Methylation and Human Disesases

Joint Graduate Seminar

The Chinese University of Hong Kong

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What is Methylation?

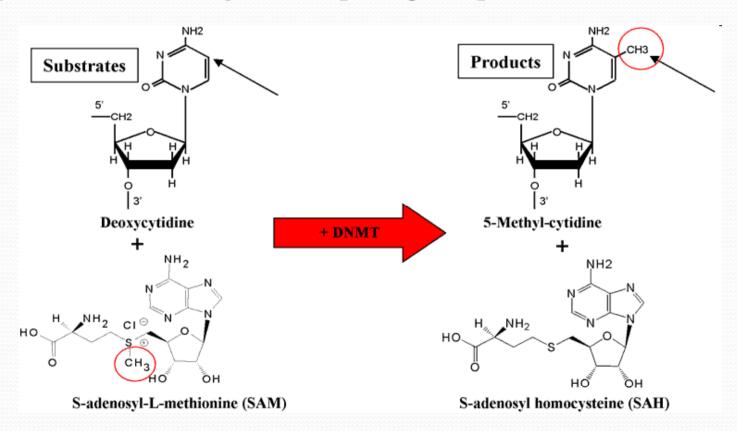
- Epigenetics
- Heritable, stable, reversible
- Catalyzed by DNA Methyltransferases (DNMTs)

DNMT3A DNMT3B Maintenance DNMT De novo DNMTs

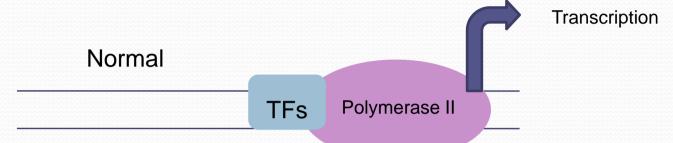
DNA methyltransferases

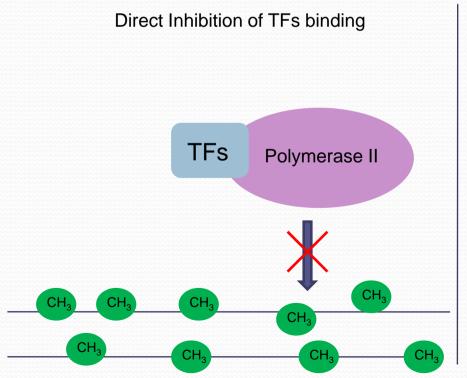
Where Does Methylation Occur?

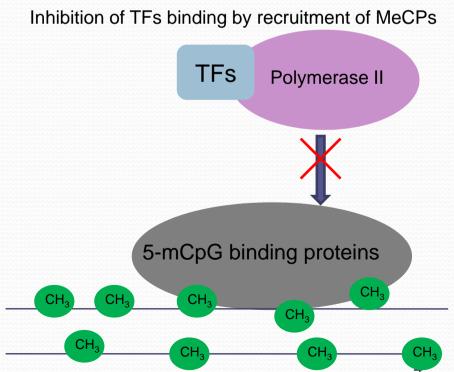
- At cytosine residues of 5'-CpG-3'
- CpG islands mostly encompass gene promoters



Silencing Effect of Methylation







Roles of Methylation

- Essential in normal development
 - Regulates gene expression and chromatin structure
 - Genomic imprinting
 - X-inactivation in females
 - Defense mechanism
- Deletion of any DNMTs is lethal in mice

Genetic Diseases Caused by Methylation Defects

- Mutations in proteins involved:
 - ICF
 - Rett Syndrome
- Defects of imprinted genes on chromosome **15q11-13**:
 - Prader-Willi Syndrome
 - Angelman Syndrome
- Loss of imprinting:
 - Beckwith-Wiedemann Syndrome

ICF Syndrome

(Immunodeficiency, Centromeric instability, Facial anomalies)

