

Comparative Genomics and Visualisation

BS32010

1.Introduction



The James
Hutton
Institute

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Types of genome comparison

Course outline



What Is Comparative Genomics?

Comparative genomics is. . .

The combination of genomic data, and comparative and evolutionary biology, to address questions of genome structure, evolution, and function.



Evolution is the central concept

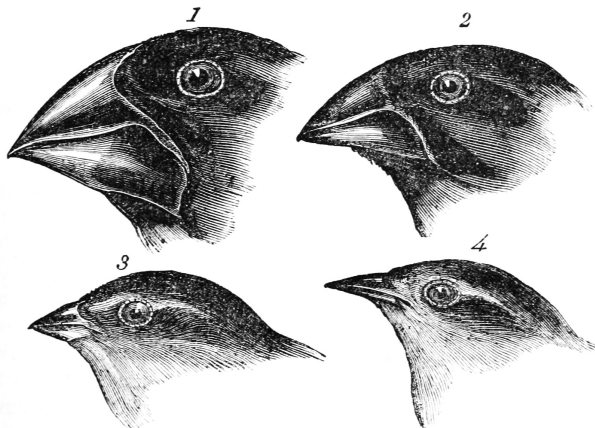
**“NOTHING IN BIOLOGY MAKES SENSE EXCEPT
IN THE LIGHT OF EVOLUTION.”**

THEODOSIUS DOBZHANSKY



Comparison of physical features

How do we determine that features are related, and evolved?



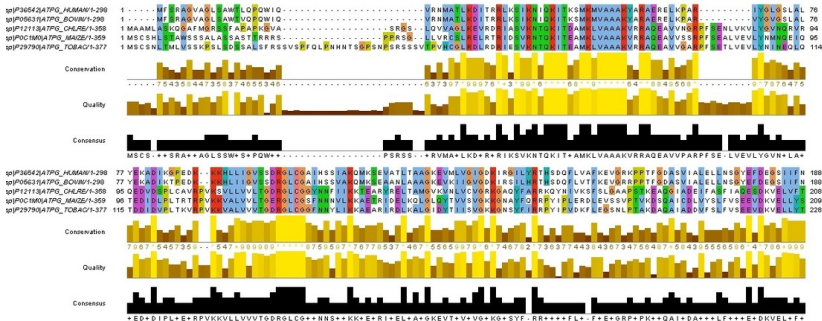
1. *Geospiza magnirostris*.
3. *Geospiza parvula*.

2. *Geospiza fortis*.
4. *Certhidea olivacea*.



Comparison of sequence features

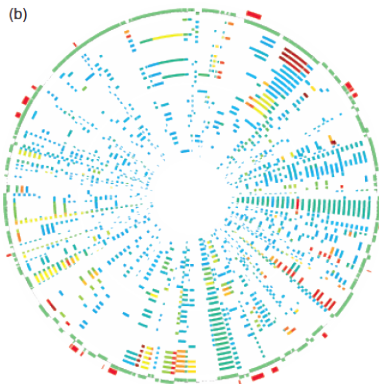
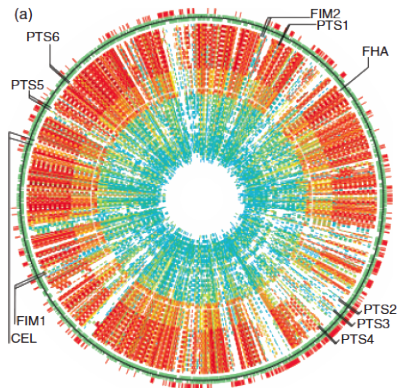
Multiple alignment of ATP synthase





