



Polimorfismi a singolo nucleotide e tossicità tardiva radioindotta



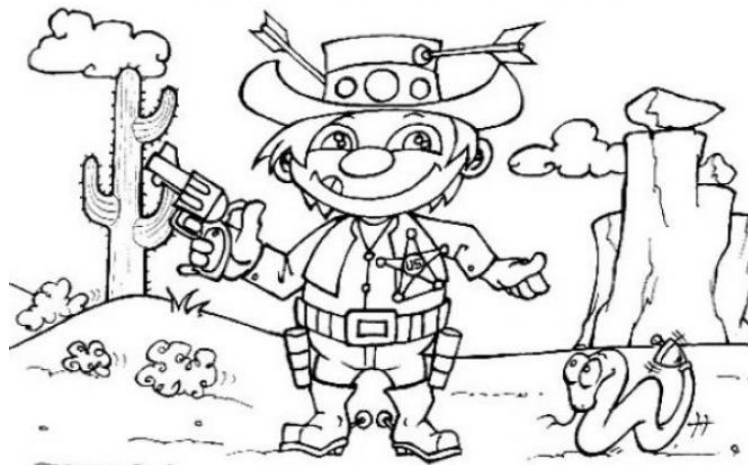
Monica Mangoni
Università di Firenze

Not all the patients react the same way to radiotherapy





Find the differences





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Radiotherapy and Oncology 64 (2002) 131–140

RADIOTHERAPY
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THEORETICAL RADIOLOGY AND ONCOLOGY

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Does variability in normal tissue reactions after radiotherapy have a genetic basis – where and how to look for it?

Christian Nicolaj Andreassen*, Jan Alsner, Jens Overgaard



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hyper-responsive clinical phenotype



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Prediction of normal tissue radiosensitivity from polymorphisms in candidate genes

Christian Nicolaj Andreassen^{a,*}, Jan Alsner^a, Marie Overgaard^b, Jens Overgaard^a

Radiotherapy and Oncology 99 (2011) 356–361



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Single Nucleotide Polymorphisms

Clinical radiobiology

Association between single nucleotide polymorphisms in the XRCC1 and RAD51 genes and clinical radiosensitivity in head and neck cancer

