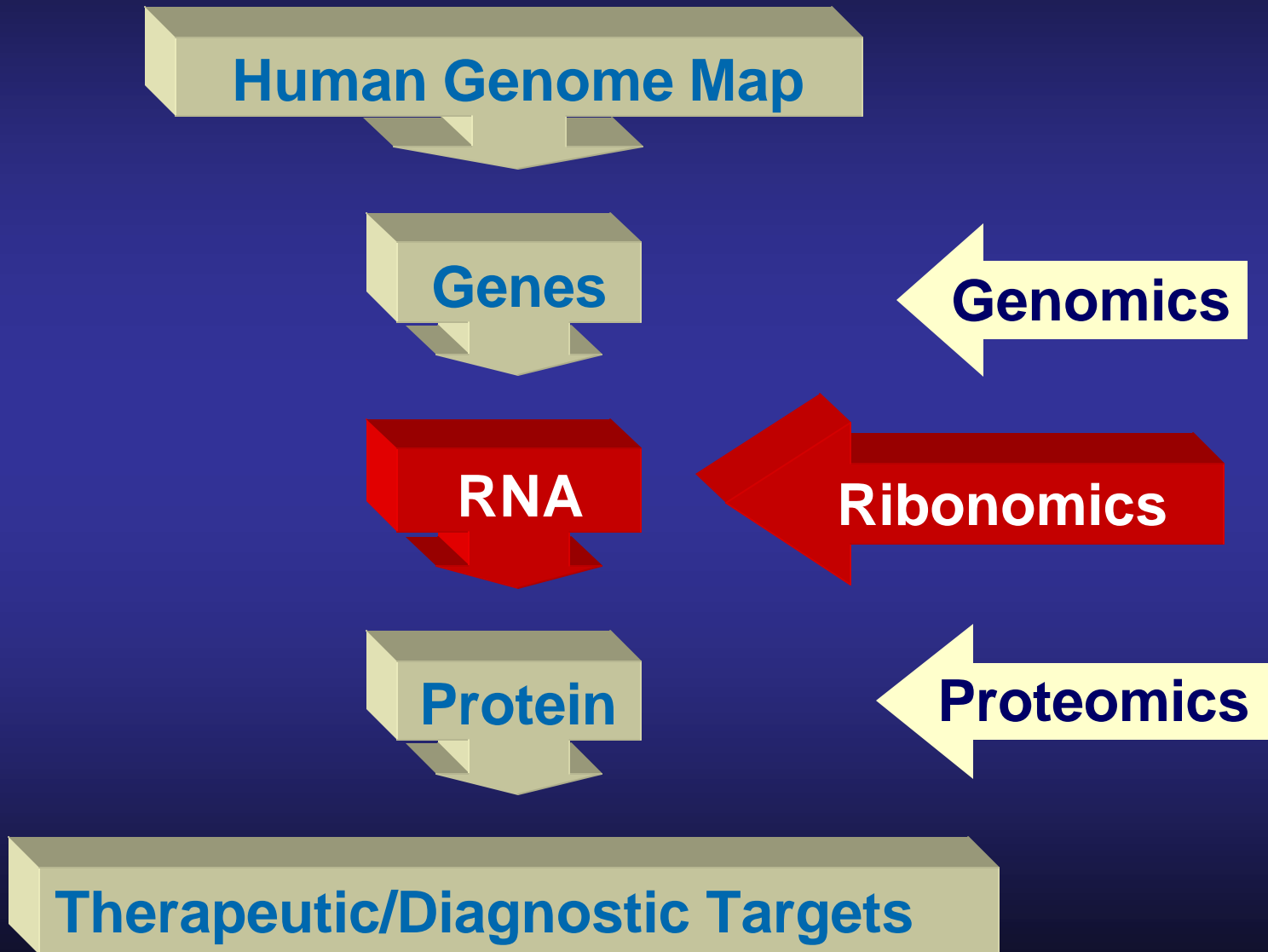
Mission

Ribonomics elucidates cellular pathways that regulate gene expression to discover novel targets for the diagnosis and therapy of human disease.