Comparison of genome features

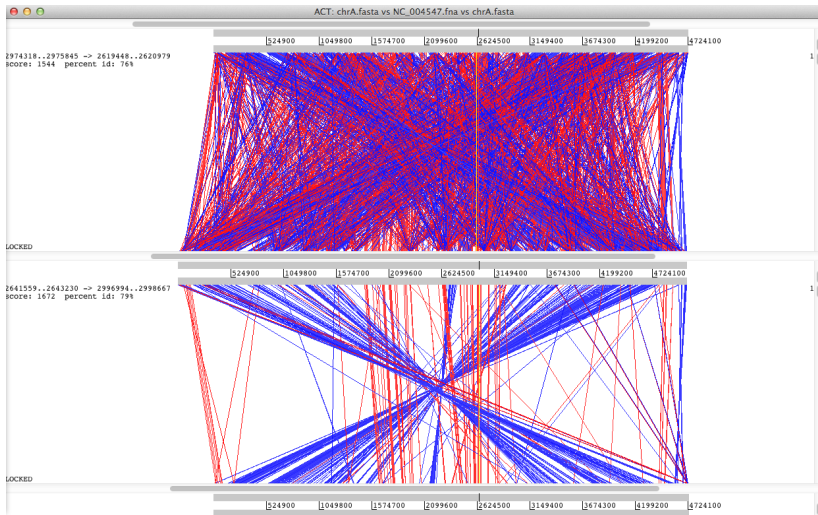
Sequence similarity of individual features





Comparison of genome features

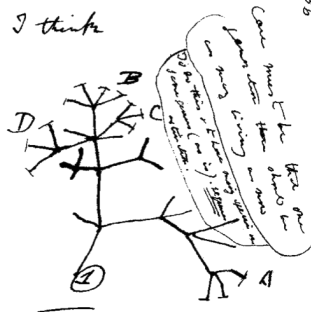
Genome structural rearrangements





Why comparative genomics?

- Genomes describe heritable characteristics
- Related organisms share ancestral genomes
- Functional elements encoded in genomes are common to related organisms

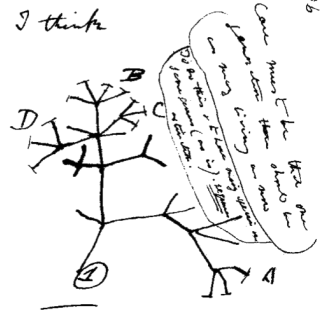


then between A & B. Various
sort of relation. C & B. The
first predation, B & D
rather greater distinction
then genus would be
formed. - binary relation



Why comparative genomics?

- Transfer of functional understanding from model systems (*E. coli*, *A. thaliana*, *D. melanogaster*) to non-model systems
- Genome comparisons can be informative, even for distantly-related organisms

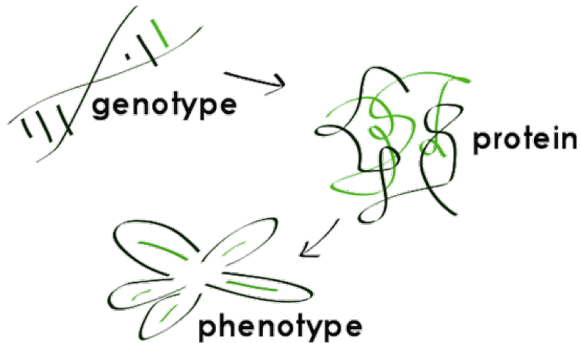


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Genomes are informative, but...

CONTEXT: epigenetics, tissue differentiation, mesoscale systems, etc.



PHENOTYPIC PLASTICITY: responses to temperature, stress, environment, etc.



Genomes to systems

Functional Genomics

- Genomic differences can underpin phenotypic (morphological or physiological) differences.
- Where phenotypes/other organism-level properties are known, comparison of genomes can give mechanistic or functional insight into differences (e.g. GWAS).
- Genomic changes reveal evolutionary processes and constraints.





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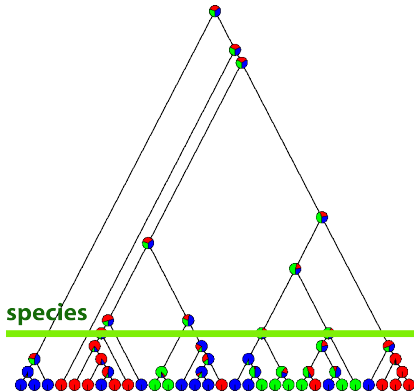
Course outline



Types of comparison

Within species

- e.g. isolate-level (and even within individuals)
- which genome features may account for unique characteristics of organisms/cell-types (e.g. tumours)?
- what epigenetic changes occur in an individual?