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CLINICAL INVESTIGATION

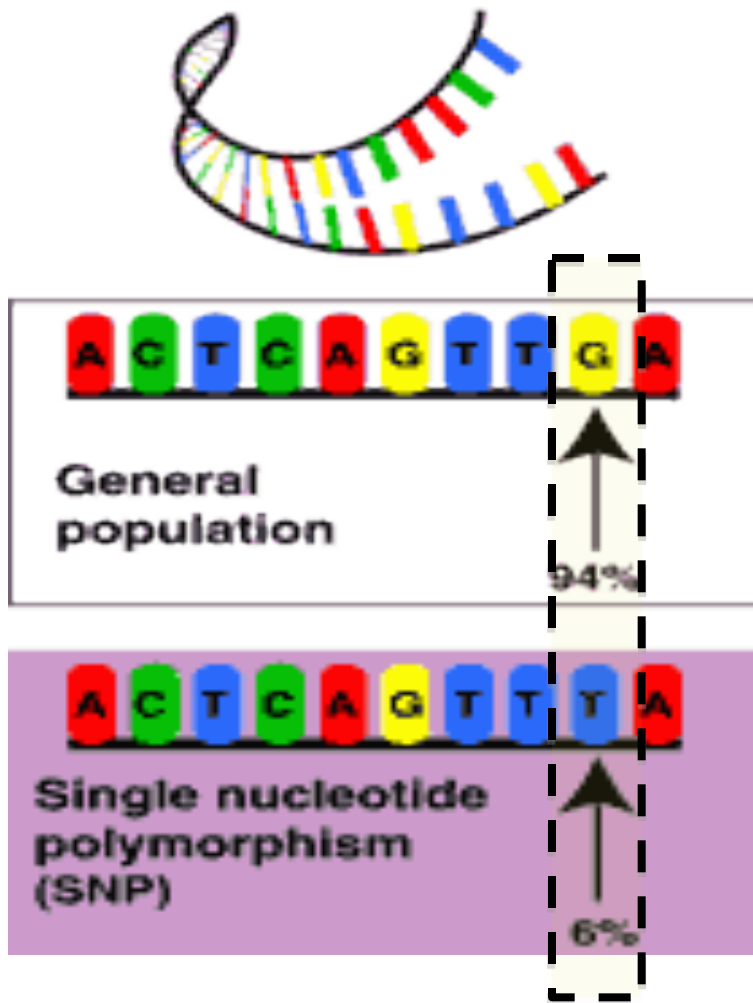
Breast

ASSOCIATION BETWEEN GENETIC POLYMORPHISMS IN THE XRCC1, XRCC3, XPD, GSTM1, GSTT1, MSH2, MLH1, MSH3, AND MGMT GENES AND RADIOSENSITIVITY IN BREAST CANCER PATIENTS

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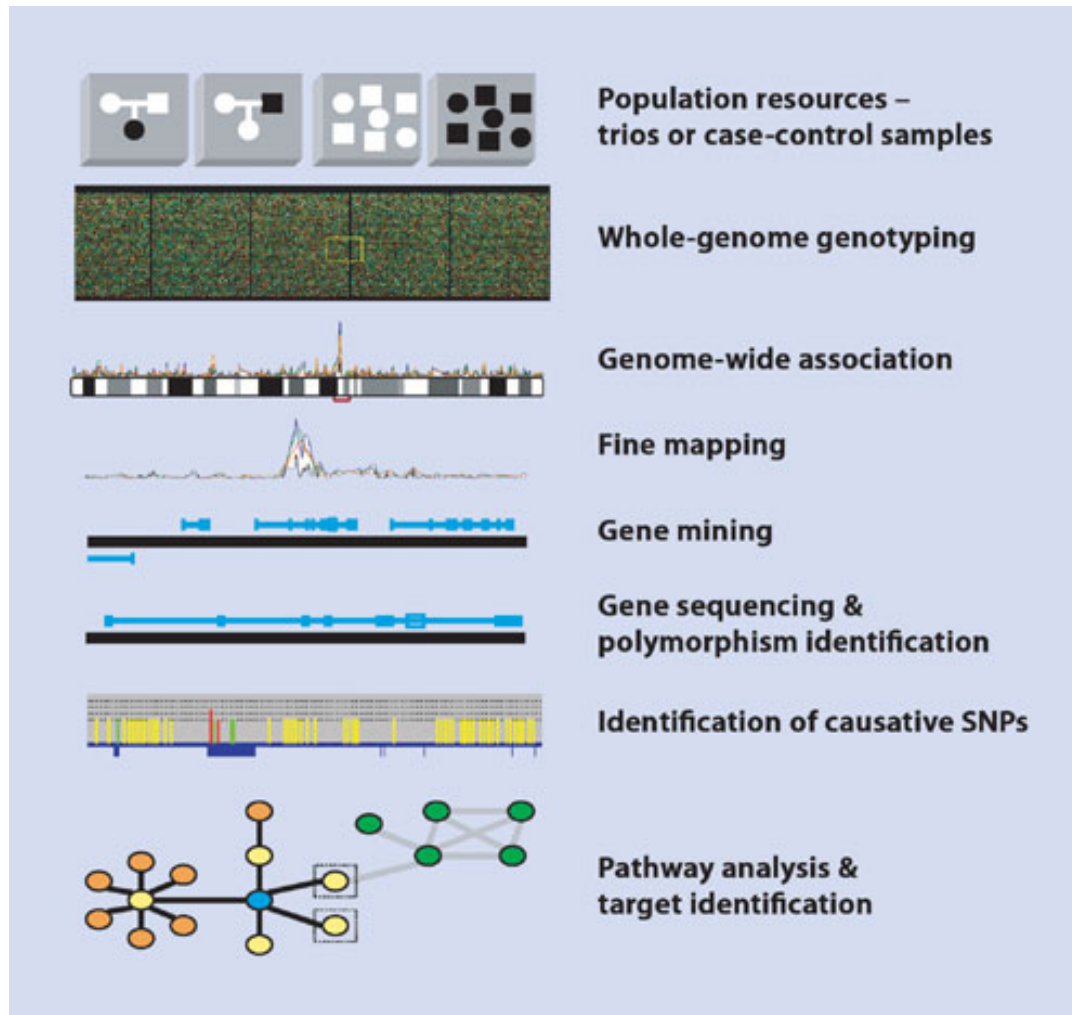


