

Basics in cancer cell biology

- DNA repair

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OUTLINE

DNA damage response and DNA repair pathways
DNA repair and cancer
DNA repair biomarkers in breast cancer
DNA repair as anticancer target
DNA repair and targeted therapies
Perspectives and conclusions

Cells are continuously submitted to DNA damage

- High fidelity of DNA replication machinery : = 10^{-10} spontaneous mutations/base
- Endogenous metabolic activities and environmental factors are continuously affecting DNA's integrity (oxidative stress, alkylating agents, UV radiation ...)
- Lesions: chemical modification of DNA bases, abasic sites, base mismatches, chemical alkylation (adducts), intra-and inter-strand cross links, single-and double-strand breaks

Table 1. DNA Lesions Generated by Endogenous and Exogenous DNA Damage

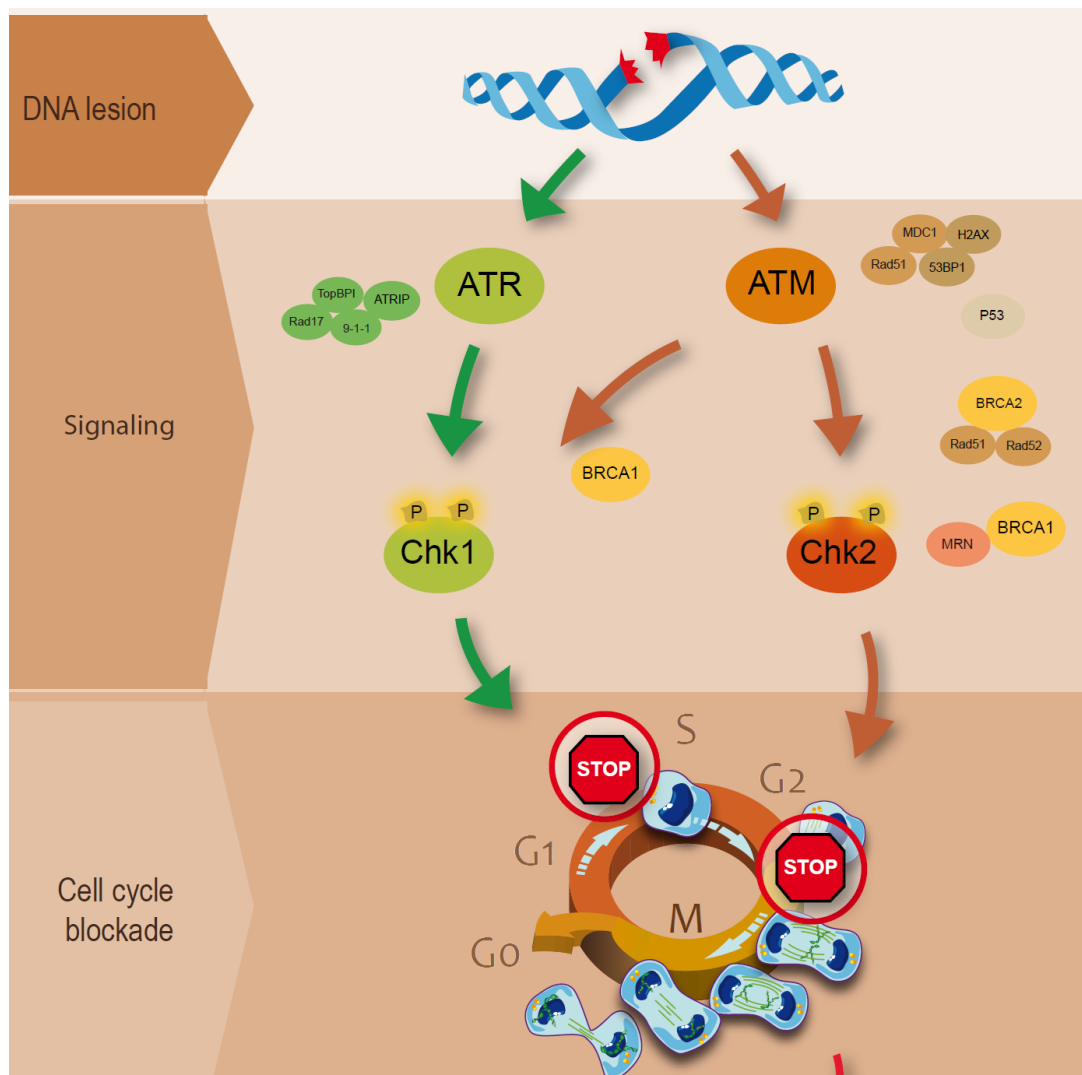
Endogenous DNA Damage	DNA Lesions Generated	Number Lesions/Cell/Day
Depurination	AP site	10000 ^a
Cytosine deamination	Base transition	100–500 ^a
SAM-induced methylation	3meA	600 ^a
	7meG	4000 ^a
	O ⁶ meG	10–30 ^b
Oxidation	8oxoG	400–1500 ^c

- Exogenous origins of DNA damage
- DSBs more dangerous and less frequent

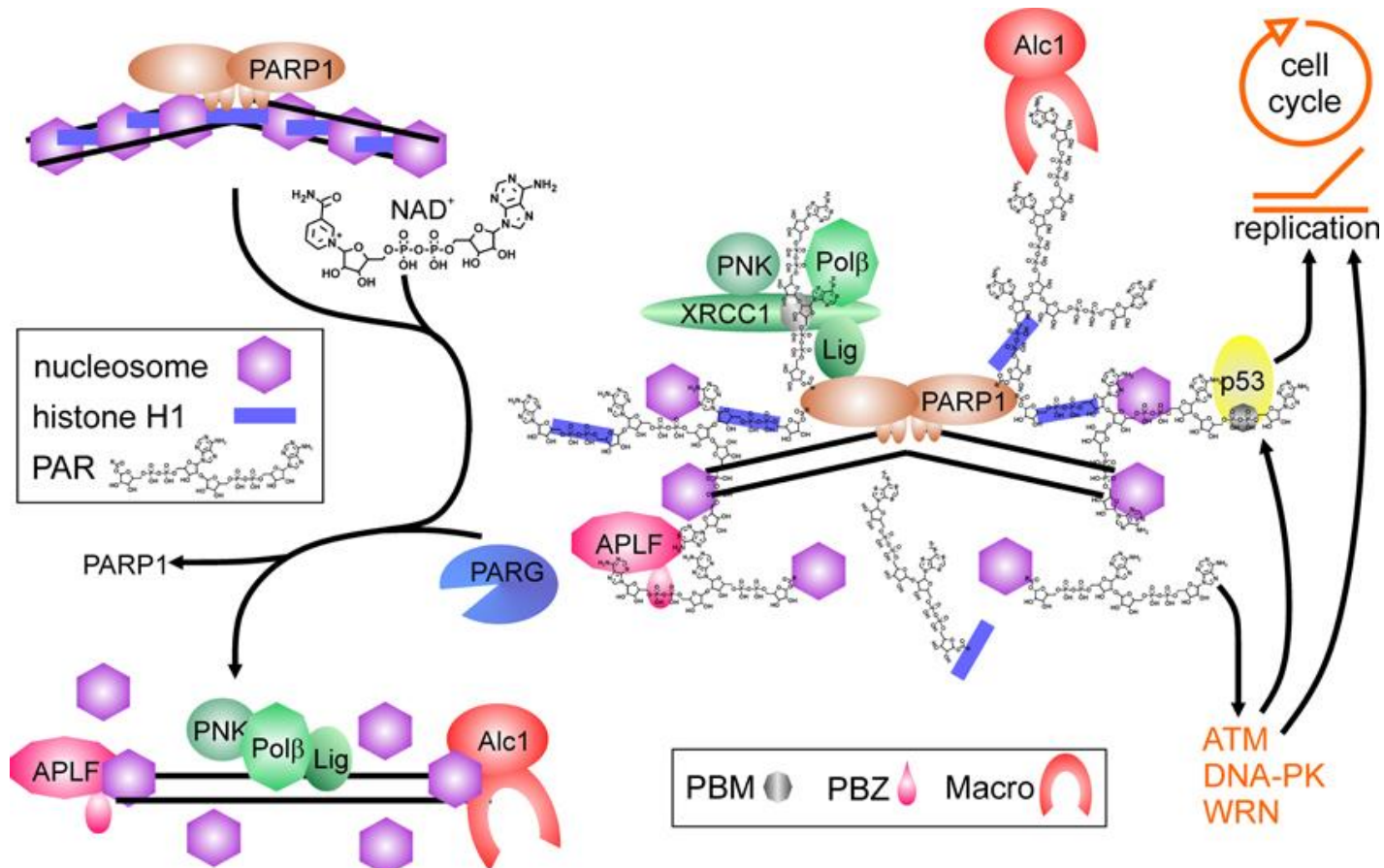
			<u>Lesions/day</u>
Exogenous DNA Damage	Dose Exposure (mSv)	DNA Lesions Generated	Estimated Number
Peak hr sunlight	—	Pyrimidine dimers, (6–4) photoproducts	100,000/day ^d
Cigarette smoke	—	aromatic DNA adducts	45–1029 ^e
Chest X-rays	0.02 ^{f,g,h}	DSBs	0.0008 ⁱ
Dental X-rays	0.005 ^{f,g,h}	DSBs	0.0002 ⁱ
Mammography	0.4 ^{f,g,h}	DSBs	0.016 ⁱ
Body CT	7 ^f	DSBs	0.28 ⁱ
Head CT	2 ^{f,g}	DSBs	0.08 ⁱ
Coronary angioplasty	22 ^h	DSBs	0.88 ⁱ
Tumor PET scan (¹⁸ F)	10 ^h	DSBs	0.4 ⁱ
¹³¹ I treatment	70–150 ^h	DSBs	2.8–6 ⁱ
External beam therapy	1800–2000 ^j	DSBs	72–80
Airline travel	0.005/hr ^f	DSBs	0.0002/hr ⁱ
Space mission (60 days)	50 ^k	DSBs	2 ⁱ
Chernobyl accident	300 ^l	DSBs	12 ⁱ
Hiroshima and Nagasaki atomic bombs	5–4000 ^k	DSBs	0.2–160 ⁱ



