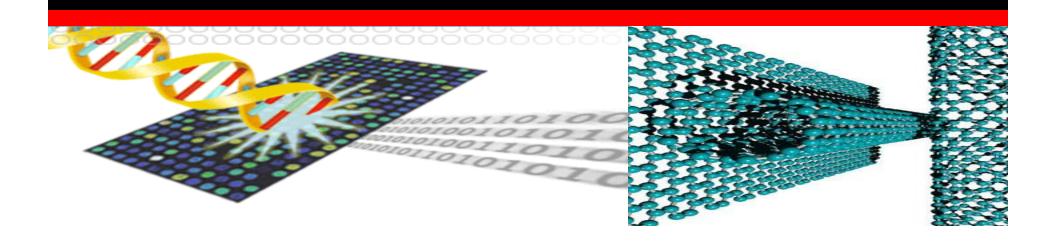


Convergence of Transcription, Metabolomics and Wireless Nano-Sensor Networks

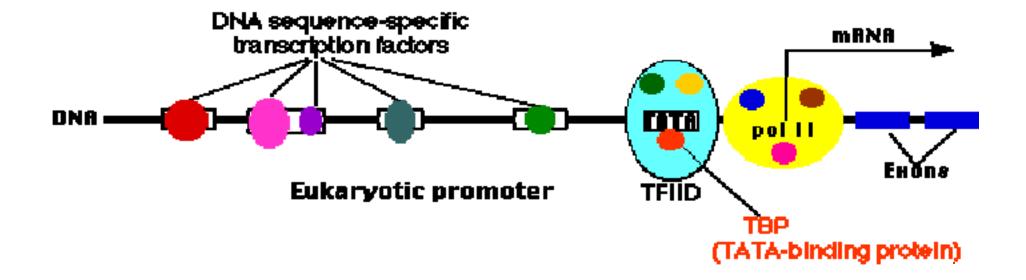
Shoumen Datta

Massachusetts Institute of Technology



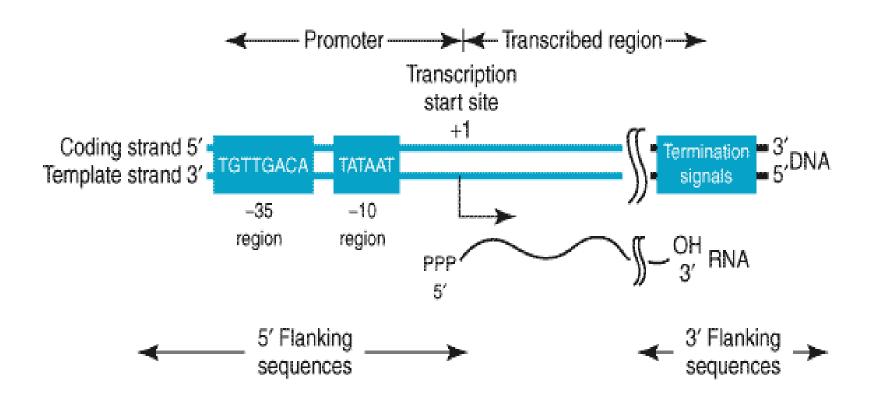


Gene Expression: Regulation of Transcription



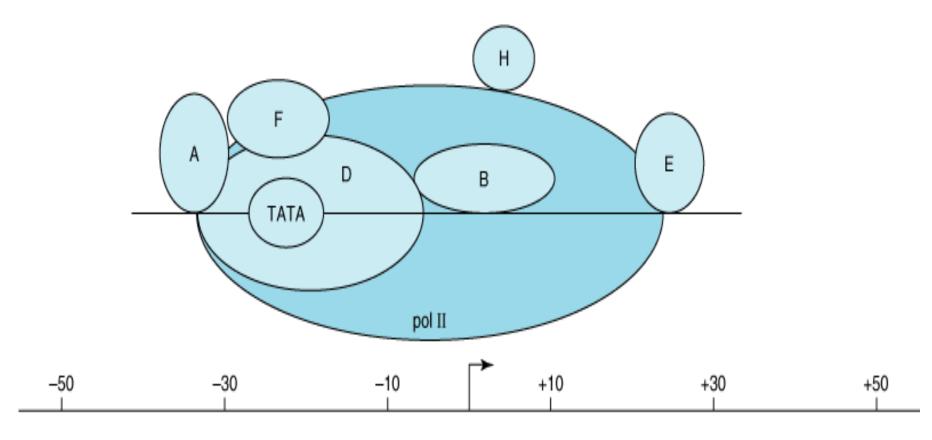


Transcription Unit



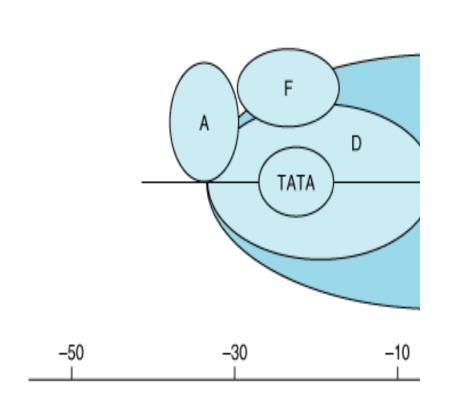


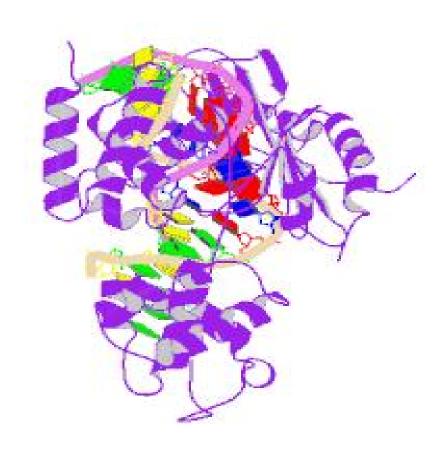
Initiation Complex: RNA Polymerase II





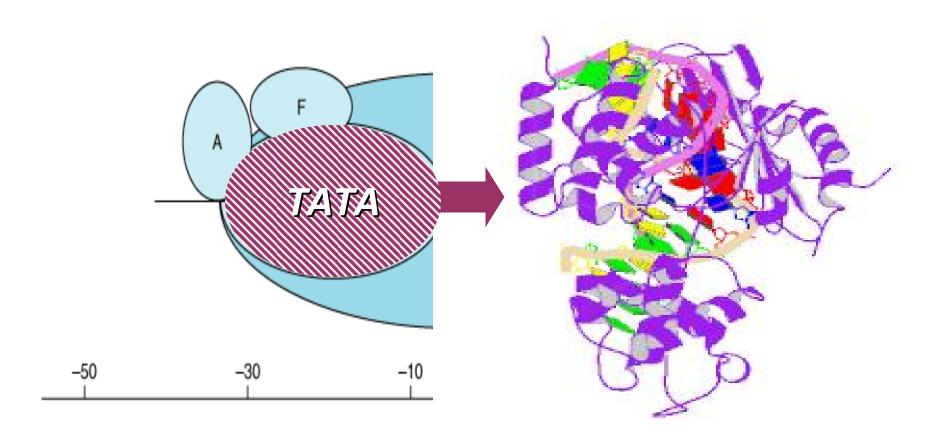
Initiation Complex: TATA Binding Protein