Single Nucleotide Polymorphisms (SNPs)



Genetic variants in which an alternate base pair is present at a particular nucleotide location

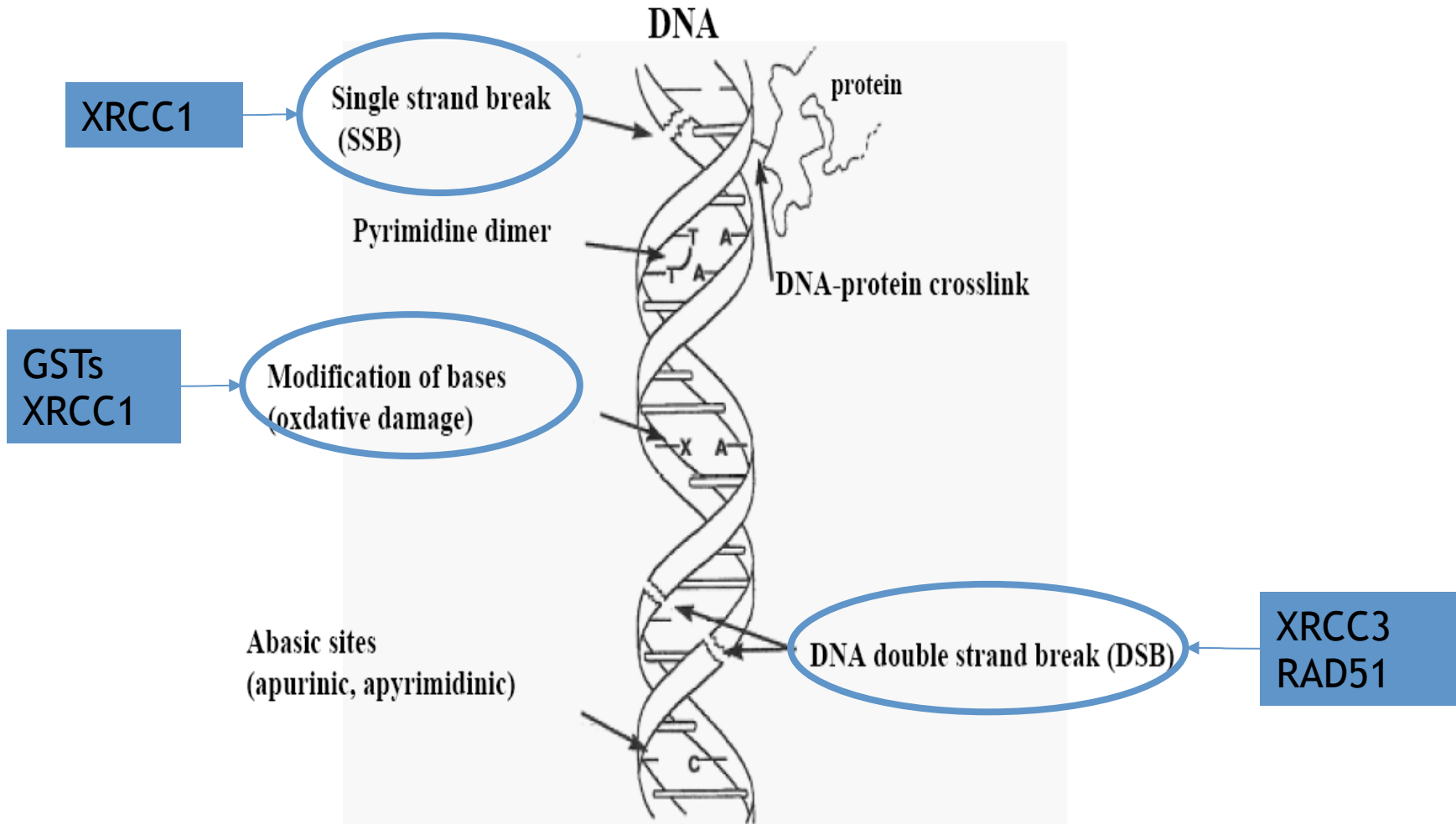
Human Genome-Wide Association Studies



GWAS permit a comprehensive scan of the genome in an unbiased fashion and thus have the potential to identify totally novel susceptibility factors.

where to look for it?