Mutation in DNMT3B

- Combined immunodeficiency
- Developmental delay
- Mental retardation

Rett Syndrome (RTT)

- Mutation in MeCP2
- X-linked neurodevelopmental disorder

- Mental retardation
- Autistic behaviour
- Stereotypical hand movements
- Seizures





Prader-Willi Syndrome (PWS)

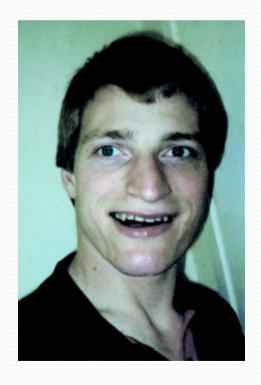
- Mental retardation
- Slow moving
- Obesity
- Hypogonadism
- Small hands and feet

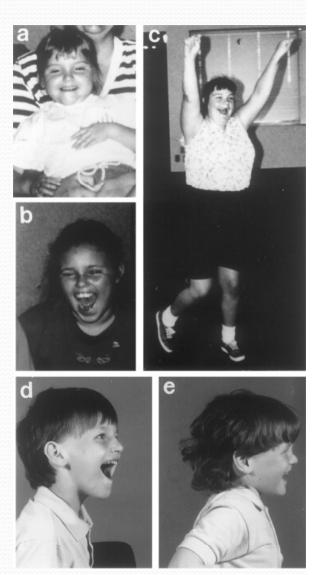


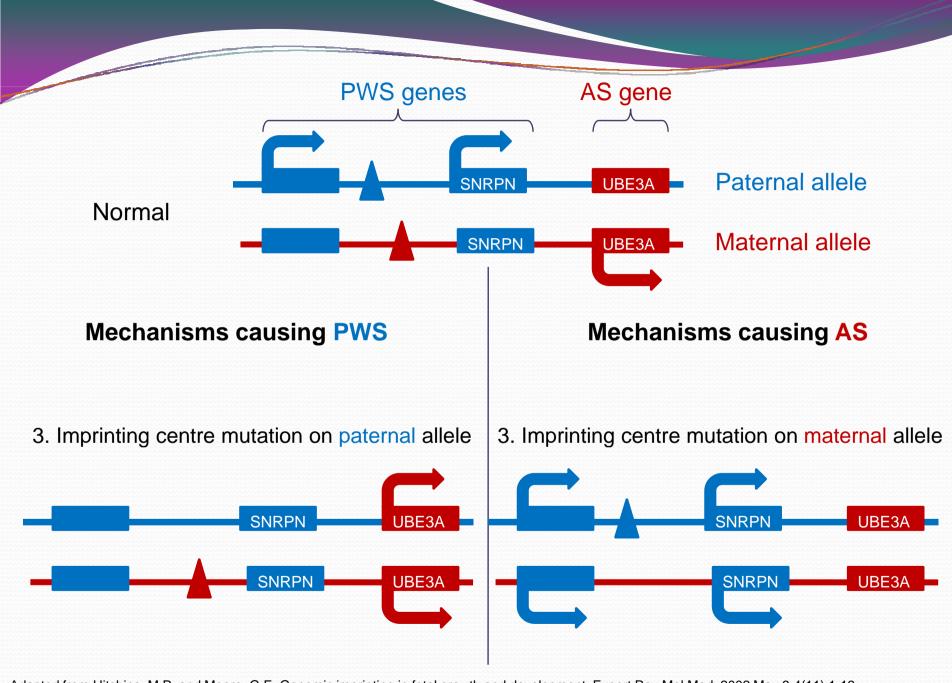
http://medgen.genetics.utah.edu/

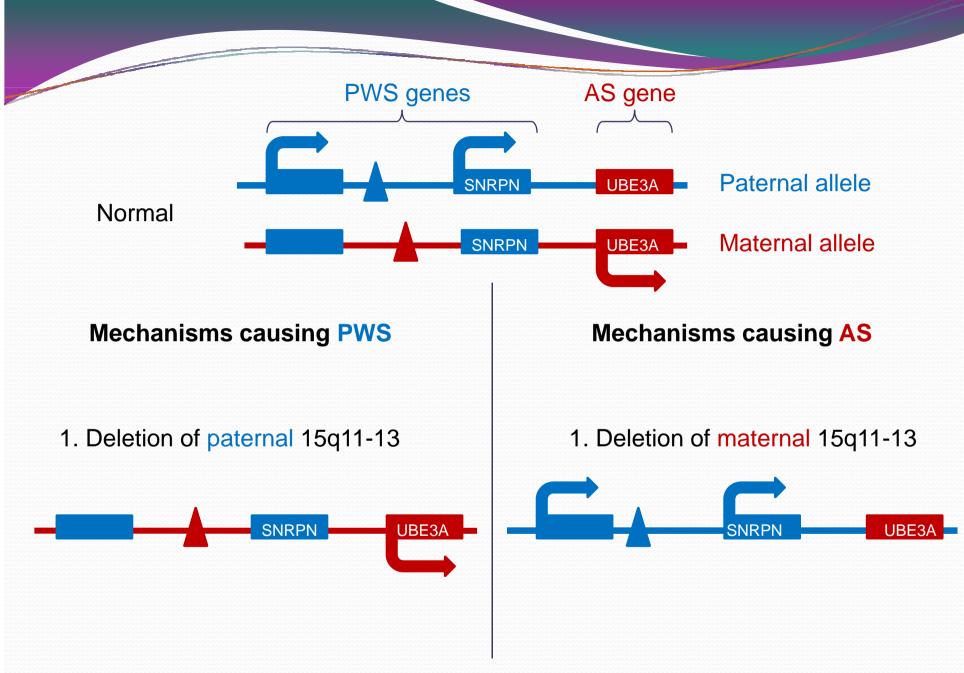
Angelman Syndrome (AS)

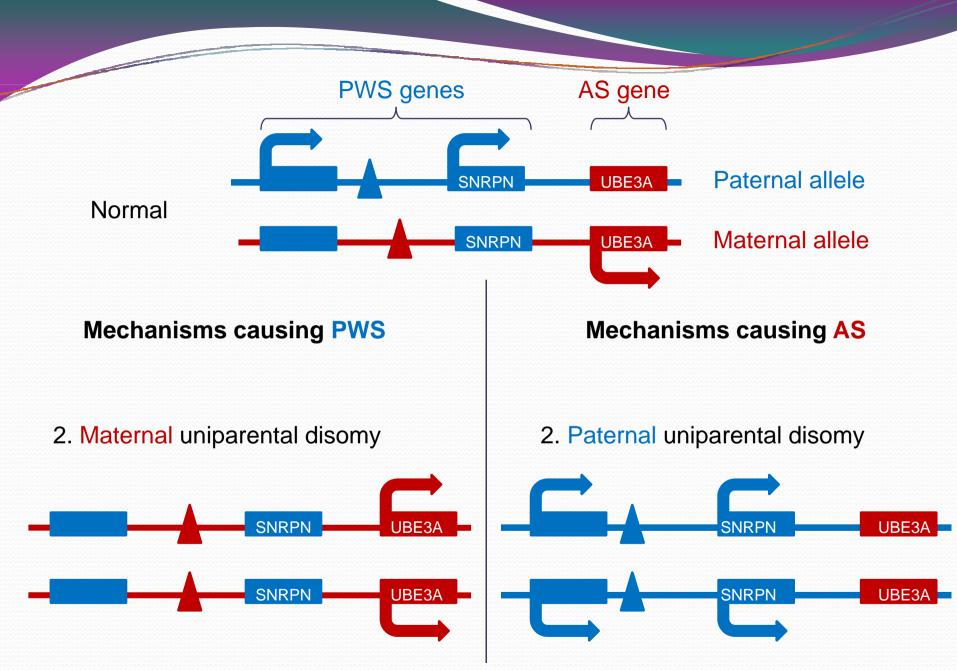
- Severe mental retardation
- Hyperactive
- Disordered movement
- Uncontrolled laughter
- Dysmorphic facial features