Initiation Complex: TATA Binding Protein

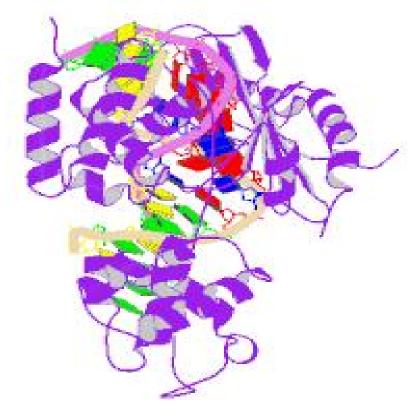




TFIID – TBP: TATA Binding Protein

PROLINE
$$(\beta_1)_{50}^{y,h}$$





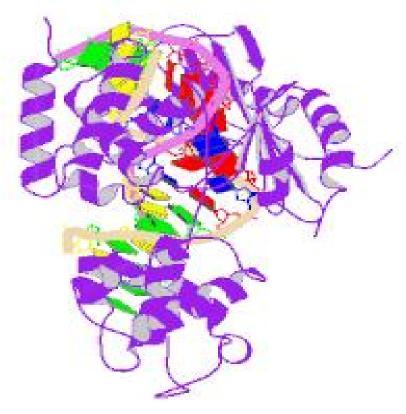


TBP: Yeast, humans and malaria parasite



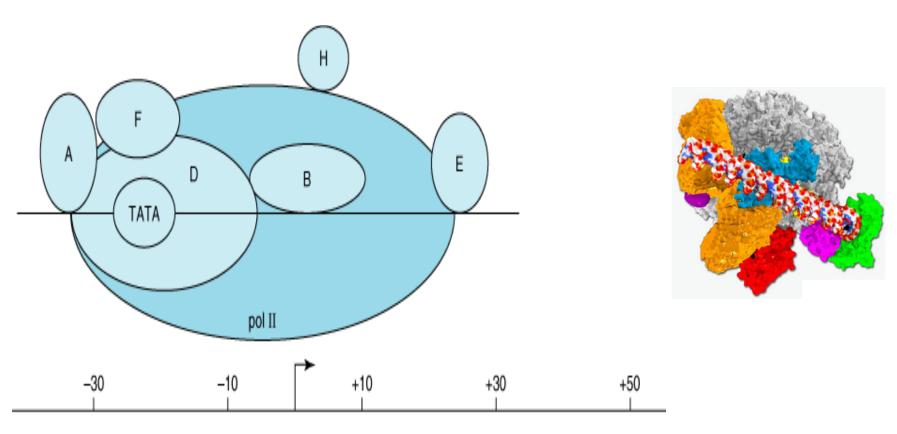
PROLINE $(\beta_1)_{50}^{y,h}$







Initiation Complex: RNA Polymerase II

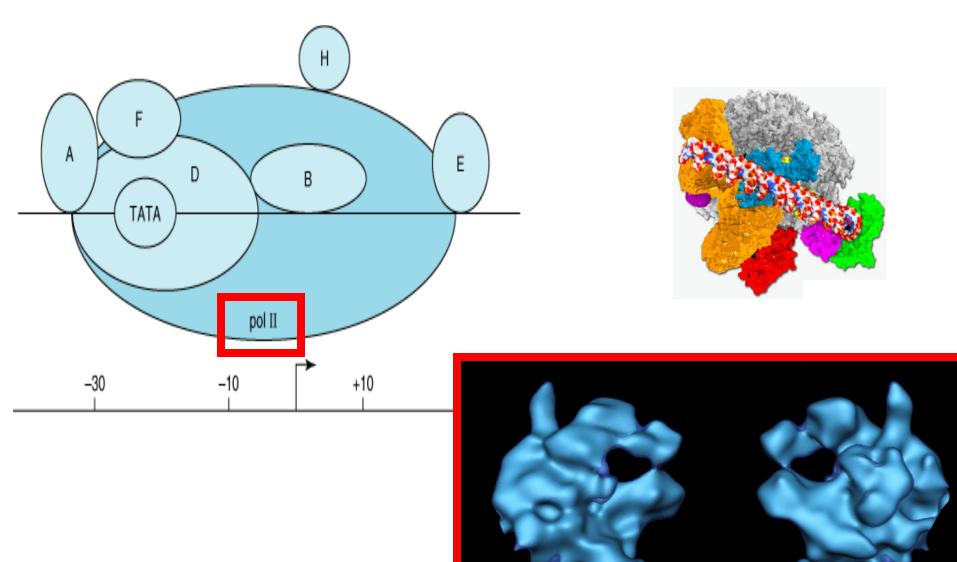




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Dr Shoumen Datta, MIT <shoumen@mit.edu>

