Genome Stability, Genetic Diseases and Cancer

- Lecture 1 DNA damage. Damage Reversal. Base excision repair. Mismatch repair
- Lecture 2 Nucleotide excision repair: cellular and clinical aspects Nucleotide excision repair: genes and proteins
- Lecture 3 Replication of damaged DNA. Mutagenesis and carcinogenesis

Learning outcomes (Lecture 1a)

Understanding:

- Different types of DNA damage
- Three examples of ways in which cells can reverse DNA damage in situ
- Basic mechanism of Base Excision Repair



DNA Damage

	UV	lonizing Radiation	Monofuctional Chemicals	Bifunctional Chemicals
Non distorting chemical damage	-	+	+	+ -
Minor distorting chemical damage	+	+	+	+ -
Major distorting chemical damage	+	-	+	+
Interstrand cross links	-	-	-	+
Strand breaks	-	+	+	+ -



Methylated purines



7-methylguanine

NH2 Т CH₃ 3-methyladenine







1-methyladenine





O-6-methylguanine

1.3

Aspects of DNA repair

- 1. Initial damage
- 2. Repair of damage
- 3. Genes involved
- 4. Mechanism of action of gene products
- 5. Replication of unremoved damage. Cell cycle progression.
- 6. Biological consequences of damage, repair and failure to repair.

Damage reversal

1. Photoreactivation





Friedberg et al, 2005 DNA Repair and Mutagenesis

Photolyase mechanism



Damage reversal 2. Repair of O6-methylguanine





Mispairing of O-6-methylguanine with thymine

Dual activities of Ada methyltransferase



Friedberg et al, 2005 DNA Repair and Mutagenesis



Ogt gene is not inducible

1.10

Alkyltransferases in mammalian cells

- Similar mechanism to *E. coli,* but for O-6-meG alone, like Ogt, not inducible.
- K/o mouse constructed, very sensitive to carcinogenesis by methylating agents.
- Conversely transgenic mice bearing MGMT gene are more resistant.
- Many cancer cell lines are Mex⁻. MGMT silenced by methylation in about 50% of tumours.
- Mex⁻ cells are sensitive to killing and mutagenesis by alkylating agents.
- Many cancer therapy drugs are alkylating agents, eg temozolomide.
- Patrin2 binds MGMT and depletes it. Currently in clinical trials together with temozolomide.

Damage reversal 3. Oxidative demethylation (A4)



Mechanism



Friedberg et al, 2005 DNA Repair and Mutagenesis

1.12

Base Excision Repair

Deamination of bases



8-hydroxyguanine

5



DNA glycosylases						
Enzyme		Size Chromosome (aa) location of		Altered base removed from DNA		
E. coli	Human		gene			
ung	UNG2	313	12q23-q24	U and 5-hydroxyuracil (rep fork)		
	MUG	410	12q24.1	U or T opposite G, ethenocytosine		
	hSMUG1	270	12q13.1-q14	U (from G:U mismatches)		
	MBD4	580	3q21	U or T opposite G at CpG sequences		
Fpg (MutM)	hOGG1	345	3p25	8-oxo G opposite C, formamidopyrimidine		
MutY	MYH	521	1p32.1-p34.3	A opposite 8-oxo G		
Nth	hNTH1	312	16p13.2-	Thymine glycol, cytosine glycol, dihydrouracil, formamidopyrimidine		
AlkA and Tag	AAG	293	16p (near telomere)	3-MeA, ethenoadenine, hypoxanthine		
Nei	Neil 1			Oxidised pyrimidines (rep fork)		
	Neil2			Oxidised pyrimidines		
	Neil3					

3-d structures of glycosylases



FROMME AND VERDINE, FIG. 4. (continued)

Protection from 8-oxoguanine



6

Base Excision-Repair



7

3-d structures of APendonucleases



APE1

ENDO IV

Summary (Lecture 1a)

