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INSTITUT LADY DAVIS DE RECHERCHES MÉDICALES / LADY DAVIS INSTITUTE FOR MEDICAL RESEARCH

*Centre Bloomfield de  
recherche sur le vieillissement*

# Cancer and Aging: Two Faces of the Same Coin

## (3) Telomere Biology and Cancer-Part 2



*The Bloomfield Centre  
for Research in Aging*





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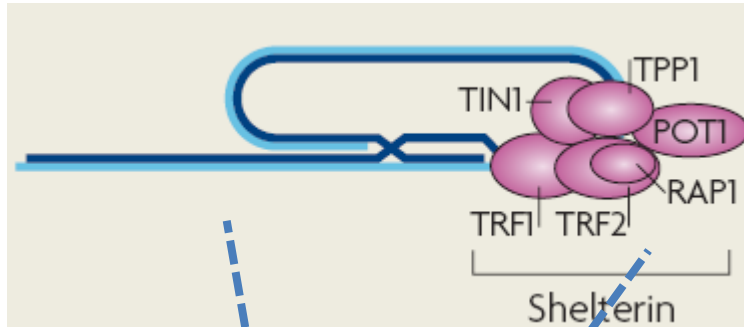


*The Bloomfield Centre  
for Research in Aging*

# Targeting Telomerase and Telomeres: Valid Anti-cancer Strategies?

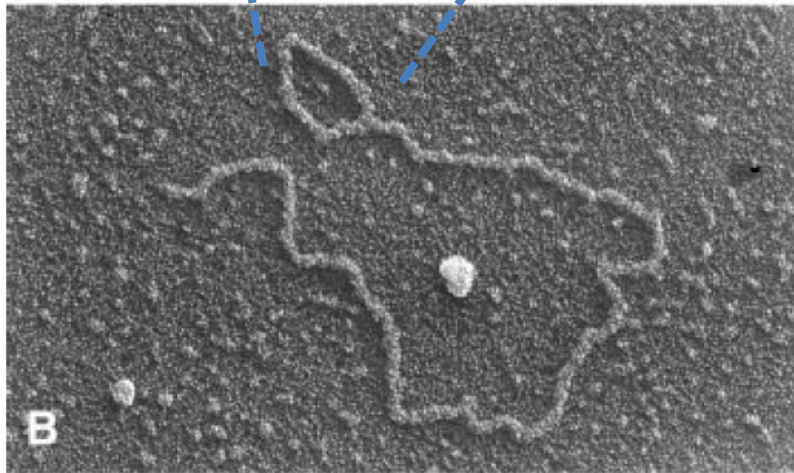
Marie Eve Brault, Johanna Mancini, Hanadi Sleiman, Chantal Autexier

# Function of telomeres: cap natural chromosome ends to make them stable structures

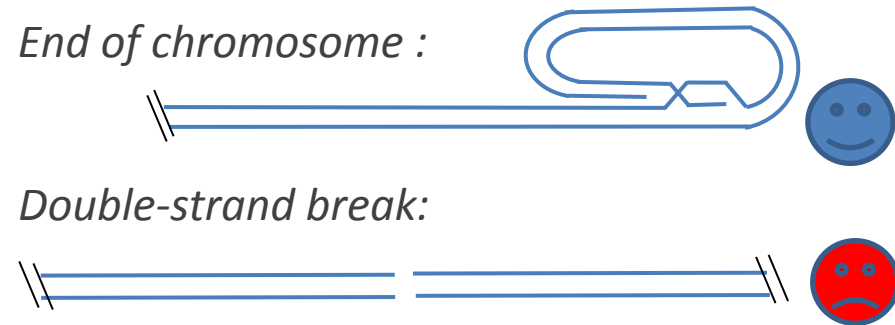


Murray, JM and Carr, AM  
Nat. Rev. Mol. Cell Biol. 2008

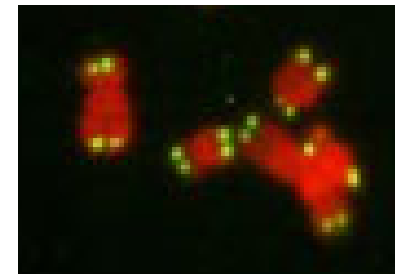
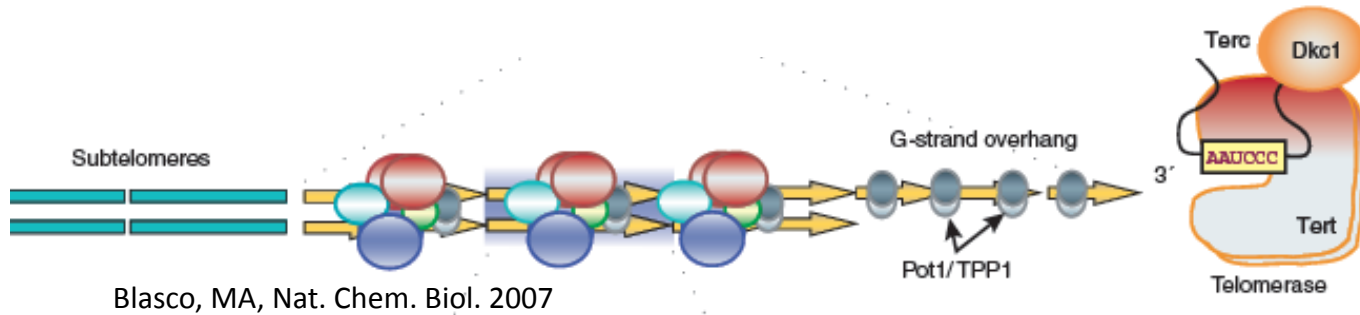
- End-to-end fusions
- Nuclease degradation
- Telomere recognized as DNA breaks
- Activation of DNA damage response
- Senescence/cell death



Griffith, JD et al., Cell 1999

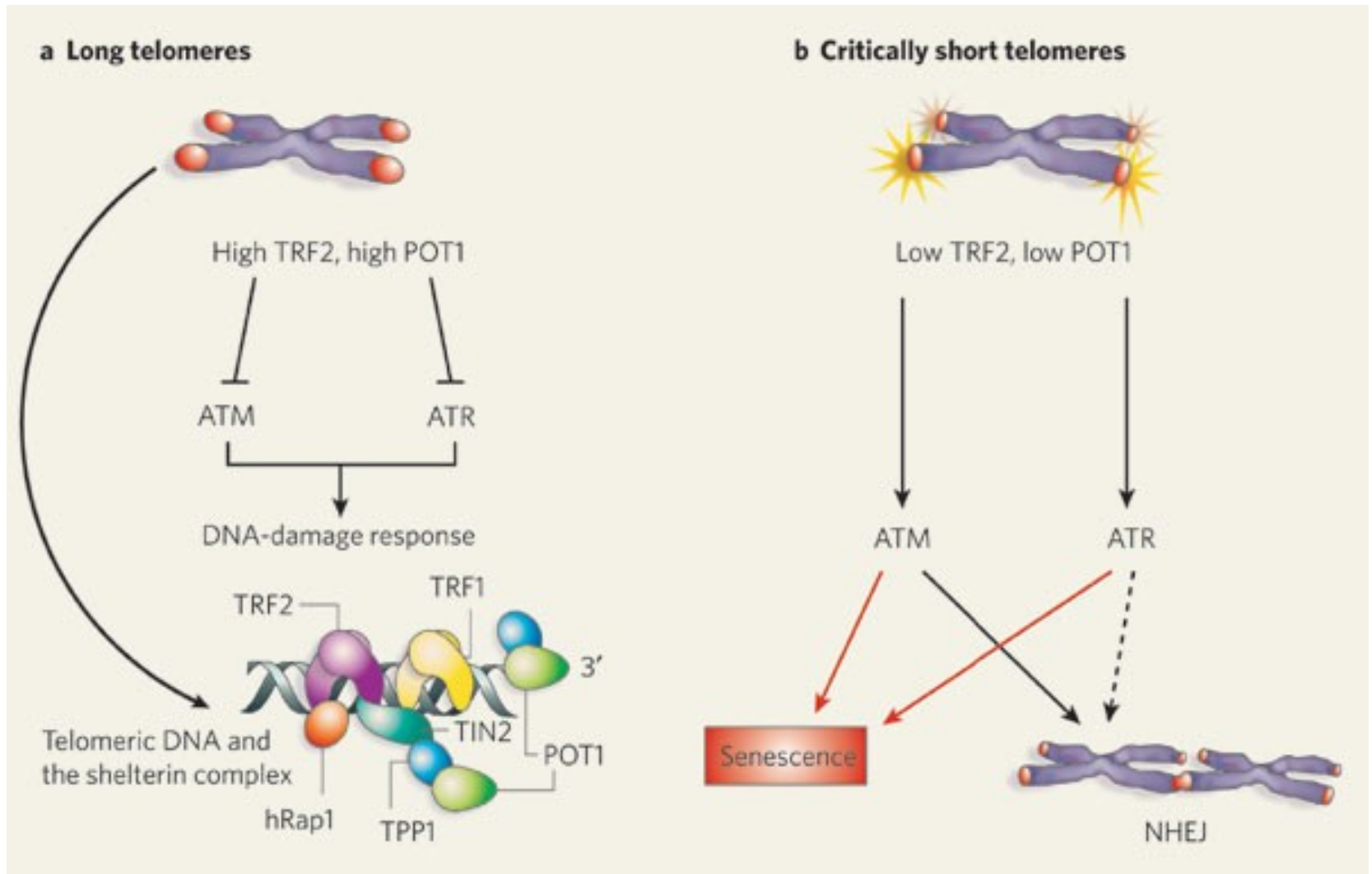


# Function of telomeres: to ensure complete DNA replication at chromosome ends via telomerase



Azzalin et al., 2008

# The T-loop structure protects telomeres

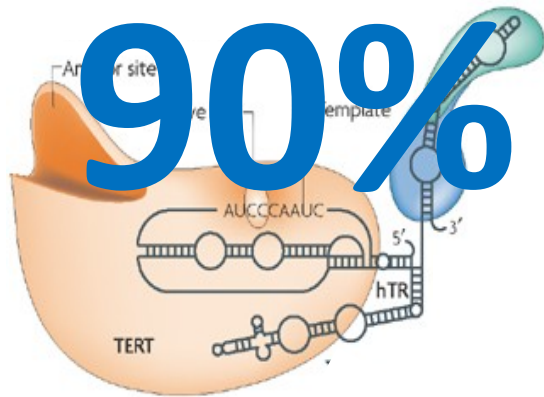


# Strategies for telomere length maintenance

a Telomerase pathway

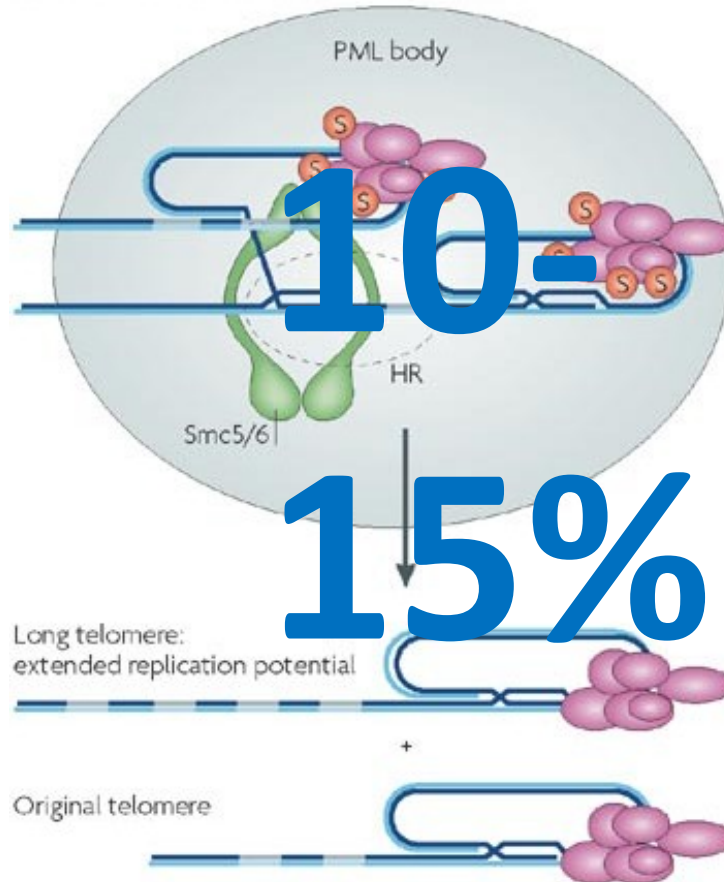


85%



90%

b Alternative lengthening of telomeres (ALT)



10%

15%

# Why does Telomerase Represent a Good Anti-cancer Target?

Universal

- 85-90% all tumors are telomerase positive

Critical

- Telomerase activity/Telomere maintenance is required for the transformed phenotype

Specific

- Most normal human somatic cells have no or very low telomerase whereas cancer cells upregulate telomerase expression

# Issues to Consider When Targeting Telomerase Function

## Lag phase

- Lag phase between the time telomerase is inhibited and the time telomeres of the cancer cells will have shortened sufficiently to produce detrimental effects on cellular proliferation

## Drug Resistance

- Telomerase inhibitors might result in the emergence of drug-resistant cancer cells (reactivation of telomerase or of the alternative lengthening of telomeres (ALT) pathway)

## ALT mechanism

- Alternative 'recombination based' mechanisms for telomere maintenance have been reported in 15% of human cancers.



# Telomerase and telomeres as potential targets

## Catherine Lauzon

Inhibition of telomere integrity  
in combination with chemotherapy



## Johanna Mancini and Hanadi Sleiman

Characterization of G-quadruplex ligands as  
telomerase inhibitors and/or disruptors  
of telomere integrity in cancer cells



## May Shawi and Raquel Aloyz

Inhibition of telomerase in combination with  
chemotherapy in chronic lymphocytic leukemia



## Marie Eve Brault , Nahid Golabi and Johanna Mancini

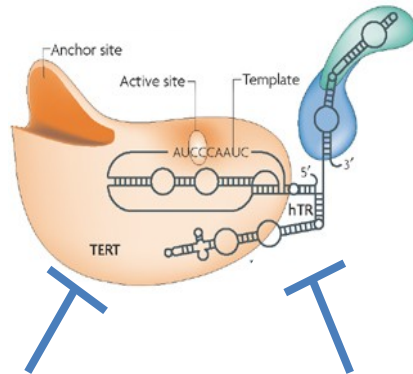
Regulation of telomere maintenance by recombination



**Telomeric recombination as a potential resistance mechanism in response to telomere dysfunction**

# Targeting the integrity of telomeres versus telomerase

Telomerase



BIBR1532

GRN163L

*Telomere erosion*

« slow » pathway

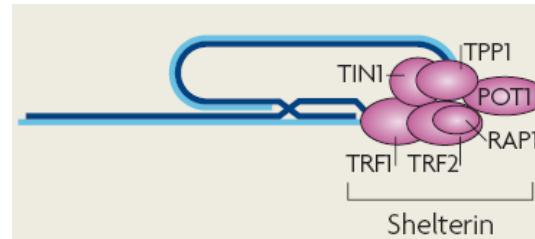
Sensescence ( $\uparrow$ p16, p21)

Telomere shortening

Decrease proliferation

Decreased tumor growth in vivo

Enhance the response to cancer drug treatment



Telomere

Q-quadruplex ligands  
ex: Telomestatin

*Telomere uncapping*

Rapid pathway

$\uparrow$  apoptosis

$\uparrow$  DNA damage response ( $\uparrow$  $\gamma$ H2AX, 53BP1, P-ATM)

Displacement of POT1 and/or TRF2

Chromosome fusions

Specific for cancer cells?

