



 Summer Institute
 in Statistical Genetics 2017

Genetics and Genomics
5a. Integrative Genomics


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The Rationale for Gene Expression Profiling

Question: What makes a muscle cell different from a skin cell different from a liver cell?
Analogy: What makes a living room different from a bathroom different from the kitchen?
Answer: What furniture and appliances and windows re placed where and when
 What genes are expressed where and when and how much

Strategy: Measure the abundance of mRNA transcripts in a bunch of samples, and use statistically rigorous approaches to identify differential expression

Methods: Prior to 1995: One gene at a time qPCR
 1995-2015: Microarrays
 Since 2015: RNAseq

Annotation of Gene Function

Gene Ontology

Biological process

- GO:0009987 - biological process
- GO:0009988 - cell cycle
- GO:0009989 - cell cycle process
- GO:0009990 - cell cycle phase
- GO:0009991 - cell cycle phase transition
- GO:0009992 - cell cycle transition
- GO:0009993 - cell cycle transition phase
- GO:0009994 - cell cycle transition phase transition
- GO:0009995 - cell cycle transition phase transition phase
- GO:0009996 - cell cycle transition phase transition phase transition
- GO:0009997 - cell cycle transition phase transition phase transition phase
- GO:0009998 - cell cycle transition phase transition phase transition phase transition
- GO:0009999 - cell cycle transition phase transition phase transition phase transition phase

Molecular function

- GO:0005488 - catalytic activity
- GO:0005489 - catalytic activity, nucleic acid dependent
- GO:0005490 - catalytic activity, nucleic acid dependent, nucleic acid dependent
- GO:0005491 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
- GO:0005492 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
- GO:0005493 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
- GO:0005494 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
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- GO:0005498 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
- GO:0005499 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent
- GO:0005500 - catalytic activity, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent, nucleic acid dependent

Cellular component

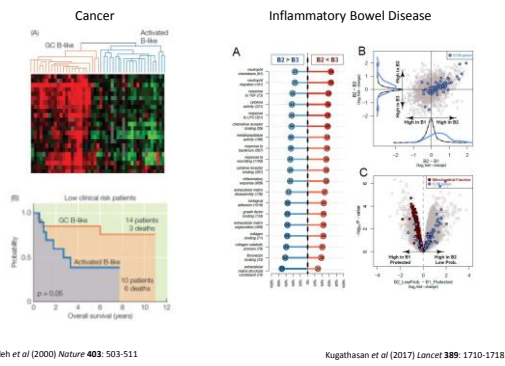
- GO:0005622 - cell
- GO:0005623 - cell body
- GO:0005624 - cell body part
- GO:0005625 - cell body part part
- GO:0005626 - cell body part part part
- GO:0005627 - cell body part part part part
- GO:0005628 - cell body part part part part part
- GO:0005629 - cell body part part part part part part
- GO:0005630 - cell body part part part part part part part
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- GO:0005643 - cell body part
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- GO:0005649 - cell body part
- GO:0005650 - cell body part

← Expression in a tissue generally implies the gene is active there

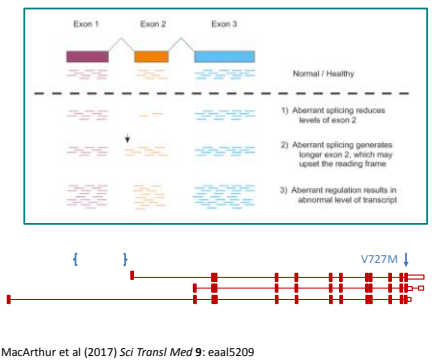
← Co-expression with similar types of gene may imply "guilt by association"

← FISH and sub-cellular imaging of proteins tells us where they act

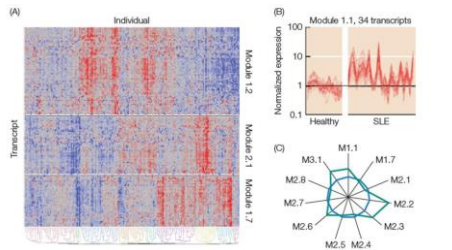
Differential Expression Analysis



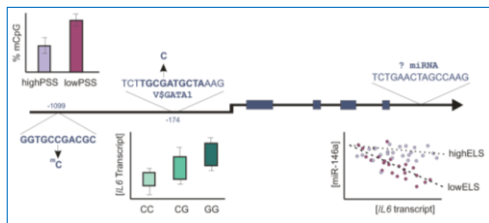
Clinical Diagnostics



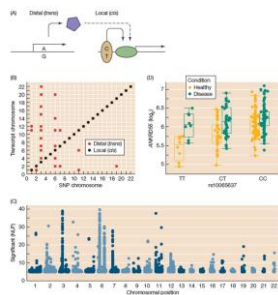
Population Profiling



Gene Regulation



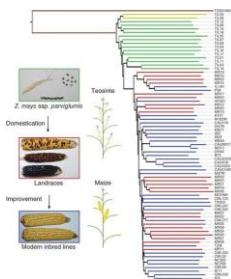
Expression QTL



Expression QTL are polymorphisms that affect the expression of a gene. They are particularly interesting when they overlap with GWAS hits.