- DNA damage can cause distortions of different severity
- With visible light, UV damage is repaired by photoreversal (not in placental mammals)
- O6-methylguanine is repaired by a specific methyltransferase
- 1-methyladenine and 3methylcytosine are repaired by oxidative demethylation
- Spontaneous lesions are removed by Base Excision Repair

Learning outcomes (Lecture 1b)

Understanding:

- Detailed mechanism of mismatch repair in *E. coli* and eukaryotes
- How mismatch repair is important both for cancer protection and cancer therapy

Mismatch Repair (A5, A6)

- DNA polymerases replicate DNA very faithfully. Accurate insertion Associated 3'-5' exonuclease for proof-reading Error rates c. 10⁻⁶ or less
- But genomes are big: *E. coli* 3x10⁶ bp, mammals 3x10⁹
- Errors can be single base mismatches or small insertions or deletions caused by base slippage
- Mismatches are repaired by the MMR system which recognises the mismatched bases
- But there's a problem

Methylation-directed mismatch repair



Dam- strains (methylation deficient) are mutators

Mismatch recognition and strand discrimination in *E. coli*

MutH, MutL and MutS⁻ strains are mutators



1.17

3-D structure of MutS



Jiricny, Current Biology, 2000



Activities of MutS

Late steps in MMR in *E. coli*



Friedberg et al, 2005 DNA Repair and Mutagenesis

Human homologues of MutH,L,S

MutS:	
Msh2	MMR
Msh3	MMR
Msh4	Meiosis
Msh5	Meiosis
Msh6	MMR

MutL:	
Mlh1	MMR
Mlh2	?
Mlh3	MMR
Pms1	?
Pms2	MMR (= Pms1 in yeast)

MutH:

No homologues Neither yeast nor Drosophila has methylated DNA

Strand discrimination based on nicks/ends in daughter DNA MMR proteins interact with PCNA at replication fork

Mismatch Repair in eukaryotes



1.21

Mismatch Repair in eukaryotes



Modified from Jiricny, Nature Rev Mol Cell Biol 2006

Microsatellite instability in tumour tissue from HNPCC (Hereditary non-polyposis colon carcinoma)



Aaltonen, et al. Science 260, 812 (1993)



2-hit tumour suppressor model

Most HNPCC result from mutations in hMsh2 or hMlh1

Extracts of **tumour cells** are deficient in MMR of dinucleotide loops and single base mismatches



1.24

Microsatellite instability results from loss of Mismatch Repair



- Microsatellite instability is a useful diagnostic tool. It's not the cause of the cancers
- Cancers arise from high rate of single-base mismatches during replication
- These lead to high frequency of somatic mutations
- Why only in colon? Not known

Damage reversal 2. Repair of O6-methylguanine





Mispairing of O-6-methylguanine with thymine

MMR and resistant tumours

- In many tumour cells MGMT is silenced. So alkylating agents are good for therapy.
- But often develop resistance.
- Select for alkylation-resistance in cells.
- MGMT not restored. O6-MeG remains in DNA. Instead cells have lost one of the MMR genes.
- Implies MMR somehow sensitises cells to alkylation damage.
- Result of futile cycles. O6-MeG:C and O6-MeG:T both recognised as mismatches.
- C or T opposite O6-MeG removed by MMR and replaced with C or T. Futile cycles.
- Results in cell cycle arrest or apoptosis

Loss of MMR protects against MNNG apoptosis



MMR and cancer

MMR deficiency

- Increases cancer susceptibility (HNPCC)
- Results in resistance to cancer therapy

Summary (Lecture 1b)

- Mismatches are repaired by the Mut(H),L,S system
- Mismatches are recognised by MutS and its homologues
- Strand discrimination is brought about by methylation in *E. coli* and nicks/ends in daughter strands in eukaryotes
- MMR deficiency leads to HNPCC and is detected by microsatellite instability
- Loss of MMR results in resistance to alkylating agents