Adapted from :

[Kelland L.](#) Clin. Cancer. Res., 2007

Harley, C.B. Nat. Reviews Canc., 2008

Murray, JM and Carr, AM

Nat. Rev. Mol. Cell Biol. 2008

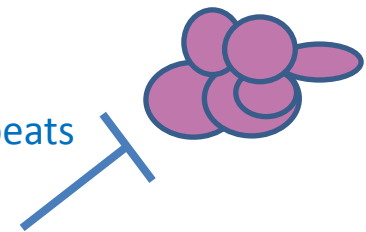
# Telomere disturbance through the expression of a telomerase RNA with a point mutation in the template region (MuA-hTR)

Telomerase-based anti-cancer approach:

**Wild-Type telomerase RNA** template 5'-CUAACCCUAA-3' specifies TTAGGG repeats



**Mutant telomerase RNA** template 5'-CAAACCCAAA-3' specifies TTTGGG repeats



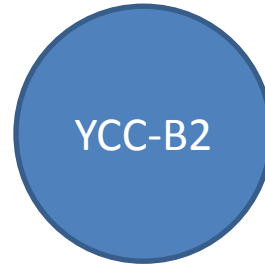
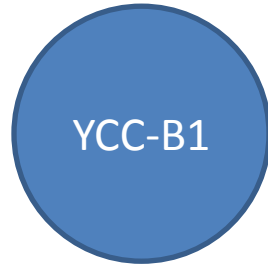
- ↓
- Tumor specific (telomerase-dependent)
  - No lag phase
  - General

# Anticancer strategies targeting telomere function by expression of template-mutated hTR

- Growth inhibition
- Altered cell cycle
- Apoptosis
- Senescence
- No effect on telomerase-negative lung fibroblasts
  
- Loss of tumor growth in xenografts
  
- Dependent on assembly of an active enzyme
- DNA damage response at telomeres
- Chromosome fusions
- Anaphase bridges
  
- Independent of p53 status
- Can be ATM dependent

Marusic L. et al. (1997) *Mol. Cell. Biol.* 17: 6394-6401; Guiducci, C. et al. (2001) *Oncogene* 20: 714-725; Kim MM. et al. (2001) *Proc. Natl. Acad. Sci.* 98: 7982-7987; Li S. et al. (2004) *Cancer Res.* 64:4833-4840; Goldkorn, A. and Blackburn, E.H. (2006) *Cancer Res.* 66: 5763-5771; Cerone, MA et al. (2006) *Oncogene* 25: 7411-7420; Stohr, B.A. and Blackburn, E.H. (2008) *Cancer Res.* 68: 5309-5317; Mahalingam, D. et al. (2011) *FEBS J.* 278, 3724-3738.

# Generation of cancer cell lines with various telomere lengths expressing template-mutated hTR

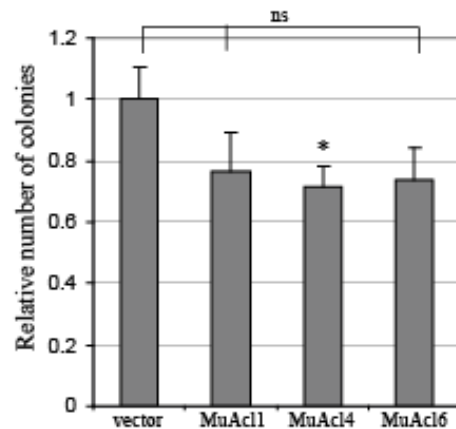


Telomere length (kb)	3.2 (short)	11 (long)	7 (intermediate)	>20 (ALT)
Telomerase	+	+	+	-

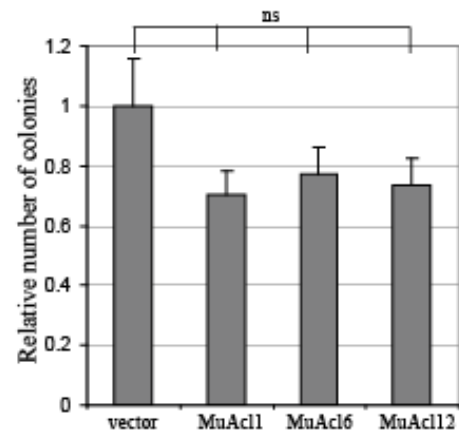
Determine effects on cell viability and proliferation upon treatment with chemotherapeutic drugs

# MuA-hTR has a mild effect on cell viability and proliferative ability

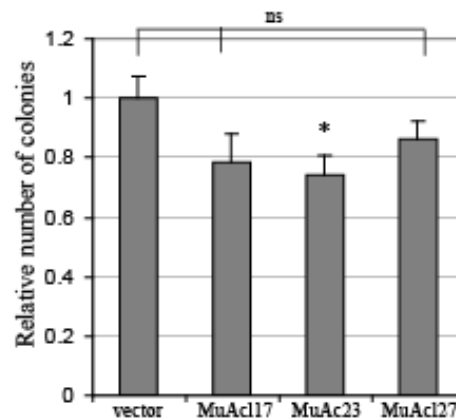
**YCC-B1 cells**



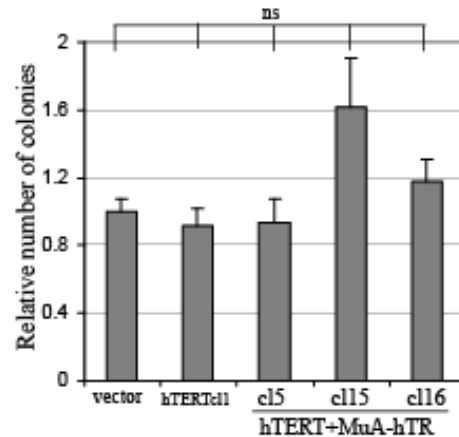
**MCF-7 cells**



**YCC-B2 cells**

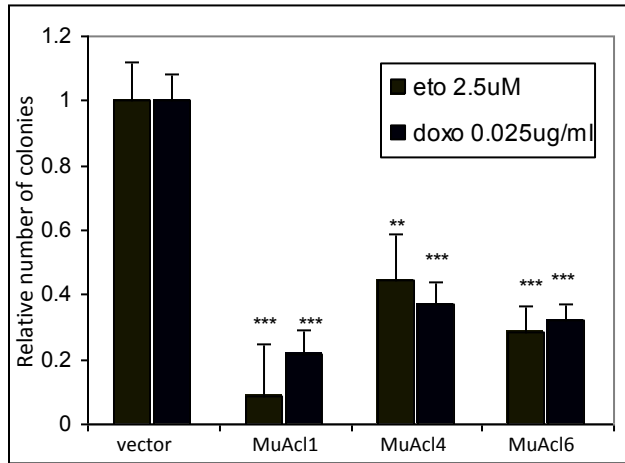


**GM847 cells**



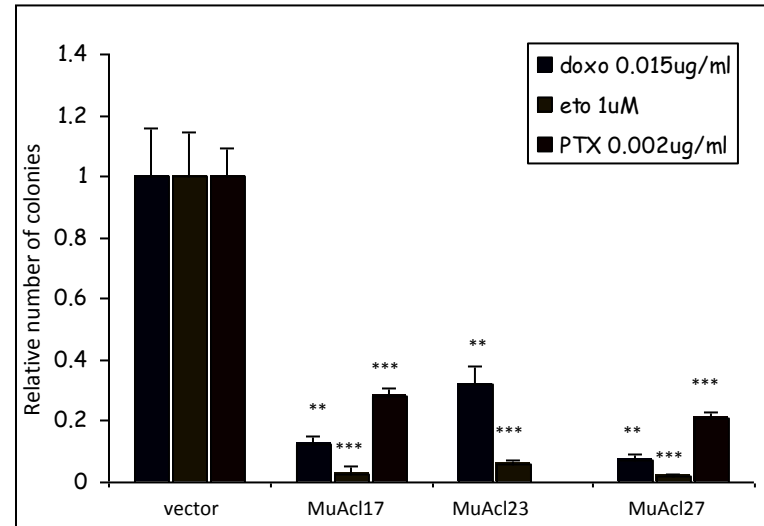
# MuA-hTR expression increases the sensitivity of cancer cells to chemotherapeutic drugs independently of telomere length and initial telomerase status

YCC-B1 (short TRF)



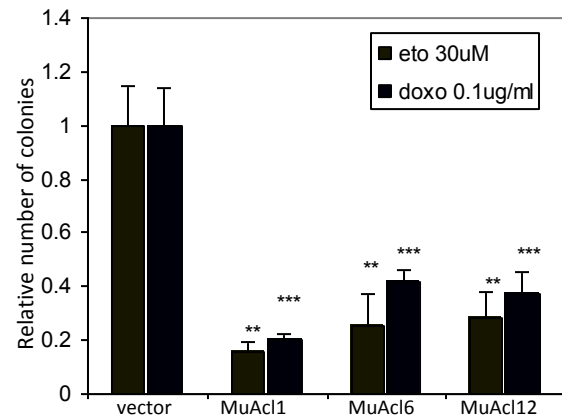
\*\*p<0.01; \*\*\*p<0.0001

YCC-B2 (long TRF)



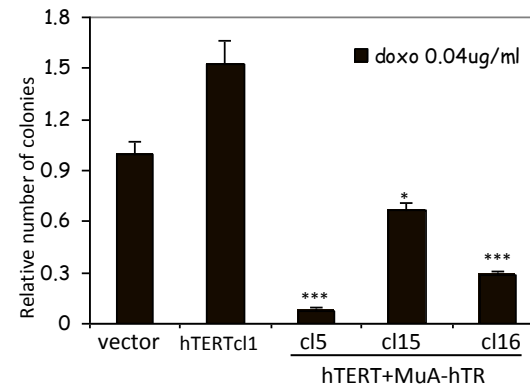
\*\*p<0.01; \*\*\*p<0.0001

MCF-7 (intermediate TRF)



\*\*p<0.01; \*\*\*p<0.0001

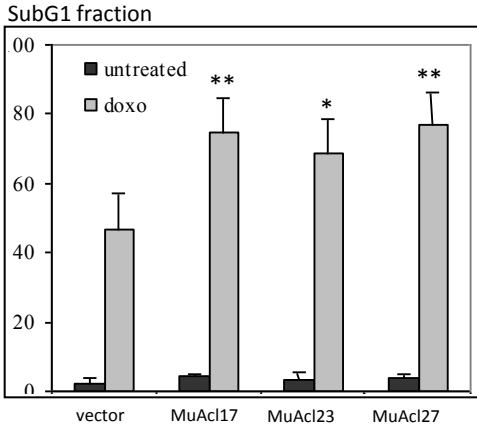
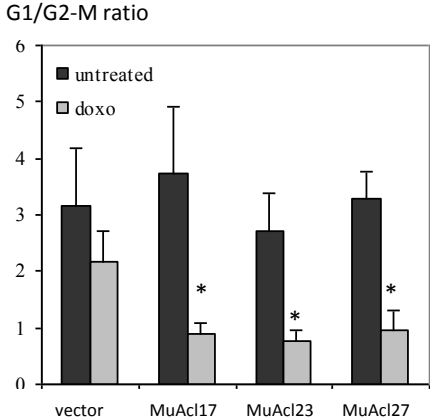
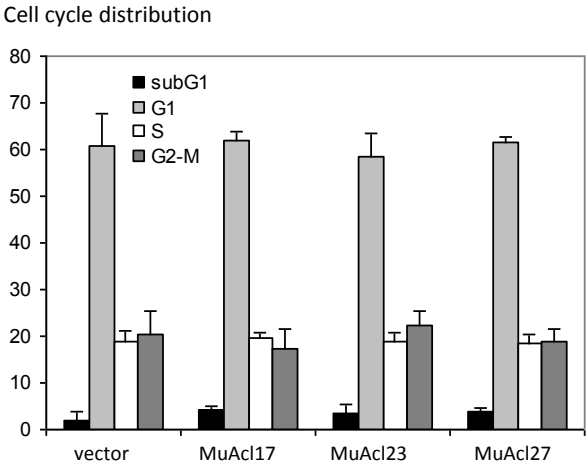
GM847 (ALT)



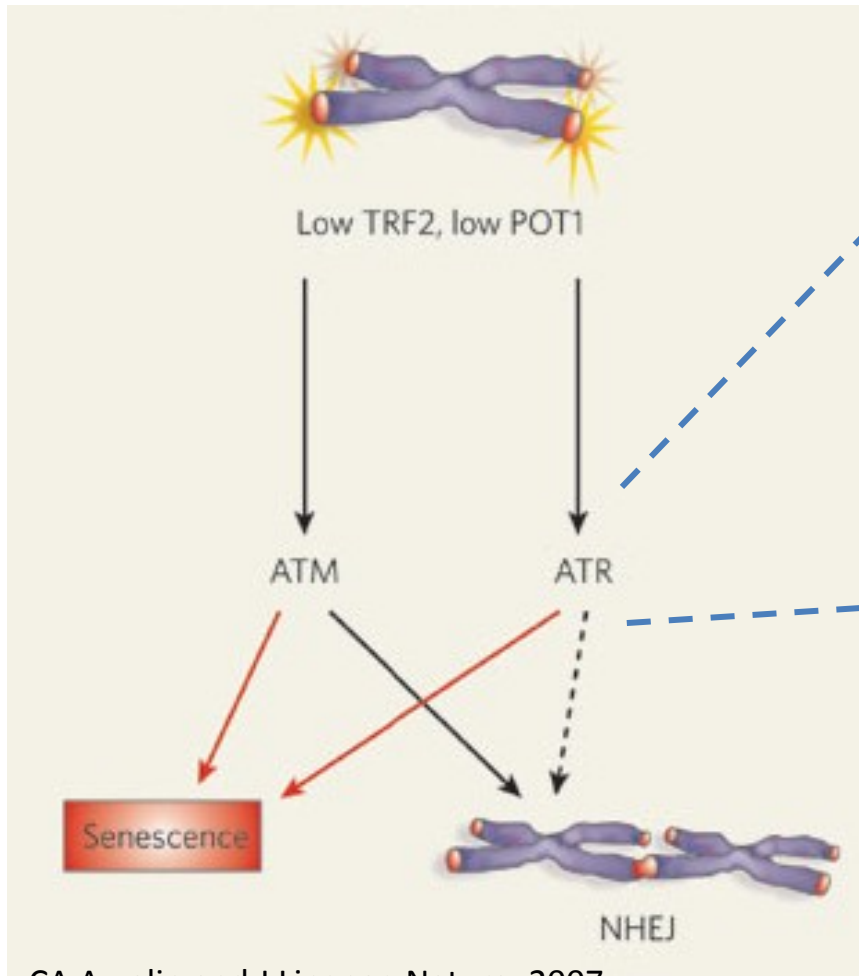
\*p<0.05; \*\*\*p<0.001



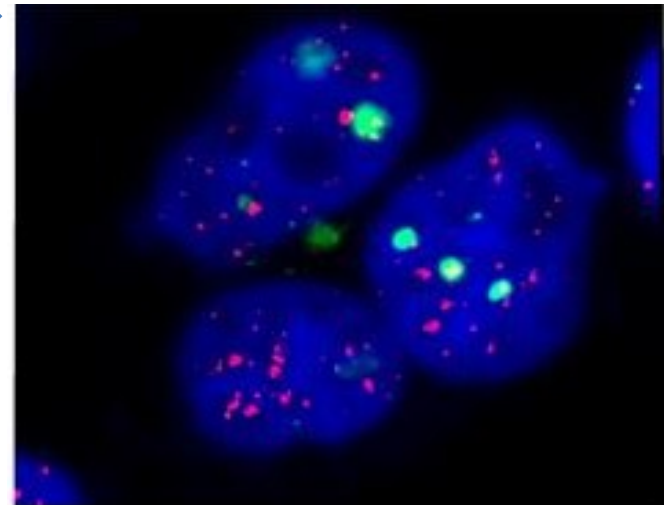
# MuA-hTR expression alters the cell cycle profile of YCC-B2 cells after drug treatment perhaps leading to the increased sensitization to drugs