Overview - Ribonomics, Inc.

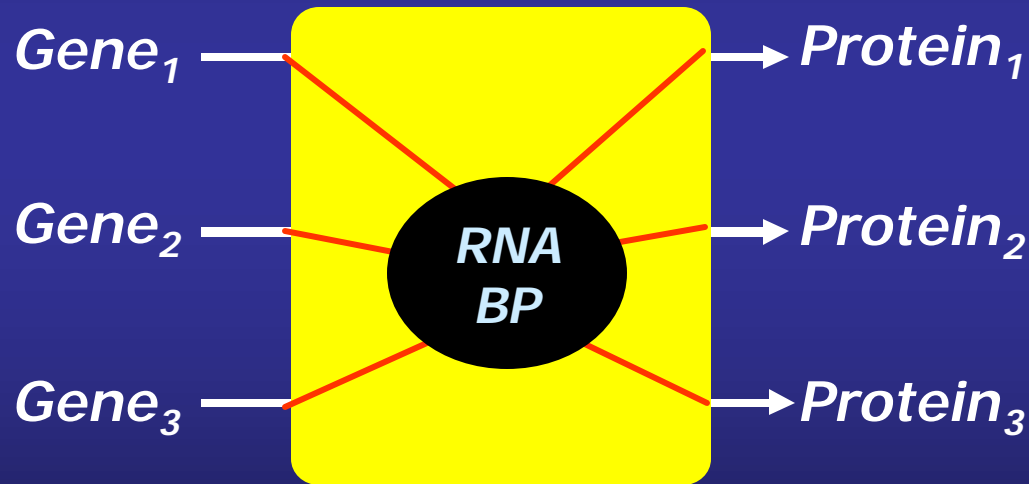
- ❑ Technology spun out of Duke University Medical School
- ❑ Operations launched in June 2001
- ❑ 7000 ft² state of the art lab facility in Research Triangle Park, NC
- ❑ 14 employees (9 Scientists/4 PhD)
- ❑ Series A funding provided by a single corporate partner:
MBL Co., Ltd (Japan)
- ❑ Series B projected Q2/Q3 2002

From Genes to Targets



RNA Binding Proteins

the Flow of Information from the Genome to the Proteome



Ribonomic Cluster: A group of mRNAs that are defined by association with a specific RNA binding protein

Ribonomics: The Proprietary Analysis of Coordinately Regulated RNA Clusters

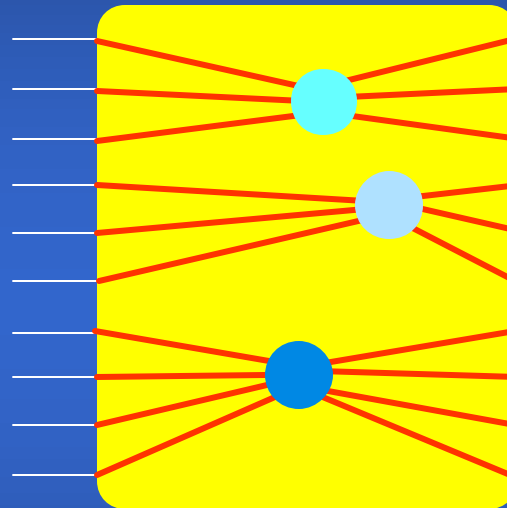
Genome



Transcriptome

mRNA₁
mRNA₂
mRNA₃
mRNA₄
mRNA₅
mRNA₆
mRNA₇
mRNA₈
mRNA₉
mRNA₁₀

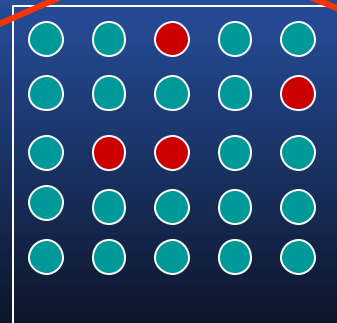
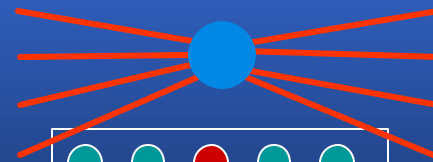
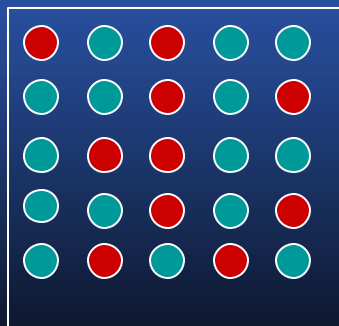
Ribonome