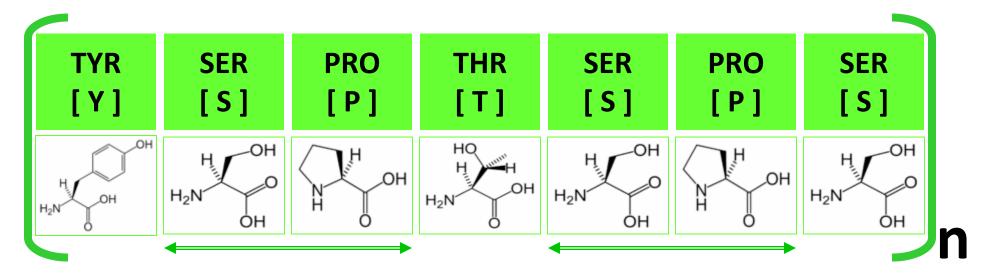
Eukaryotic RNA Polymerase II Subunits



© Darst

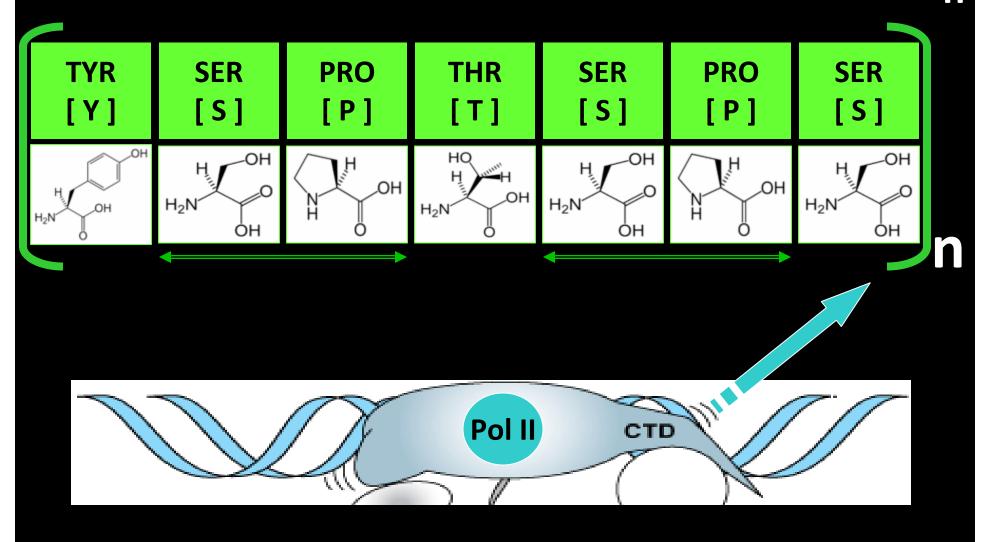
* SACHUSE TITO * A STATE OF TECHNOLOGY

RNA Polymerase II Subunit 1: RPB1 C Terminal Domain (CTD): [YSPTSPS]_n



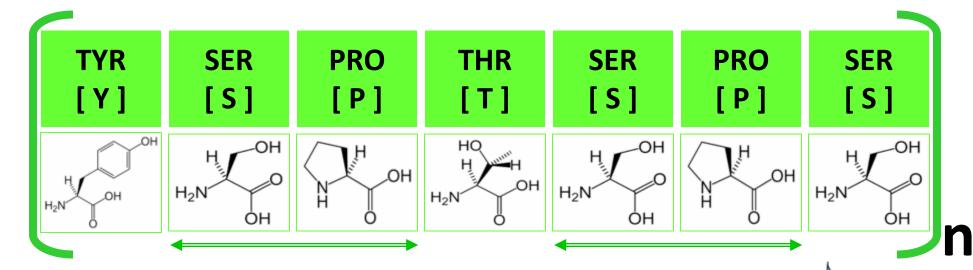


RNA Polymerase II Subunit 1 : RPB1 C Terminal Domain (CTD): [YSPTSPS]_n



SSACHUSE II. SPACHUSE III. SPACHUSE III.

RNA Polymerase II Subunit 1: RPB1 C Terminal Domain (CTD): [YSPTSPS]_n



Yeast

yRPB1-CTD: [YSPTSPS]₂₆

Human

hRPB1-CTD: [YSPTSPS]52

Malaria

pRPB1-CTD: [YSPTSPS] + 63 aa

Pol II

Trypanosome

tRPB1-CTD: [YSPTSPS]

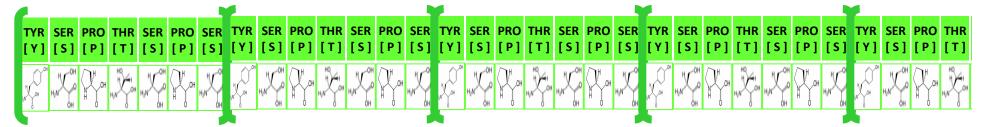




Host-Parasite Interactions: CTD

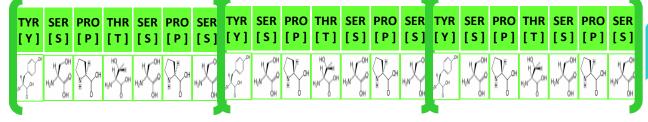
Human

hRPB1-CTD: [YSPTSPS]52



Malaria

pRPB1-CTD: [YSPTSPS]₁₅ + 63 amino acids



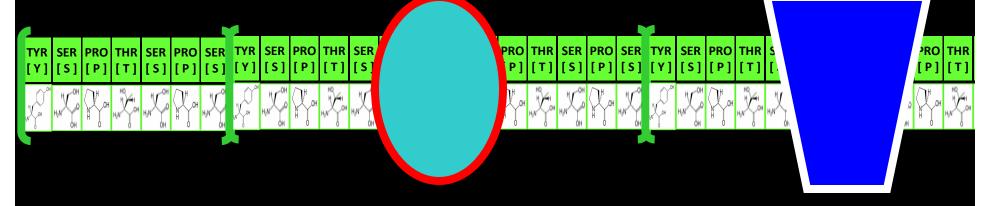




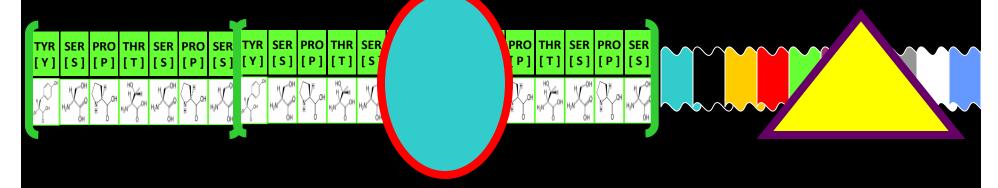
Host-Parasite Interactions

Unique CTD Proteins Regulating Malaria Gene Expression in Infected Humans

Human hRPB1-CTD: [YSPTSPS]52



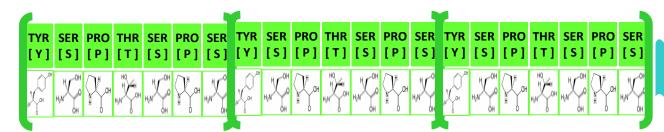
Malaria pRPB1-CTD: [YSPTSPS] 15 + 63 amino acids

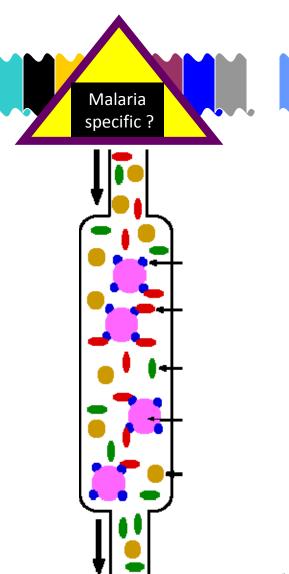




Identification of Targets for RNA Immunity?

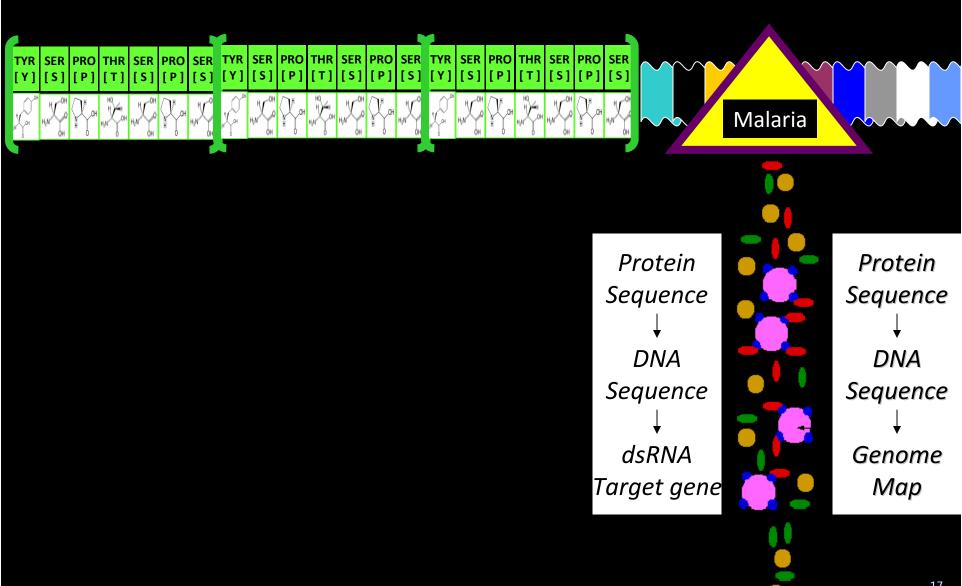
Are Unique CTD Proteins Regulating Malaria Gene Expression in Infected Humans?