# DNA damage response at the telomeres : Telomere Dysfunction-Induced Foci (TIFs)

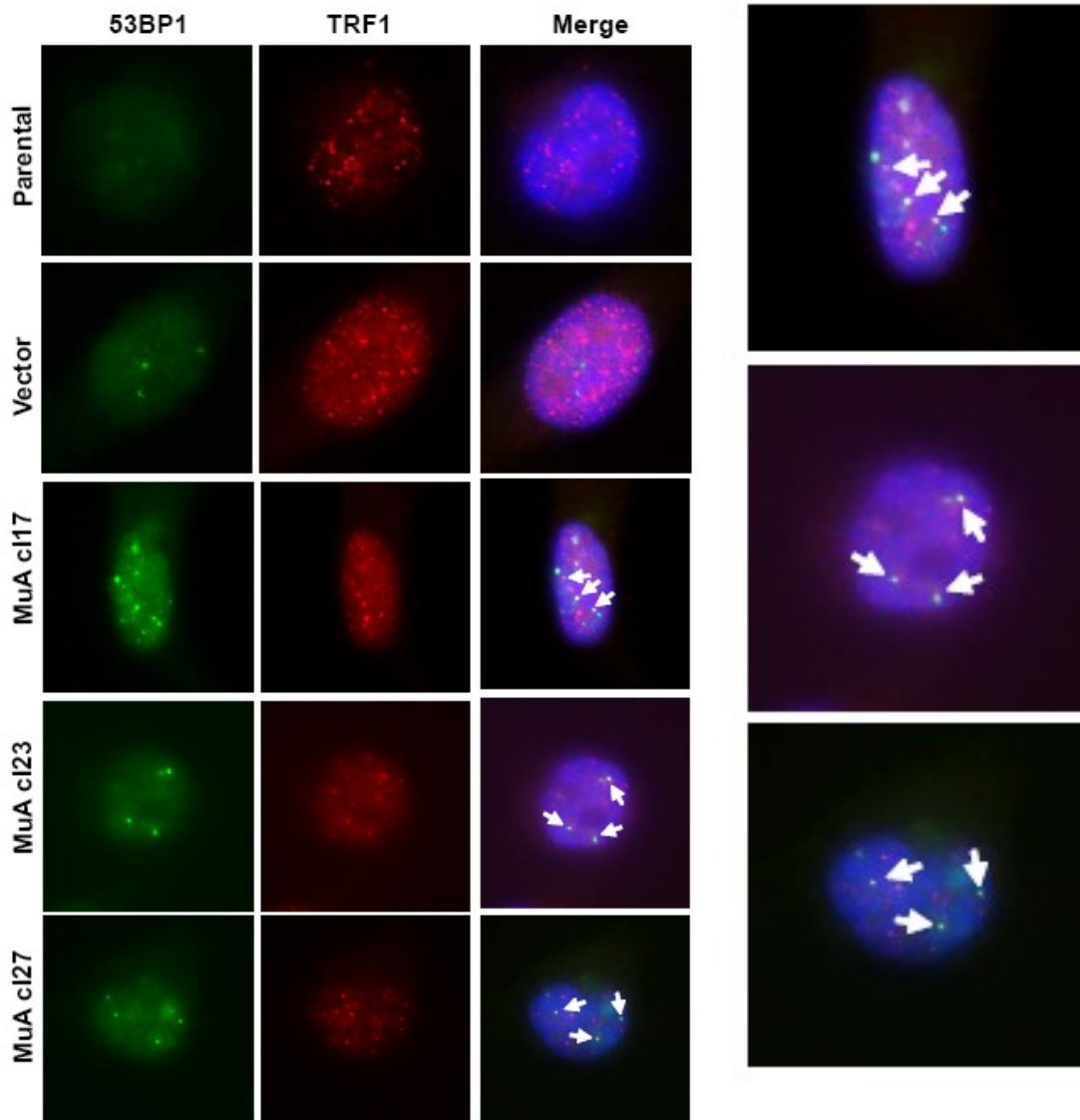


*Telomere dysfunction induced-foci (TIFs)*

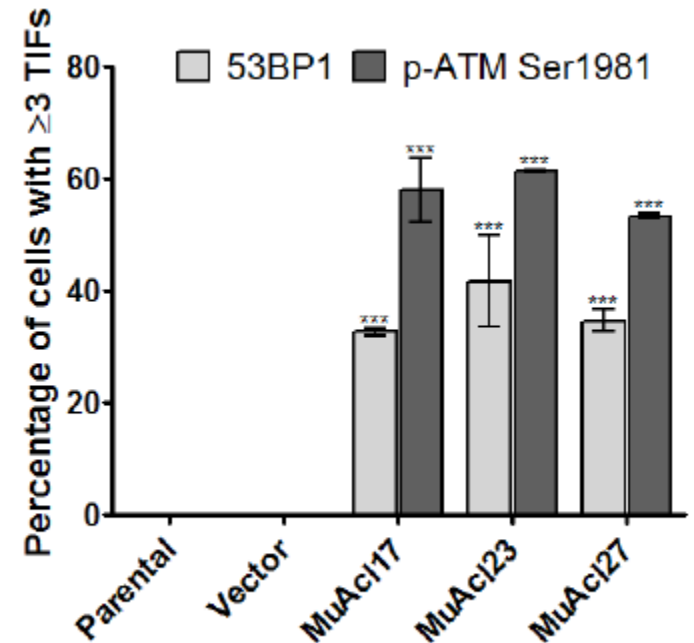
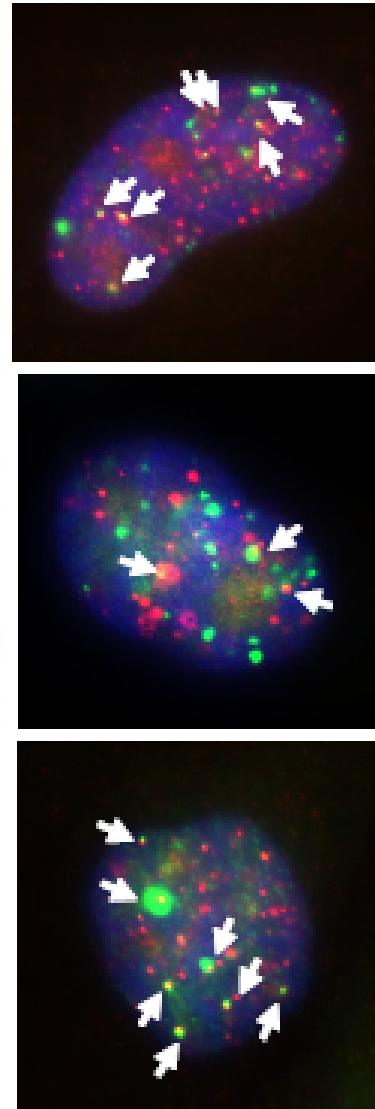
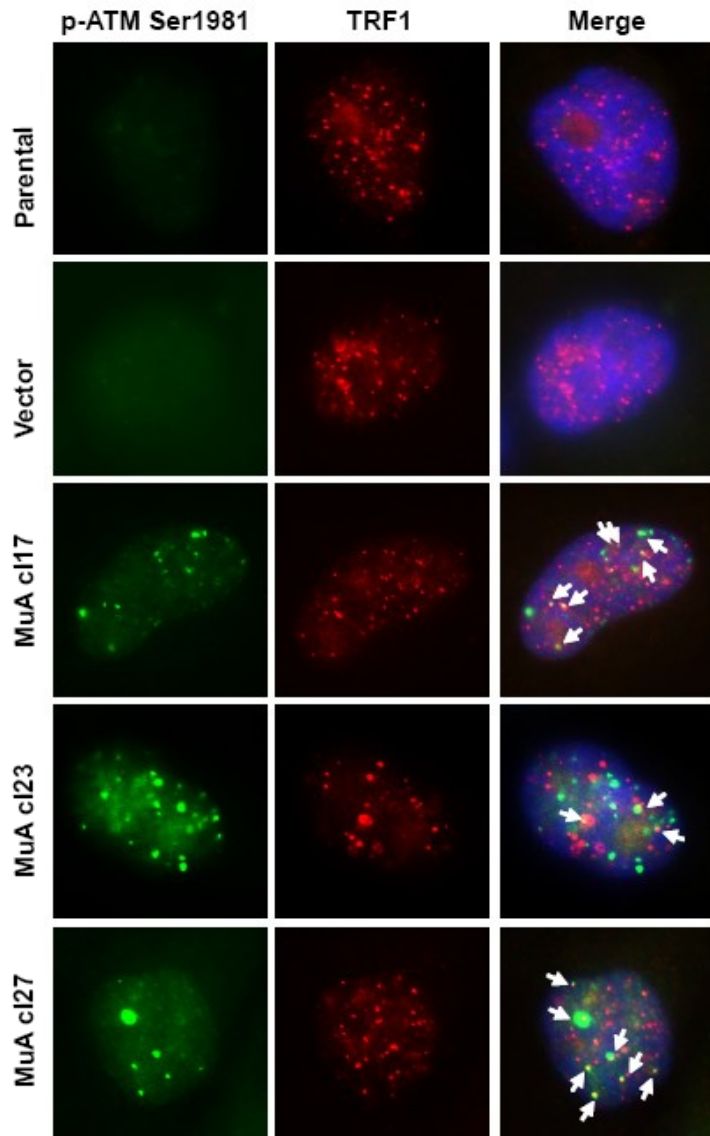


53BP1  
 $\gamma$ H2AX  
p-ATM  
MRN  
MDC1

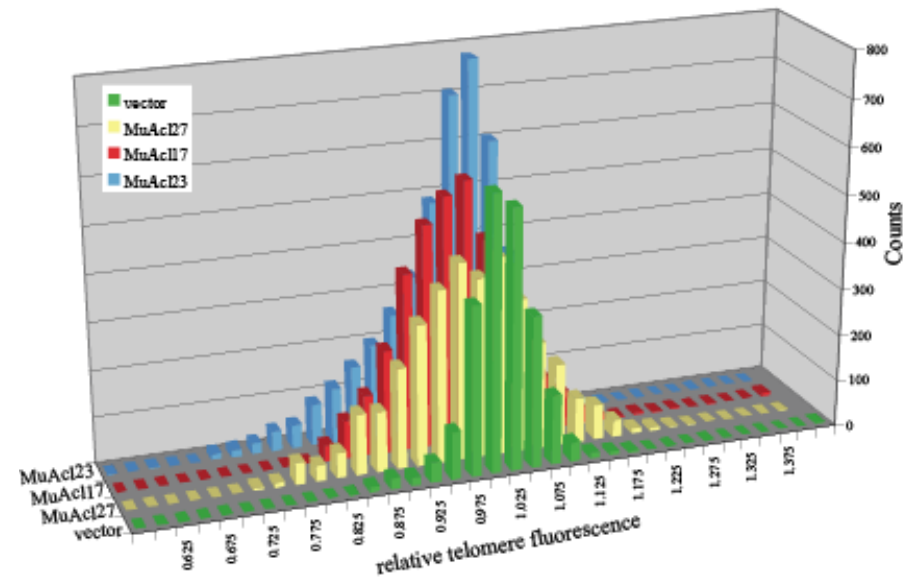
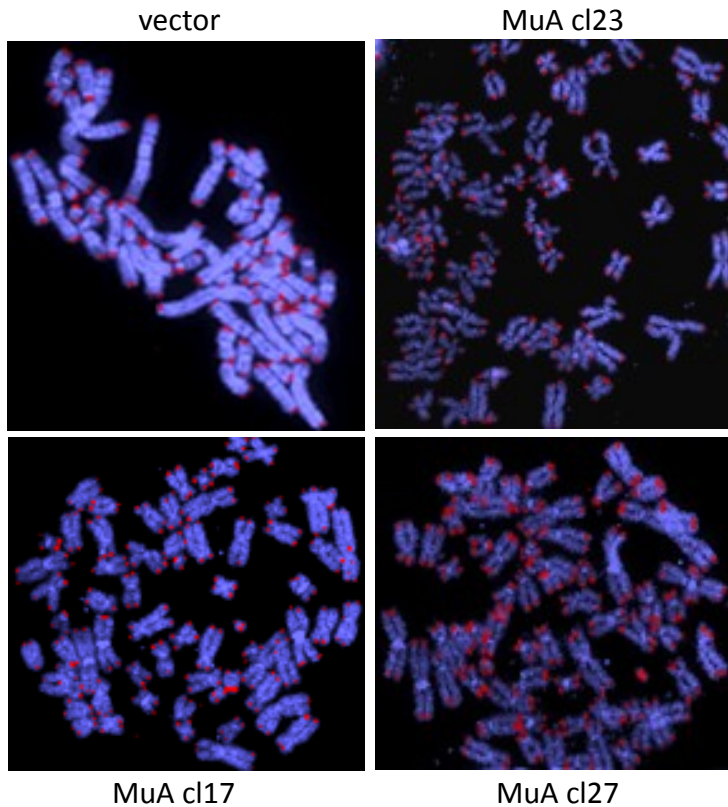
# MuA-hTR induces the formation of TIFs containing TRF1 and 53BP1 in YCC-B2 cells



# MuA-hTR induces the formation of TIFs containing TRF1 and P-ATM Ser 1981 in YCC-B2 cells



Observed a broader distribution of relative telomere lengths within cells carrying the mutant RNA, suggesting possible telomeric recombination events



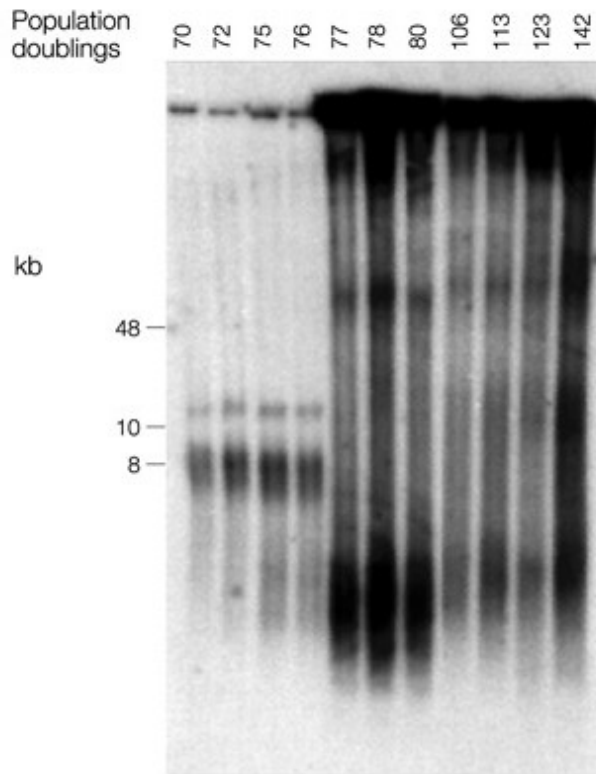
# Characteristics of ALT cells

Homologous recombination (HR) mediated events leading to:

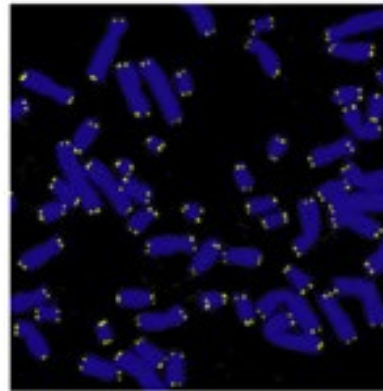
1. exceptionally long and heterogeneous telomeres, ranging from <2kb to >50kb
2. High levels of telomeric sister chromatid exchange (T-SCE)
3. Altered pq-ratios
4. extrachromosomal telomeric DNA, of circular (termed t-circles) forms

ALT-associated promyelocytic leukemia bodies (APBs) which may be the sites of recombination

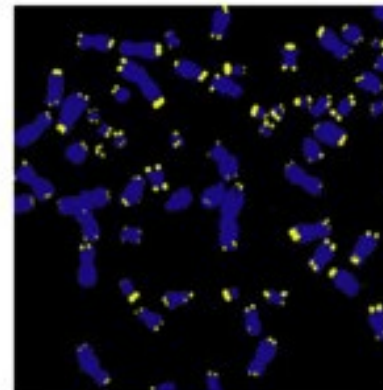
# The telomeres in ALT cells are highly heterogeneous and extremely long



