

Drug design: genomics and proteomics

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Learning outcomes

- Gene therapy
- Gene therapy: inherited disorders
- Gene therapy: cancer
- The impact of biotechnology on drug discovery
- Proteomics: studying proteomics

Gene therapy

- Ex-vivo techniques
 - Explant cells, insert gene, reimplant cells
- *In-vivo* techniques
 - Insert gene already inside the host (need for useful vessels)

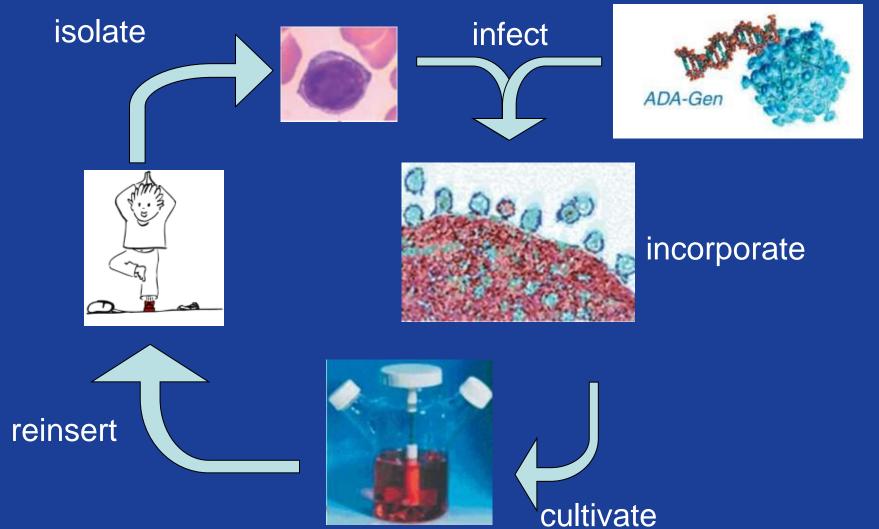
The 'ideal' gene delivery system should...

- 1) Take up also larger DNA/RNA-pieces
- 2) Be easily handled and manufactured in concentrated form
- 3) Target specific cells
- 4) Not be inactivated (=long term activity)
- 5) Not exhibit toxic properties
- 6) Not exert immunogenic responses

Vectorial systems

- A) Plasmids
 - Used in cystic fibrosis
- B) Viral vectors
 - i. Adeno viruses
 - ii. Retroviruses
 - iii.Lentiviruses

ADA (adenosine-deaminase deficiency)



Cystic fibrosis

- Caused by a mutation in a gene called the cystic fibrosis transmembrane conductance regulator (CFTR-del508)
- CFTR = chloride channel → trafficking deficit
- Gene therapy attempts to place a normal copy of the CFTR gene into affected cells
- Only 5–10% of the normal amount of CFTR gene expression is needed
- "There is a realistic prospect of a clinically effective treatment in the next 10 years" (Tate & Elborn, 2005)

Cystic fibrosis

 Caused by a mutation in a gene called the cystic fibrosis transmembrane conductance regulator

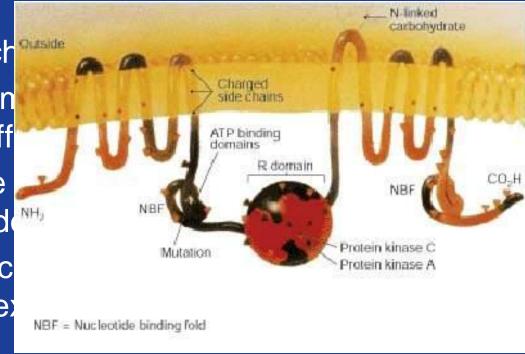
(CFTR-del508)