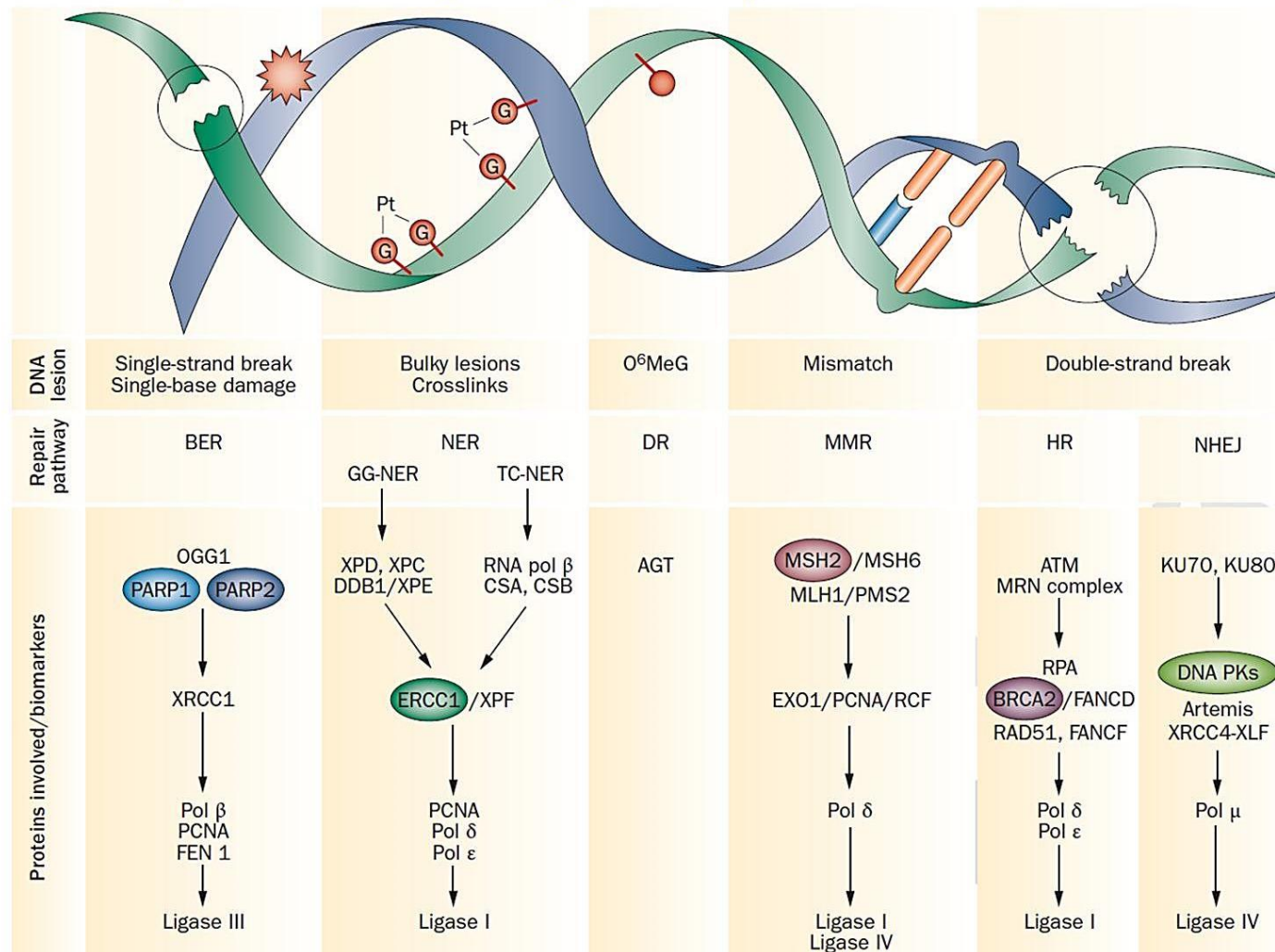
DNA Damage Response (DDR)



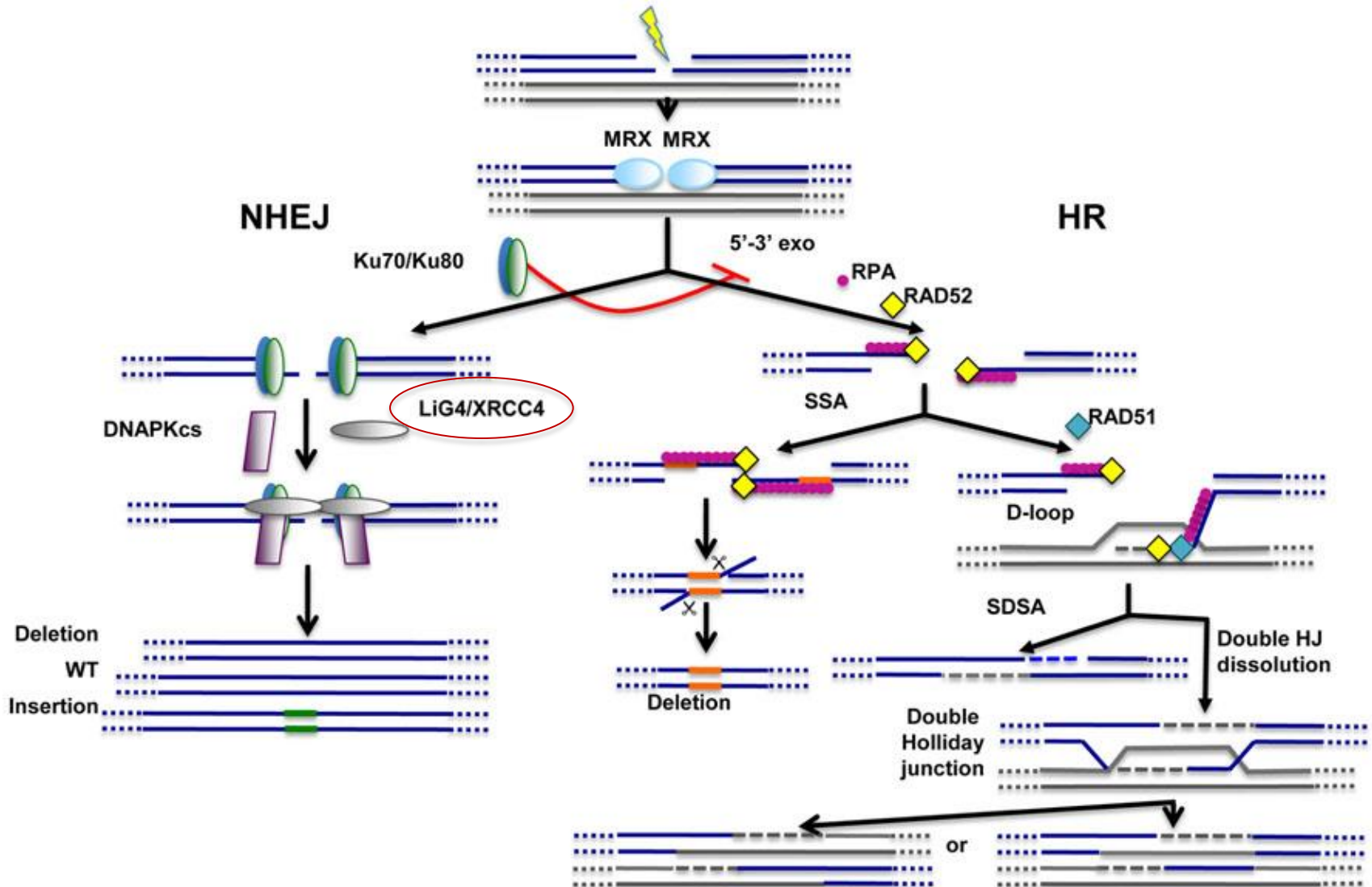
BER pathway depends on the automodification (poly(ADP-ribosyl)ation (PAR) of PARP1 to release DNA from nucleosomes and recruit repair proteins



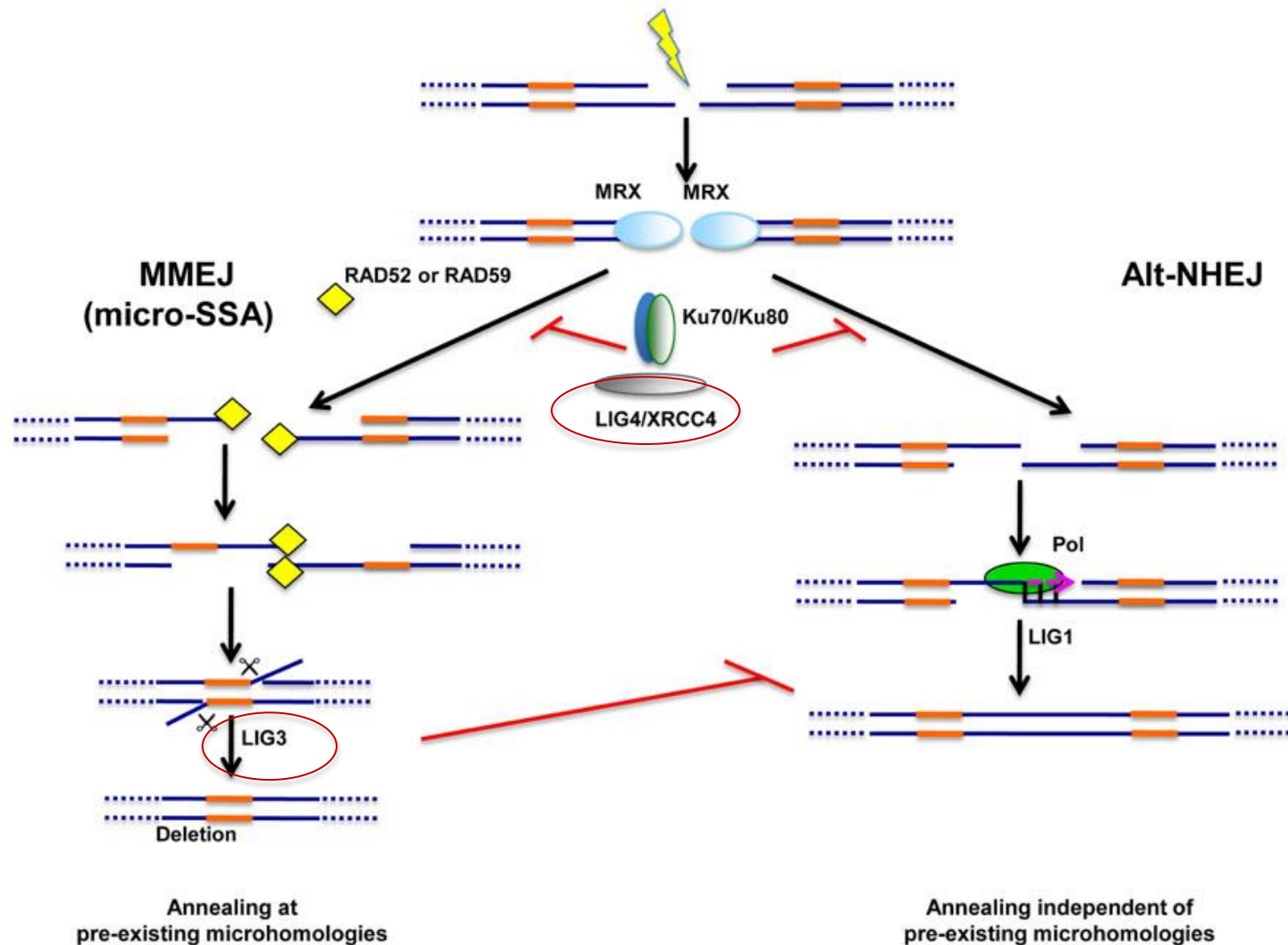
Major DNA repair pathways and biomarkers