Proteome

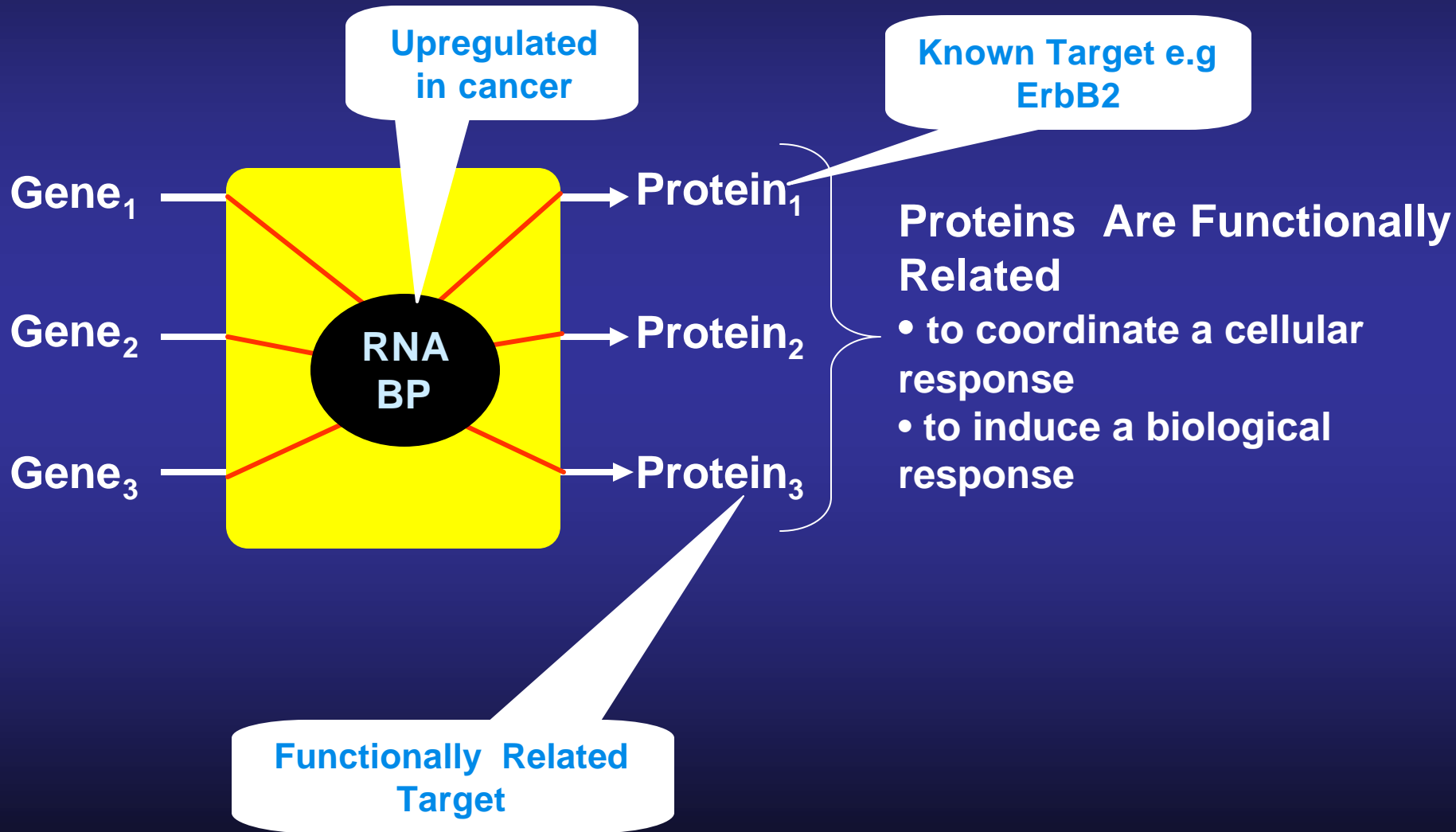
Protein₁
Protein₂
Protein₃
Protein₄
Protein₅
Protein₆
Protein₇
Protein₈
Protein₉
Protein₁₀