Telomerase

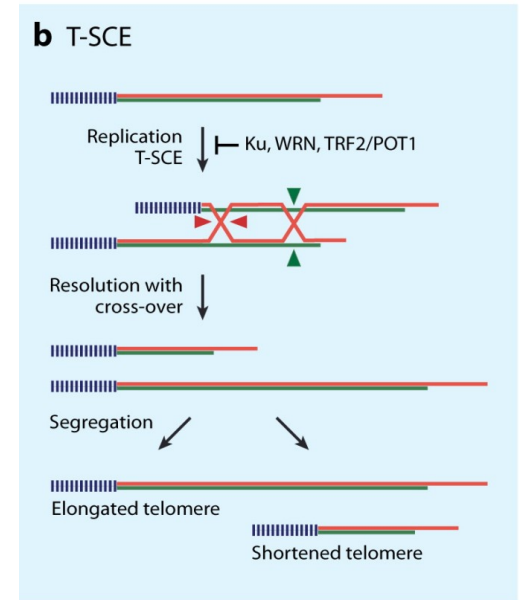
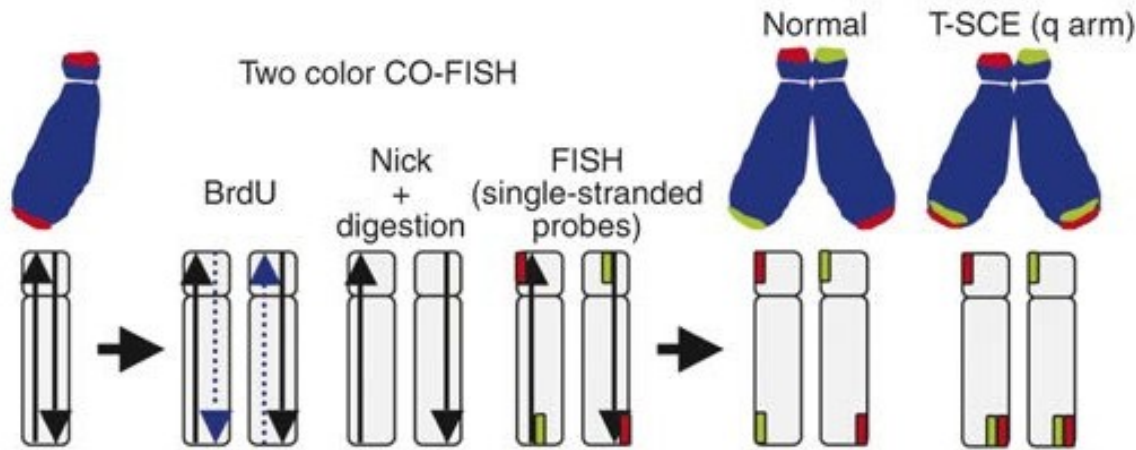


ALT





# Measuring Telomere-Sister Chromatid Exchanges (T-SCEs) : the CO-FISH Technique



Palm, W. and de Lange, T. Annu. Rev. Genet. 2008.

Lagging

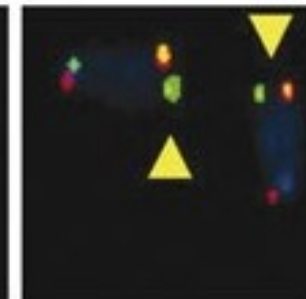
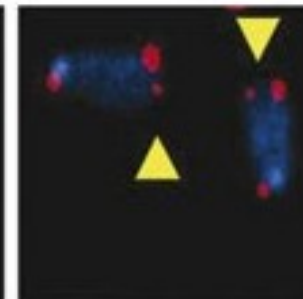
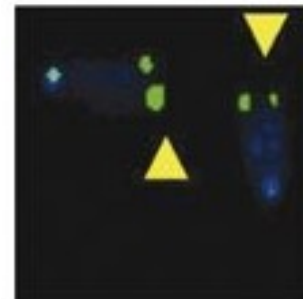
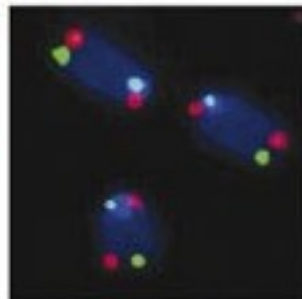
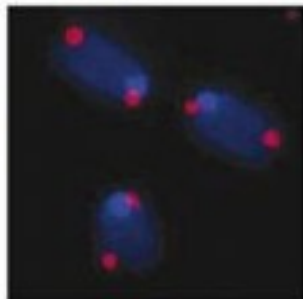
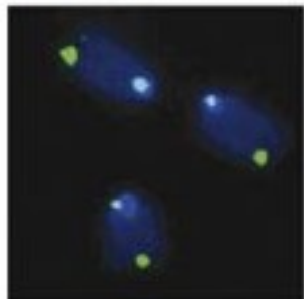
Leading

Combined

Lagging

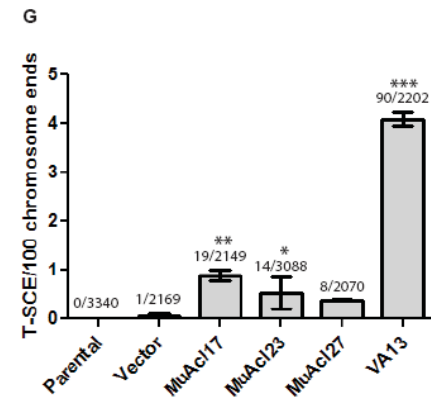
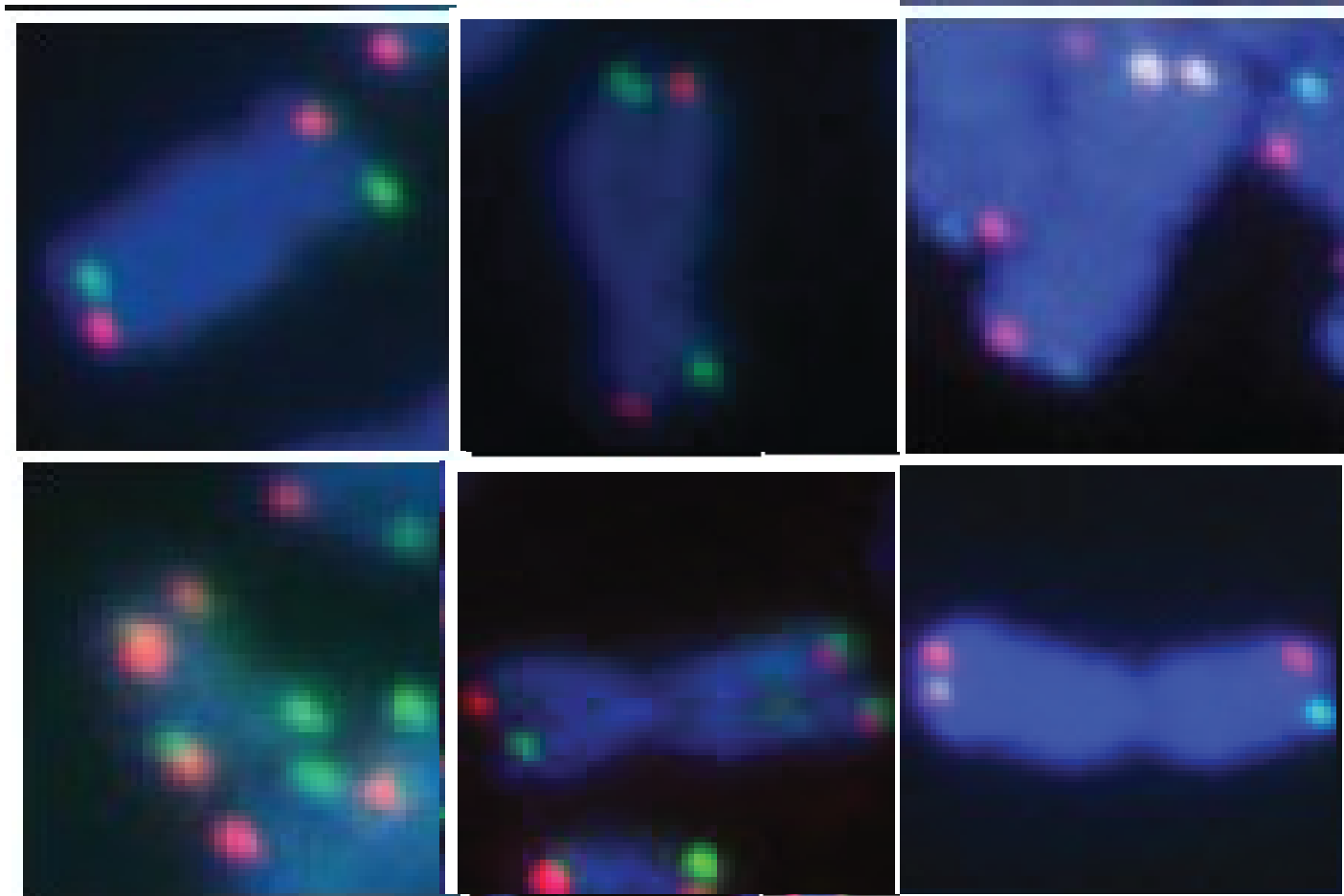
Leading

