



# Clinical epigenetics and its relation to ART

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## Epigenetics and ART

- Basics of epigenetics
- Genomic imprinting
- Human imprinting disorders
- Review data on imprinting disorders in ART children



ESHRE workshop PORTO



16-9-2010

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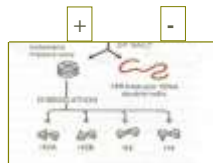
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## What is epigenetics?



Chromatin structure:  
histones and DNA



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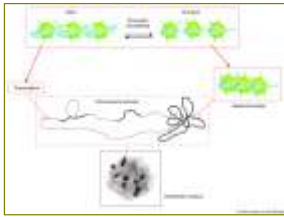
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# What is epigenetics?



gene expression      gene silencing




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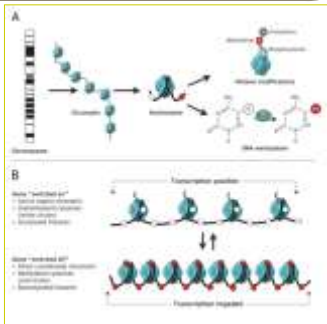
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# Basic epigenetic mechanisms

“epigenetics”:  
refers to modifications on top of DNA and histones that will influence gene expression patterns without changes in the DNA code




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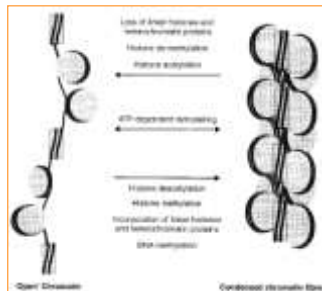
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# Basic epigenetic mechanisms

euchromatin      heterochromatin



gene expression      gene silencing




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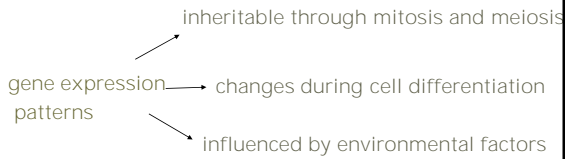
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## What is epigenetics?

### “epigenetics”:

study of changes in gene expression that occur without changes in the DNA code



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## Genomic imprinting

most genes: biallelic expression or silencing  
expression of maternal **and** paternal alleles



imprinted genes: monoallelic expression  
expression of maternal **or** paternal alleles



Maternal copy is expressed

Paternal copy is expressed



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## Genomic imprinting

- > 80 imprinted genes (0.1 - 1% of all genes)
- Eutherian mammals, marsupials, higher plants
- Key role in embryonic growth and placental function, cognition and maternal behaviour
- Growth regulation: paternally expressed genes enhance growth, maternally expressed genes repress growth
- Defective imprinting involved in carcinogenesis and in human genetic diseases



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# Genomic imprinting

Mouse imprinting genes, regions and phenotypic effects



<http://www.mgu.har.mrc.ac.uk/research/imprinting/>

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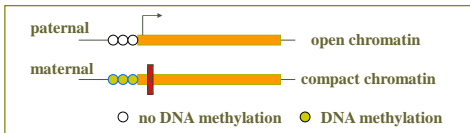
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# Genomic imprinting

- Imprint Control Regions (ICR) control imprinting of the clusters
- Maternal and paternal alleles carry different epigenetic modifications (“imprints”)




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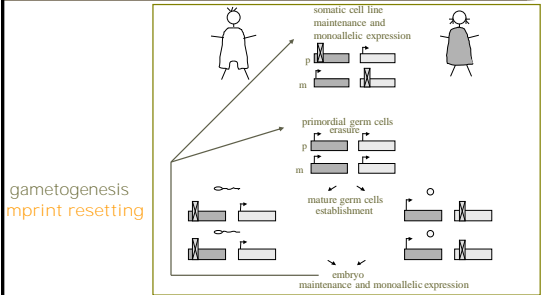
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# Genomic imprinting



gametogenesis  
mprint resetting

early embryonic development  
imprint maintenance

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## Epigenetics and ART

Does ART interfere with

- epigenetic reprogramming (imprint maintenance) in the early embryo?
- epigenetic reprogramming (imprint resetting) during gametogenesis?