DSB repair pathways



Alternative end-joining pathways

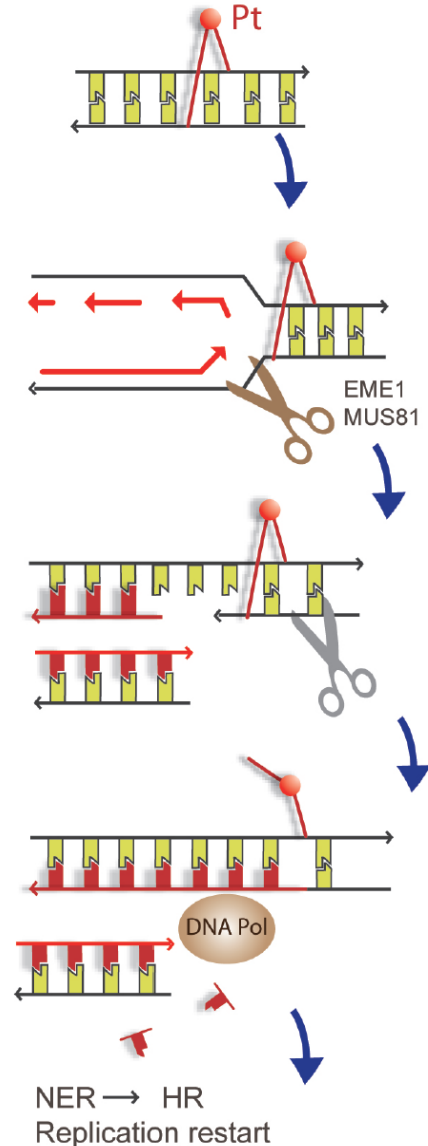
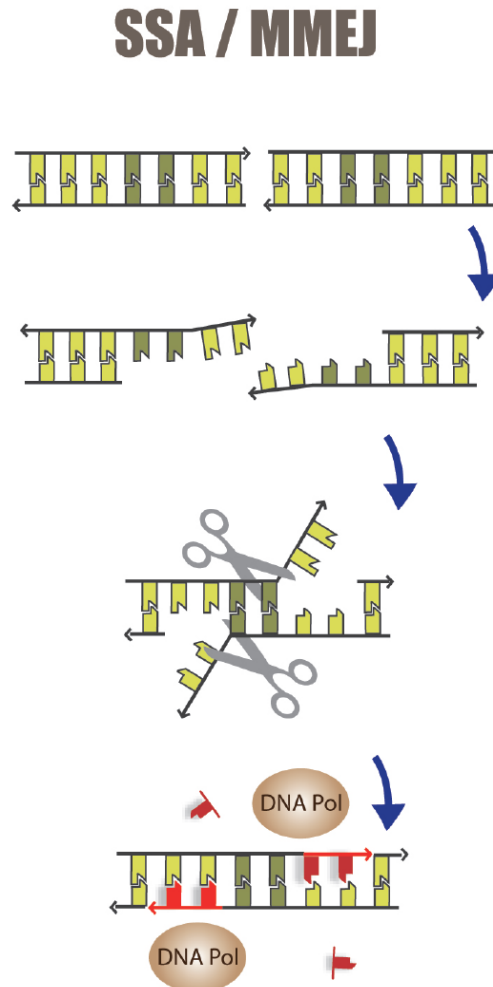
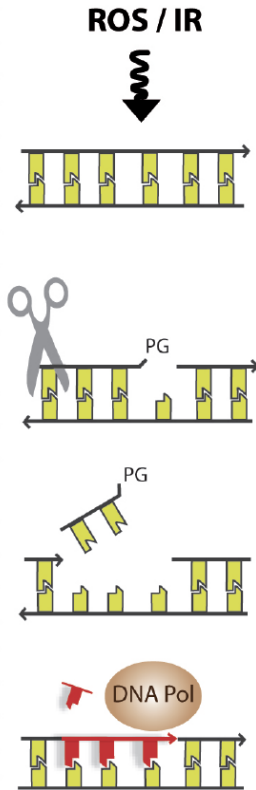
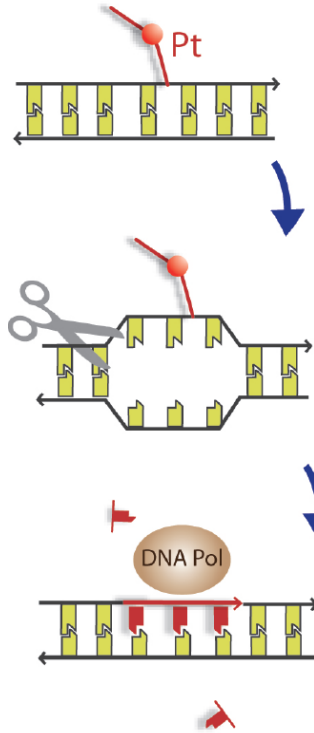


