ERROR CORRECTION 5 Oct 15

ANNOUNCEMENTS

The 2015 – 2016 Saul O Sidore Memorial Lecture Series

Personal Genomic **Medicine:**

Social, ethical, and scientific implications

With the success of the Human Genome Project and advances that permit individuals to have their genetic code determined, the era of personal genomics is already upon us. Leading scholars representing multiple areas of human genome and microbiome research will outline and navigate the current state of knowledge. The series will explain how the genomic revolution will affect our lives, and will stimulate debate of the scientific, medical, ethical, legal, and societal implications of sequencing human genomes.

The Saul O Sidore Memorial Lecture Series was established in 1965 in memory of Saul O Sidore of Manchester, New Hampshire. The purpose of the series is to offer the University community and the state of New Hampshire programs that raise critical and sometimes controversial issues facing our society. The University of New Hampshire Center for the Humanities sponsors the programs



http://cola.unh.edu/center-humanities/sidore

Image of Smithsonian e:

14 October

Gut Feelings: How the Microbiome Influences Behavior Dr. Jane A. Foster, Brain-Body Institute, McMaster University

7 December

Personalized Medicine: Using Integrative Omics to Analyze Complex Disease and Manage Health Dr. Michael Snyder, Director, Stanford Center for Genomics and Personal Medicine

10 February

Implementing 'Precision' Medicine: Ethical Concerns in a Postgenomic World Dr. Barbara Koenig, Institute for Health and Aging, UCSF

9 March

The Dog Genome: Shedding Light on Human Diseases Dr. Elaine Ostrander, Head of Comparative Genetics, NIH

13 April

The Invisible Influence of the Human Microbiome Dr. Jack A. Gilbert, University of Chicago

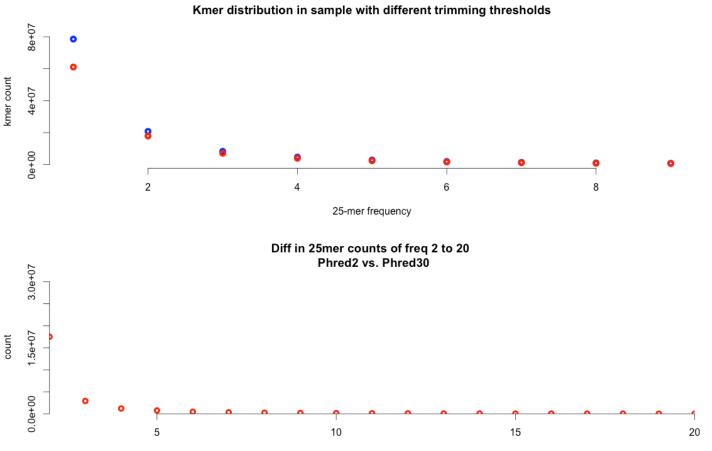
27 April

Probing Human Ancestry with Ancient DNA

Dr. John Hawks, University of Wisconsin

All lectures held at 3 PM in the University of New Hampshire Memorial Union Building Theaters

