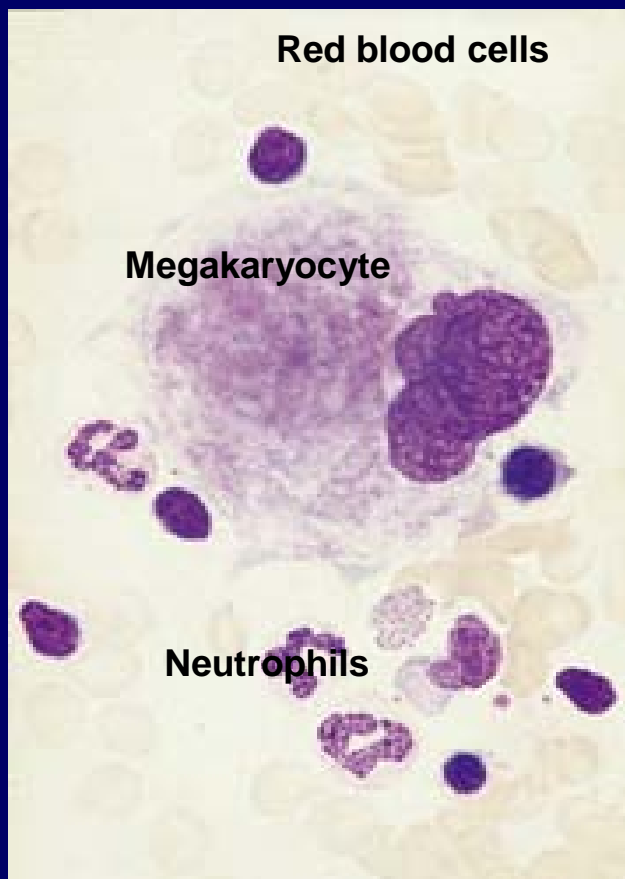
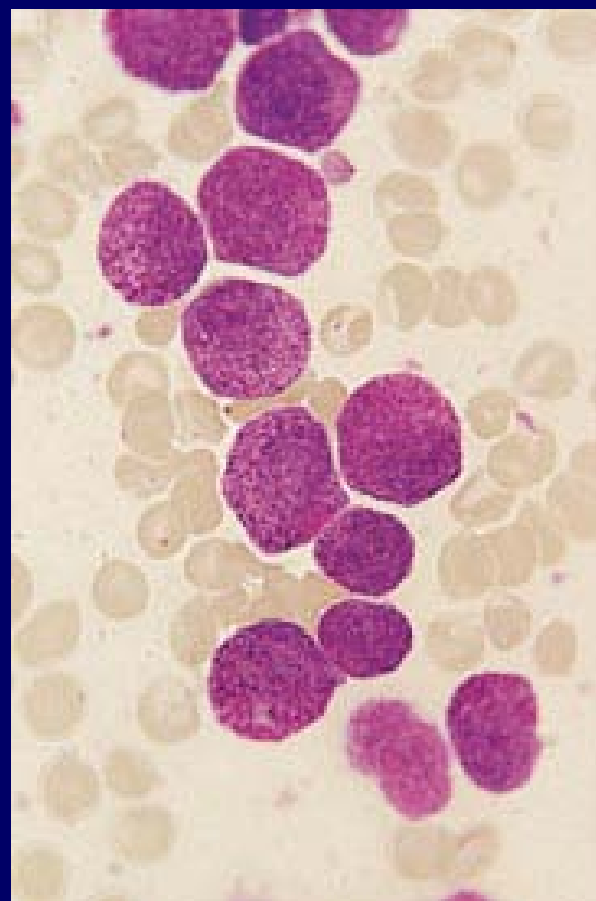


Acute promyelocytic leukaemia (APL)