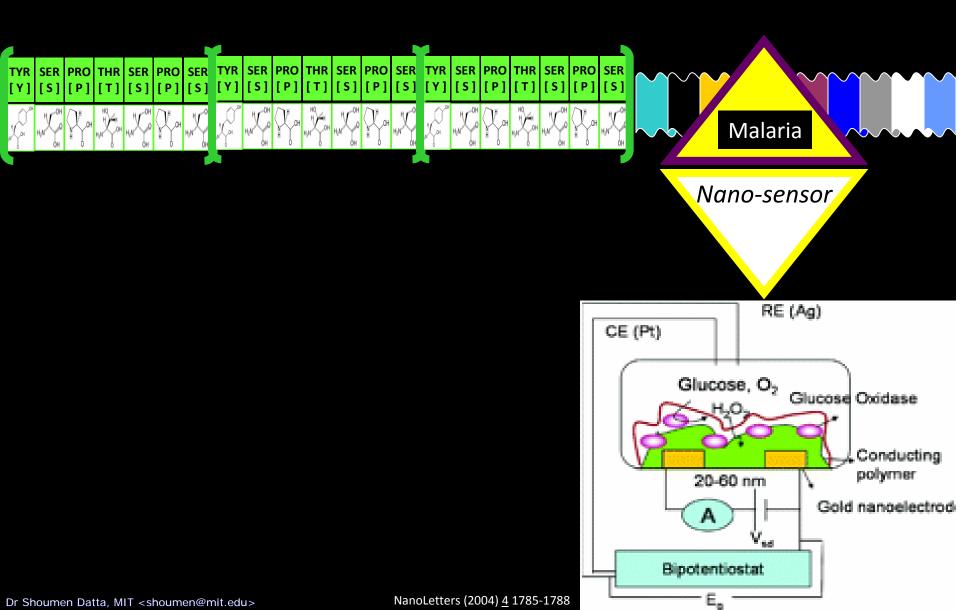


If target protein is specifically parasite induced: A Potential for Development of Malaria Therapy



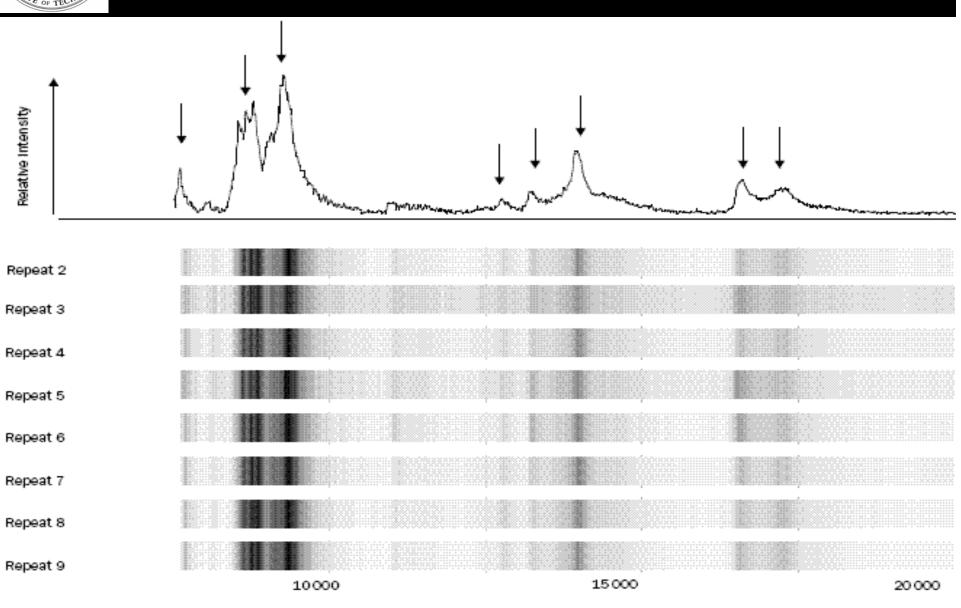


Detection of Target Human Protein: Constitutive Expression? Early detection of infection by Malaria parasite *P. falciparum*?





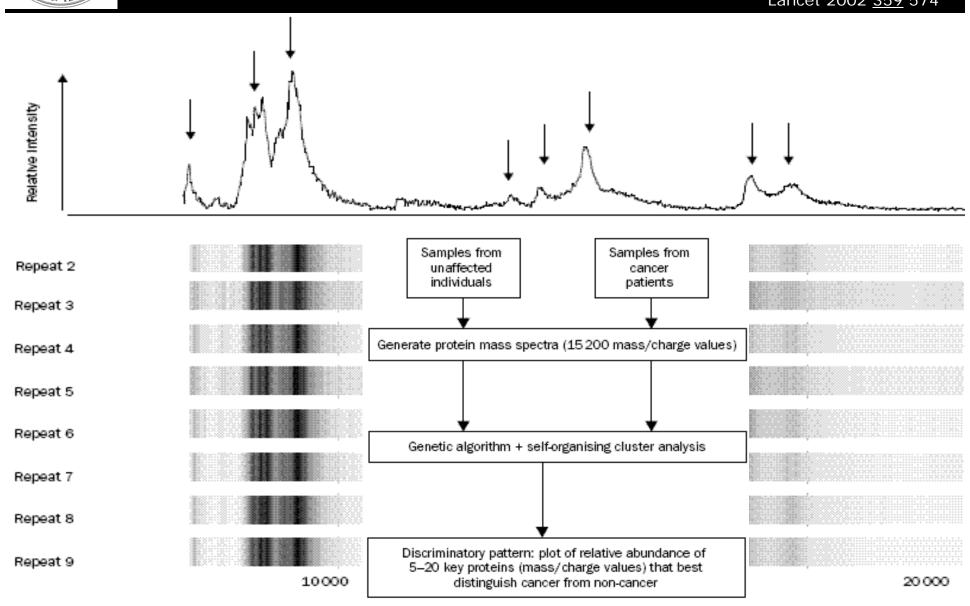
Other Tools to Identify Unique Target Proteins: Mass Spec Analysis

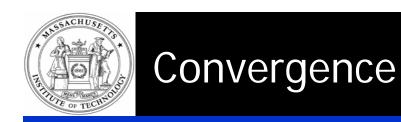




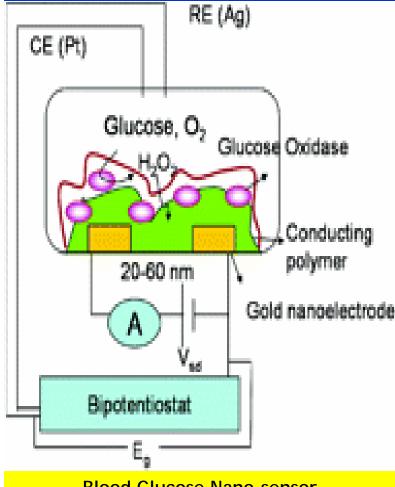
Mass Spectrometric Analysis of Proteomes in Ovarian Cancer