Combined

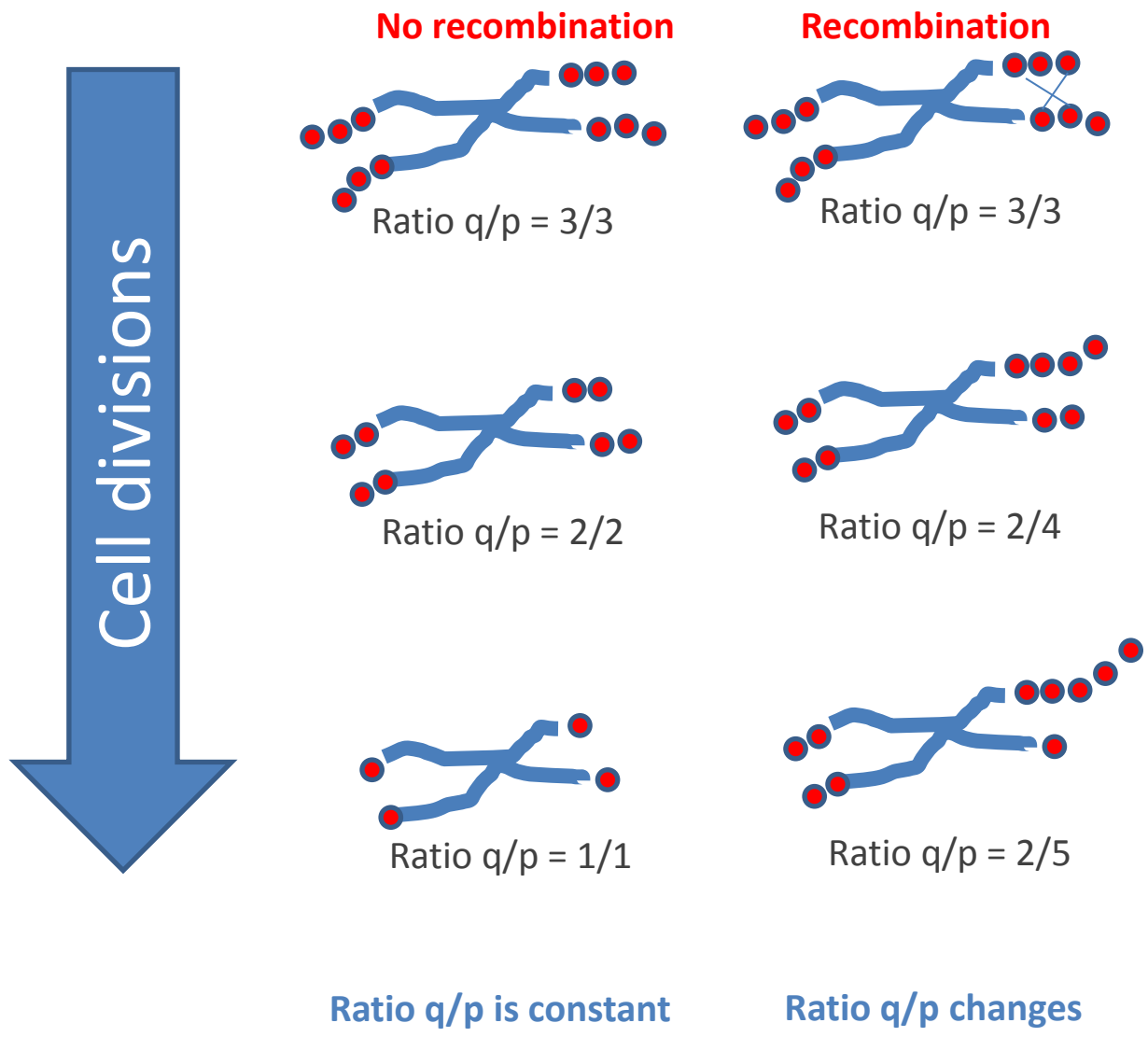




# Increased frequency of T-SCEs in the YCC-B2 cells after MuA-hTR expression

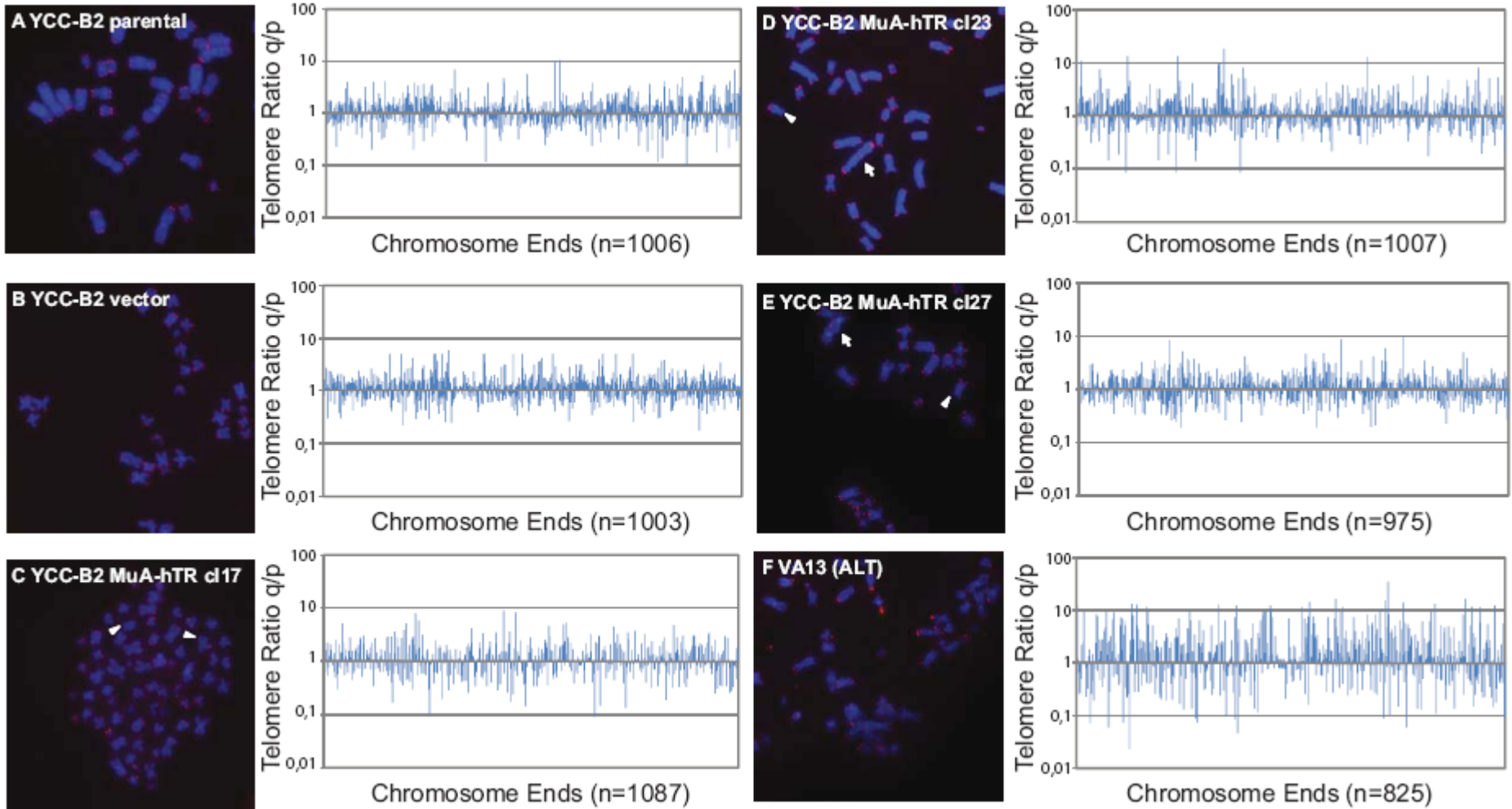


# Changes in pq-ratios indicate telomeric recombination

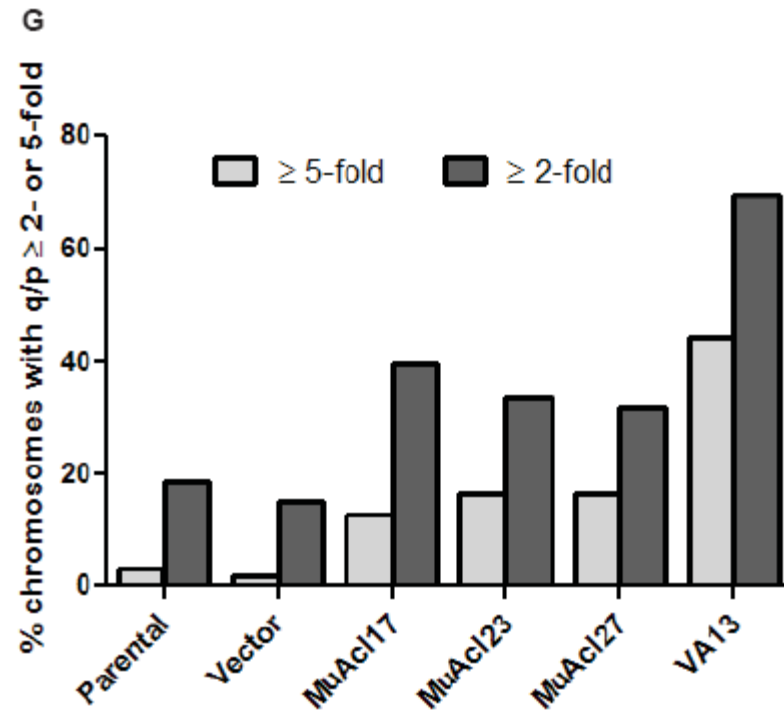


# Mutant telomerase RNA expression is associated with changes in telomere pq-ratios

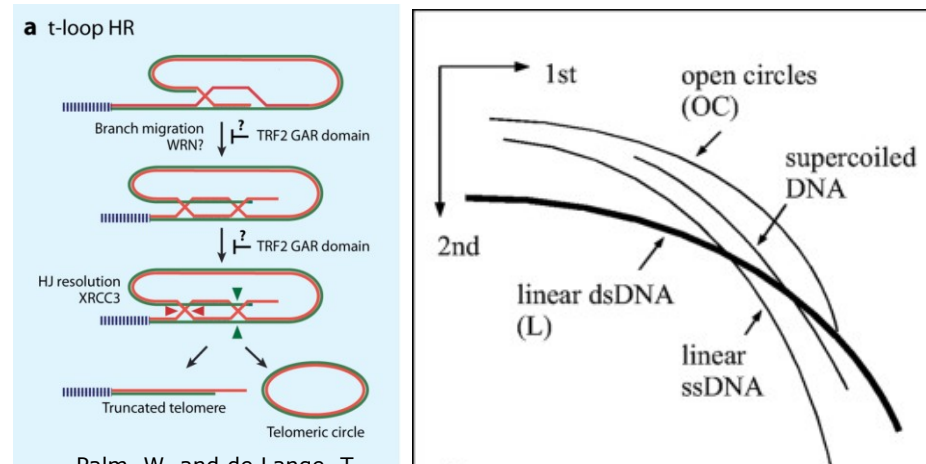
Figure 3



# Mutant telomerase RNA expression is associated with changes in telomere pq-ratios

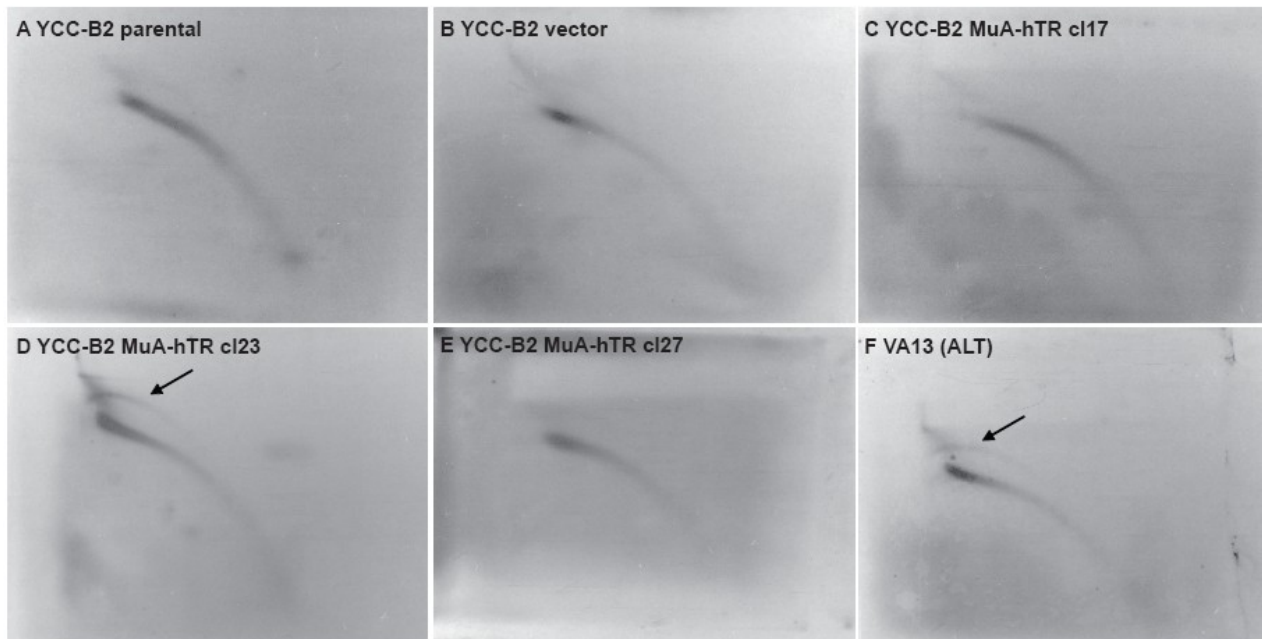


# MuA-hTR expression results in the accumulation of circular extrachromosomal telomeric DNA

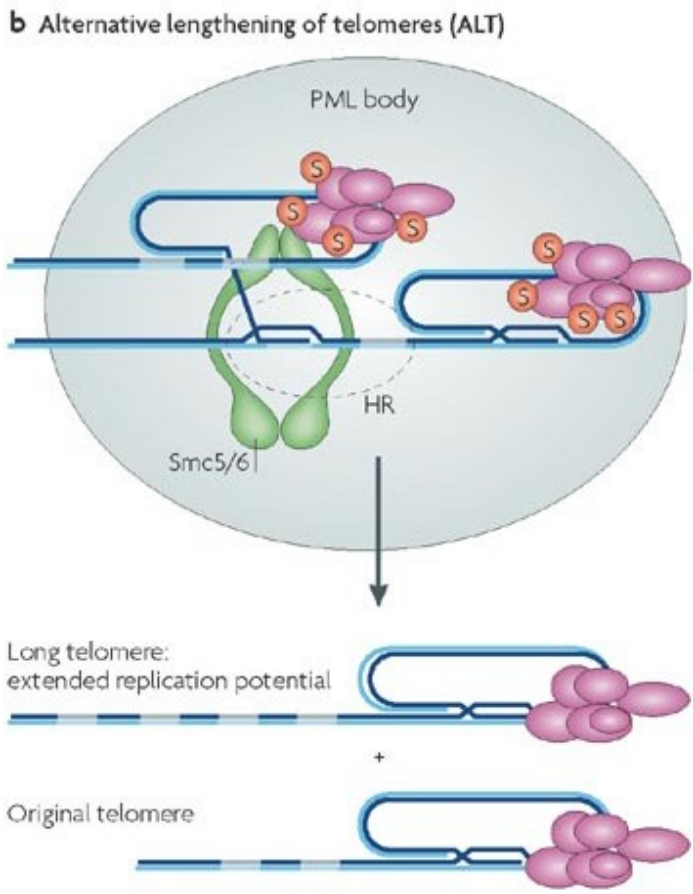


Palm, W. and de Lange, T.  
Annu. Rev. Genet. 2008.

Figure 4

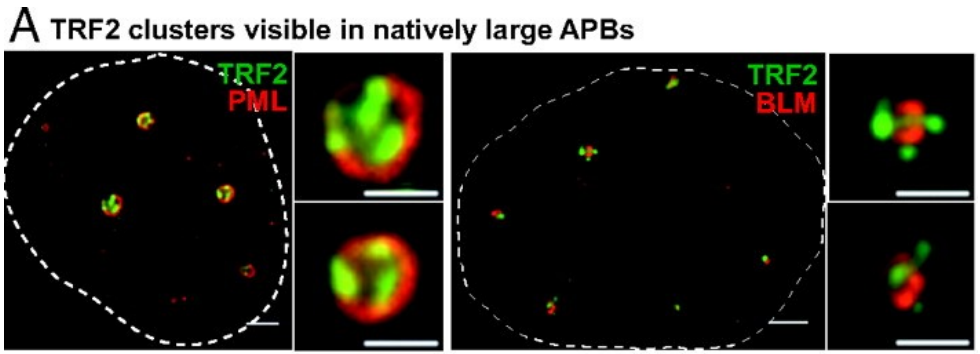


# ALT cells contain a high level of ALT associated promyelocytic leukemia bodies (APBs)



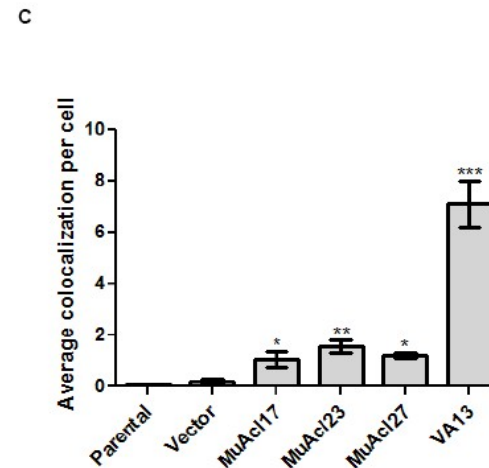
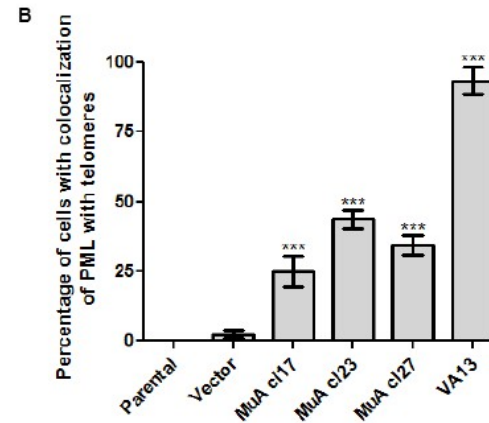
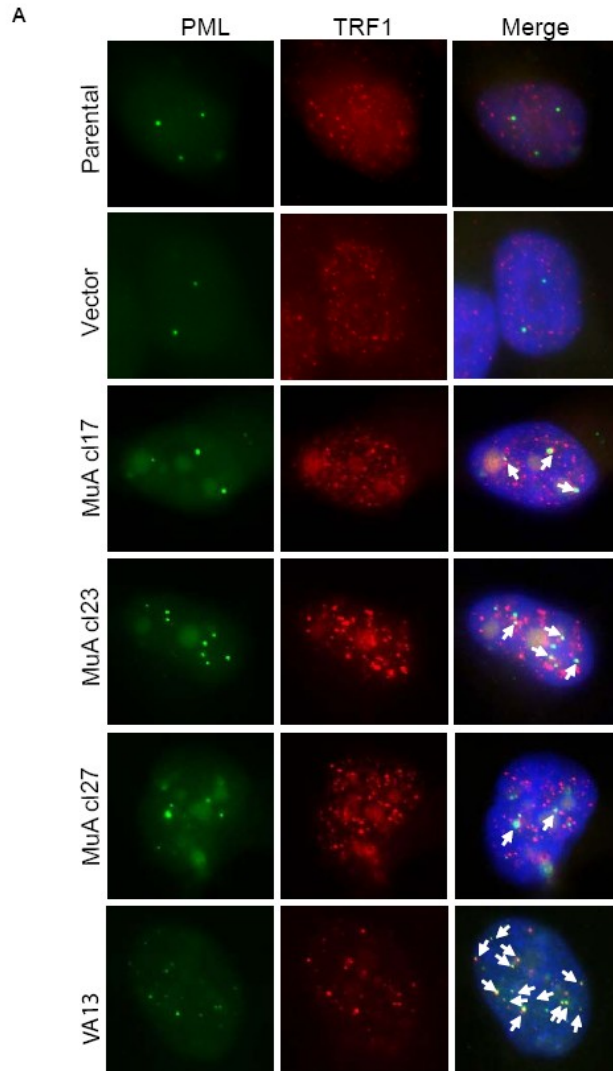
Nature Reviews | Molecular Cell Biology

Murray, JM and Carr, AM. Nat. Rev. Mol. Cell Biol. 2008



Irena Draskovic et al., PNAS, 2009.