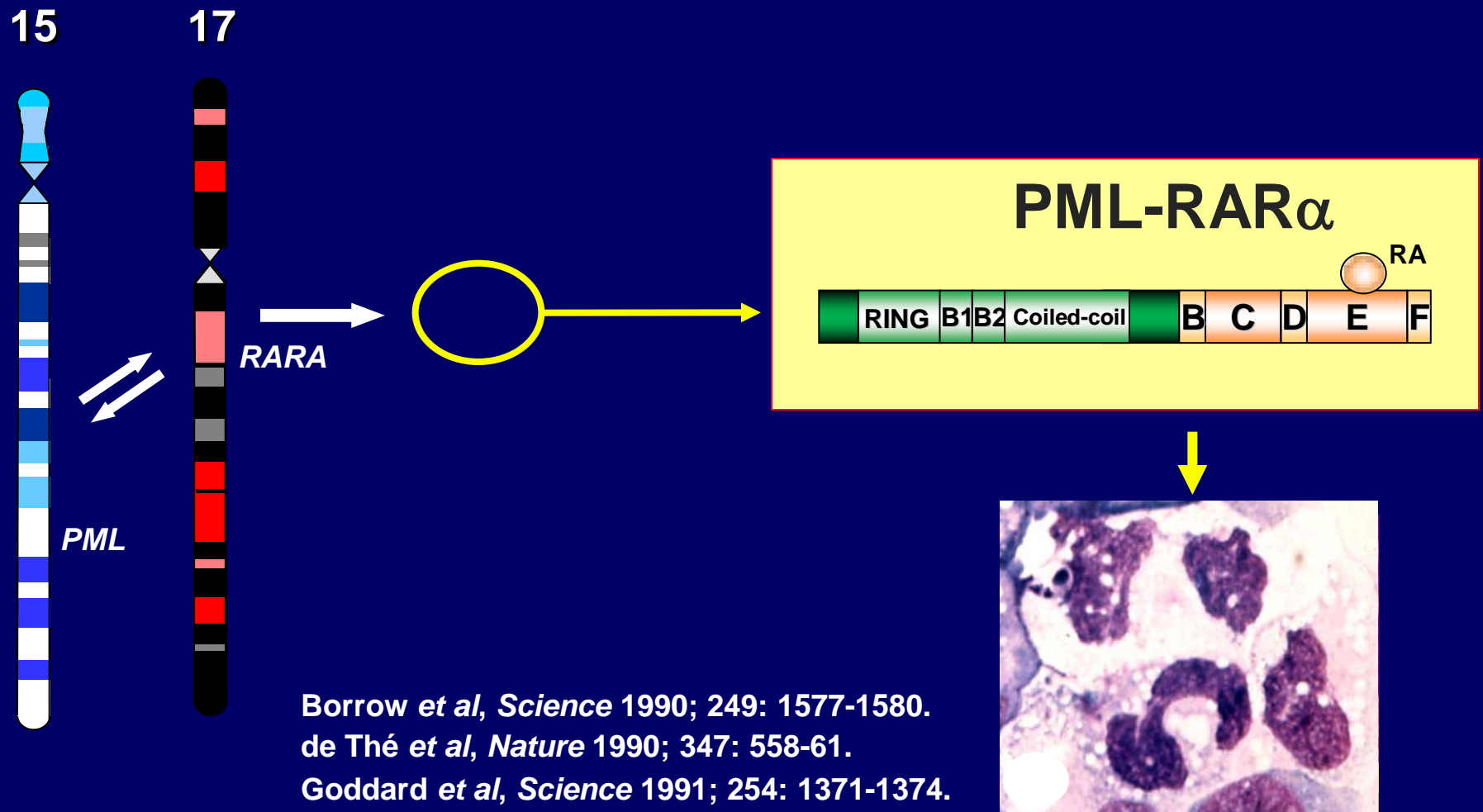
Normal bone marrow



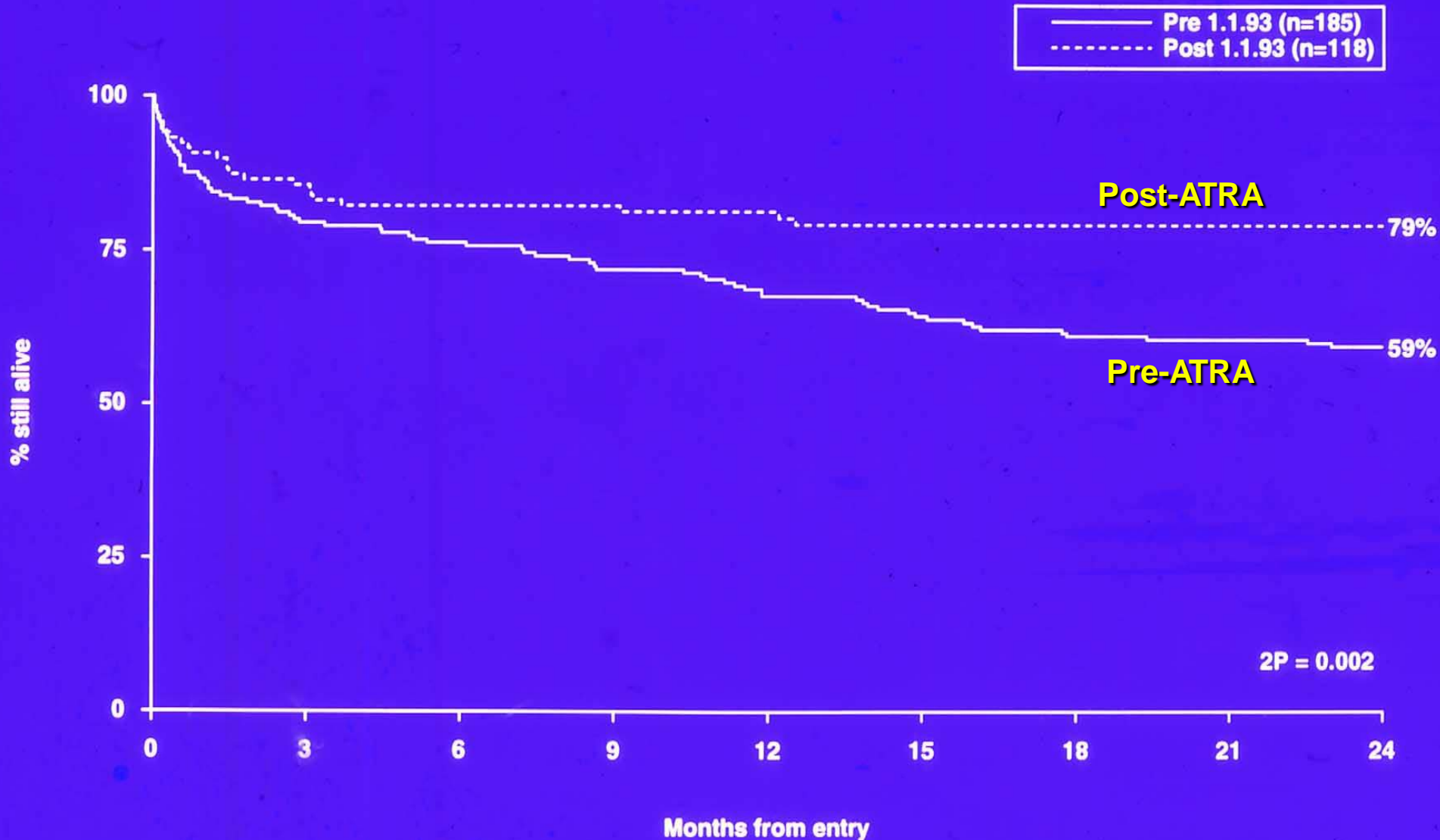
Leukaemic bone marrow (APL)