Lancet 2002 359 574

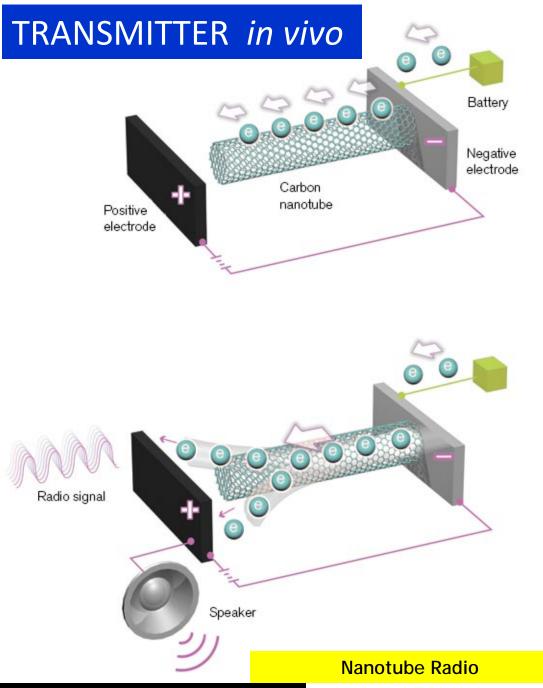




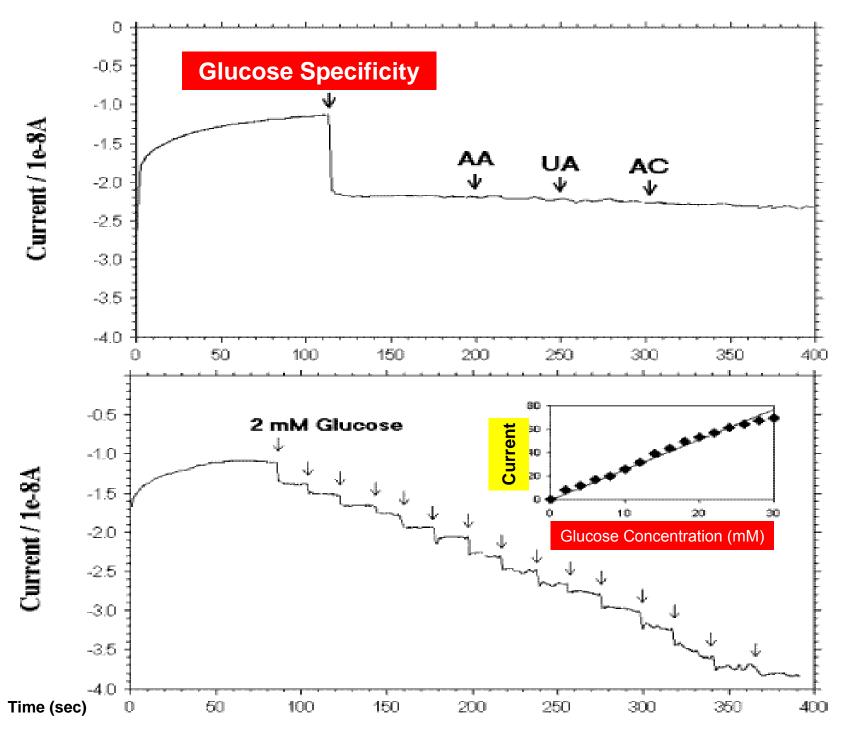
MALARIA SENSOR DATA



Blood Glucose Nano-sensor







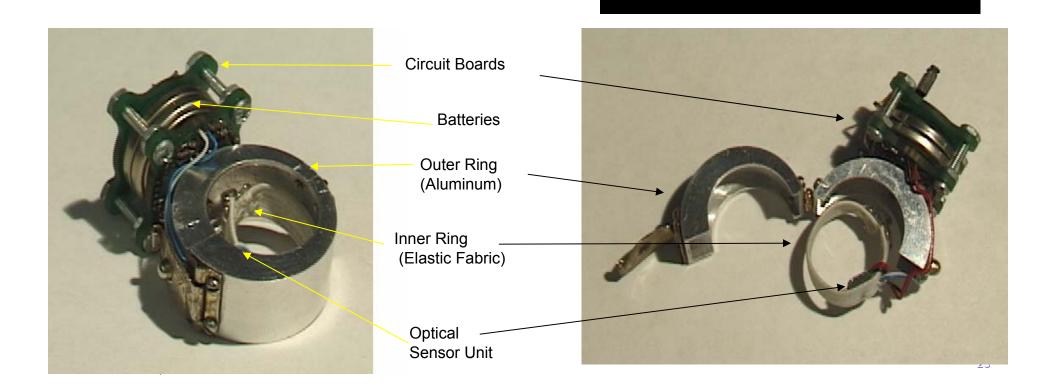


Sensors: Not necessarily in vivo, Not necessarily nano

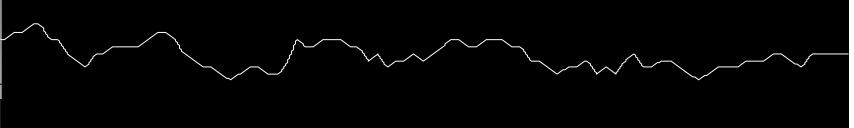
Ring Sensors

Sokwoo Rhee, Ph.D. thesis, MIT

www.sokwoo.com







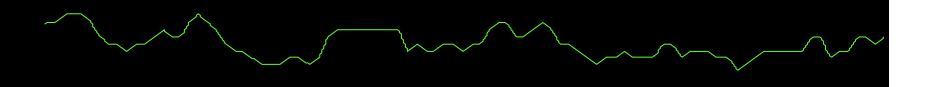
(sec

(a) Heart Rate (beats/min) by Electrocardiogram (ECG)