# Increased formation of PML bodies associated with telomeric DNA in telomerase-positive cells expressing mutant telomerase RNA



# Telomeric recombination induced by dysfunctional telomeres

- Elevated DNA damage response located at the telomeres (Telomere dysfunction-induced foci), suggesting that the incorporation of mutant repeats disturbs the telomere cap
- Increased frequency of Telomere Sister Chromatid Exchange (T-SCEs), elevated pq-ratios and extrachromosomal telomeric DNA fragments were observed in the MuA-expressing cells, which are consequences of homologous recombination between telomeres
- First report of telomere dysfunction inducing telomeric recombination in mammalian cells without telomerase inhibition and of endogenous telomerase and telomeric recombination pathways coexisting spontaneously in cancer cells

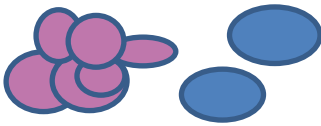
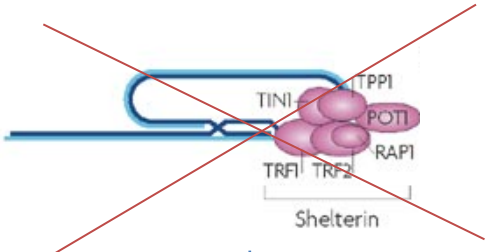
## Conclusion

*Our results suggest that after telomere destabilization, there is a strong selection pressure for the emergence of resistant cells with increased telomeric recombination and telomeric recombination could be a potential resistance mechanism in response to telomere dysfunction*

*However, the use of chemotherapeutic drugs decreases the proliferation of the MuA-expressing cells, despite the presence of a recombination-based mechanism for telomere maintenance*

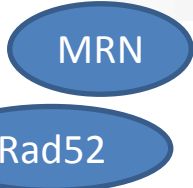
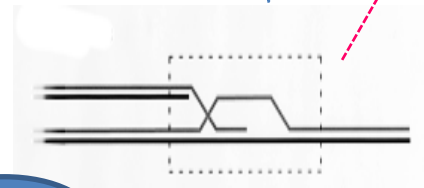


# Model



TRF1, TRF2 and POT1 cannot bind to mutant sequences

Telomere maintenance independent of telomerase and survival

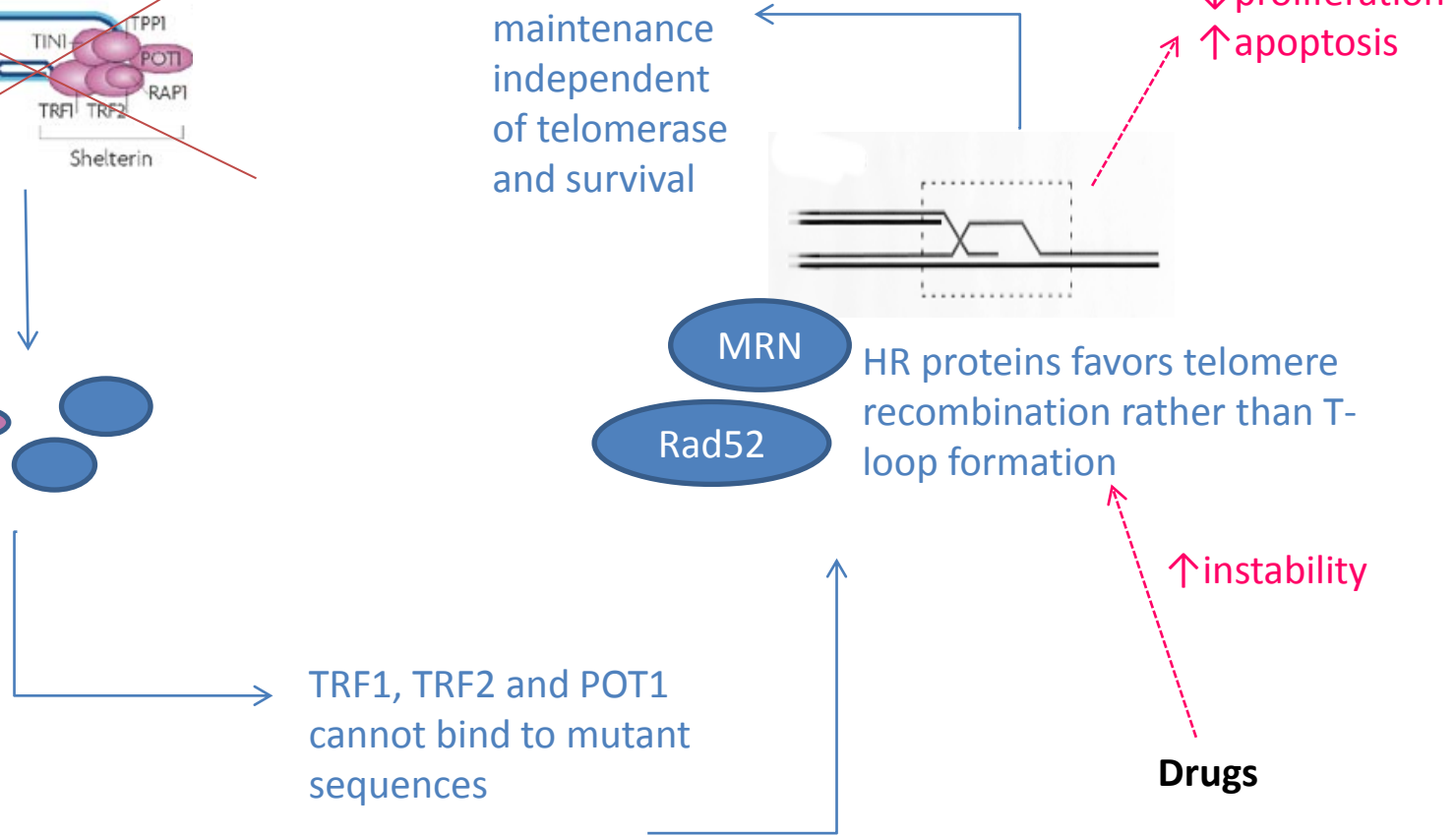


HR proteins favors telomere recombination rather than T-loop formation

↓ proliferation  
↑ apoptosis

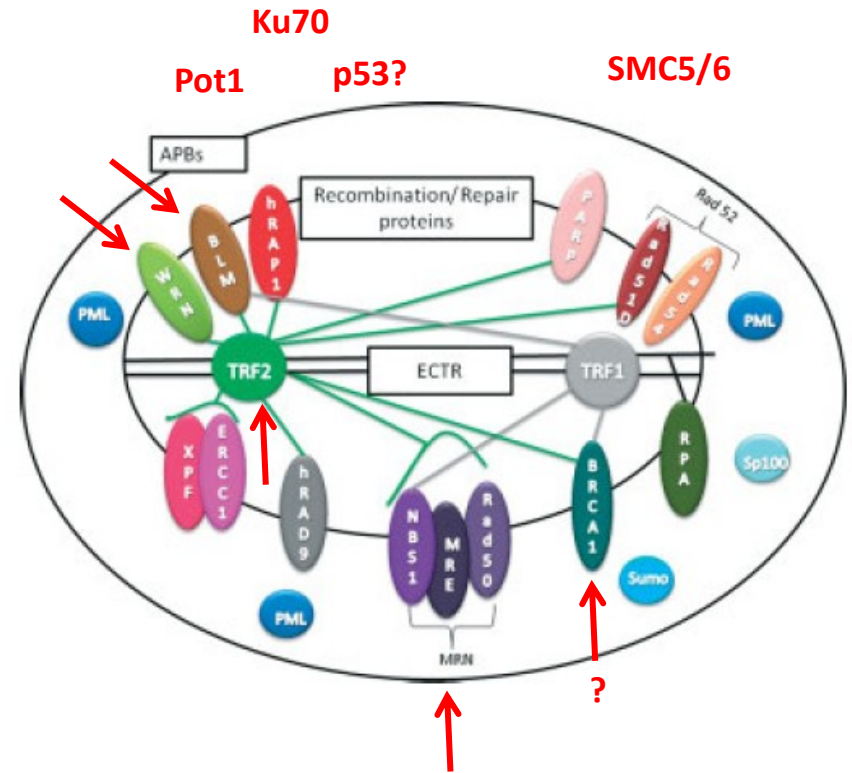
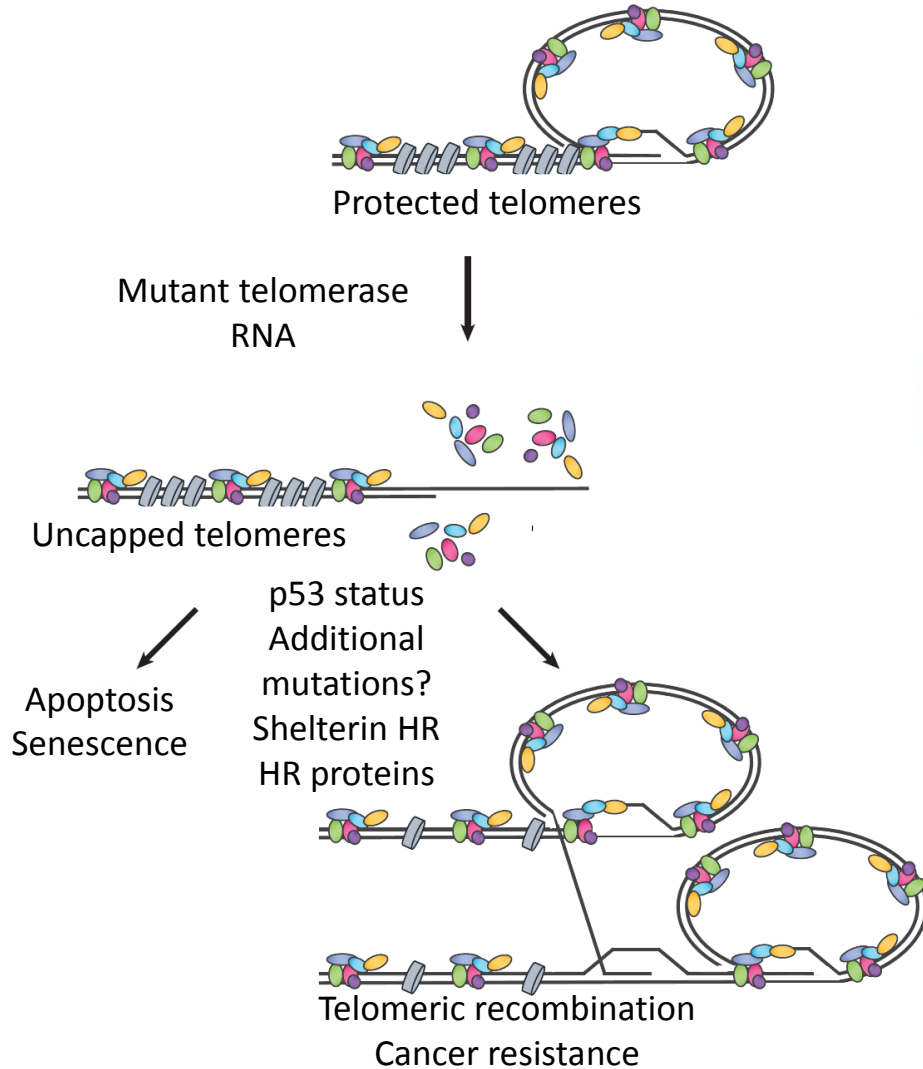
↑ instability

Drugs



# Perspective

The factors involved in resistance to induced-telomere dysfunction could include the impaired p53 status of YCC-B2 and additional proteins which regulate or may regulate ALT or telomeric recombination



Adapted from De Boeck, G. et al. J. of Path. 2008.

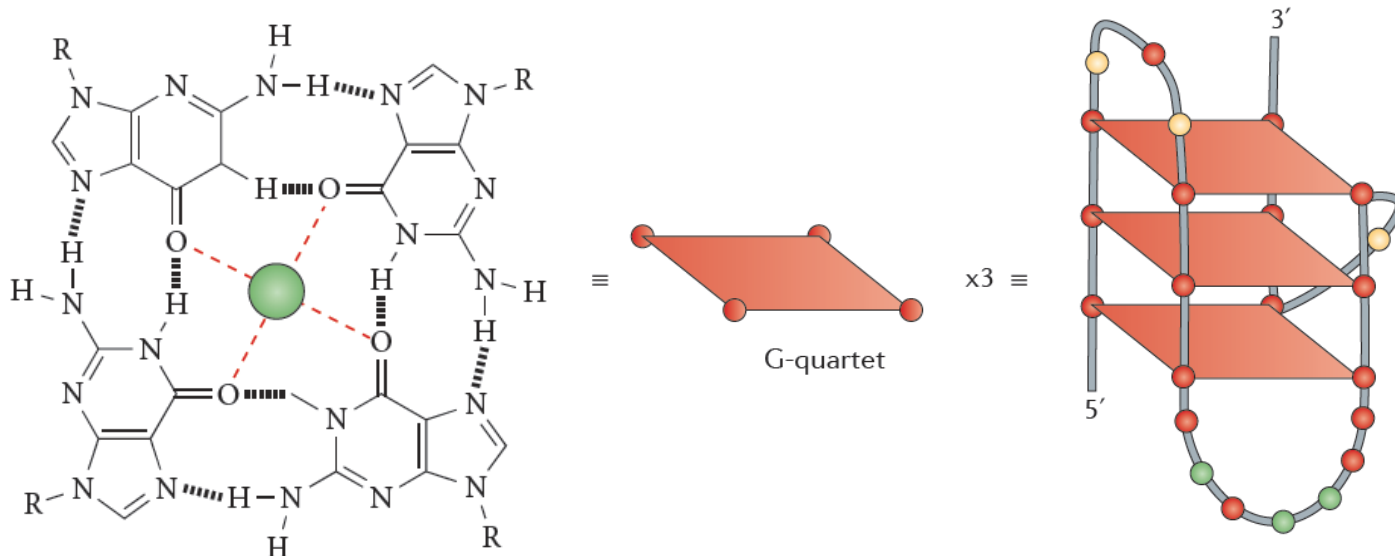
Adapted from Brault and Autexier, Cell Cycle, 2011



**Inhibiting telomerase and telomere function in cancer cells  
with G-quadruplex ligands**

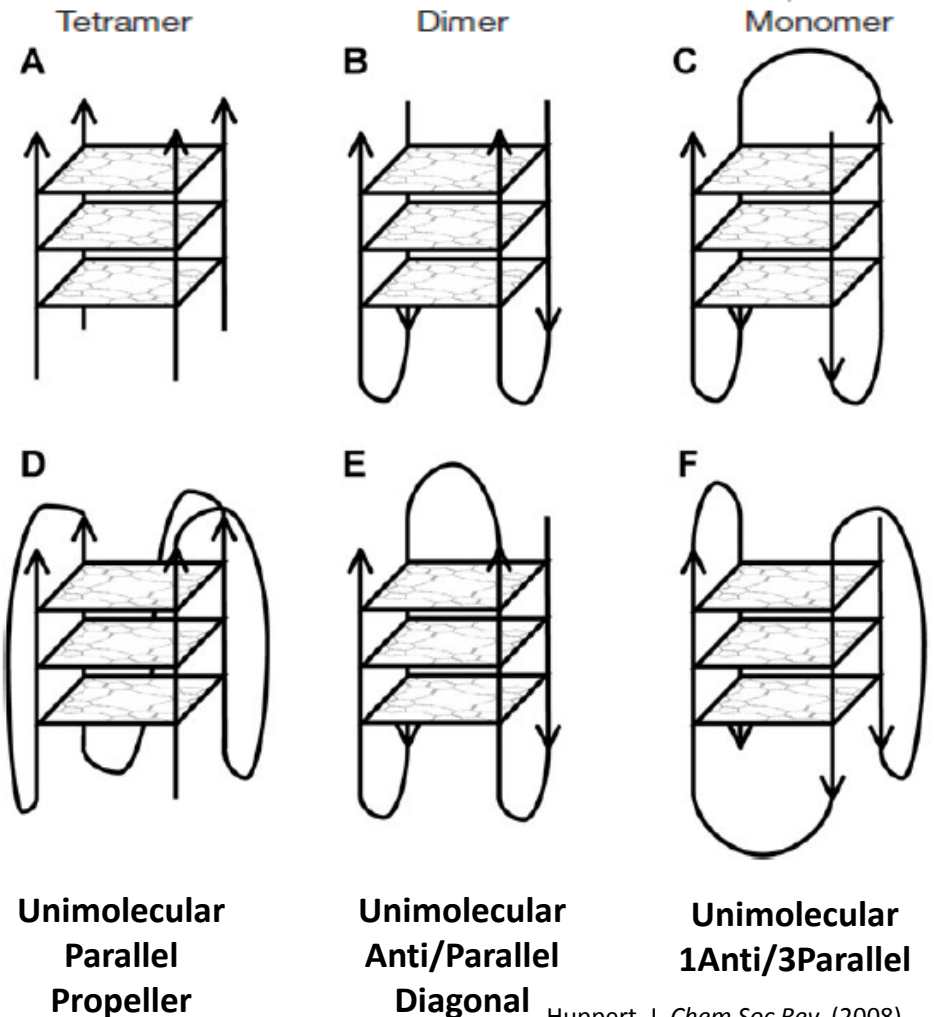
# G-quartet

- Bang 1910
- Gellert *et al* 1962
- 4 Guanines
- Stabilized by Hoogsteen Hydrogen bonds
- Stacked = G-quadruplex
- Stabilized by  $K^+$  and  $Na^+$



