

Need to Transform Health and Medicine in the 21st Century		
207/02/2	20 th Century	21 st Century
3	Treat disease when symptoms appear and normal function is lost	Intervene before symptoms appear and preserve normal function for as long as possible
	Did not understand the molecular and cellular events that lead to disease	Understanding preclinical molecular events and ability to detect patients at risk
	Expensive in financial and disability costs	Orders of magnitude more effective
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Dan, a 50 something year XY, had his genome scanned 4 years ago by his physician and has 5 gene variants that increase his risk for heart disease 6-fold; he also has 3 genes that protect him from getting cancer. Recommendations for his diet and medications have been selected based on his profile for drug metabolizing enzyme variants.

Circa 2015

- He developed chest pain while jogging. A analysis of his serum based upon biomarkers for myocardial ischemia revealed no necrosis.
- Molecular imaging showed a 'hot' plaque in his right coronary artery.
- A targeted anti-inflammatory agent and targeted blood thinner were given to him and re-imaging showed quiescence of the 'hot' plaque.

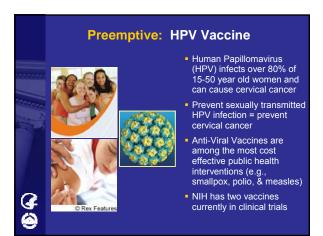


Genomic Medicine

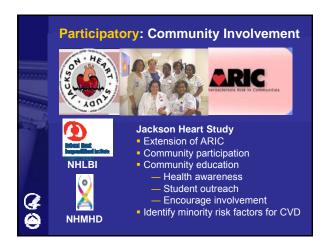
- Prediction of individual risk for various diseases
- Preemption of clinical disease through early detection
- Personalized treatments
- Participation in health maintenance

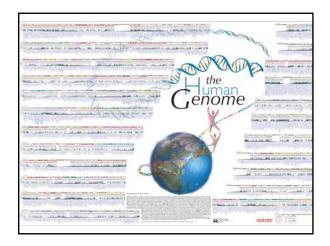


Predictive: End Stage Renal Disease • End-stage Renal Disease: - \$22.8 billion in U.S. public and private spending (2001) - In the past decade, the absolute number of ESRD patients more than doubled and the incidence rate doubled - More than 85,000 new cases per year • Apolipoprotein E (APOE) - Variation predicts kidney disease progression - Prediction independent of diabetes, race, lipid and non-lipid risk factors

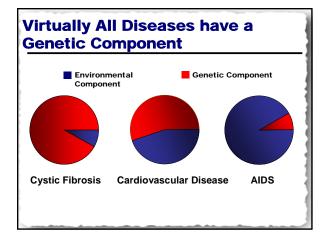


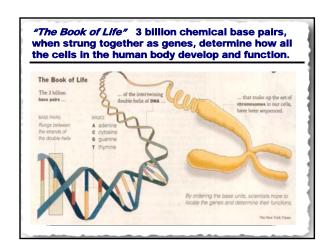




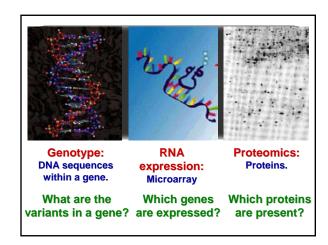


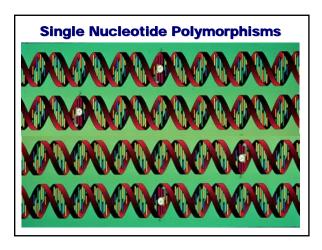


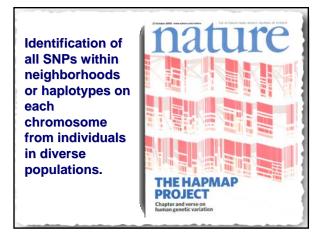


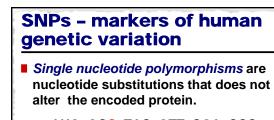


The Human Genome 3.1 billion bases of DNA sequence -97% of the genome contains coding regions 30,000 genes 100,000 proteins Mutations causing disease have been identified in >1,000 genes









AUG GCC TAC GTT CGA CCC

Met Ala Tyr Val Arg Pro

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