NER

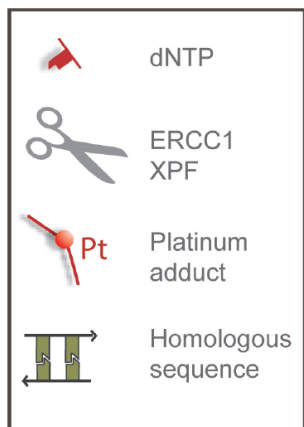
SSB-R

DSB-R

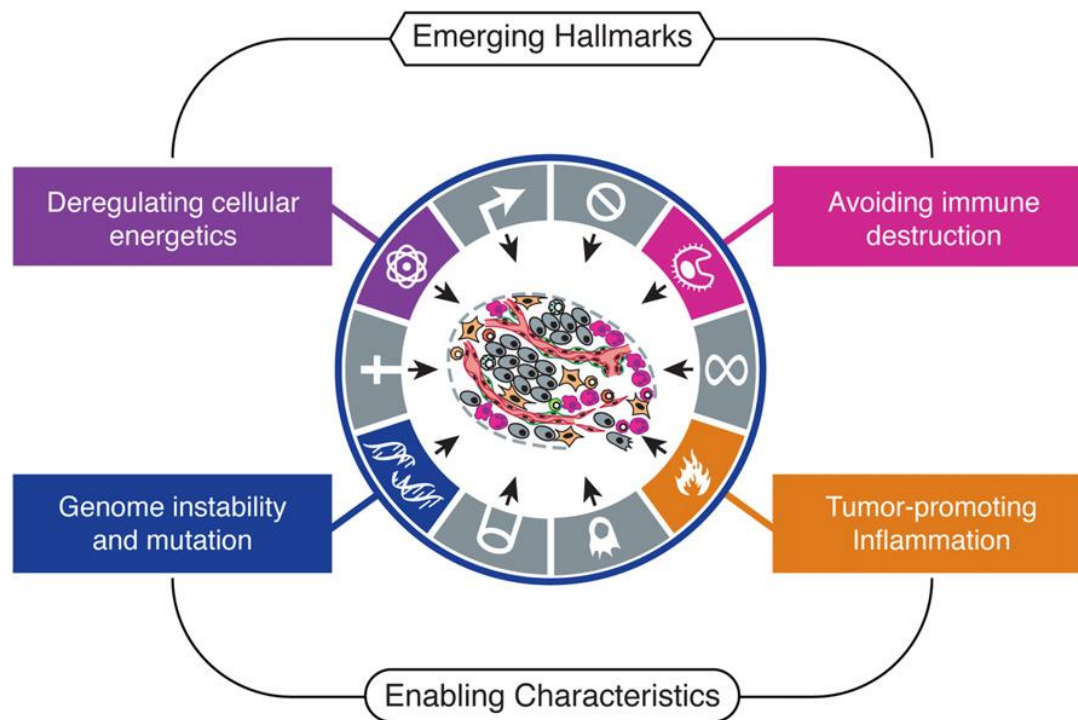
ICL-R



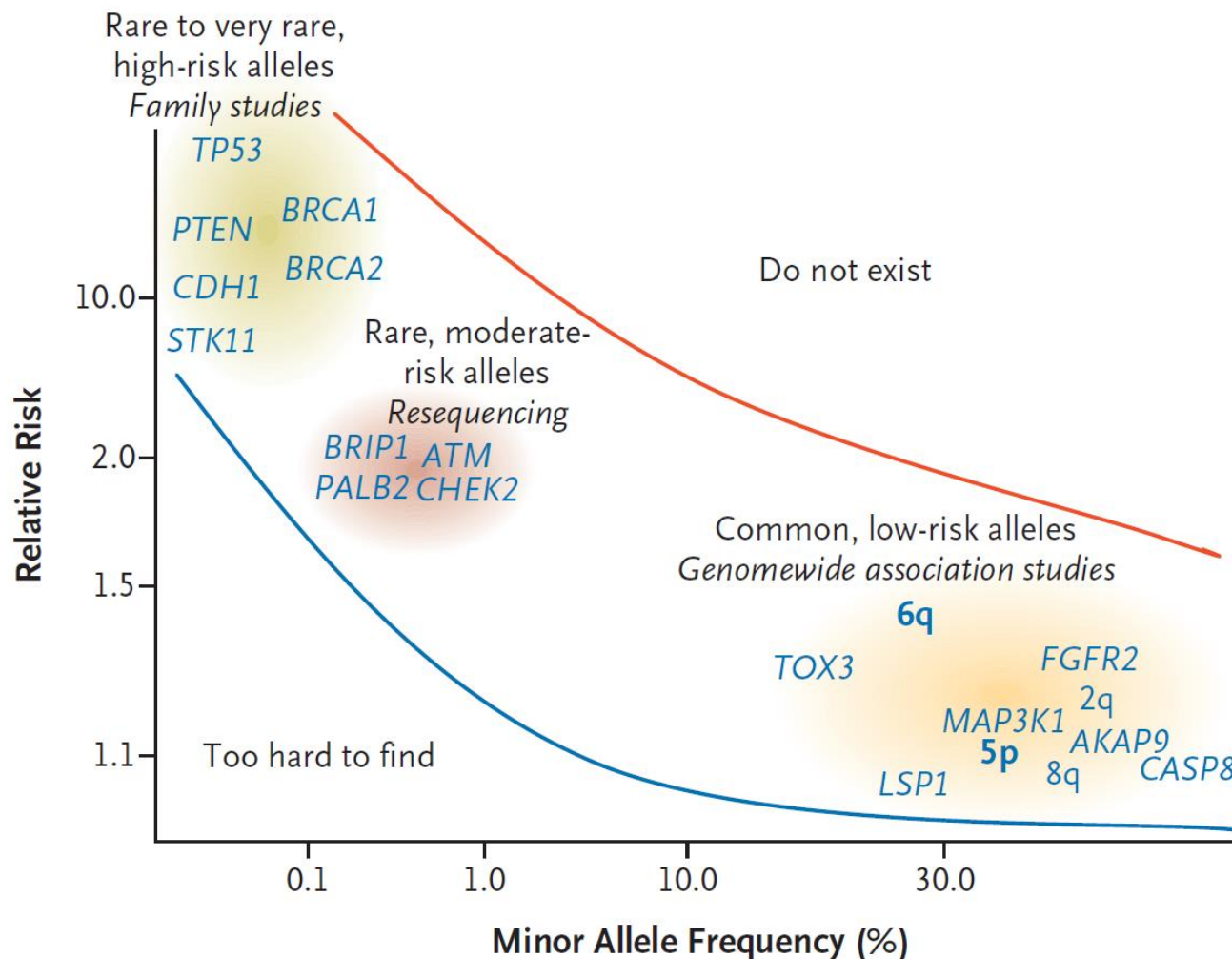
Numerous interconnexions between pathways
e.g. ERCC1 is implicated in several



« Mutability endow cancer cells with genetic alterations that drive tumor progression »

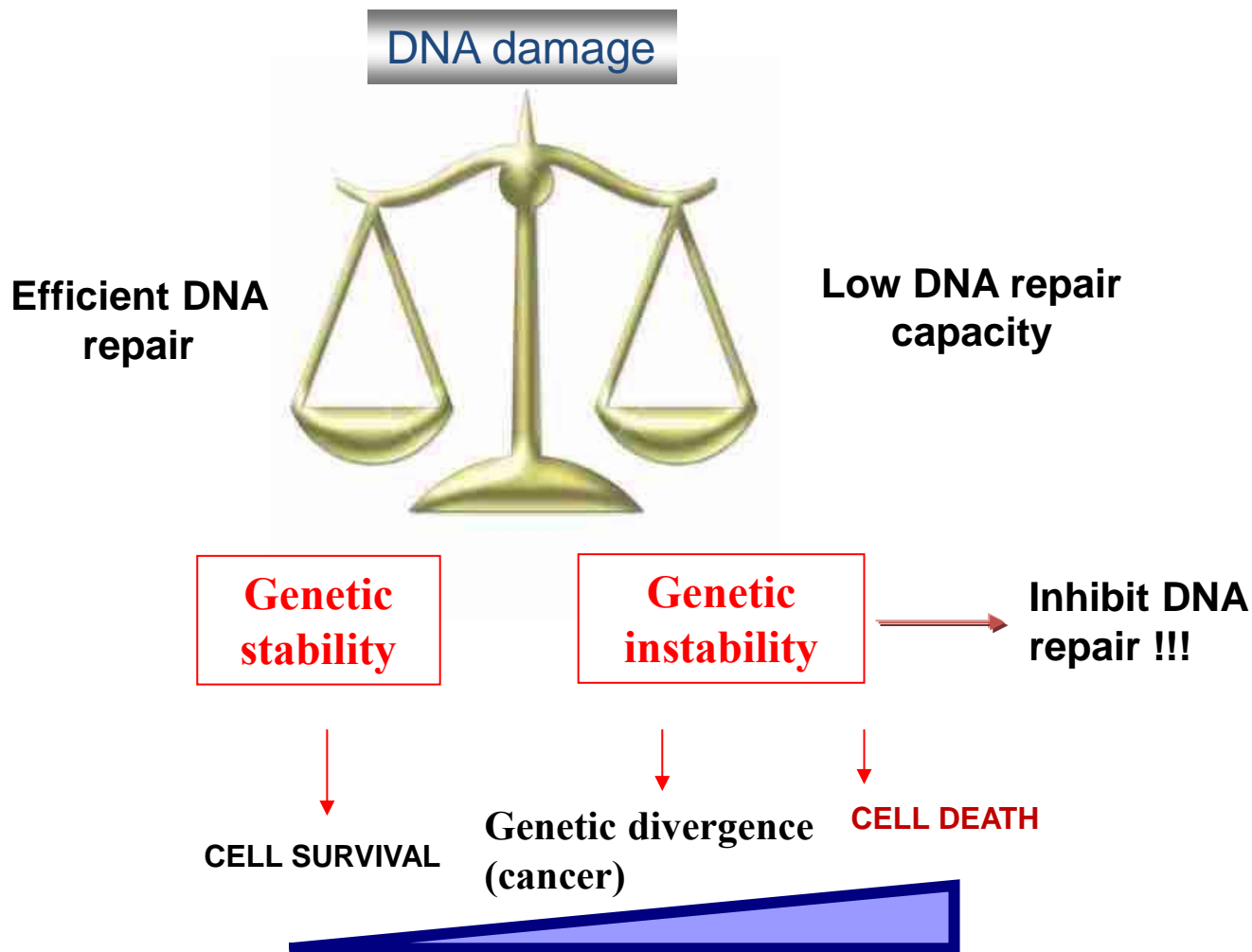


Risk alleles in breast cancer: several DNA-damage response or repair-related genes: BRCA1/2, TP53, ATM, CHEK2...

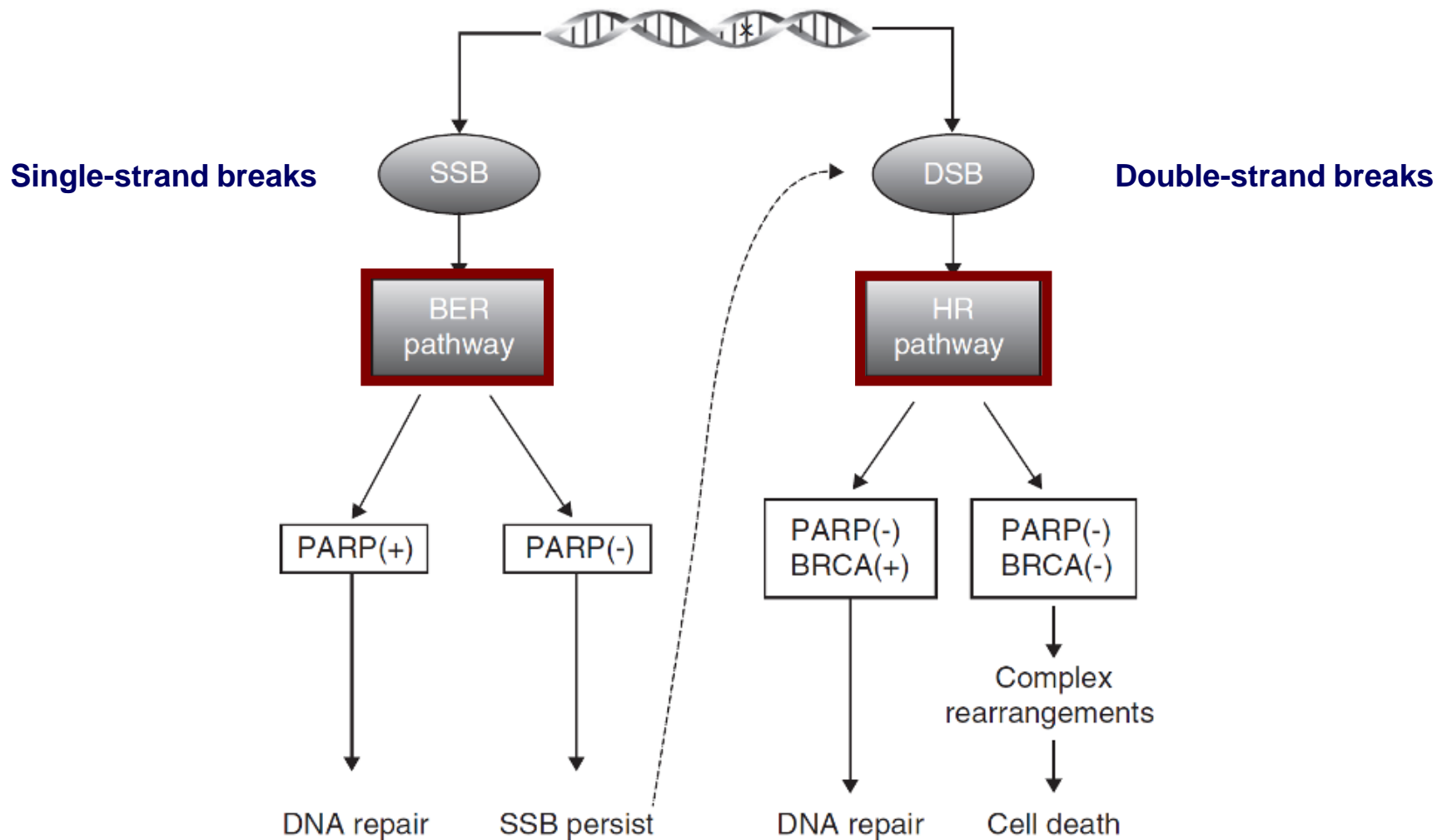




“The balance of life” theory supports that DNA repair mechanisms can be an anticancer target

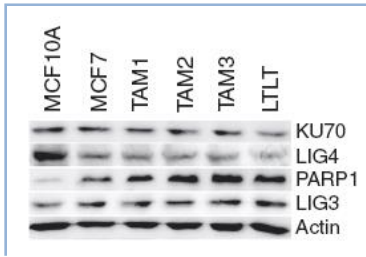


PARP inhibition is synthetic lethal in BRCA1-deficient tumors because HR pathway is a back-up pathway for unrepaired SSBs