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## Epigenetics and ART

Recent studies have presented (limited) evidence for a higher risk of imprinting disorders after ART

Imprinting disorders: altered expression of imprinted genes

- epigenetic mechanisms
- genetic mechanisms (large duplications or deletions, mutations in imprinted genes and ICRs, uniparental disomy)



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## Angelman syndrome

- OMIM 105830
- Incidence: 1/15.000
- Neurogenetic disorder
  - severe mental retardation
  - ataxia
  - "happy puppet syndrome"
  - absence of speech



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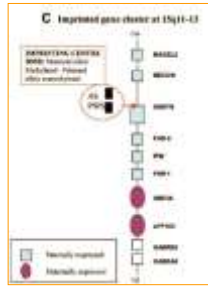
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## Angelman syndrome

### Angelman syndrome

Inheritance: most cases are sporadic

- 85-90 % genetic defects
- < 5 % epigenetic defects
- => loss of *UBE3A* expression




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## ART and AS

Syndrome	# cases	ART	country	reference
AS	2	ICSI	Germany	Cox et al., 2002
AS	1	ICSI	Norway	Orstavik et al., 2003

molecular analysis: an epigenetic defect (loss of maternal methylation at the IC of imprinted region on chr15q11-13) was found in all three cases

- + 1 case in Ludwig et al., 2005
- + 1 case in Sutcliffe et al., 2006




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## Beckwith-Wiedemann syndrome

- Incidence: 1/13700
- Overgrowth syndrome with predisposition for embryonal tumours
- Caused by genetic or epigenetic defects in an imprinted region on chr11p15
- Inheritance:
  - 15% familial cases, autosomal dominant
  - 85% sporadic cases:
    - 20 % genetic (pUPD, mutations)
    - 60 % epigenetic (aberrant methylation)
    - 20 % unknown




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## Beckwith-Wiedemann syndrome

**Beckwith-Wiedemann Syndrome**

**A Clinical Synopsis**

- Pre and/or postnatal overgrowth
- Macroglossia
- Anterior abdominal wall defect (e.g. omphalocele, umbilical hernia)
- Neonatal hypoglycemia
- Hypogenitalism
- Endocrinological tumors (~7% of patients)
- 10% familial

**B Molecular Genetic Findings**

- 2% Paternal duplication or maternal rearrangement of chromosome 1p15.5
- 20% Uniparental (paternal) disomy chromosome 11
- 7% CDKN1C mutation (40% of familial cases)
- 40% Imprinting center 2 (IC2) defect (hypermethylation, loss of maternal methylation)
- 3% Imprinting center 1 (IC1) defect (gain of maternal methylation)
- 1% Imprinting centers (IC1 or IC2) deletion

**C Imprinted gene cluster at 11p15.5**

IC1/DMR1: Maternal allele unmethylated, Paternal allele methylated

IC2/DMR2: Maternal allele methylated, Paternal allele unmethylated

Legend: Red square = Paternally expressed, Blue square = Maternally expressed

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## ART and BWS

ART/ cohort	study	#LOM/ #tested	#ICSI/ #IVF	RR	country	reference
4/3/65*	retro. prosp.	5/6	5/2	x 6	USA	DeBaun et al., 2003
6/149	retro.	2/2	3/3	x 4	UK	Maher et al., 2003
6/149	retro.	6/6	2/4	x 3	France	Gicquel et al., 2003
4/37	retro. case-contr.	3/3	1/3	x 9	Australia	Halliday et al., 2004
	retro.	24/25	13/12	-	UK	Lim et al., 2009
11/79	retro.	8/8	5/1/5HT		UK	Sutcliffe et al., 2006

RR = relative risk; HT = hormonal treatment  
LOM = loss of maternal methylation \* overlapping patients

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## ART and BWS

Association between  
ART and BWS to be considered

- epidemiological data
- molecular analysis: an epigenetic defect (loss of maternal methylation at BWSIC2) in nearly all BWS patients after ART
- general population: epigenetic defect in 50-60% of cases

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## ART and imprinting disorders

- Insufficient evidence for association between ART and other imprinting disorders (SR, PWS, global hypomethylation syndrome, RB)
- limited evidence for association between ART and AS and BWS
- absolute risk is low



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## Cause of imprinting disorders after ART

### \* in-vitro culture systems

- Khosla et al. 2001: culture (serum) of preimplantation mouse embryos reduced fetal development (lower birthweight) and expression of imprinted genes
- Young et al. 1998: after exposure *in vitro* unusually large offspring syndrome (LOS) in relation with imprinted genes
- Young et al. 2000: epigenetic changes in *IGF2R* are associated with fetal overgrowth (LOS) after sheep embryo culture

The mammalian preimplantation embryo is very sensitive to culture conditions; epigenetic changes induced at the early stages may lead to altered phenotypes at later stages