CFTR = chloride ch

 Gene therapy atten CFTR gene into aff

 Only 5–10% of the expression is need

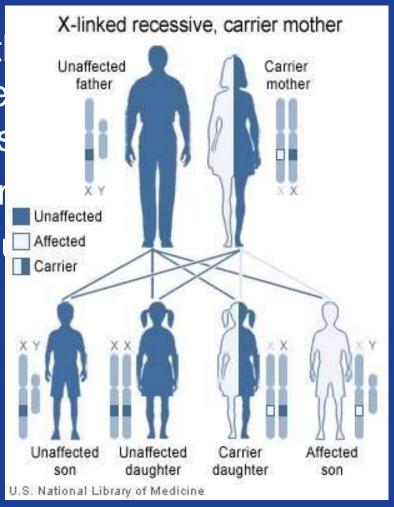
 "There is a realistic treatment in the nex



Duchenne muscular dystrophy

- Caused by mutations in tencodes dystrophin, an embrane protein in mus
- Stem cell research = pror
- U7 gene transfer techniq

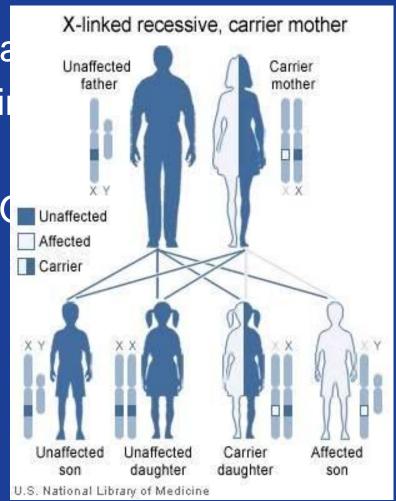
Genetic counselling



Haemophilia

- Bleeding disorder, severa
- Clotting factors are missing

Today: production in CH(

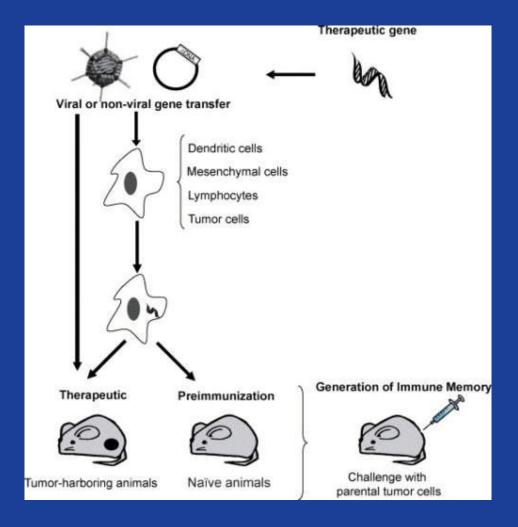


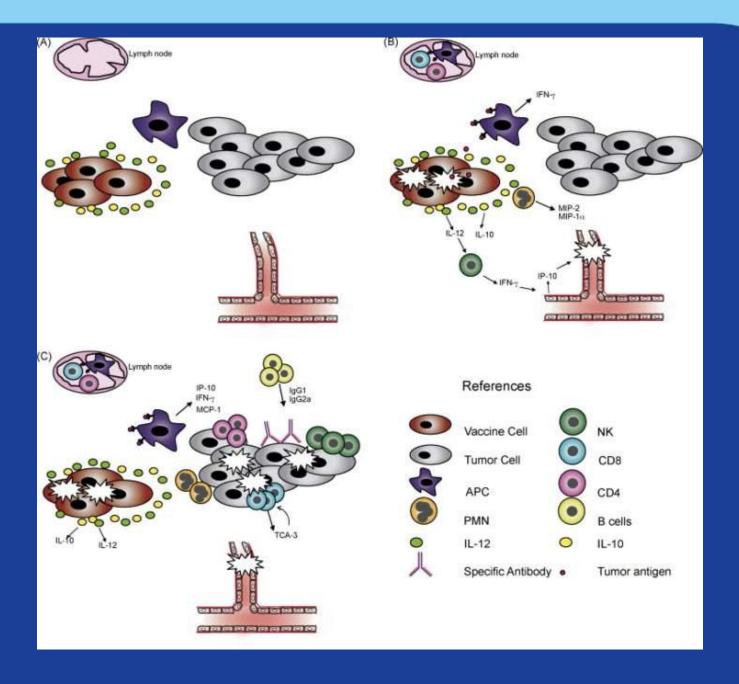
Gene therapy in cancer

 The major problems in cancer therapy with biopharmaceuticals:

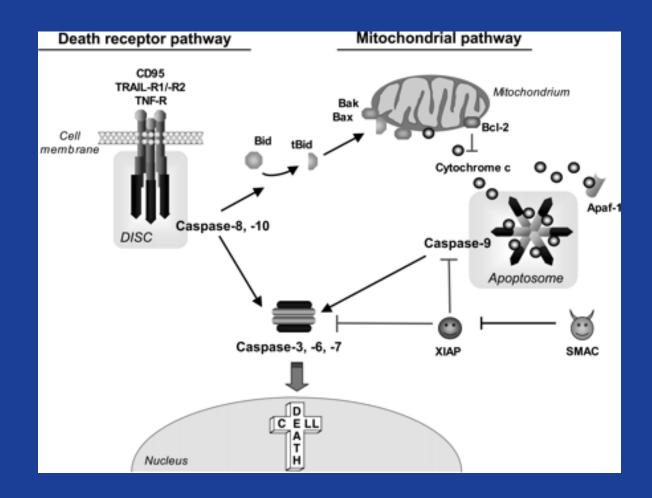
 "The immunogenicity problem, the killing problem and the target problem"

Cytokine genes

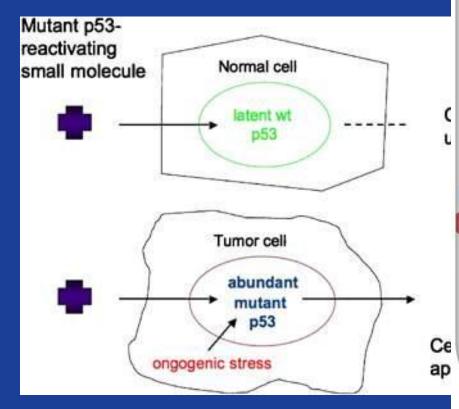




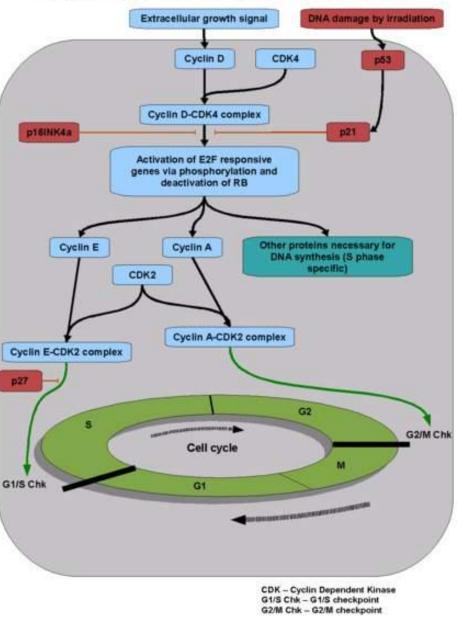
Suicide genes



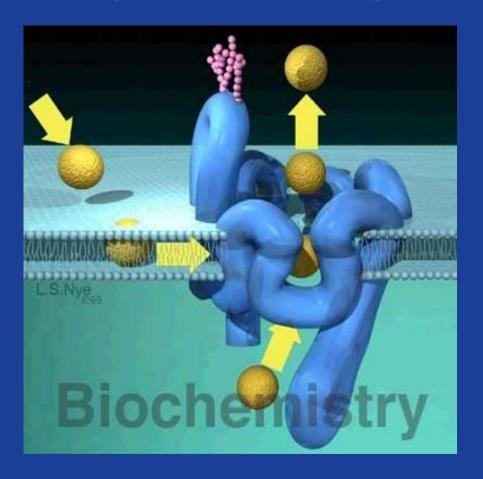
Tumour supp



Regulation of cell cycle - Schematic



Multiple drug resistance gene (MDR-i)