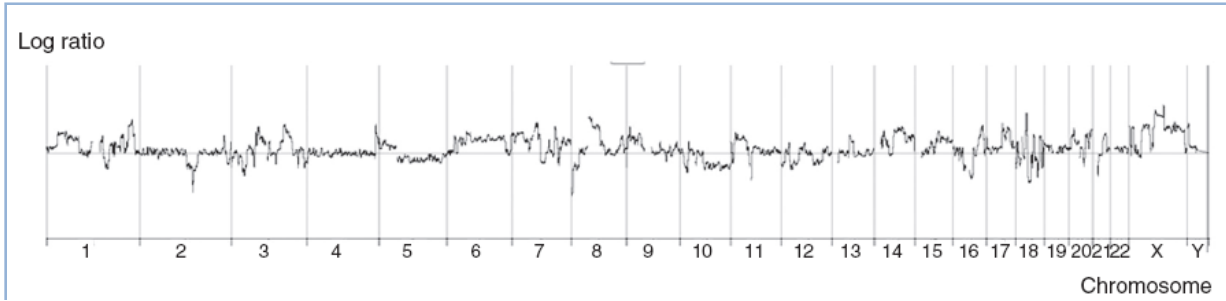


Targeting Abnormal DNA Repair in Therapy-Resistant Breast Cancers

Lisa A. Tobin¹, Carine Robert¹, Pratik Nagaria¹, Saranya Chumsri², William Twaddell³, Olga B. Ioffe³, George E. Greco⁵, Angela H. Brodie⁴, Alan E. Tomkinson⁶, and Feyruz V. Rassool¹



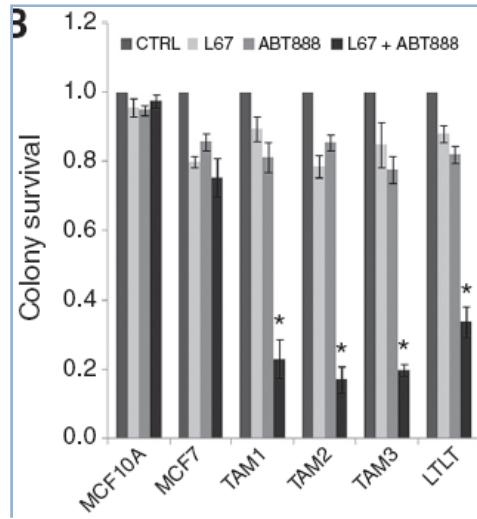
Altered expression of DNA repair proteins in breast cancer cell lines with acquired resistance to the antiestrogens tamoxifen and letrozole



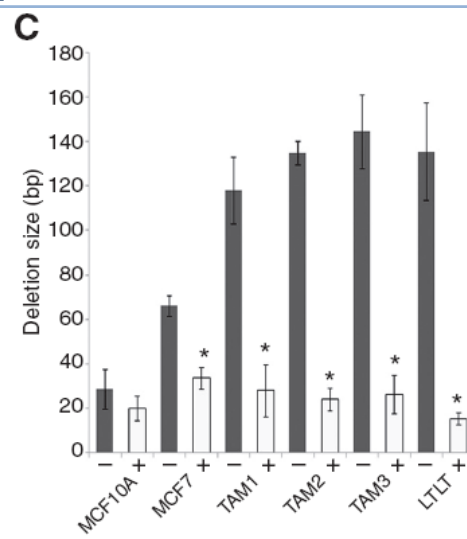
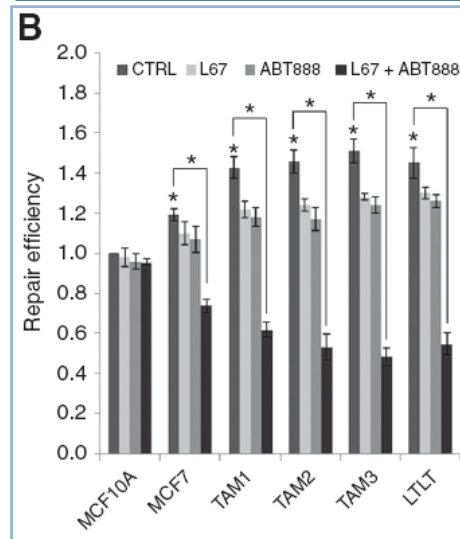
The increase of Ligase 3 favors error-prone ALT NHEJ resulting in large deletions and chromosomal translocations in therapy-resistant cells

Targeting Abnormal DNA Repair in Therapy-Resistant Breast Cancers

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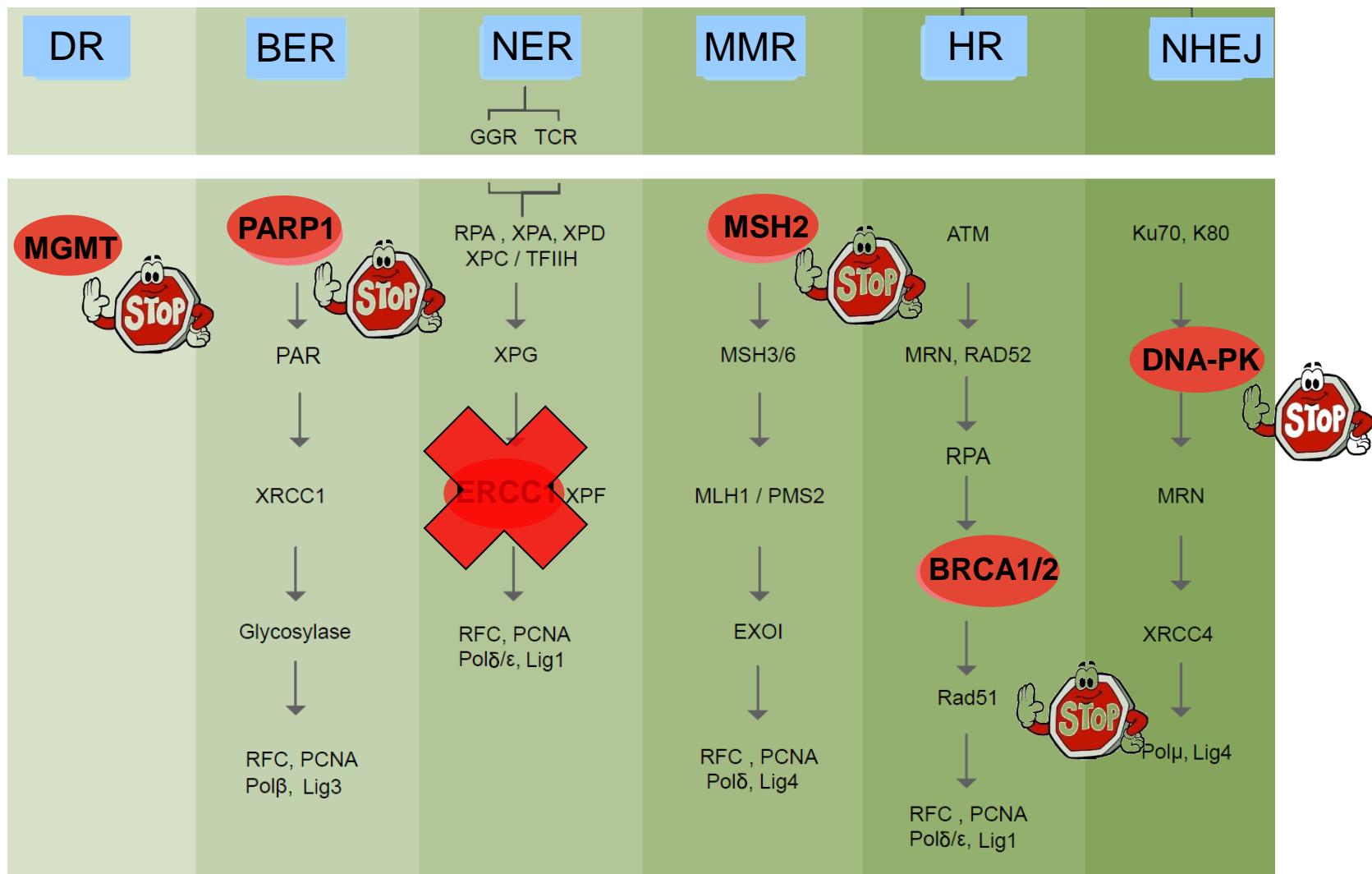


Cell lines with acquired resistance to antiestrogen therapeutics are hypersensitive to a combination of DNA ligase 3 (L67) and PARP inhibitors