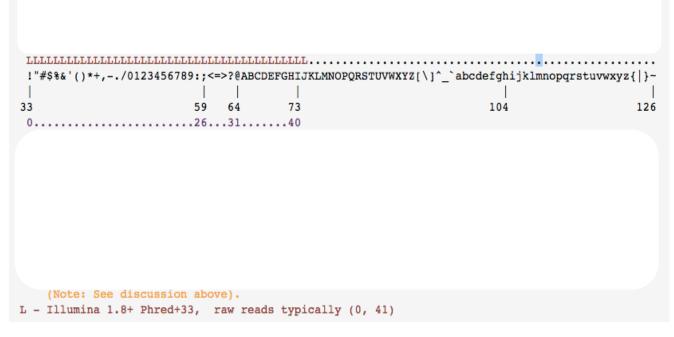
LAB REVIEW

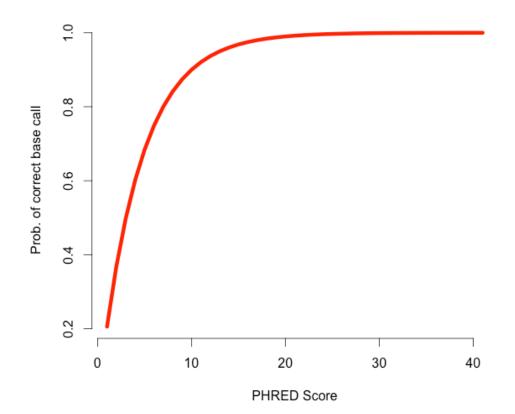


kmer frequency

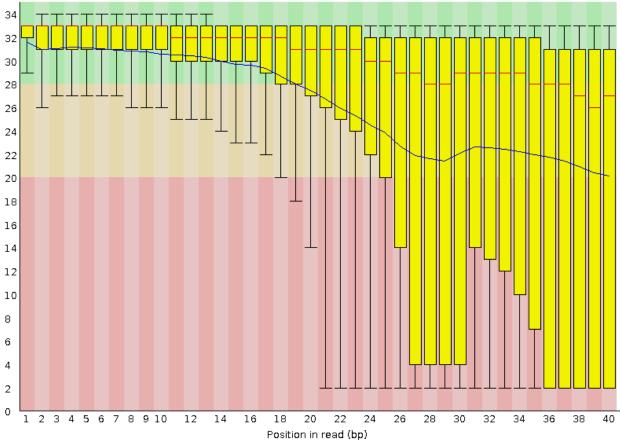
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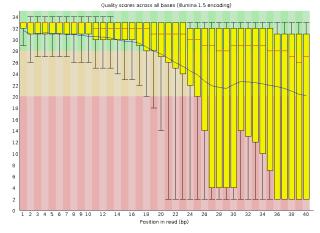
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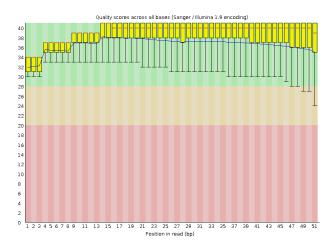




Quality scores across all bases (Illumina 1.5 encoding)







ERROR CORRECTION

MSA based

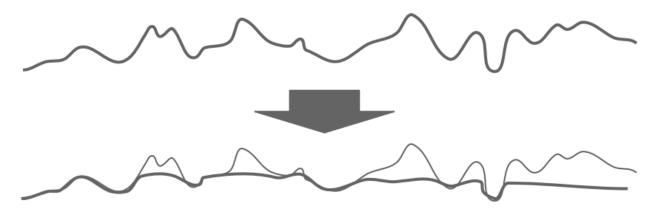
ERROR CORRECTION

Evaluation of Correction

High-throughput seq: 2 major challenges

Perfect Storm of data analysis – What to do???

Perfect Storm of data analysis – What to do???



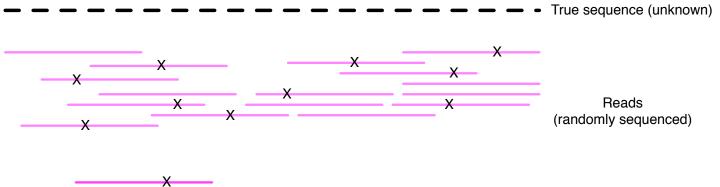
Brown 2012 arXiv:1203.4802v2

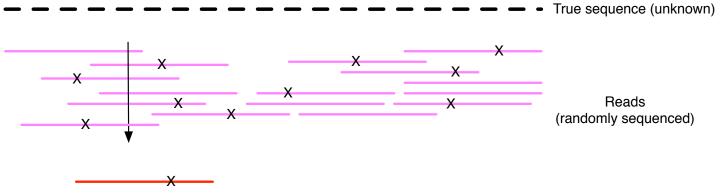
Х

- True sequence (unknown)

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Reads (randomly sequenced)





If next read is from a high coverage region - *discard*