Human genome sequence

identify gene sequences

Functional genomics (and SNPs)

identify gene function in disease of interest

Proteomics/bioinformatics

identify gene product involved in disease of interest

Protein production

produce potential protein target (new molecular site for intervention) for disease of interest

Ligand discovery

identify molecules that bind to protein target in disease of interest

Target validation, ligand bioassay, ligand optimization

prove protein target critical to disease of interest, prove ligand binding affects protein target appropriately, logically modify ligand structure to optimize potential therapeutic effect

New drug entity with novel mechanism of action

Rational drug design

Identification of target structure

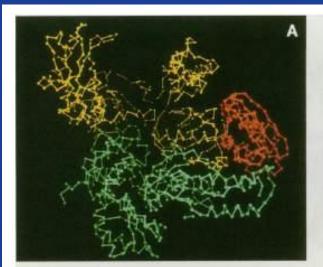
eg HIV reverse transcriptase

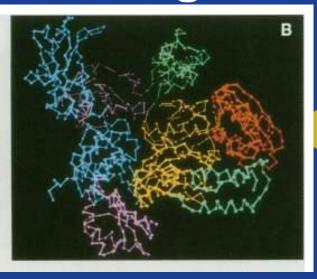
'Lead' compound

Proteomics Genomics Structural biology approach **Bioinformatics**

'Omics'

Rational drug design

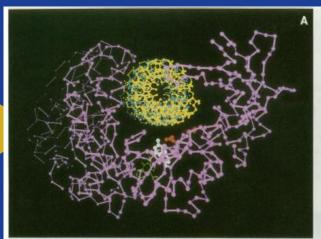


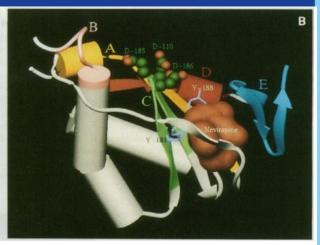


Crystalstructure

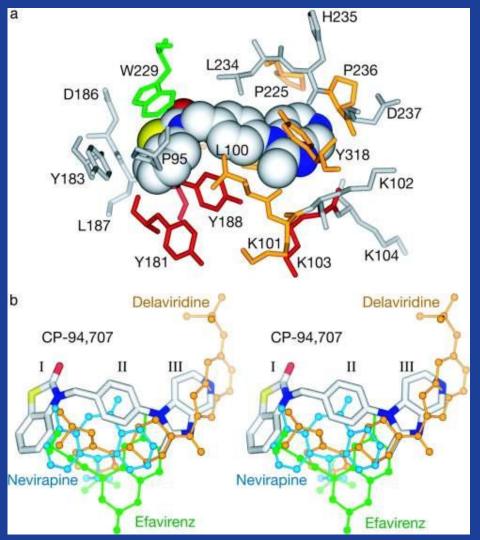
Kohlstaedt et al. Crystal structure at 3.5. A resolution of HIV-1 reverse transcriptase complexed with an inhibitor. *Science* 1992;26(256):1783–1790

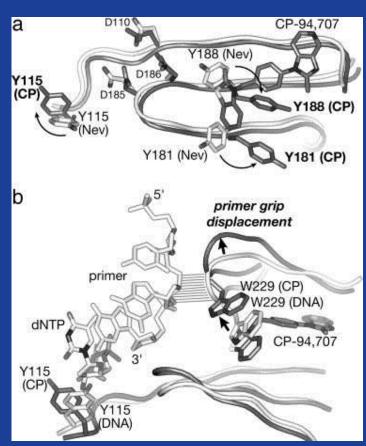
Crystalstructure + nevirapine





Rational drug design





Pata et al. Structure of HIV-1 reverse transcriptase bound to an inhibitor active against mutant reverse transcriptases resistant to other nonnucleoside inhibitors. *PNAS* 2004;101:10548–10553