Total RNA



Ribonomic Cluster

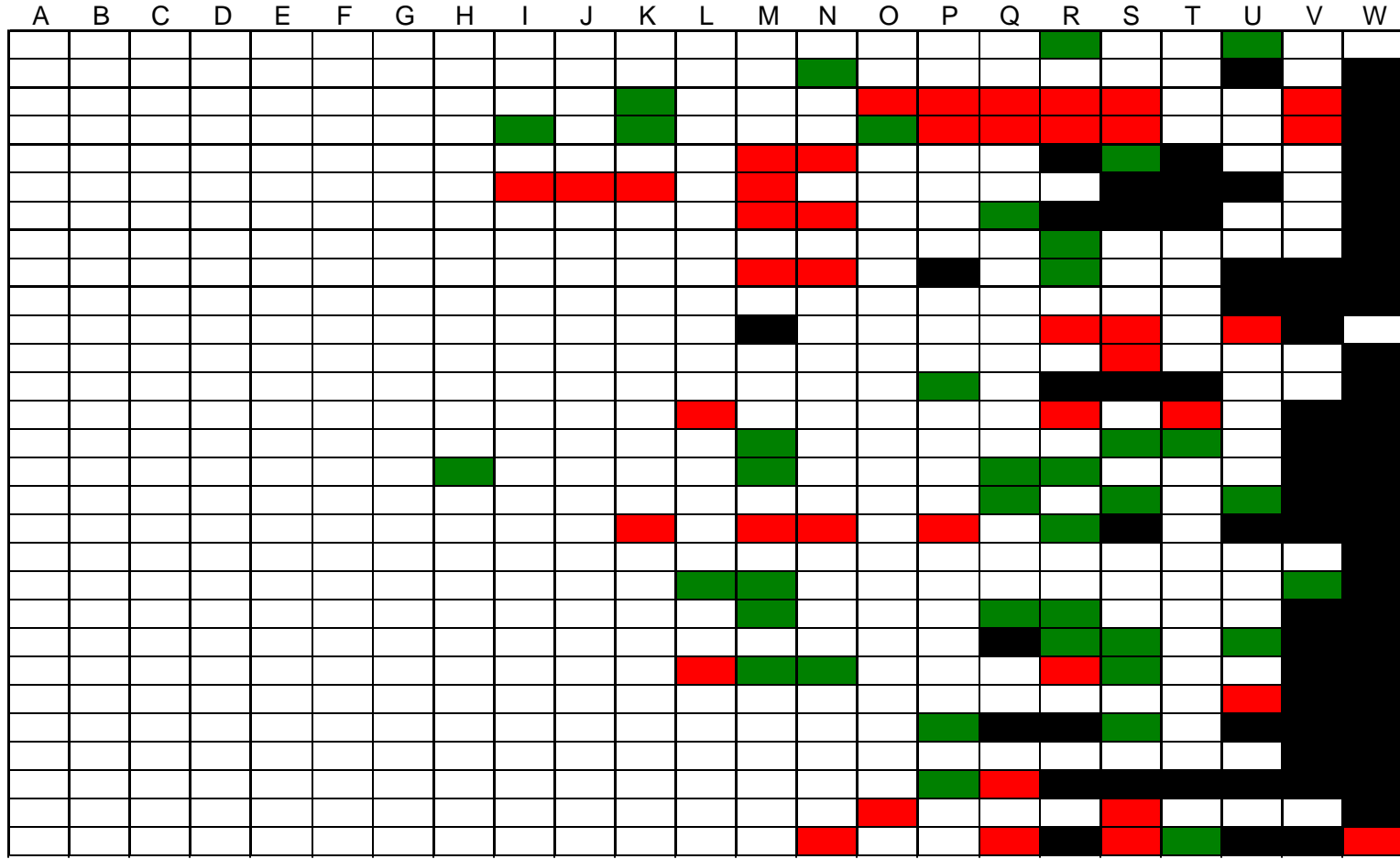
Functional Clusters



RNA BP Genes

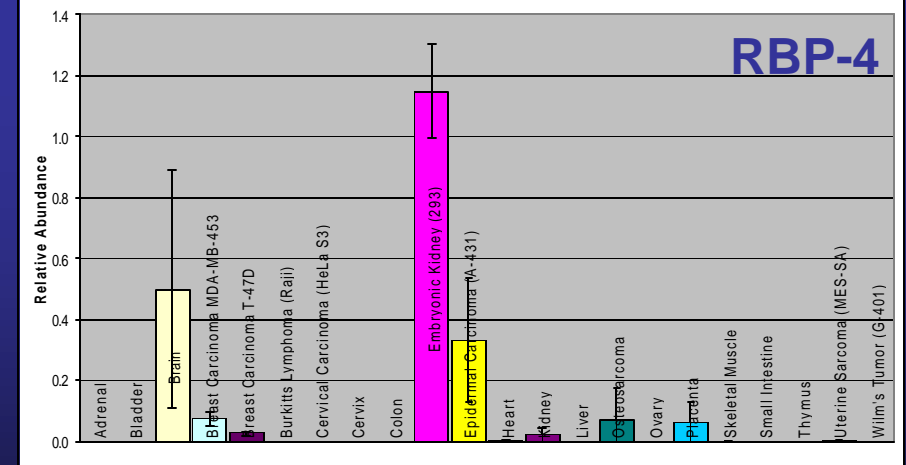