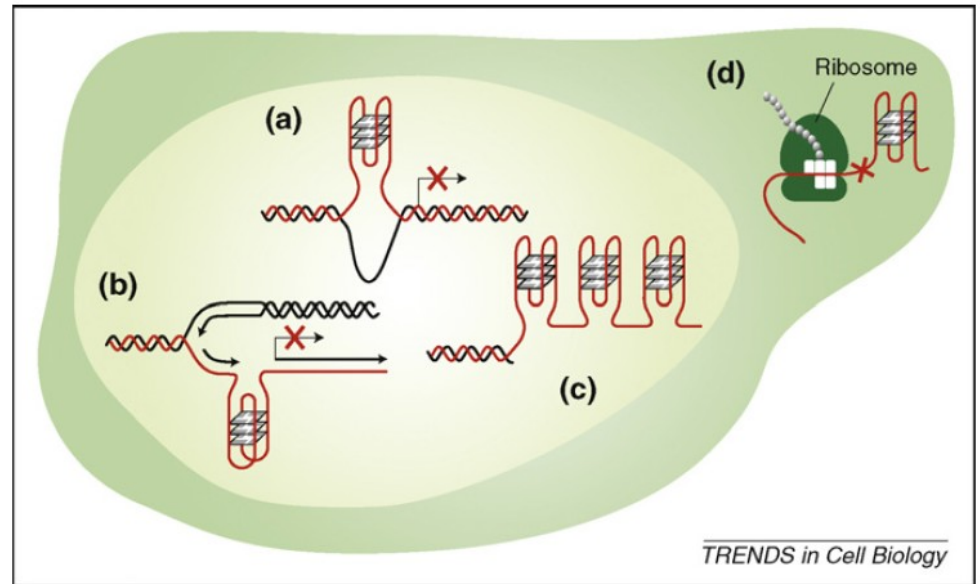
# G-quadruplex

- Forms determined by:
  - Strand number
    - Monomer, Dimer, Tetramer
  - Strand direction
    - Parallel, anti-parallel
  - Loops
    - Lateral/Edgewise
    - Diagonal
    - Chain-reversal/Propeller
  - D, E, & F formed in human telomeric repeats

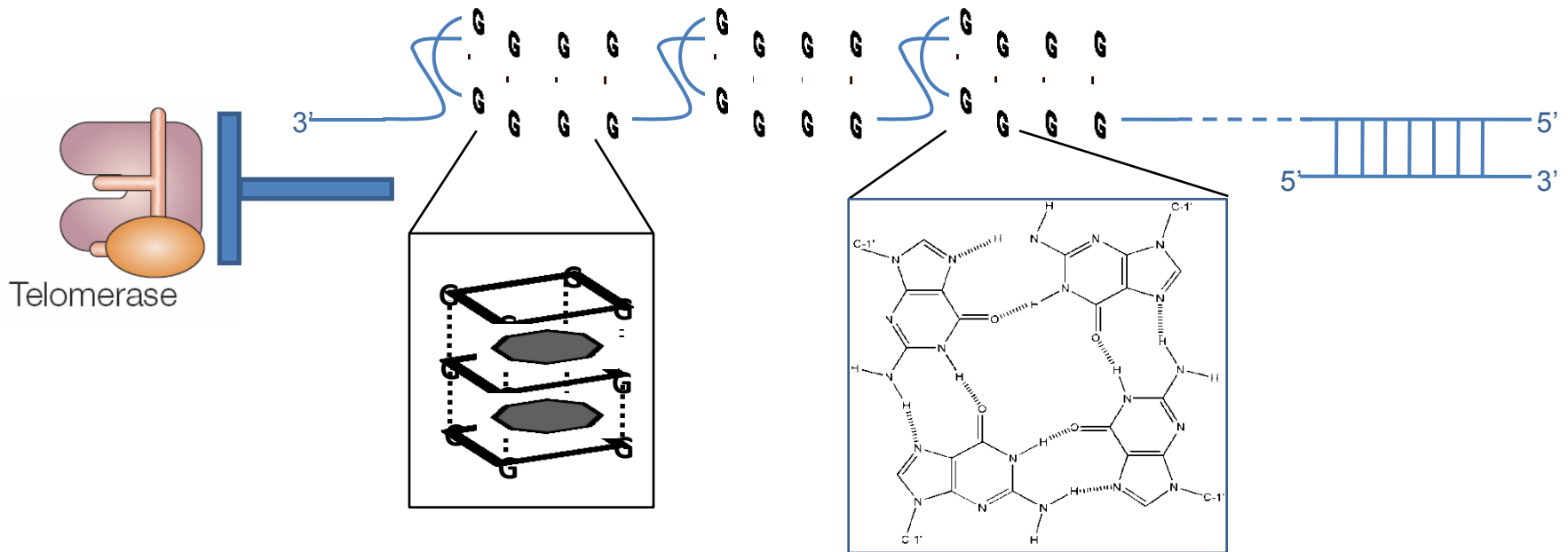


# Location of G-quadruplexes

- Genome-wide sequence analysis identified **376 000** putative sequences (*not homogenously distributed*)
- Within the nucleus:
  - a) promoter region of genes
  - b) during replication
  - c) **telomeric 3' G-overhangs**
- Outside of nucleus:
  - d) 5' UTR of mRNA



# G-Quadruplex Stabilization Leads to Telomerase Repression



Adapted from Fakhoury, J, Nimmo, G, Autexier, C. Anticancer Agents in Medicinal Chemistry, 2007

# Phenotypes elicited by G-quadruplex ligands

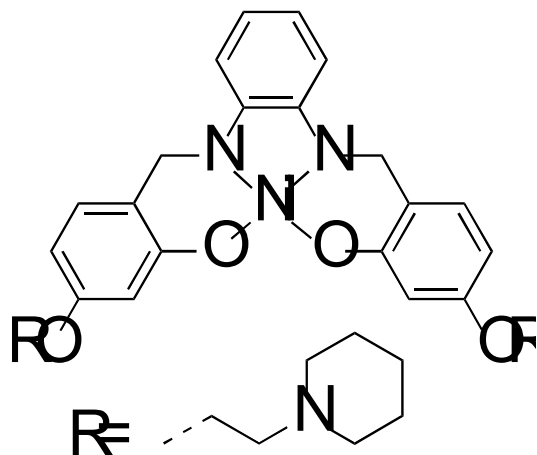
- Telomerase inhibition
- Telomere shortening
- Lag phase dependent antiproliferative response in cancer cells
  
- Rapid telomere shortening-independent antiproliferative effects
- Loss of the G-rich overhang
- Dissociation of TRF2 and Pot1 from the telomere
- Increased DNA damage foci at telomeres
- Activation of DNA damage response
- Apoptosis
- Antitumor activity in mouse tumor models
  
- Antiproliferative response restricted to cancer cells



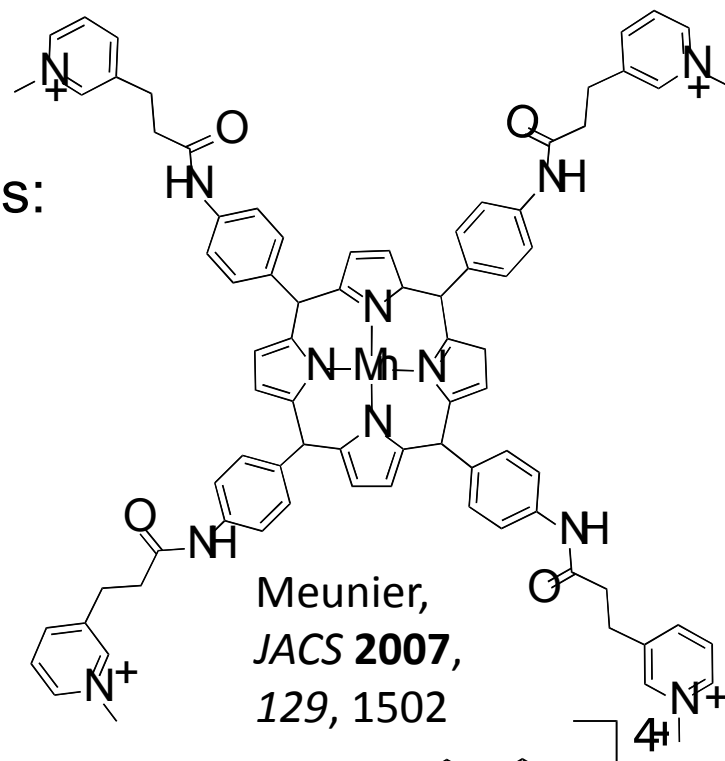
# Transition metal-based G-quadruplex binders

Many advantages to implementing metals:

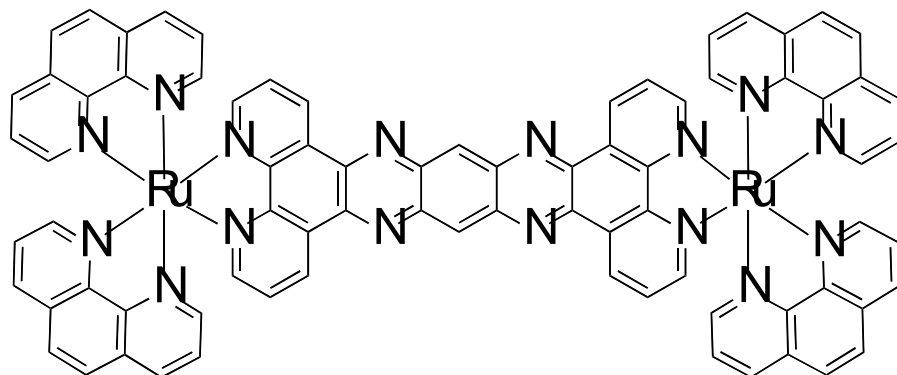
- Modular/tunable
- Multiple metal geometries/inherent positive charge
- Ease of synthesis
- Potential reduction in overall synthetic time



Neidle, Vilar, *JACS* **2006**, 128, 5992



Meunier,  
*JACS* **2007**,  
129, 1502



Haq, Thomas, *JACS* **2006**, 12, 4611

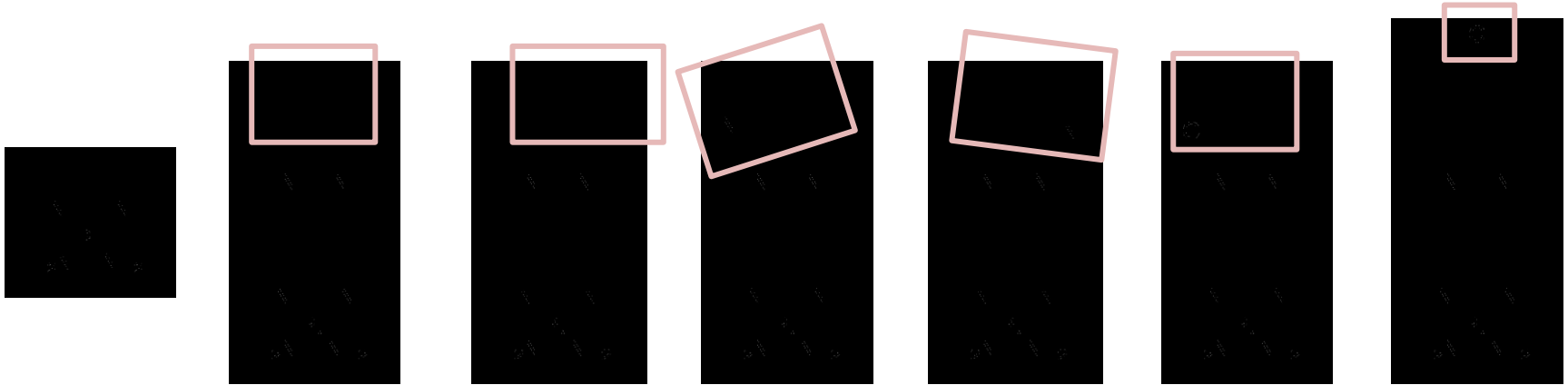
# Platinum phenanthroimidazoles as G-quadruplex binders

Challenge: Make DNA binders selective for G-quadruplexes using square planar transition metals to extend the  $\pi$ -surface



R. Kieltyka, Fakhoury J., Moitessier N., Sleiman H., *Chem. Eur. J.*, 2008, 14, 1055.

# Q-quadruplex Ligands



Q-quadruplex ligands with high affinity and selectivity to G-quadruplex over duplex DNA  
platinum(II)

- New class of Pt(II) complexes with high affinity and selectivity to G-quadruplex over duplex DNA
- Bigger role of hydrogen bonding in binding affinity
- Added a Halogen, shifting electron density towards Cl
- Increased binding affinity and selectivity

# Ligand Binding Affinity & Selectivity

- Binding Affinity assessed by FID

fluorescence intercalator displacement assay

- Compared to 17mer & 26mer

- Binding affinity measured by  $^{ds}DC_{50}/^{G4}DC_{50}$

DC 50 concentration of cplx req to give 50% decrease in fluorescence

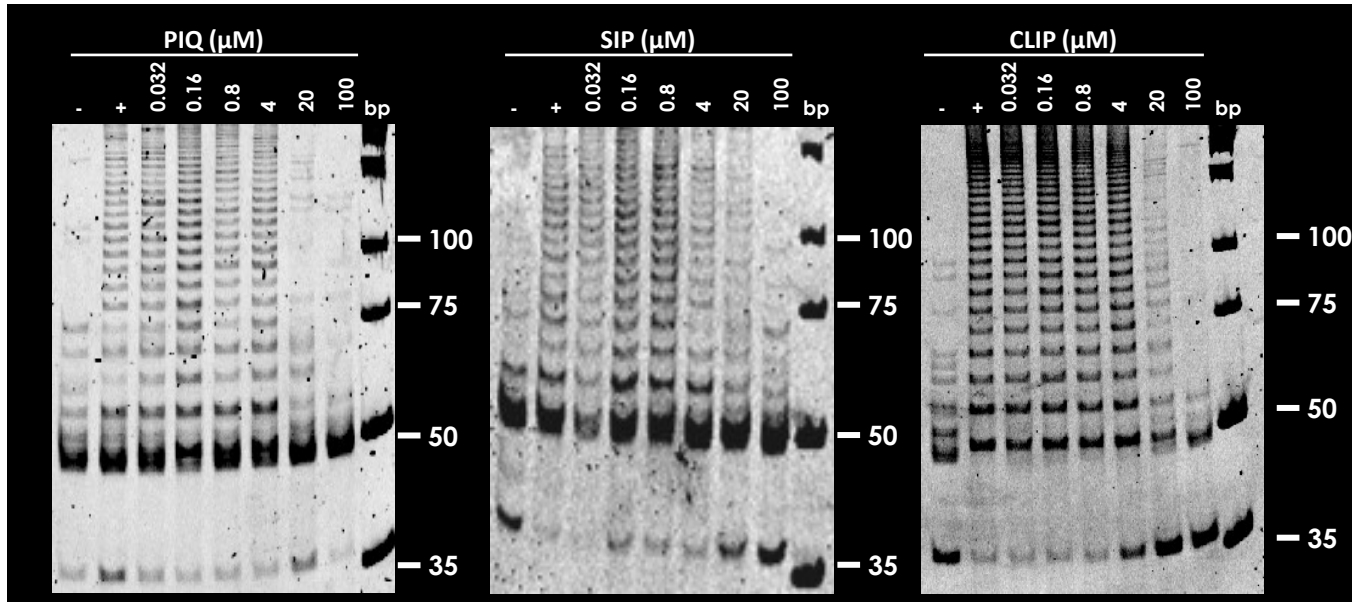
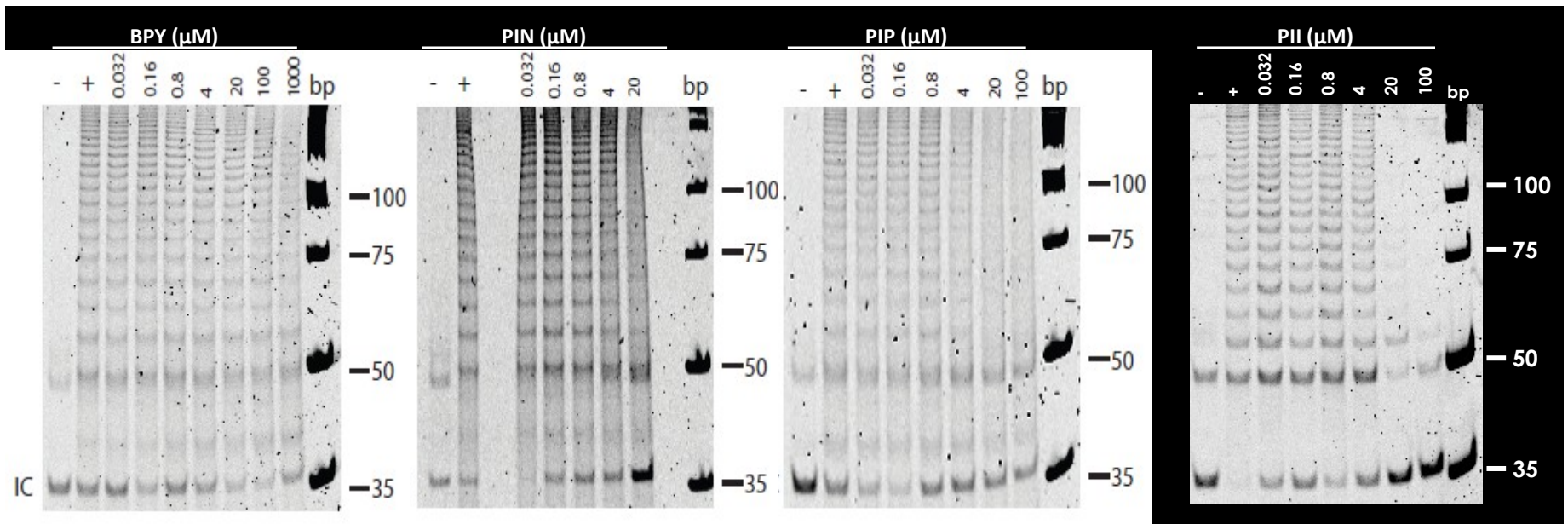
PIP  
PIN  
PII  
PIQ  
SIP  
CLIP  
BPY