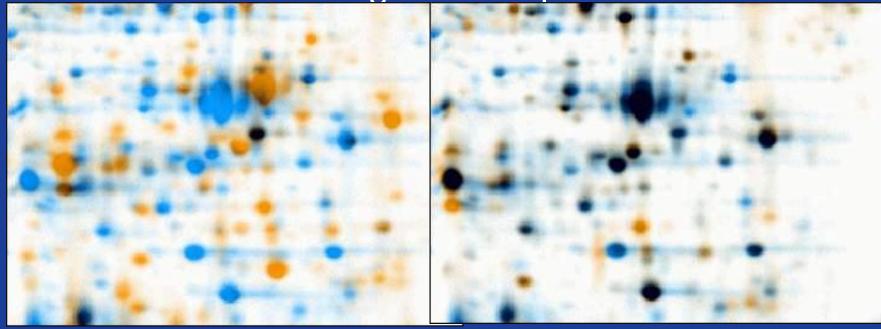
Proteomics



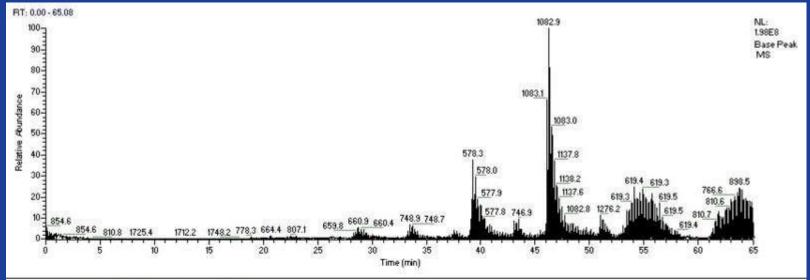
Robotic preparation of MALDI mass spectrometry samples on a sample carrier

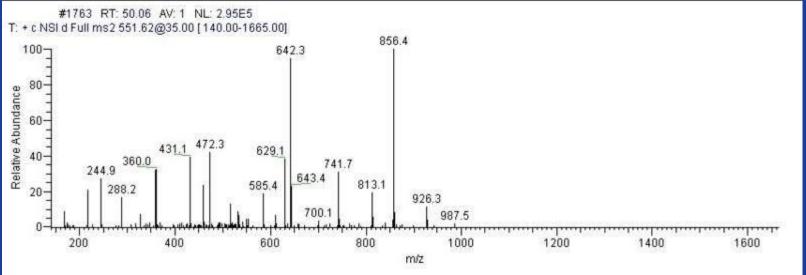
Protein separation techniques

Two-dimensional gel electrophoresis



'Warping'





Protein quantification

- By incorporation of radioactive isotopes into the sample
- Carbon (¹³C) or nitrogen (¹⁵N)
- eg SILAC (stable isotope labellings with amino acids in cell culture)

Protein sequence analysis

- Part of bioinformatics
- Searching databases for possible protein or peptide matches by different algorithms
- Databases include Mascot, PEAKS (software), OMSSA, SEQUEST and X!Tandem
- The prime goals are: functional assignment of domains, prediction of function from sequence, and evolutionary relationships of proteins

Conclusion slide

- Gene therapy
 - Promising technique to treat...
- Gene therapy: inherited disorders
 - Examples for genetically encoded disease
- Gene therapy: cancer
 - New strategies for cancer treatment
- The impact of biotechnology on drug discovery
 - From genome to drug impact of...
- Proteomics: studying proteomics
 - How to examine the structure and nature of proteins