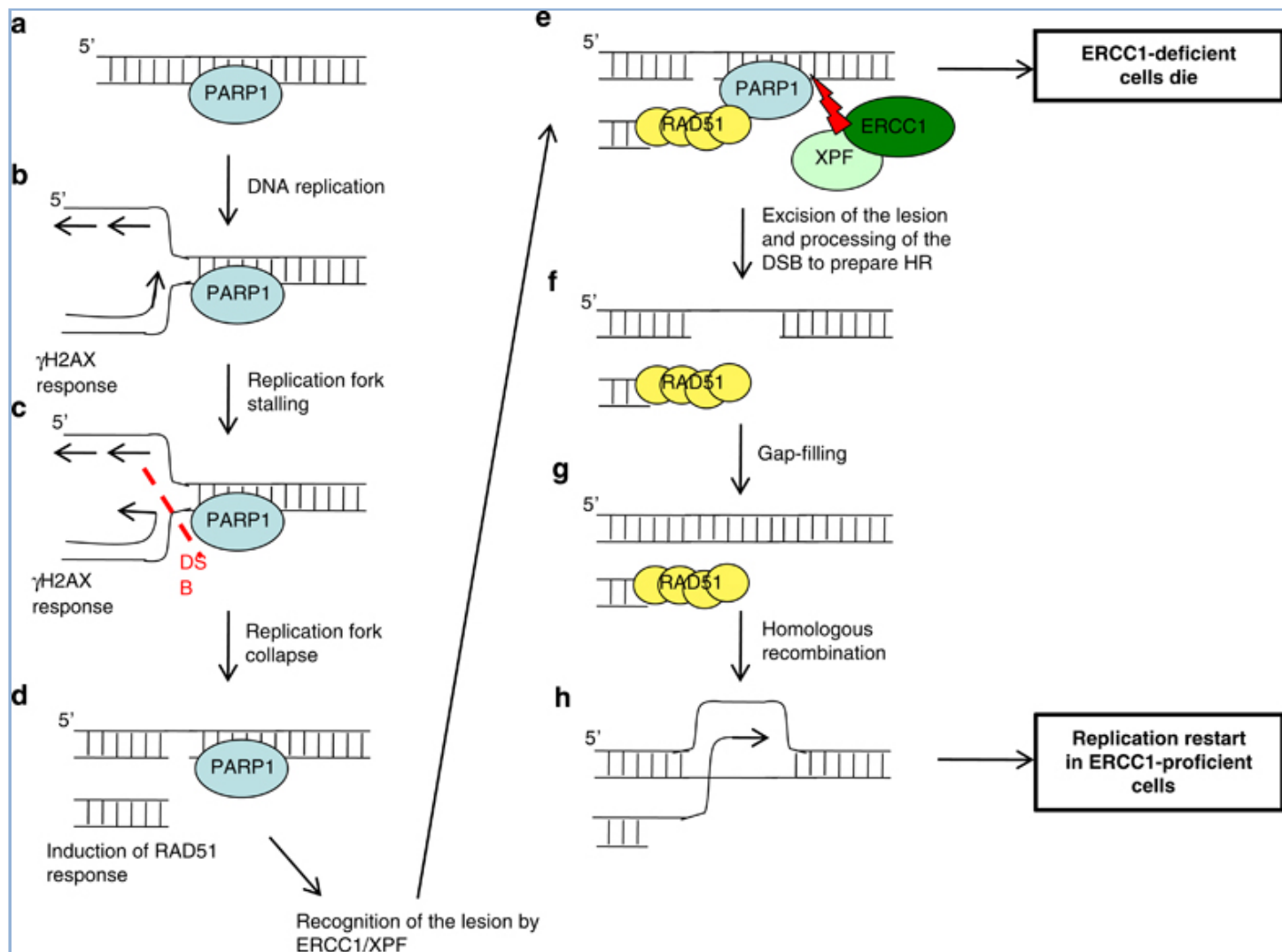
The hypersensitivity of the combination is followed by loss of efficient DSB-repair (although less error-prone)

ERCC1 deficient cell lines to explore synthetic lethality with candidate drugs/pathways



PARP1/2 inhibitors selectively kills ERCC1-deficient cells

PARP is trapped onto the DNA by the PARP inhibitor leading to replication fork stalling and collapse

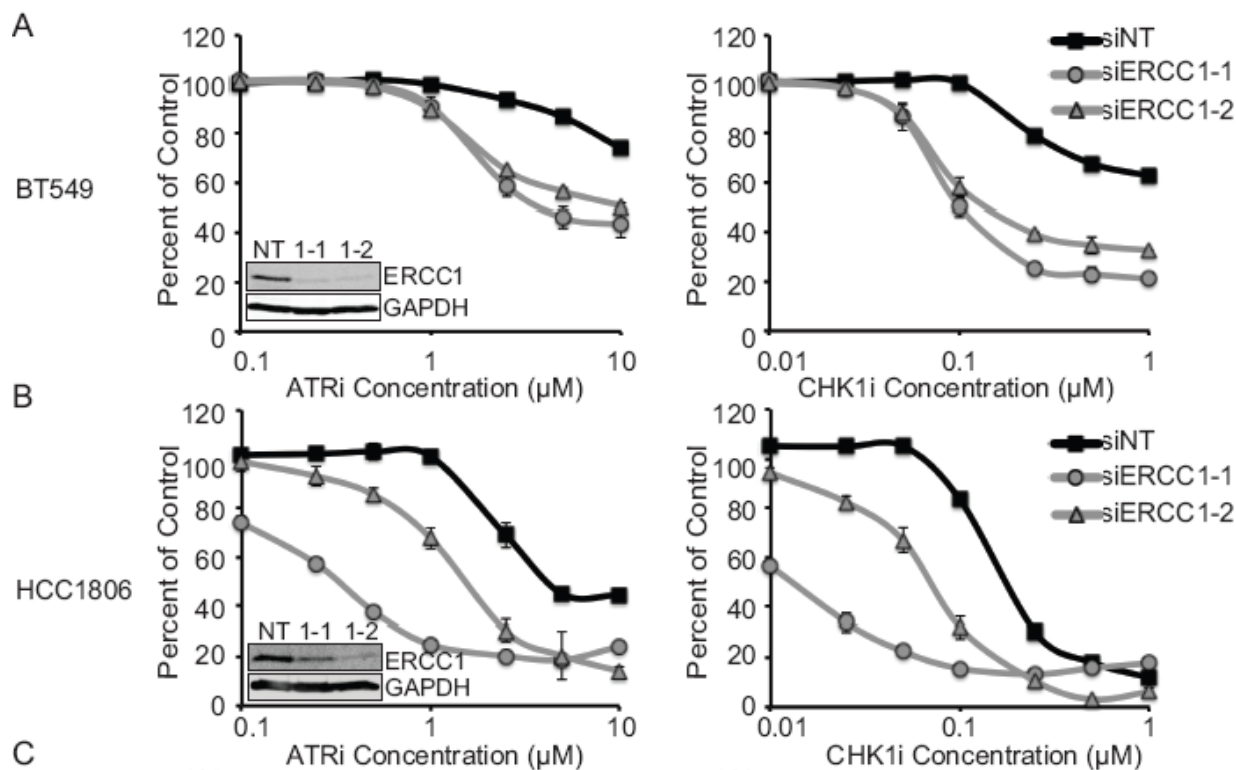


ATR pathway inhibition is synthetically lethal in cancer cells with ERCC1 deficiency

Kareem N Mohni, Gina M Kavanaugh and David Cortez

Cancer Res Published OnlineFirst March 24, 2014.

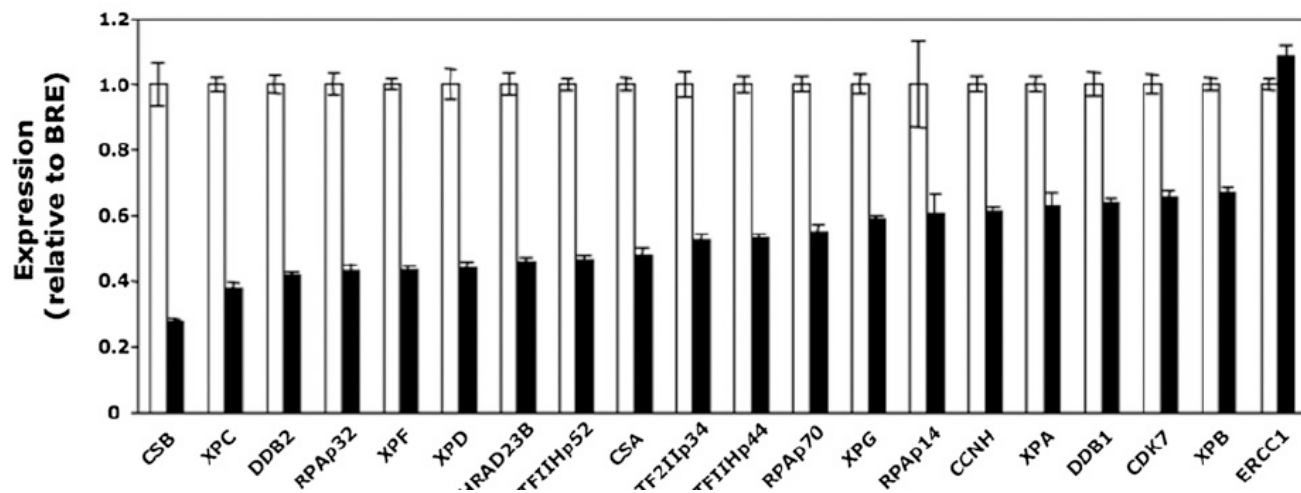
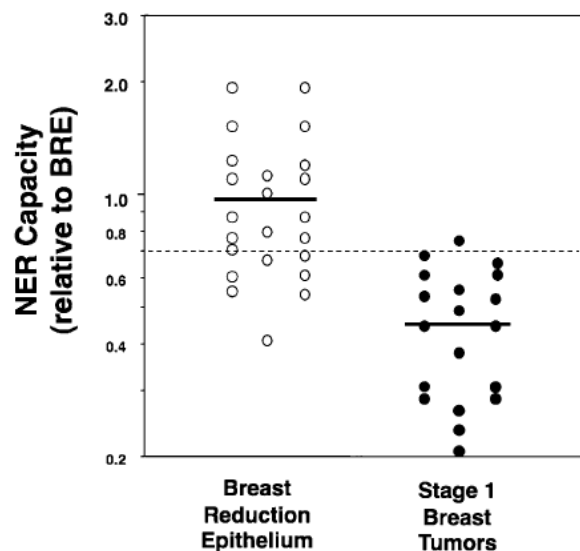
Breast cancer cell lines



Nucleotide excision repair deficiency is intrinsic in sporadic stage I breast cancer

Jean J. Latimer^{a,b,1}, Jennifer M. Johnson^{b,c}, Crystal M. Kelly^a, Tiffany D. Miles^b, Kelly A. Beaudry-Rodgers^d, Nancy A. Lalanne^b, Victor G. Vogel^{b,e}, Amal Kanbour-Shakir^f, Joseph L. Kelley^a, Ronald R. Johnson^g, and Stephen G. Grant^{b,h}

^aDepartment of Obstetrics, Gynecology and Reproductive Sciences, Magee-Womens Hospital, University of Pittsburgh Medical Center, Pittsburgh, PA 15213-3180; ^bHillman Cancer Center, University of Pittsburgh Cancer Institute, Pittsburgh, PA 15232-1301; ^cProgram in Biochemistry and Molecular Genetics, University of Pittsburgh School of Medicine, Pittsburgh, PA 15261; ^dDepartment of Human Genetics, Graduate School of Public Health, University of Pittsburgh, Pittsburgh, PA 15261; Departments of ^eMedicine, ^fPathology, and ^gSurgery, Magee-Womens Hospital, University of Pittsburgh Medical Center, Pittsburgh, PA 15213-3180; and ^hDepartment of Environmental and Occupational Health, Graduate School of Public Health, University of Pittsburgh, Pittsburgh, PA 15219-3130