Beckwith-Wiedemann syndrome (BWS)

• Loss of imprinting of IGF2 gene

- Macroglossia (large tongue)
- Macrosomia (birth weight and length >90th percentile)
- Umbilical hernia (midline abdominal wall defects)
- Ear creases / ear pits







Methylation and Cancers

- Cancers once viewed as genetic disorders
- Epigenetic events also responsible
- 2 types of methylation pattern defects:
 - Genome wide <u>hypo</u>methylation
 - Gene specific <u>hyper</u>methylation

Hypomethylation









Oncogene activation

Retrotransposons activation

Chromosome instability

Loss of imprinting

Loss of Imprinting of IGF2 in cancers

Normal

Cancers (e.g. colorectal cancer, Wilm's tumour)

Methylation

1

Imprinted

1

Expression of IGF2 from paternal allele

Hypomethylation

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Loss of imprinting

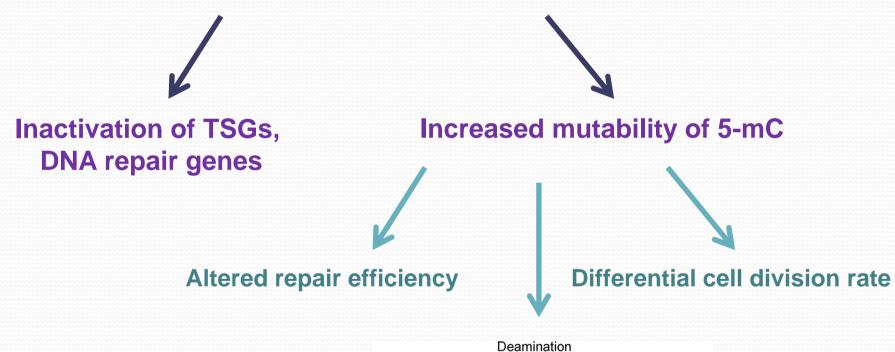
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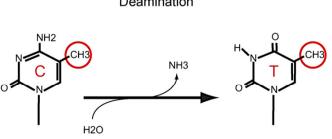
Expression of IGF2 from both alleles



Transformation

Hypermethylation





Most Frequently Reported Genes Hypermethylated in Cancers

TSGs	Function	Cancer type with hypermethylation
p16-INK4	Cell cycle arrest	Lung, liver, leukemia, lymphoma
RASSF1	Cell cycle regulation	Ovarian, thyroid, cervical, kidney
MGMT	DNA repair, drug resistance	Brain, colorectal, lung, gastric
CDH1	Cell-cell adhesion	Gastric, lung, leukemia
MLH1	DNA mismatch repair	Gastric, colorectal
APC	Apoptosis	Lung, breast, prostate

Methylation and Oncogenic Viruses

- Viral oncoproteins can silence host genes that interfere with cancer progression
 - E.g. LMP1 of EBV activates DNMTs and downregulates E-cadherin
- Hypermethylation identified in viral genomes
 - E.g. EBV, HPV, HBV
- Possible roles:
 - Evasion of host immune system
 - Establishment of latent infections

Methylation Patterns as Cancer Biomarkers

- Defects found in almost all cancer types
- Patterns specific in different tumours
- Possible biomarker for early detection of cancers
- Predict metastatic potential and therapy sensitivity

Epigenetic Therapy

- Reversibility of epigenetic events can be targets of therapy in cancer
- DNMT inhibitors:
 - 5-Aza-cytidine
 - 5-Aza-2'-deoxycytidine
- In vitro, diminish activities of DNMTs and induce demethylation of TSGs

The End Thank You!