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## Cause of imprinting disorders after ART

### \* use of hormonal ovarian stimulation

- mouse: comparison of methylation in 4 imprinted genes in normal vs superovulated GVs: no differences for maternally methyl. genes, but gain of methylation for H19 after superovulation  
*Sato et al., 2007*
- human: superovulated GV & MI oocytes show gain of H19 methylation (2/6) and loss of maternal methylation at PEG1 (6/16)  
*Sato et al., 2007*
- human: methylation analysis of KvDMR1 in GV, MI & MII superovulated oocytes shows loss of methylation in 1/16 oocytes  
*Geuns et al., 2007*



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## Cause of imprinting disorders after ART

### \* parental infertility

- Doornbos *et al.*, 2007: correlation between ART, fertility problems and epigenetic defects (cohort 220 AS, BWS, PWS patients)

	cohort	general population
Born after ART	6.4%	2.1%
Fertility problems (no ART)	6.8%	3.5%




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## Cause of imprinting disorders after ART

### \* parental infertility

subfertile couples are predisposed to epigenetic defects  
imprinting defects and subfertility have a common cause

- AS cohort study: subfertile couples have an increased risk of conceiving a child with an imprinting defect  
TTP > 2 years, no therapy RR 6.25  
TTP > 2 years, treatment RR 12.5 *Ludwig et al., 2005*




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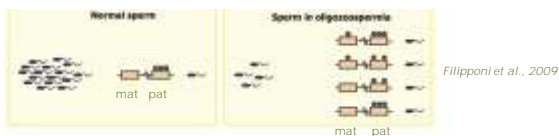
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## Cause of imprinting disorders after ART

### \* parental infertility

Link between imprinting errors and disruptive spermatogenesis



- Incomplete establishment of imprints (incomplete DNA methylation of *H19*) in sperm from men with fertility problems  
*Marques et al., 2004 & 2008 & 2009*




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## Cause of imprinting disorders after ART

### \* parental infertility

Link between imprinting errors and disruptive spermatogenesis

- Aberrant DNA methylation at imprinted loci in men with fertility problems *Kobayashi et al., 2007*

- Aberrant DNA methylation at imprinted loci in 17/78 paired DNA samples (conceptus and father)  
the DNA methylation errors were inherited from the father (7/17)  
they were associated with DNA sequence variants in DNMT3L and more prevalent in oligozoospermic men *Kobayashi et al., 2009*



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## Epimutations associated with ART

- in-vitro culture systems
- hormonal stimulation
- parental infertility

imprinted genes: vulnerable

- functionally haploid
- differential epigenetic modifications
- embryonic and placental development



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## Iceberg theory

epigenetic defects



- ↳ imprinting disorders (congenital syndromes) (tip of iceberg)
- ↳ neurodevelopmental delay
- ↳ cancer predisposition
- ↳ IUGR
- ↳ increased risk for cardiovascular diseases



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## Iceberg theory

- No higher risk of childhood cancer  
Doyle et al. 1998, (UK n= 2057), Bruinsma et al. 2000, (Australia, n = 5249), Klip et al. 2001, (the Netherlands, n = 9484), Lerner-Geva et al. 2000, (Israel n = 332)
- Lidegaard et al., 2005:
  - systematic follow up of 6052 ART singletons (1995-2001) till 2002 vs a control group of 442.349 non-ART
  - the incidence rate of childhood cancer, mental diseases, imprinting syndromes and developmental disturbances was equal in the two groups



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## Iceberg theory

- *Ceelen et al., 2008*: follow-up of IVF children vs controls born to subfertile couples
  - IVF children show higher blood pressure levels
- ART is associated with an increased incidence of **Low Birth weight**  
maybe as a consequence of embryonic/fetal epigenetic programming in response to early adverse environmental factors and stress???



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## Iceberg theory

- *Katari et al., 2009*: changes in DNA methylation and gene expression patterns of imprinted and non-imprinted genes in cord blood and placenta of ART children (n=10) versus spontaneously conceived children (n = 13)
  - > phenotype?
  - > long term: increased risk for metabolic and cardiovascular diseases?



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## Conclusion and perspectives

- Epidemiological data: increased relative risk for AS and BWS after ART but absolute risk is low
- need for long-term follow up
- Prospective multi-centre studies and record data on infertility history, hormonal stimulation, *in vitro* culture conditions
- Need for basic epigenetic and imprinting research
  - in animal models
  - human gametes and embryos



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