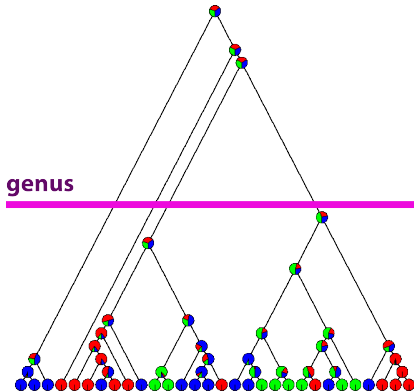




Types of comparison

Within genera/between species

- what genome features show evidence of selective pressure?
- which species are under selective pressure for which functions?



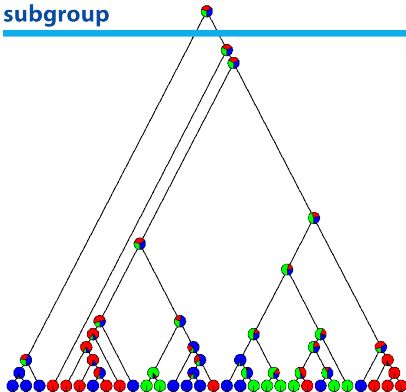


Types of comparison

Between subgroups

- what are the core set of genome features that define a subgroup or genus?
- what functions are present/absent between groups?

subgroup



E. coli LTEE *a b c*



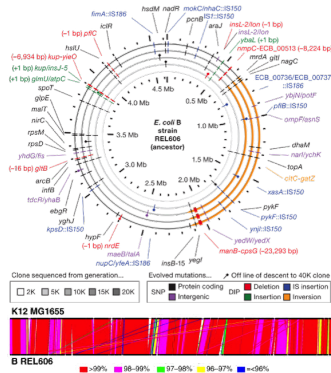
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^aJeong *et al.* (2009) *J. Mol. Biol.* doi:10.1016/j.jmb.2009.09.052

^b Barrick et al. (2009) *Nature* doi:10.1038/nature08480

^cWiser et al. (2013) *Science* doi:10.1126/science.1243357

- Run by the Lenski lab, Michigan State University since 1988 (<http://myxo.css.msu.edu/ecoli/>)
- 12 flasks, citrate usage selection
- >50,000 generations of *E. coli*!
 - Cultures propagated every day
 - Every 500 generations (75 days), mixed-population samples stored
 - Mean fitness estimated at 500 generation intervals





Comparative genomics in the news ^{a b c}

^aWashington Post 23/2/2015

^bIbarra-Laclette *et al.* (2013) *Nature* doi:10.1038/nature12132

^cCarretero-Paulet *et al.* (2015) *Mol. Biol. Evol.* doi:10.1093/molbev/msv020

- *Utricularia gibba*: carnivorous bladderwort
- Genome: 82Mbp, 17,324 genes (wheat: 17bn bases, \approx 94-96k genes)
- Intergenic region contraction (3% repeat elements; most plants: 10-60% repeat elements)
- Genomic context for flowering plants does not require “hidden regulators” (cf. ENCODE)

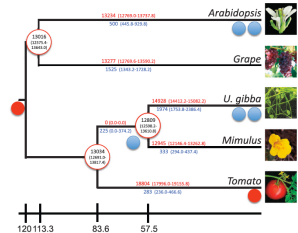




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Levels of comparison

Bulk Properties

- chromosome/plasmid counts and sizes, nucleotide content, etc.

Whole Genome Sequence

- sequence similarity
- organisation of genomic regions (synteny), etc.

Genome Features/Functional Components

- numbers and types of features (genes, ncRNA, regulatory elements, etc.)
- organisation of features (synteny, operons, regulons, etc.)
- complements of features
- selection pressure, etc.



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