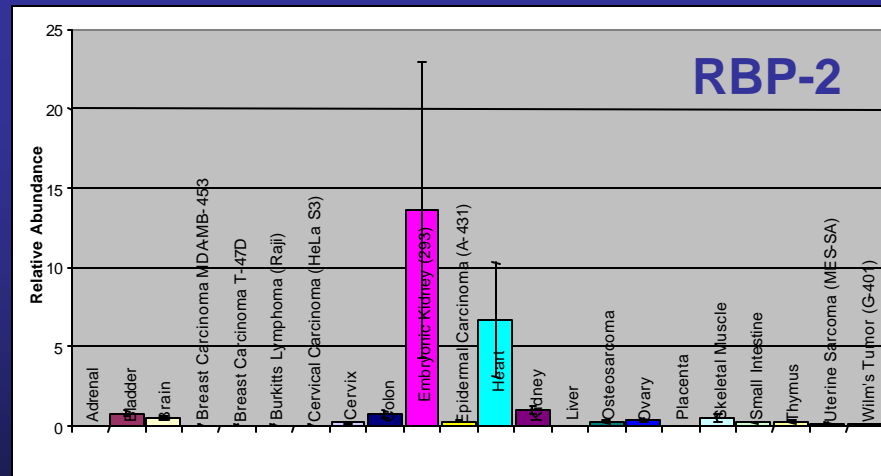
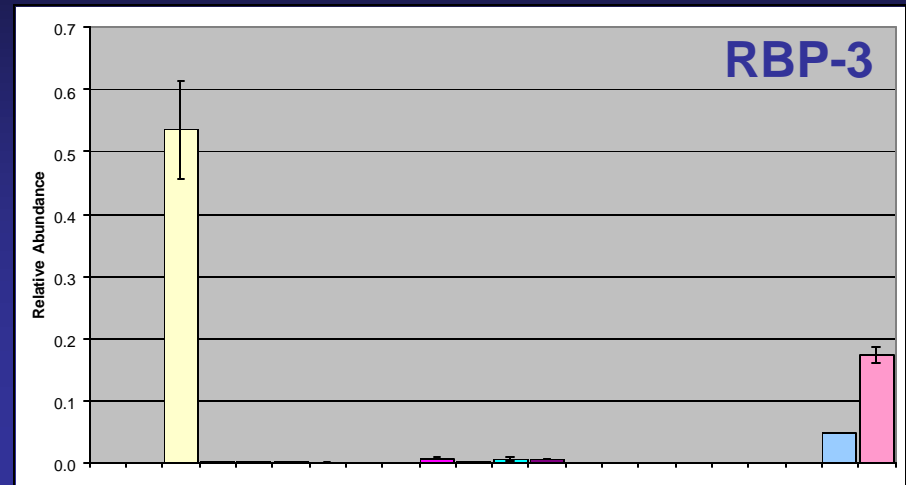
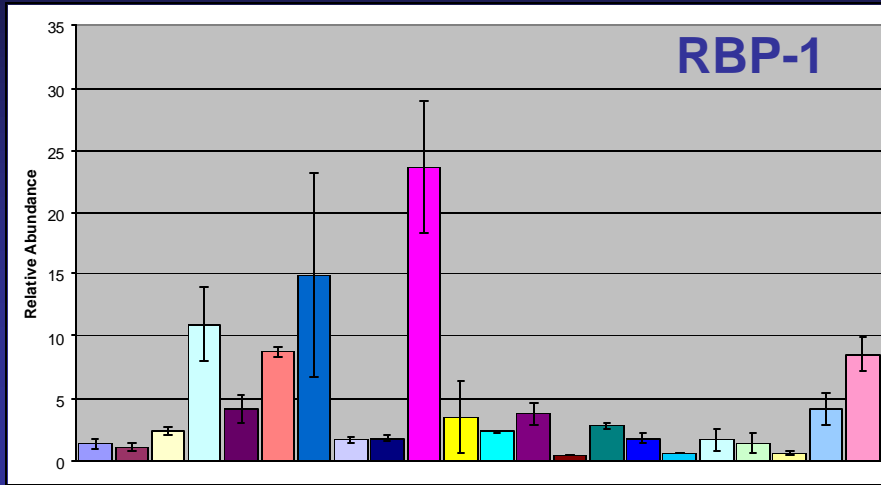
Tissues