**Table 1.**  $DC_{50}$  values determined from FID assays.

Complex	$^{G4}DC_{50}$ [ $\mu M$ ]		$^{ds}DC_{50}$ [ $\mu M$ ]		Selectivity <sup>[b]</sup>
	Na <sup>+</sup>	K <sup>+</sup>	17mer	26mer	
1	0.66	0.68	1.11	0.84	1.2–1.7
2	1.30	1.71	2.12	1.58	0.92–1.6
3	0.53	1.55	1.37	1.03	0.66–2.6
4	0.70	1.09	1.65	1.37	1.3–2.4
5	0.53	0.70	1.60	1.10	1.6–3.0
6	0.31	0.66	1.47	1.20	1.8–4.7
7	> 2.5	ND <sup>[a]</sup>	> 2.5	> 2.5	–

[a] Not determined: given the inability of complex 7 to bind strongly to G4 DNA, no thiazole orange displacement was observed, and therefore no  $DC_{50}$  value could be determined with the G4 structures formed in potassium-containing buffer. [b] Selectivity range determined from dividing the lowest  $^{ds}DC_{50}$  by the highest  $^{G4}DC_{50}$  and the highest  $^{ds}DC_{50}$  by the lowest  $^{G4}DC_{50}$  for each complex.

# *In vitro* Telomerase Inhibition



Ligand	TRAP-Lig $\text{IC}_{50}$ ( $\mu\text{M}$ )
BPY	ND
PIP	0.46 0.29
PIN	6.81 5.84
PII	0.93 1.78
PIQ	2.48 1.49
SIP	11.6 1.78
CLIP	5.71 1.55
BRACO-19	6.30
Telomestatin	0.60

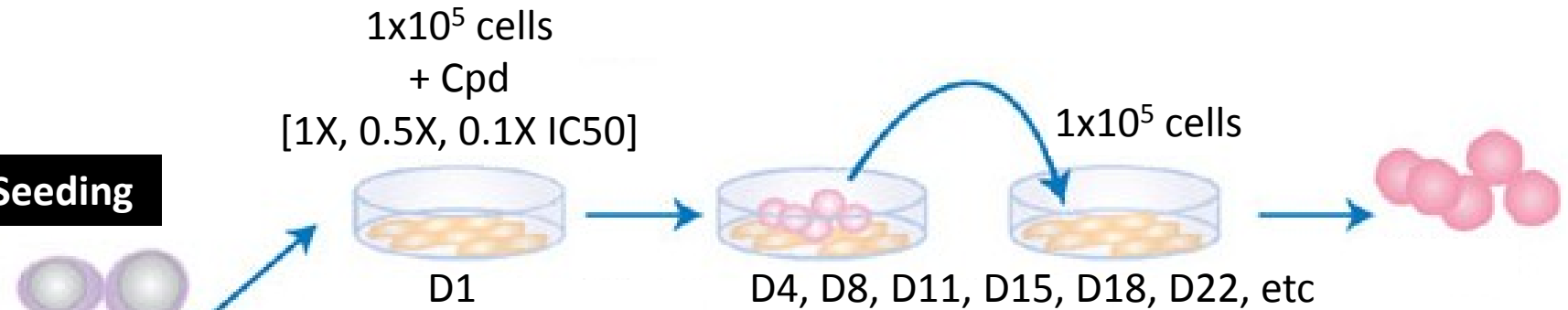
# Short term cytotoxicity assay suggest cancer cell specific effects

MTS (72h) – Proliferation		IC <sub>50</sub>			
Cell Type	Cell Line	BPY (μM)	PIP (μM)		CLIP (μM)
Cancer Telomerase +	A549	N/A	21.91	1.20	13.41 1.23
	HUH7	N/A	11.45	1.09	16.70 1.14
	MCF7	N/A	42.91	1.17	18.37 1.11
Cancer, ALT Telomerase -	GM847	N/A	27.73	1.04	9.33 1.02
Normal Primary	MRC-5	N/A	50.00		57.61 2.85
	WI-38	N/A	70.85		53.63

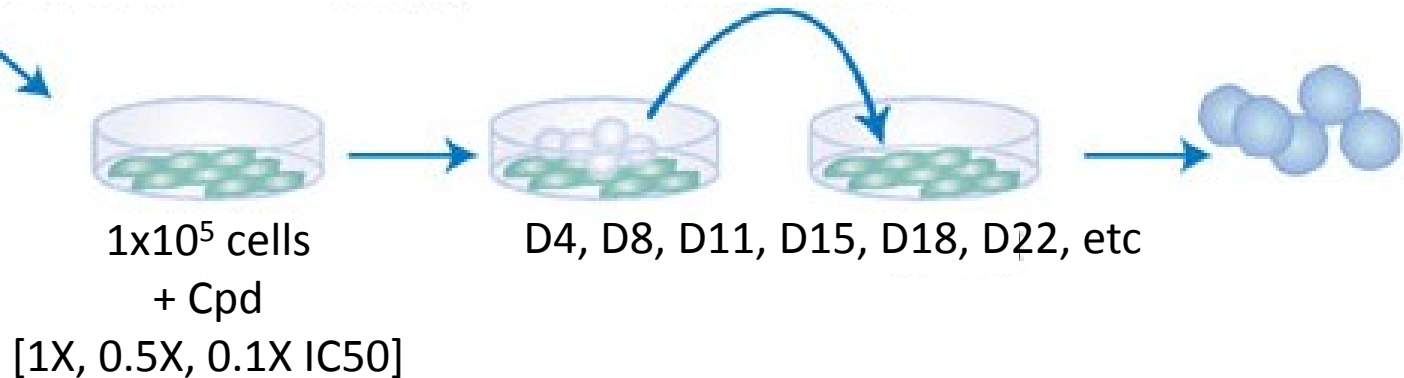
**MTS: Metabolic activity assay**

# Cell Studies: Experimental Design

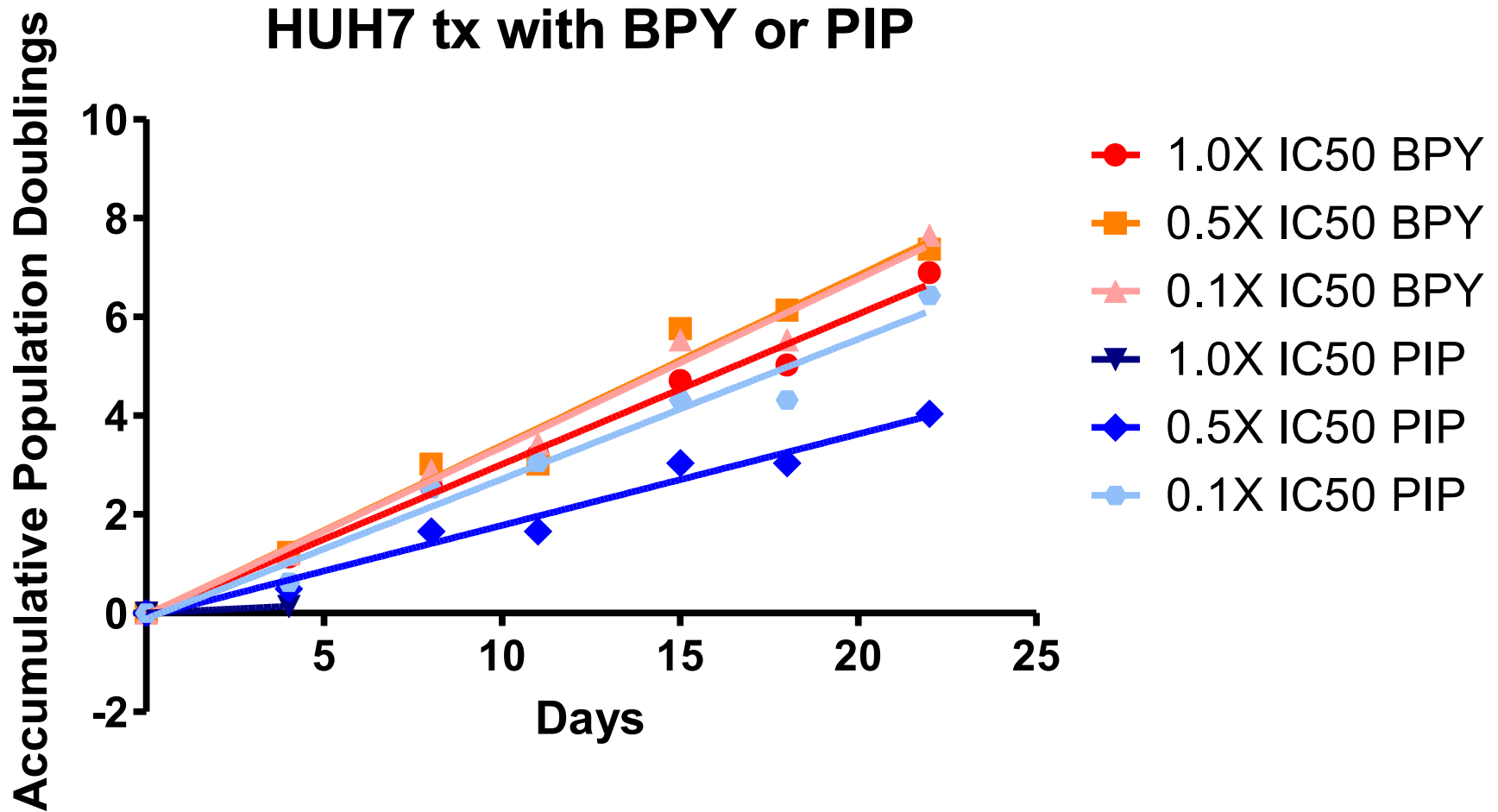
## 1) Seeding



## 2) Promoter

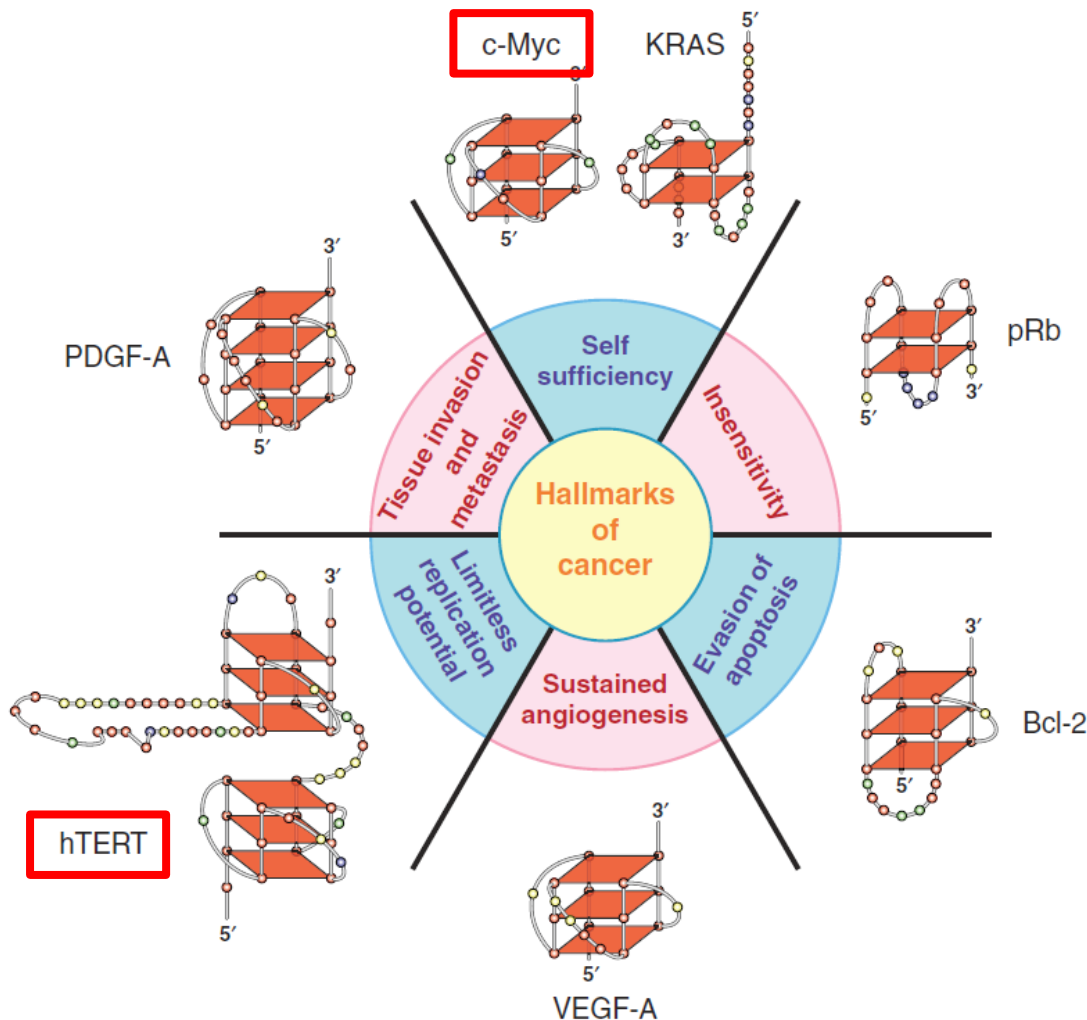


# Cell Studies: Seeding Experiment





# Cell Studies: qPCR



- **c-myc**

- most studied G4-forming promoter region
- upregulated in many cancers
- is a regulator of hTERT

- **hTERT**

- forms 2 different G4 in the promoter region
- expression is required for telomerase activity

# Conclusions and perspectives

- Molecules with distinct structural features that target G-quadruplex can be generated using supramolecular self-assembly
- phenanthroimidazole platinum(II) complexes are G-quadruplex stabilizers and telomerase inhibitors
- We will further evaluate improved ligands with increase binding affinity and selectivity to the G-quadruplex substrate (telomeric or promoter)
- Ligands are currently being tested for cancer cell specific antiproliferative effects
- Mechanism of action to be determined: telomere length dependent or independent, effect on G-quadruplex containing promoters relevant to telomerase (c-myc, hTERT)
- Best molecule will be tested for antitumor activity in xenograft mouse model

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CIHR IRSC

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Sanjida Khondaker  
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