candidate gene approach





Genes related to the endogenous oxidative stress defense

- GSTM1, GSTT1, GSTA1, GSTP1
- SOD2, MPO, eNOS

- **SOD2 (Val16Ala) (rs4880 T/C)**: severe **skin fibrosis** in breast cancer

(Andreassen CN, Radiother Oncol 2003; 69:127-135)

but other studies haven't confirmed this observation

(Green H, Radiother Oncol 2002; 63: 213-216)

(Andreassen CN, Radiother Oncol 2005; 75: 18-21)

- **GSTP1 Ile105Val (rs1695)** (variant with low activity): **pleural thickening**

(Edvardsen H, IJROBP 2007; 67: 1163-71)

- **variant form of the GSTA1**: **increased teleangiectasia**
- **eNOS T allele (lower activity)**: **lower risk of teleangiectasia**

(Kuptsova N, Int J cancer 2008; 122: 1333-1339)



Genes involved in the inflammatory response and in cytokine activity related to fibrosis



- **TGFβ1 509 C>T** in the promoter region: **elevated TGF β1 serum levels**
- **TGFβ1 869 T>C** and **TGFβ1 915 G>C** (amino acid changes): **fibrosis in breast cancer**

(Andreassen CN, Radiother Oncol 2003; 69: 127-35)

(Andreassen CN, Radiother Oncol 2005; 75: 18-21)

but data haven't been confirmed in another study

(Andreassen CN, Int J Radiat Biol 2006; 82: 577-86)

nor for other types of tumors (cervical, endometrial, prostate tumour)

(De Ruyck k, IJROBP 2006, 65: 1240-8)

(Damaraju D, Clin Canc Res 2006; 12: 2545-54)

- **TGFβ1 509 C>T: decline in erectile function and rectal bleeding in prostate cancer**

(Peters CA, IJROBP 2008; 70: 752-9)



Genes contributing to DNA damage signalling and cell cycle control



- **ATM 5557G>A: fibrosis and telangiectasia** in breast cancer
(Andreassen CN, Radiother Oncol 2005; 75: 18-21)
however results were not confirmed in other cohorts
(Andreassen CN, Int J Radiat Biol 2006; 82: 577-586)
(Ho AY, IJROBP 2007; 69: 677-84)
- **ATM 4258C>T (rs1800058): pleural thickening and lung fibrosis**
(Edwardsen H, Radiat Oncol 2007: 2:25)
- **ATM sequence alterations: proctitis and decline of erectile function**
with 125I brachytherapy
(Cesaretti JA, IJROBP 2005; 61: 196-202)



DNA repair genes

- **BER**: XRCC1, APEX1, OGG1, LIG3
- **NER**: ERCC2/XPD, ERCC4/XPF, RAD9A
- **HR and NHEJ**: RAD51, RAD52, XRCC3, XRCC2, NBN, LIG4, BRCA1, BRCA2

- **XRCC3 18067 C > T**: severe **fibrosis**

(Andreassen CN, Radiother Oncol 2003; 69: 127-135)

but other findings could not be confirmed in the other studies

- **XRCC3 17893 A>G**: **late** adverse effects in cervical and endometrial carcinoma

(De Ruyck K, IJROBP 2006; 65: 1240-8)

- **LIG4 Asp568Asp**: late **bladder or rectal toxicity** in prostate cancer

(Damaraju D, Clin Cancer Res 2006; 12: 2545-54)



DNA repair genes

- **BER**: XRCC1, APEX1, OGG1, LIG3
- **NER**: ERCC2/XPD, ERCC4/XPF, RAD9A
- **HR and NHEJ**: RAD51, RAD52, XRCC3, XRCC2, NBN, LIG4, BRCA1, BRCA2

- **XRCC1 28152G**: severe **fibrosis** and **telangiectasia** in breast cancer
(Andreassen CN, Radiother Oncol 2003; 69: 127-35)
(Gitopoulos G, Br J Cancer 2007; 96: 1001-7)
- **XRCC1 26304T**: **late** adverse effects in breast cancer
- **XRCC1 28152G in combination with XRCC1 26304T**: side effects
(Moullan N, Cancer Epidemiol Biomarkers Prev 2003; 12: 1168-74)

In one study with only a small number of patients: no effects of XRCC1 SNPs

(Andreassen CN, Radiother Oncol 2005; 75: 18-21)