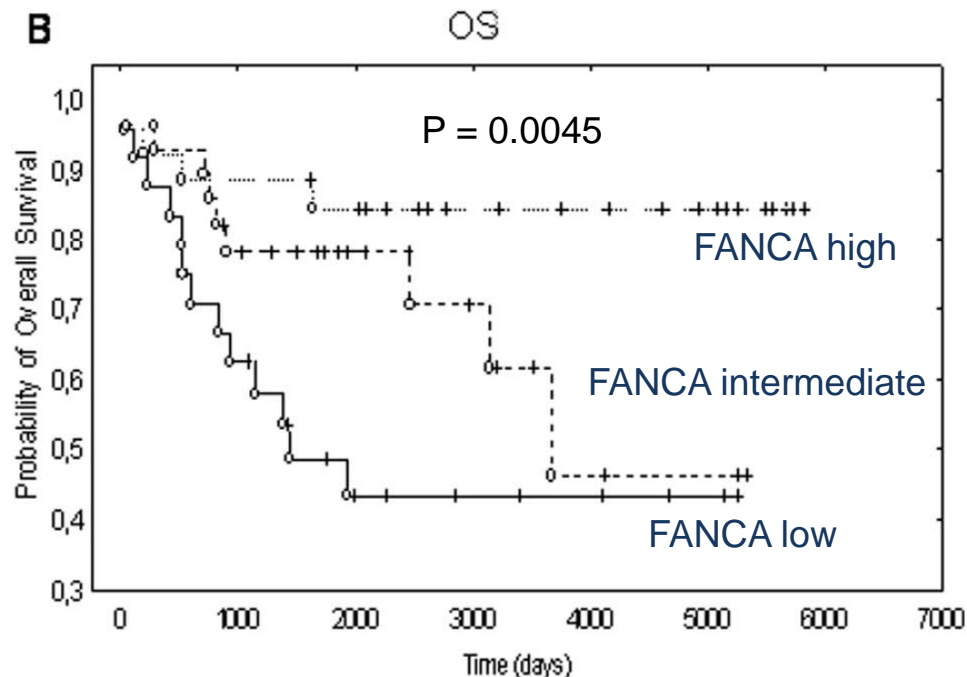


Remarkable loss of DNA repair proteins in TNBC compared to LABC

Table 3. Normalized and calibrated values (mean \pm SD and median) of the different DNA repair genes in tumor samples.

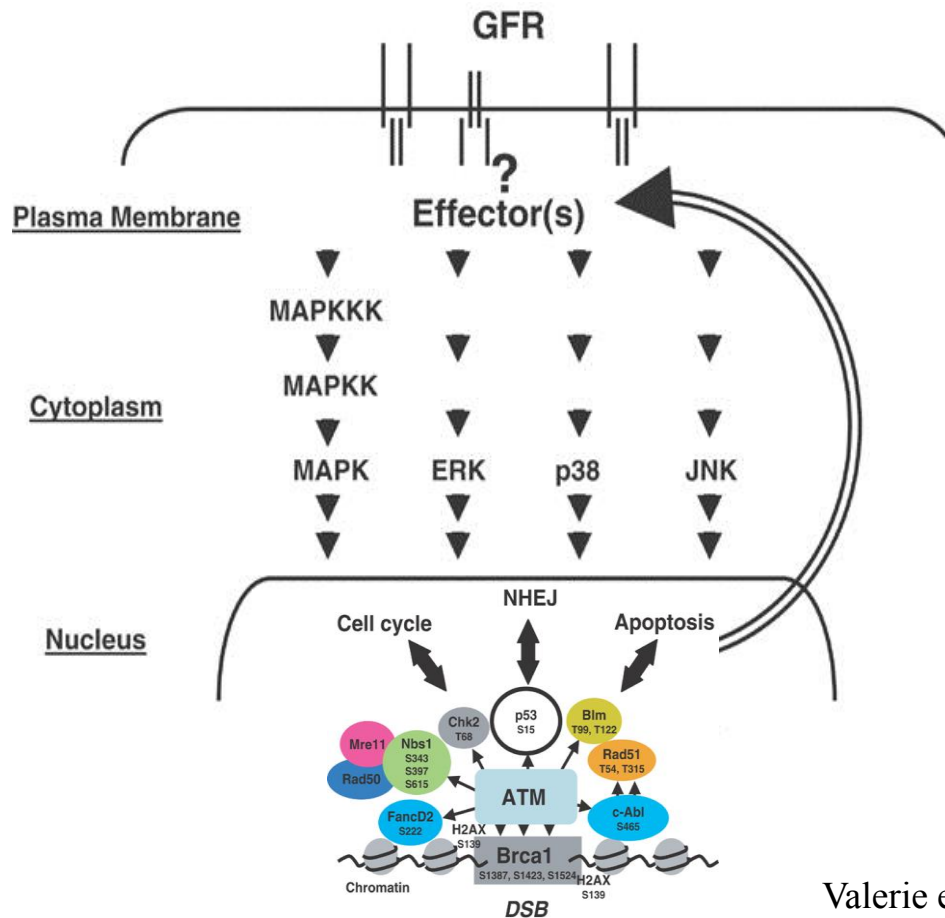
PATH WAY	GENE	TNBC (N = 80)		LABC (N = 70)		p value*
		Mean \pm SD	Median	Mean \pm SD	Median	
BER	<i>PARP1</i>	10.87 \pm 10.92	8.250	7.05 \pm 5.46	5.221	0.0002
NER	<i>ERCC1</i>	0.71 \pm 0.66	0.633	6.87 \pm 20.77	1.040	<0.0001
	<i>XPA</i>	0.10 \pm 0.08	0.080	0.16 \pm 0.21	0.104	0.0309
	<i>XPF</i>	23.35 \pm 55.61	5.108	68.93 \pm 107.05	21.996	<0.0001
	<i>XPG</i>	0.66 \pm 0.71	0.468	1.51 \pm 3.52	0.287	0.1534
	<i>XPD</i>	0.276 \pm 0.360	0.186	6.68 \pm 10.26	3.662	<0.0001
FA	<i>BRCA1</i>	0.028 \pm 0.025	0.020	0.83 \pm 3.36	0.062	<0.0001
	<i>FANCA</i>	0.138 \pm 0.279	0.057	0.27 \pm 0.71	0.094	0.8293
	<i>FANCC</i>	0.005 \pm 0.007	0.003	0.002 \pm 0.050	0.003	0.3627
	<i>FANCD2</i>	0.08 \pm 0.124	0.054	4.45 \pm 23.32	0.150	<0.0001
	<i>FANCF</i>	0.026 \pm 0.027	0.018	0.451 \pm 1.170	0.031	<0.0001
	<i>PALB2</i>	0.453 \pm 0.563	0.306	5.91 \pm 23.03	2.010	0.0006
SEN SOR	<i>CHK1</i>	0.179 \pm 0.300	0.108	11.51 \pm 46.44	1.779	<0.0001

FANCA mRNA expression in TNBC



Ribeiro E et al., PLoS ONE, 2013

DNA repair and targeted therapies



Valerie et al., Oncogene 2003

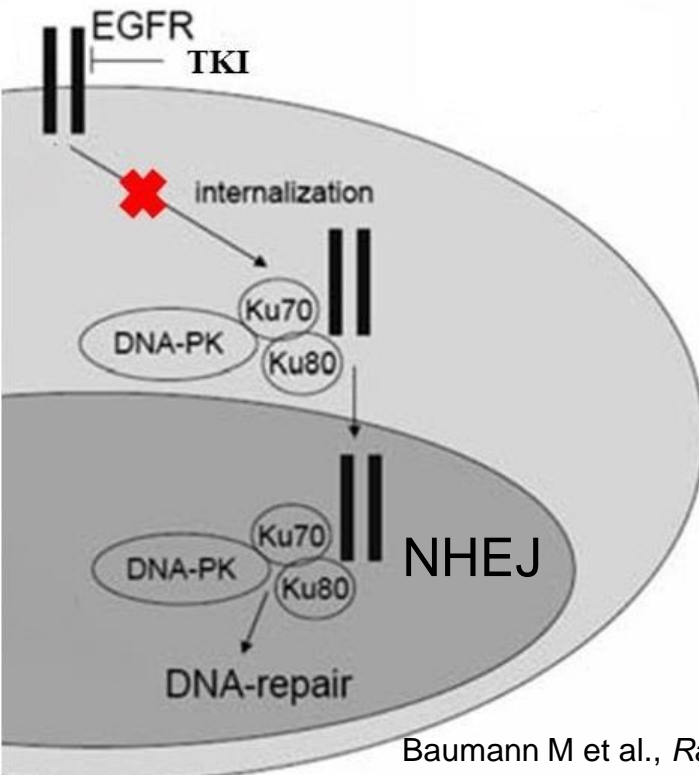
Recent example: EGFR cooperates with glucose transporter SGLT1 to enable chromatin remodeling in response to ionizing radiation. Dittmann K, et al. Radiother Oncol. 2013

Inhibiting PI3K/AKT and Ras/ERK pathways

Intracellular distribution of DNA repair proteins

Transcription of DNA repair genes

Phosphorylation of DNA repair proteins

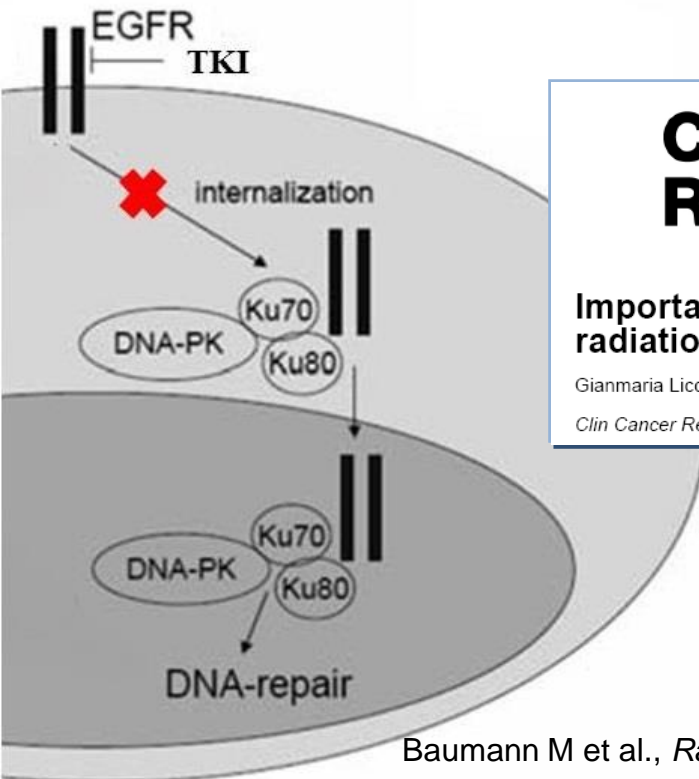


Inhibiting PI3K/AKT and Ras/ERK pathways

**Intracellular distribution
of DNA repair proteins**

Transcription of DNA
repair genes

Phosphorylation of DNA
repair proteins



Clinical Cancer Research

ACR

Importance of EGFR/ERCC1 interaction following radiation-induced DNA damage

Gianmaria Liccardi, John A. Hartley and Daniel Hochhauser

Clin Cancer Res Published OnlineFirst April 29, 2014.