Ecology and Evolution

Comparative population genomics of maize domestication and improvement

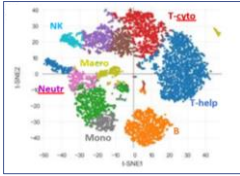


~1000 Domestication and Improvement loci identified with $s > 0.015$. Improvement candidates tend to be more highly expressed in multiple tissues, suggesting selection targeted cis-regulatory sites.

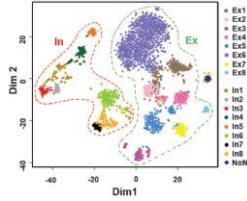
Hufford et al (2012) Nature Genetics 44: 808-811

Single Cell RNASeq

Peripheral Blood Monocytes



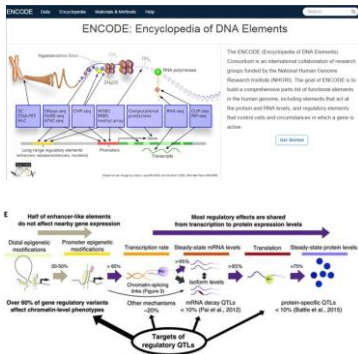
Neuronal nuclei



Zheng et al (2017) Nature Comm 8: 14049

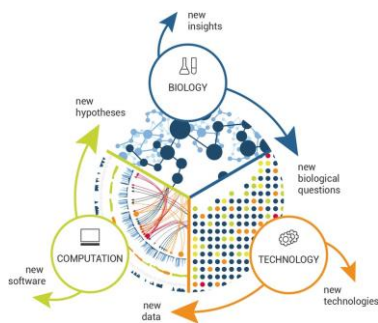
Lake et al (2016) Science 352: 1586-1590

Additional Tiers of Gene Regulation



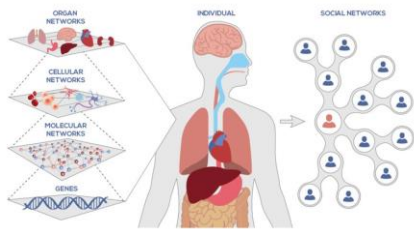
Li et al (2016) Science 352: 600-604

Systems Biology



<http://www.omicscouts.com/en/disease-and-systems-biology.html>

Human Systems Biology



<https://www.systemsbio.org/about/what-is-systems-biology/>

First genome sequencing success story

Diagnosed at age 5 with dopa-responsive dystonia

Worsening respiratory and neuromuscular disease not responsive to dopamine precursor therapy

WGS shows mutation in SPP "sepiapterin reductase" gene

5-HT serotonin precursor supplementation had immediate impact



The Beery twins

Bainbridge, et al. (2011) *Sci. Transl. Med.* 3, 87re3

Some Personal Genomics Companies



Some Public Initiatives

The image shows three screenshots of public initiatives related to genomics. The first is the 'Personal Genome Project' website, which describes a collaborative effort to sequence personal genomes. The second is a 'Live better, together!' website with a navigation menu including 'learn', 'connect', and 'track'. The third is the 'genomes unzipped' website, which provides information about the 100,000 Genomes Project.

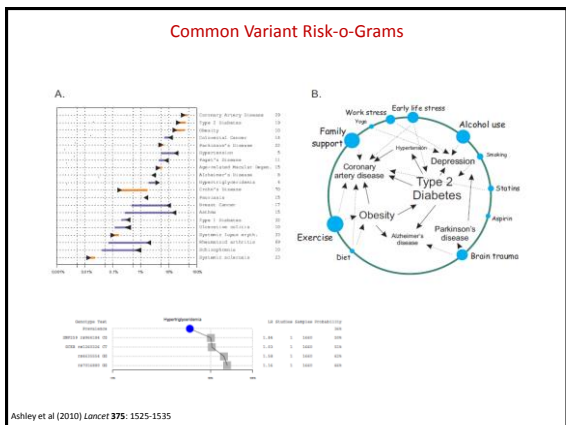
Precision Medicine and Predictive Health

Precision Medicine is molecular pathology based on a patient's genome sequence. It is about finding the mutation or perturbed genetic pathway that is largely responsible for a congenital birth defect, or for a specific cancer.

Predictive Health is about using your own clinical and genomic profile to make better decisions about wellness in an effort to prevent the onset of chronic disease.

Personalized genomic medicine encompasses both, and essentially captures the idea that each person's individual genome sequence will eventually be part of their own medical care.

Common Variant Risk-o-Grams



Then why don't we all get our PGS?

- 1. Cost: \$500 for the lite version, \$3000 the whole
- 2. Privacy: What if WikiLeaks releases it to the world
Can providers sell information to companies?
- 3. Insurance: The Genetic Non-Discrimination Act protects from workplace and health discrimination only
- 4. Anxiety: Will I learn something I don't want to know?
And will there be anything I can do about it?
- 5. Consent: How can I provide informed consent to something I don't really understand? Can I trust my doctor to help me decide?