- Adrenal
- Bladder
- Brain, Fetal
- brain, Adult
- Breast Adenocarcinoma
- Breast Carcinoma
- Breast Carcinoma
- breast
- Burkitts Lymphoma
- Burkitts Lymphoma
- Cervix
- Colon
- Epidermal Carcinoma
- Heart
- Kidney
- Liver
- Lung
- Osteosarcoma
- ovary, 47 yrs Female
- Ovary
- Pancreas
- Placenta
- Skeletal Muscle
- Small Intestine
- Spleen
- Thymus
- Uterine Sarcoma
- Uterus
- Wilm's Tumor



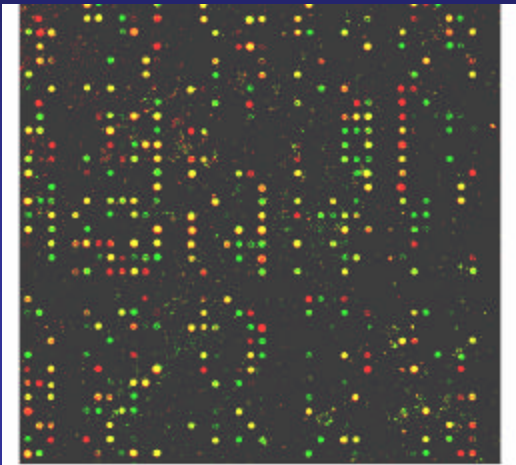
Average +/- 10X
 10X > Average
 10X < Average
 No Expression Detected

RiboMap (QPCR)