**Error-prone
Alt-NHEJ**

Inhibiting PI3K/AKT and Ras/ERK pathways

↓

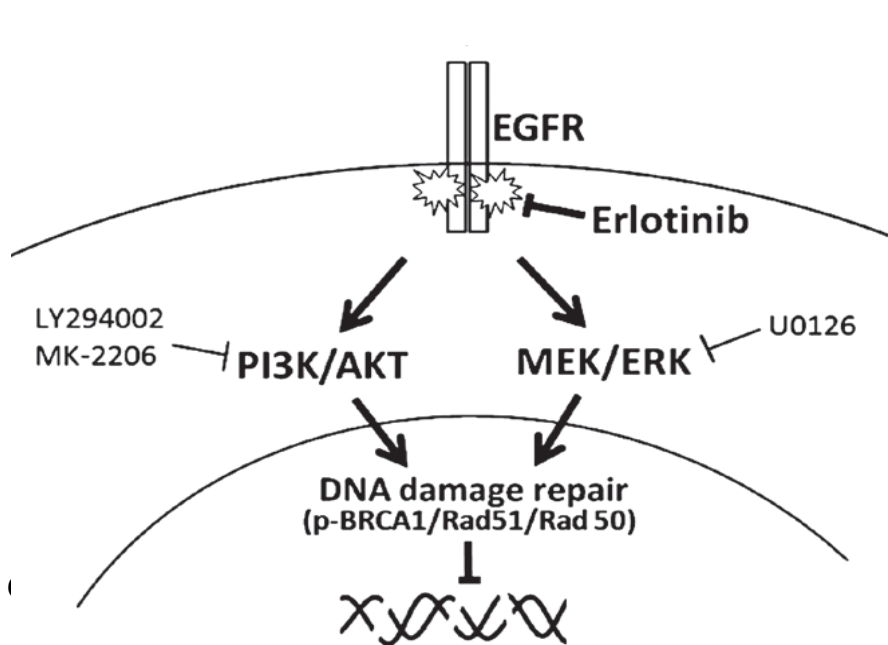
Intracellular distribution
of DNA repair proteins

↓

**Transcription of DNA
repair genes**

↓

Phosphorylation of DNA
repair proteins



↓ Rad51/Rad50 expression

↓ HRR

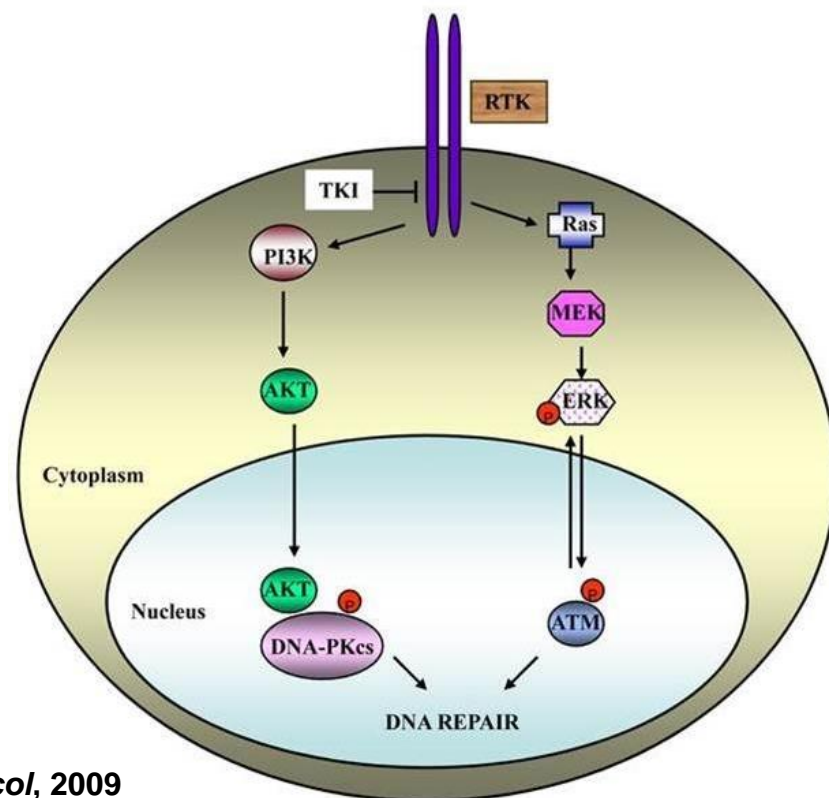
↑ DSBs

Inhibiting PI3K/AKT and Ras/ERK pathways

↓
Intracellular distribution
of DNA repair proteins

↓
Transcription of DNA
repair genes

↓
**Phosphorylation of DNA
repair proteins**



↓
AKT and
PI3K

↓
DNA-PKcs phosphorylation
(NHEJ)

↓
ERK

↓
ATM phosphorylation
(NHEJ and HRR)

PI3K inhibition increases indicators of DNA damage such as γ -H2AX, but decreases Rad51 foci and BRCA1 expression allowing synthetic lethality approaches (with PARPi) in HRR proficient cells

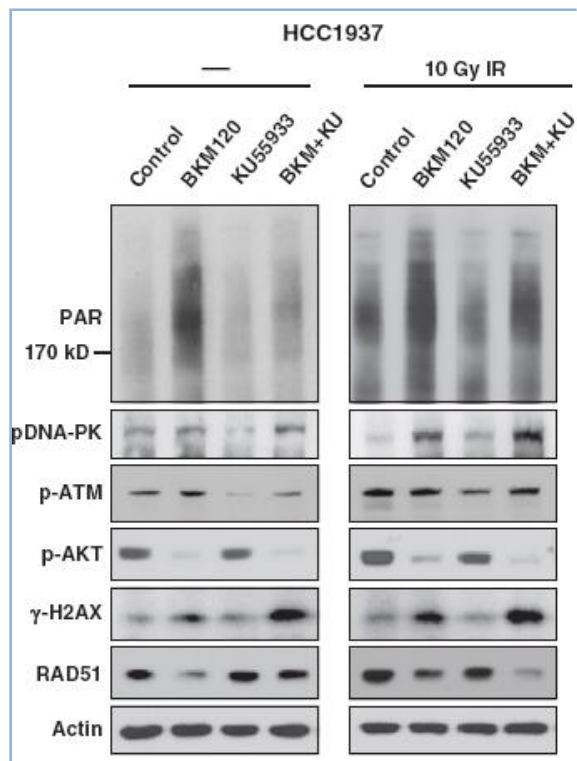
CANCER DISCOVERY

ACR

Combining a PI3K Inhibitor with a PARP Inhibitor Provides an Effective Therapy for BRCA1-Related Breast Cancer

Ashish Juvekar, Laura N. Burga, Hai Hu, et al.

Cancer Discovery 2012;2:1048-1063. Published OnlineFirst August 22, 2012.



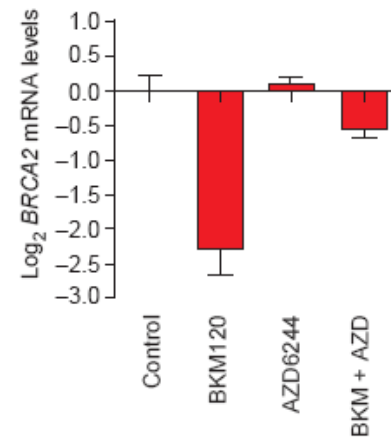
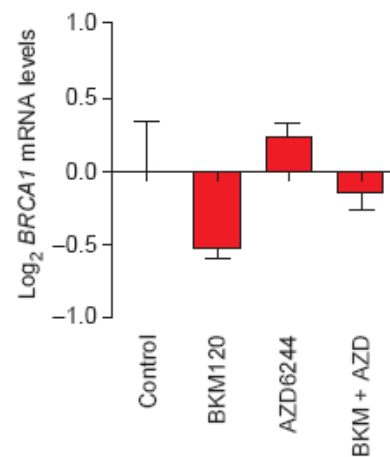
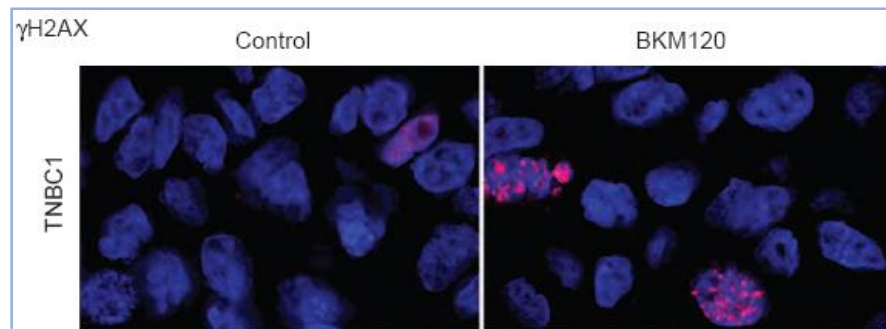
CANCER DISCOVERY

ACR

PI3K Inhibition Impairs BRCA1/2 Expression and Sensitizes BRCA-Proficient Triple-Negative Breast Cancer to PARP Inhibition

Yasir H. Ibrahim, Celina Garcia-Garcia, Violeta Serra, et al.

Cancer Discovery 2012;2:1036-1047. Published OnlineFirst August 22, 2012.



Conclusive remarks

- DNA repair pathways important in cancer development and risk
- Complex interactions between DNA repair pathways give new therapeutic opportunities (synthetic lethality)
- A more robust technology to select patients with DNA repair dysfunctionality is to be developed
- TKIs can change expression, intra-cellular distribution and activation of DNA repair proteins
- This indirect modulation inhibition of DNA repair might be combined with direct modulators (PARPi...)
- Better understanding of fundamental DNA repair mechanisms to be encouraged

Thank you !



Jean-Charles Soria
Fabrice André
Luc Friboulet
Sophie Postel-Vinay
Tony Sourisseau
Julien Adam
Florence Ponsonnailles
Angélique Robin
Nicolas Dorvault
Mei-Shiue Kuo-Cassin