Epigenetics of methylation BIOL 426/626 Approaches to Molecular Biology

Class 20: Epigenetics



Learning Goal

 To understand the dynamics of DNA modification during development and the methods used to analyze the modification state genome wide with single nucleotide accuracy

Learning Objectives

- Explain the role of CpG island methylation in regulating gene expression globally in the genome
- Describe how bilsuphite sequencing is used to identify sites of DNA methyaltion
- Summarize how patterns of methylation can vary based on sex, developmental stage or tissue type
- Discuss the role of DNA methylation in suppressing transposition by retroelements in the genome
- Explain how DNA methylation is used for the epigenetic control system called imprinting

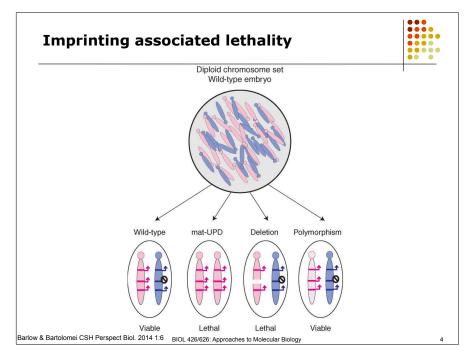
• Reading assignment:

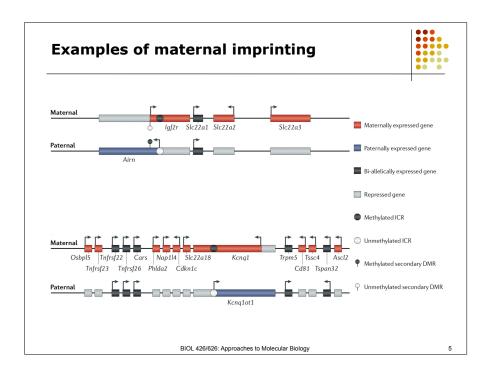
• Smith et al. A unique regulatory phase of DNA methylation in the early mammalian embryo. Nature. 2012 484:339-344.

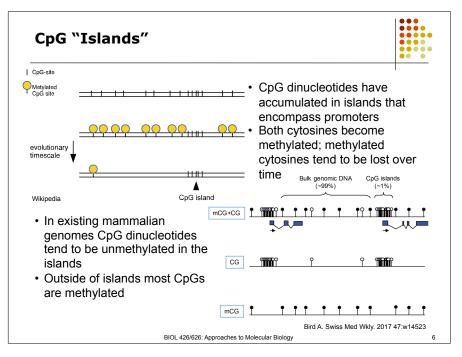
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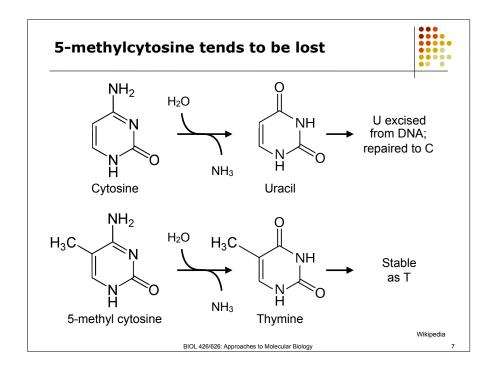
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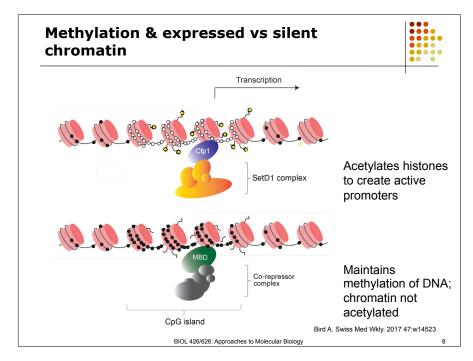