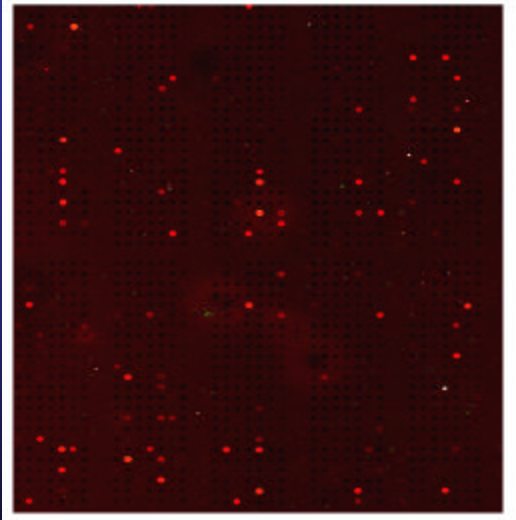


The Ribonomic Advantage

**Total
RNA**



**Cluster
RNA**



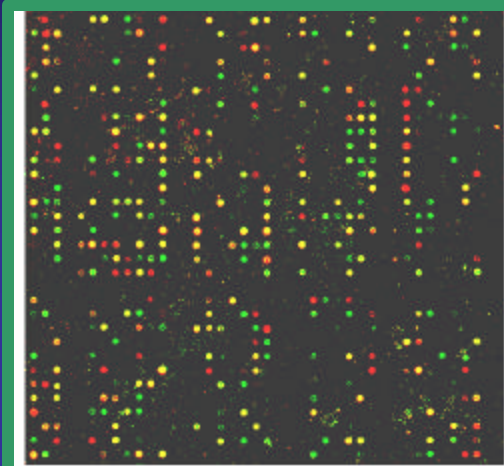
1. Novel growth factor receptor mRNA, 3' cds
2. HSPC010 mRNA
3. cp19 homeobox from HOX-3 locus
4. DKFZp434I068_r1 434 (synonym: htes3)
5. DKFZp434I1820
6. PAX-9 mRNA
7. Transitional epithelia response (TERE1) mRNA
8. IMAGE:3124866
9. Cyclooxygenase (COX-2)
10. E2-25K ubiquitin conjugating enzyme

1. Transitional epithelia response (TERE1) mRNA
2. LUPUS KU AUTOANTIGEN
3. KERATIN, TYPE II CYTOSKELETAL 8
4. PAX-9 mRNA
5. IMAGE:2364179
6. clone 24566
7. nuclear factor NF-IL6
8. H_DJ0555L14.2
9. ATF6
10. IMAGE:2980306

RAS

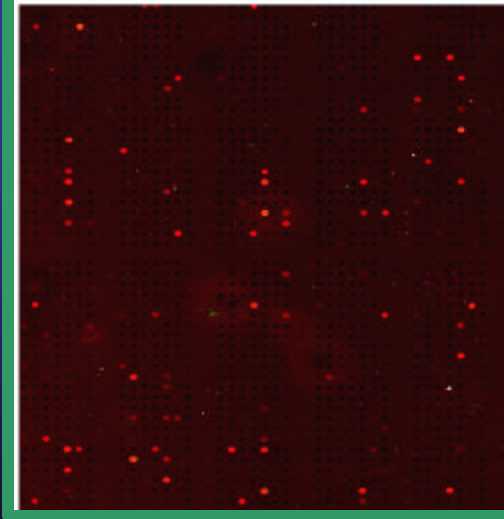
Separating the wheat from the chaff

Total RNA



- Global view of transcriptome
- Low sensitivity
- Complex Data Sets

RBP-1



- Enriched subset of mRNAs
- Higher Sensitivity
- Focus--Simplifies Data sets
- Are coordinately regulated
- Many in same pathway, i.e. 'functionally related'
- Identity of Novel Components

The Ribonomic Value

- Enables use of the cell's own organization to analyze and simplify gene expression
- Enables the discovery of novel targets by elucidating functional relationships between genes
- Bridges the gap from Genomics to Proteomics

Business Model

Ribonomics, Inc. is pursuing novel therapeutic and diagnostic targets in

- Cancer**
- Metabolic Disease**
- CNS**

Ribonomics seeks to develop strategic technology partnerships and focused research collaborations to extend our technology platform into key scientific areas.



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