- **XRCC1 polymorphisms**: no significant effect in gynecological and in prostate cancer
(De Ruyck K, IJROBP 2006; 65:1240-8)
(Damaraju D, Clin Cancer Res 2006; 12: 2545-54)



Single Nucleotide Polymorphisms and radiation-induced late toxicity in prostate cancer

Cipressi et al, AIRO 2011

Case-control study on 46 patients affected by prostate cancer

GSTP1 A313G

XRCC3 c.722C>T
RAD51 c.3429G>C
RAD51 c.3392G>T

- **GSTP1c313A>G**: late rectal and bladder toxicities
- **XRCC1 c.1196A>G**: late rectal and bladder toxicities

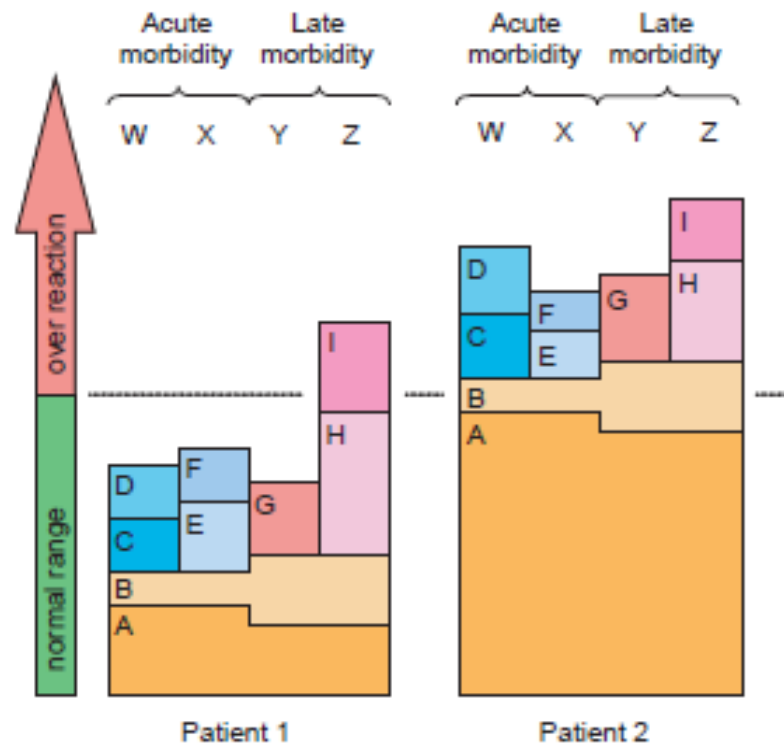
≠ XRCC1 polymorphisms: no significant effect in prostate cancer
(De Ruyck K, IJROBP 2006; 65:1240-8)
(Damaraju D, Clin Cancer Res 2006; 12: 2545-54)



Bias of the available studies

- ✓ candidate gene approach is insufficient: the physiology underlying a given phenotype is very complex or poorly understood
- ✓ investigate haplotypes rather than individual SNPs (tag SNPs)
- ✓ different normal tissue damage endpoints used to evaluate radiosensitivity (basic vs complex endpoints, acute vs late vs effects considered together)
- ✓ numerous confounding factors
 - ✓ comorbidities
 - ✓ different treatments: External beam vs brachiterapy vs fractionation (BED) vs carbon ions (EBR)

Clinical normal tissue radiosensitivity: complex trait



Andreassen CN, Acta Oncologica, 2005; 44: 801-15



Conclusions

- ✓ We are still far from having a comprehensive understanding of the sequence alterations that may underlie variability in clinical normal tissue radiosensitivity.
- ✓ On the other hand, the available data seem to provide a reasonable proof of principle that the risk of radiation induced normal tissue complications among unselected cancer patients is influenced by genetic factors



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QUANTEC

QUANTEC: VISION PAPER

BIOMARKERS AND SURROGATE ENDPOINTS FOR NORMAL-TISSUE EFFECTS OF RADIATION THERAPY: THE IMPORTANCE OF DOSE-VOLUME EFFECTS

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GENEPI
Europe

REPORT

ESTABLISHMENT OF A RADIOGENOMICS CONSORTIUM

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REPORT

GENETIC PREDICTORS OF ADVERSE RADIOTHERAPY EFFECTS: THE GENE-PARE PROJECT

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