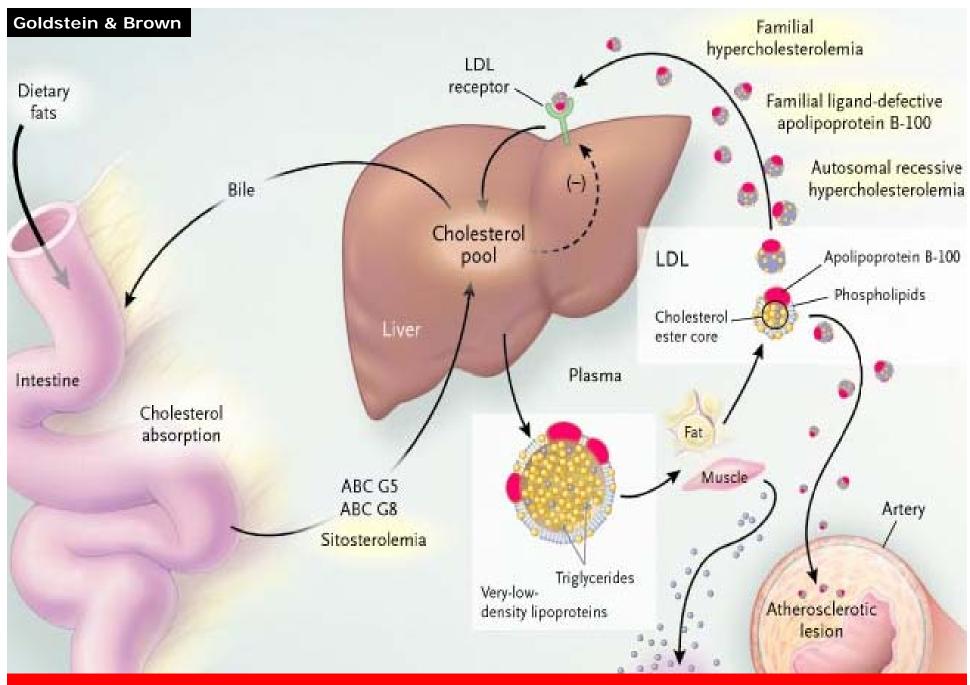
(se

(b) Heart Rate (beats/min) by Fingertip Photoplethymograph (PPG



S

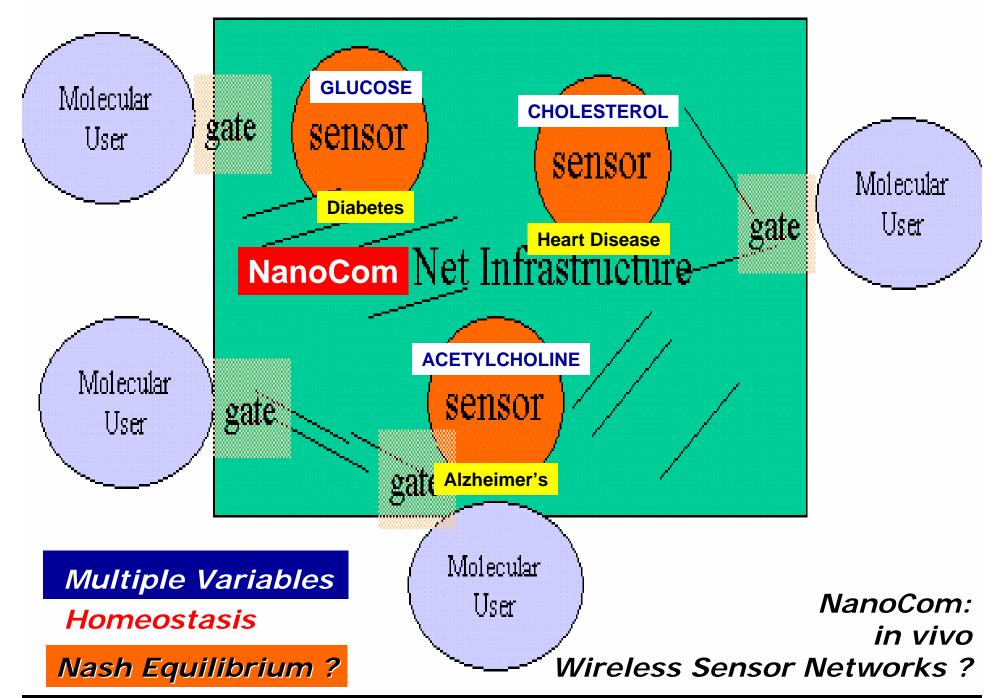
(c) Heart Rate (beats/min) monitored by Wireless Ring Sensor shown in Figure



: Prime Target for Sensor-based Early detection & RNA-based Immunity

Human	LYR	PQG	R	CACPI	G F E L
Chimp	LYR	PQG	L R	CACP!	G F E L
Monkey	LYR	P Q G	L R	CACPI	G F E L
Mouse	LYR	P Q G	R R	CACP	G F E L
Rat	LYR	PQG	R R	CACP	GFEL
Dog	LYR	PQG	L R	CACPI	G F E L
Cow	LYR	P Q G	L R	CACP	G F E L
Opposum	FR	P Q G	L R	CACPI	G F E L
Chicken	LYR	PQG	R R	CACP	GLEL
Frogs	L F R	PQG	PR	CACPI	GLEL

Stuart Lipton et al Mutation in LRP6 cosegregates with early CAD and high LDL Science (2007) 315 1278 -1282





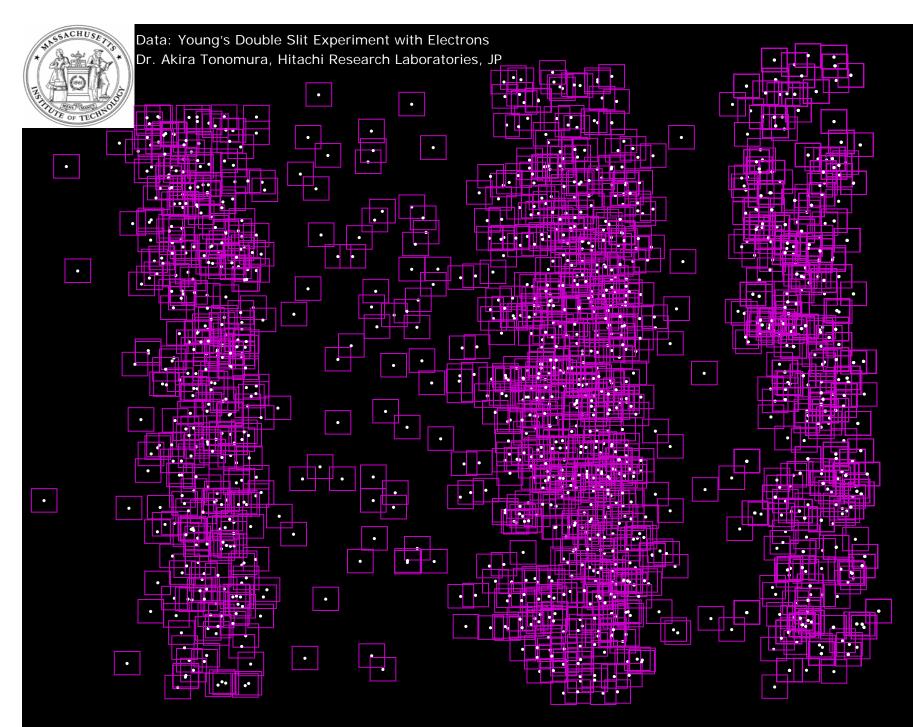




More data points ...

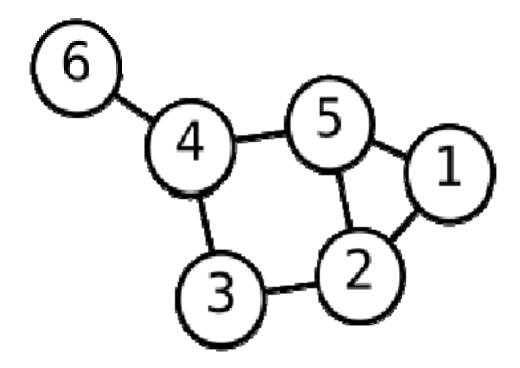


Data shows emerging pattern



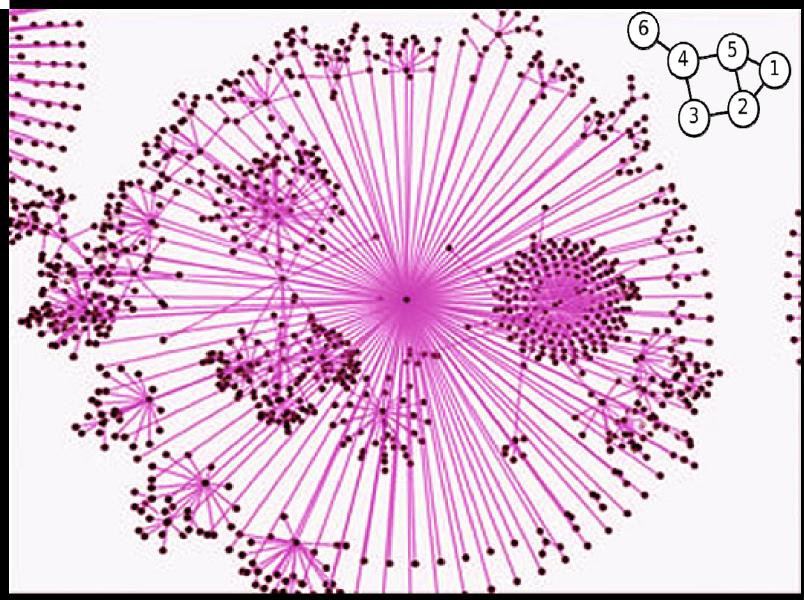


Directed Graph: Today's Network





Social Networking





Sense of Distributed Data Analytics

n = 10; p = 1,000 10 locations 1,000 lags

Estimate Coefficients:

10,000 **Φ**

10.000 for x's

20,000 per stage

or

200,000 for n=10 (excluding constants

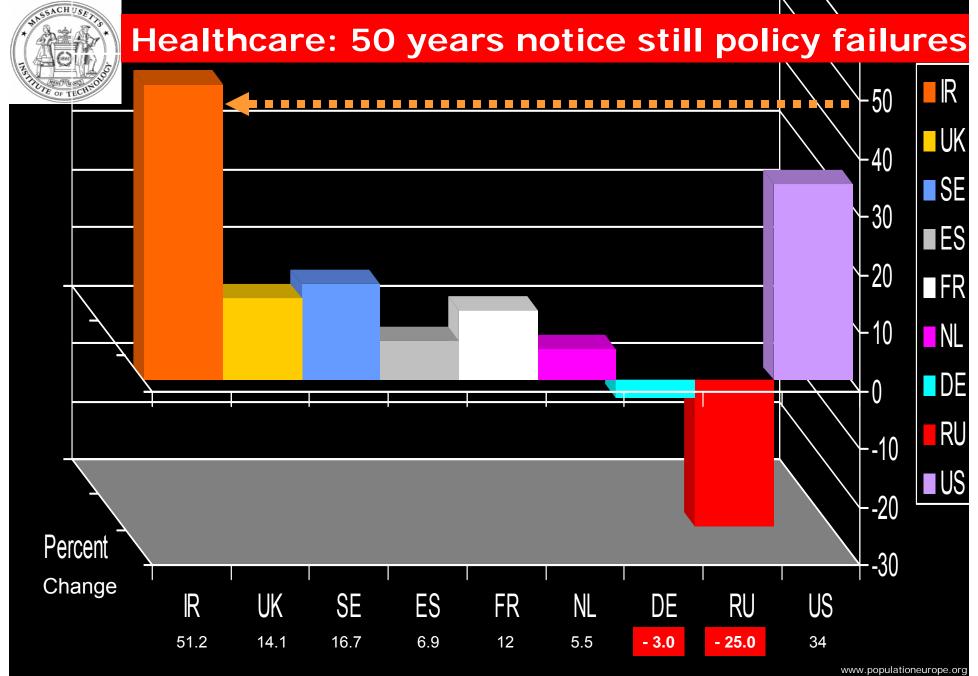
and error coefficients)

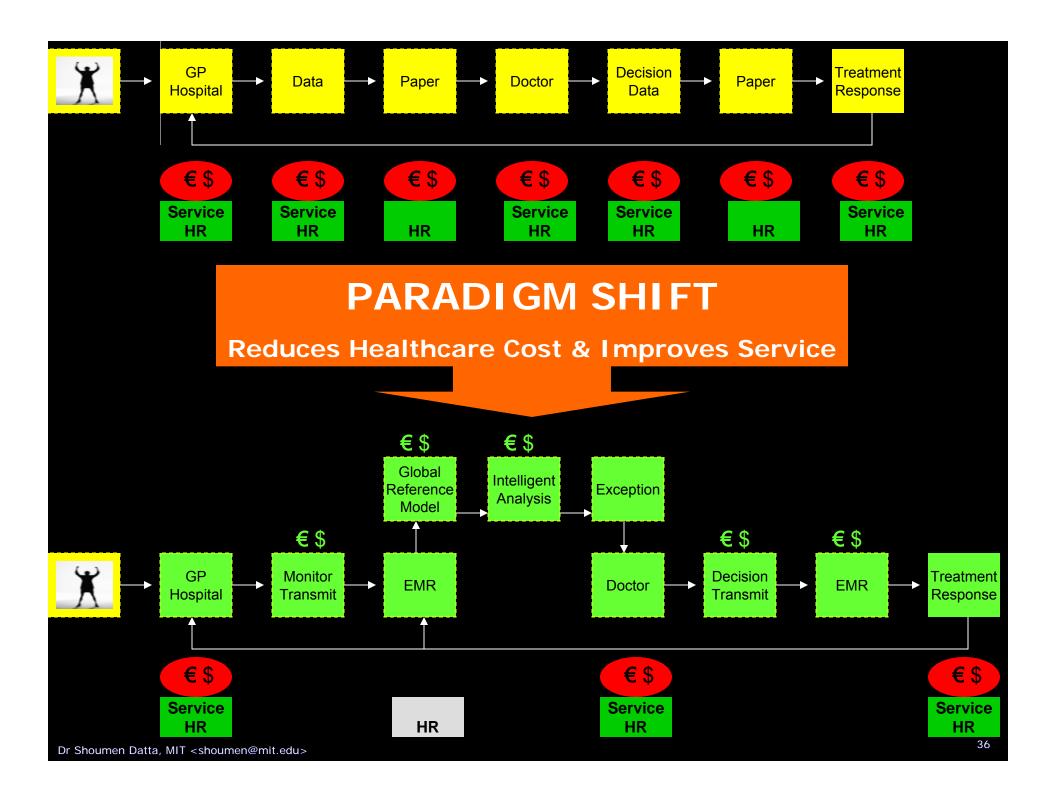
$$\mathbf{y_{1t}} = \beta_0 + \sum_{k=1}^{K} \sum_{i=1}^{N_{\mathbf{x_{kt}}}} \alpha_{ki} \mathbf{x_{kt-i}} + \boldsymbol{\phi}_{11} \mathbf{y}_{1t-1} + \boldsymbol{\phi}_{12} \mathbf{y}_{2t-1} + \boldsymbol{\varepsilon}_{1t}$$

$$\mathbf{y_{2t}} = \beta_0 + \sum_{k=1}^{K} \sum_{i=1}^{N_{\mathbf{x}_{kt}}} \alpha_{ki} \mathbf{x}_{kt-i} + \phi_{21} \mathbf{y}_{1t-1} + \phi_{22} \mathbf{y}_{2t-1} + \mathbf{\epsilon}_{2t}$$

$$\sigma_{1t}^2 = \theta_0 + \sum_{i=1}^q \theta_i \varepsilon_{1t-i}^2 + \sum_{j=1}^p \tau_j \sigma_{1t-j}^2$$

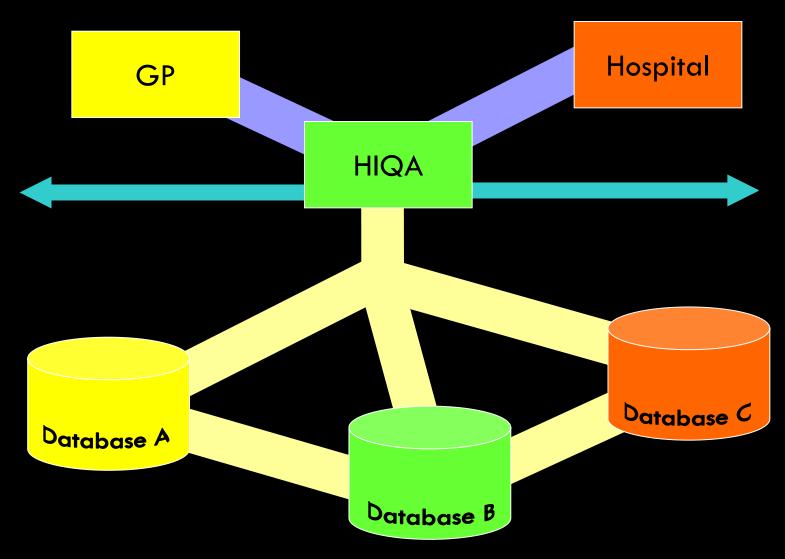
$$\sigma_{2t}^2 = \theta_0 + \sum_{i=1}^q \theta_i \epsilon^2_{2t-i} + \sum_{j=1}^p \tau_j \sigma^2_{2t-j}$$