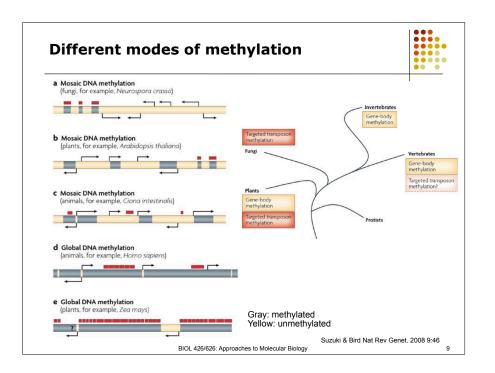


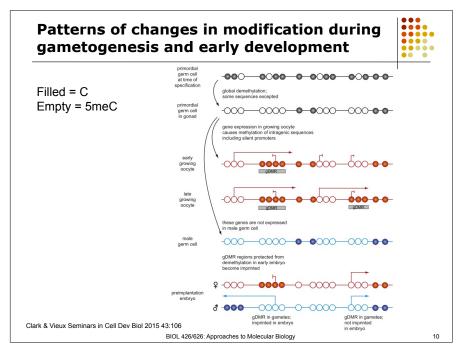


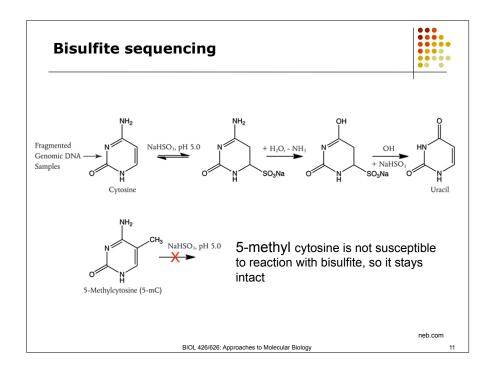


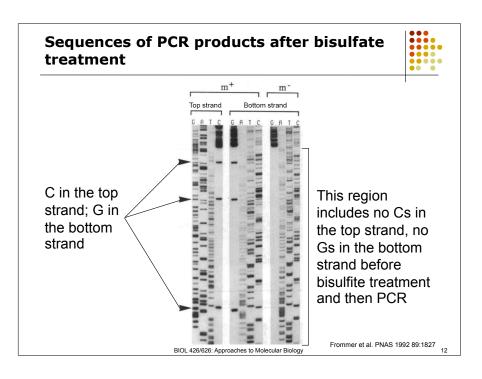


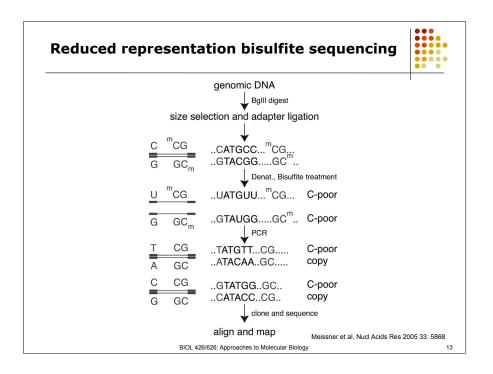


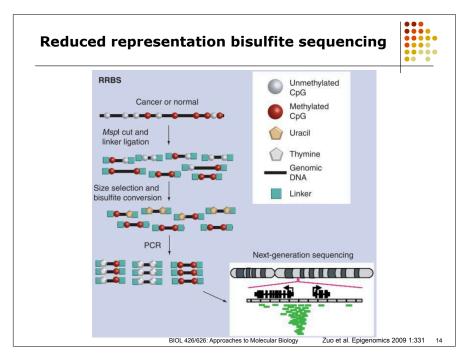




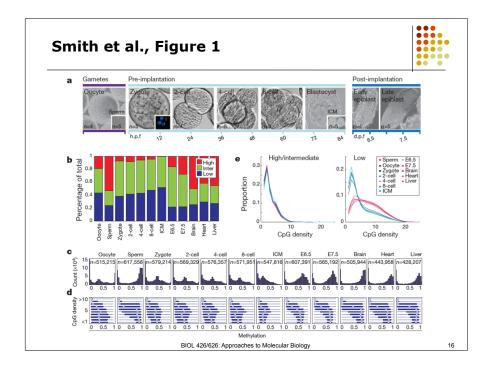


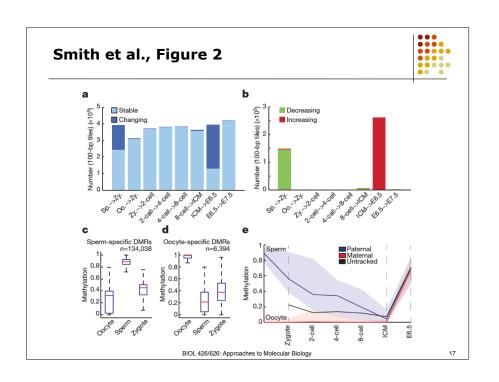


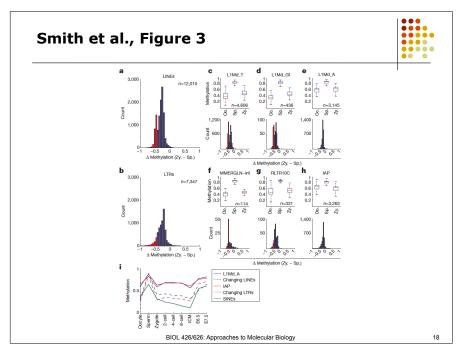


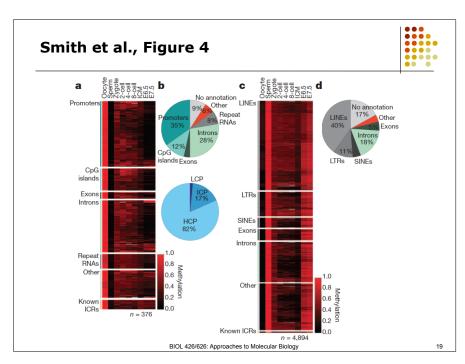


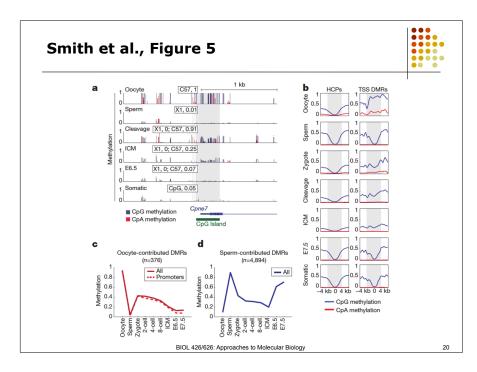












Reading for next time:



- Dale et al. From Genes to Genomes, Chapter 11
- Rydning et al. Conditional transgenic technologies. J. Endocrinol. 2001. 171, 1–14

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