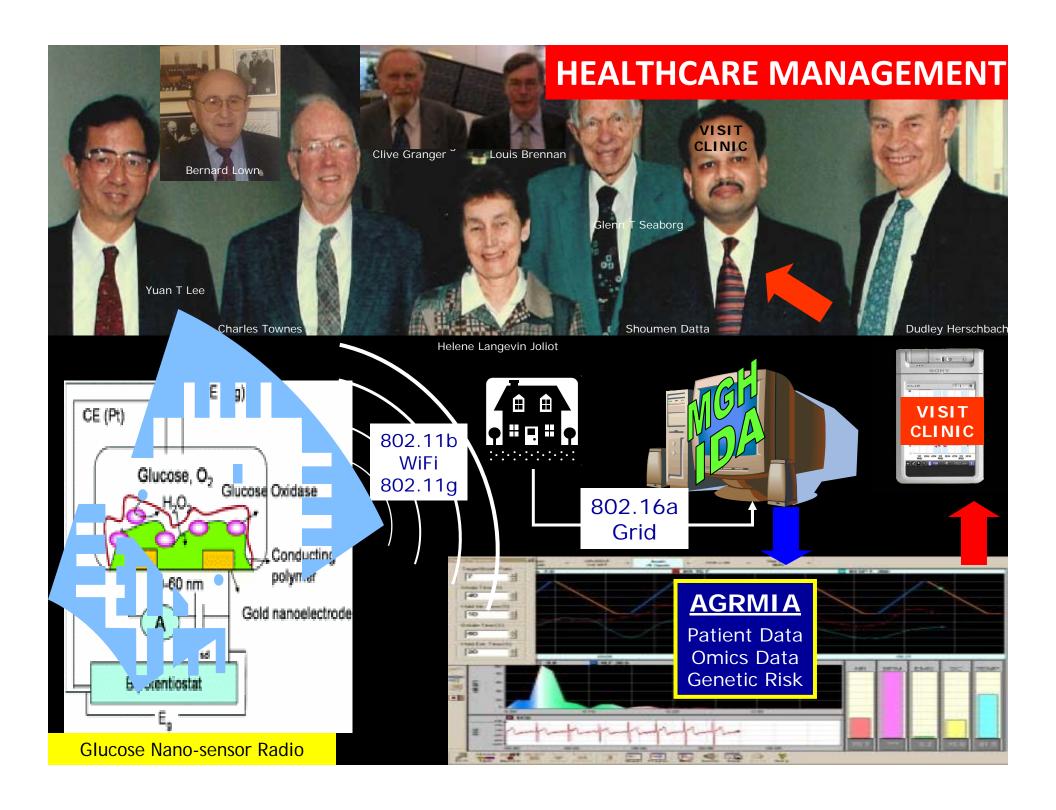






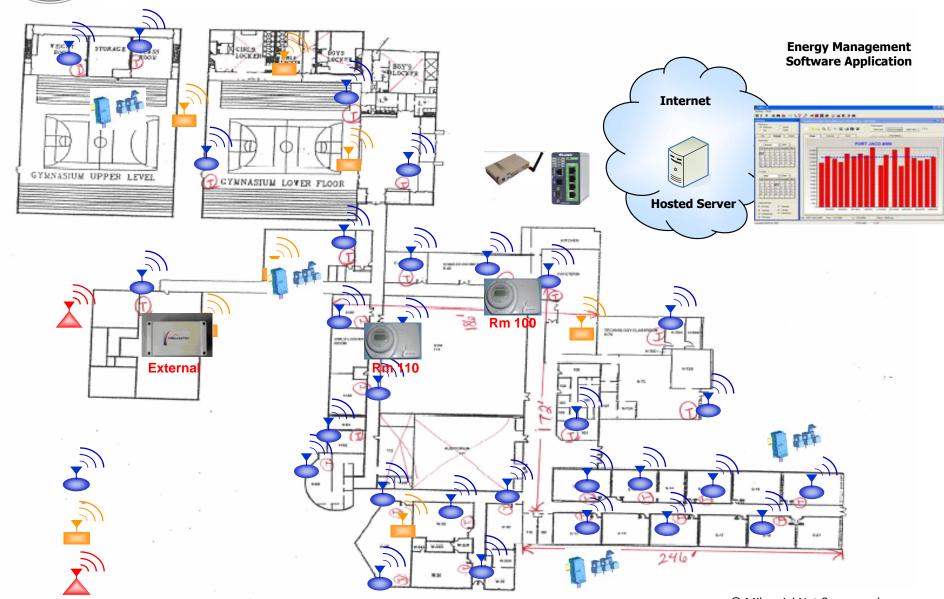
Data & Information Asymmetry: Isolated DDT Systems





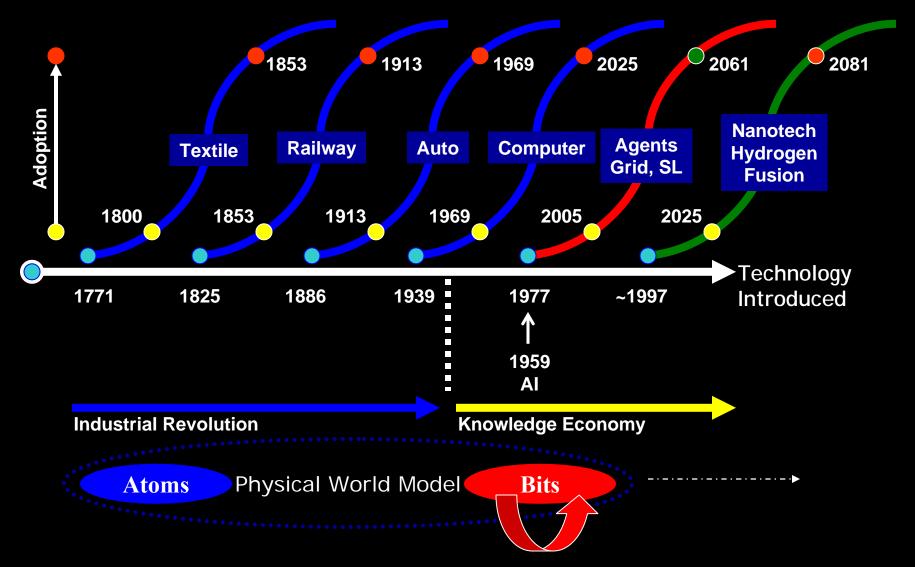


ENERGY MANAGEMENT





Conceptual Convergence and the Wealth of Nations





"Did not entail being right all the time. It was rather to dare, to propose new ideas, and then to verify them and to know how to admit errors."

Professor Pierre-Gilles de Gennes* (1932-2007) after receiving the 1991 Nobel Prize for Physics

Support research ...

"Research is four things: brains with which to think, eyes with which to see, machines with which to measure and fourth, money."

Albert Szent-Gyorgyi de Nagyrolt Nobel Prize in Medicine (1937)