First success for molecularly targeted therapy:

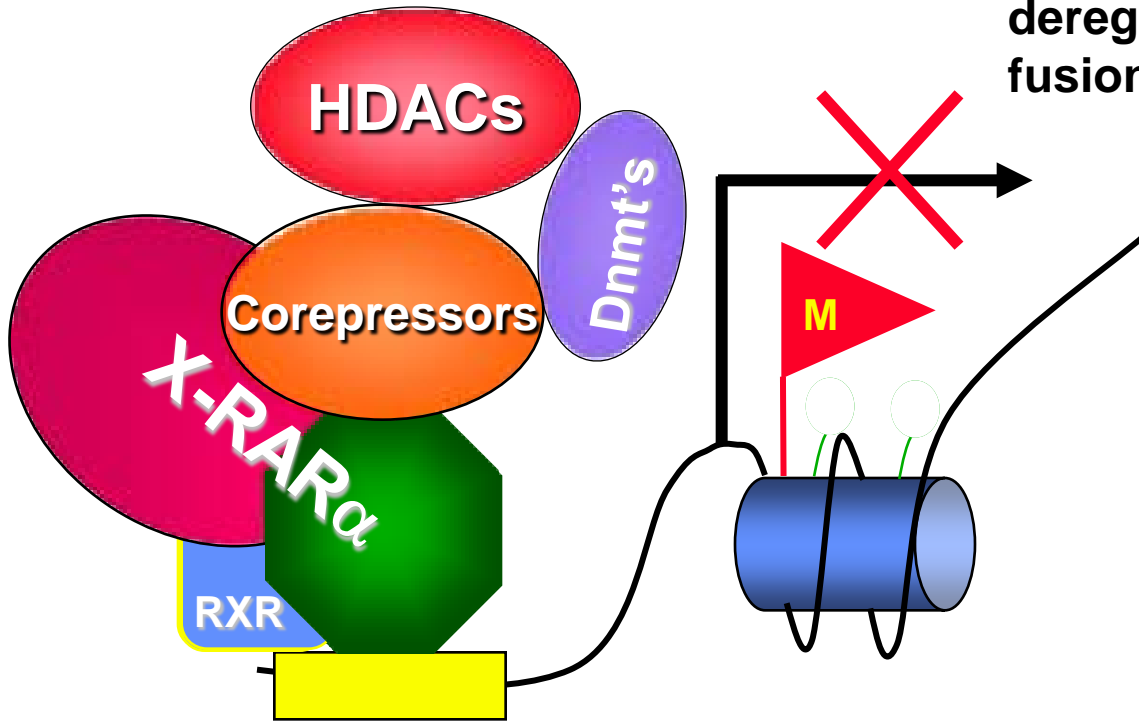
Targeting of PML-RAR α oncoprotein formed by t(15;17) in APL



Impact of molecularly targeted therapy – all *trans*retinoic acid (ATRA) on outcome in APL



Repression of target genes implicated in myeloid differentiation as a common mechanistic theme in APL

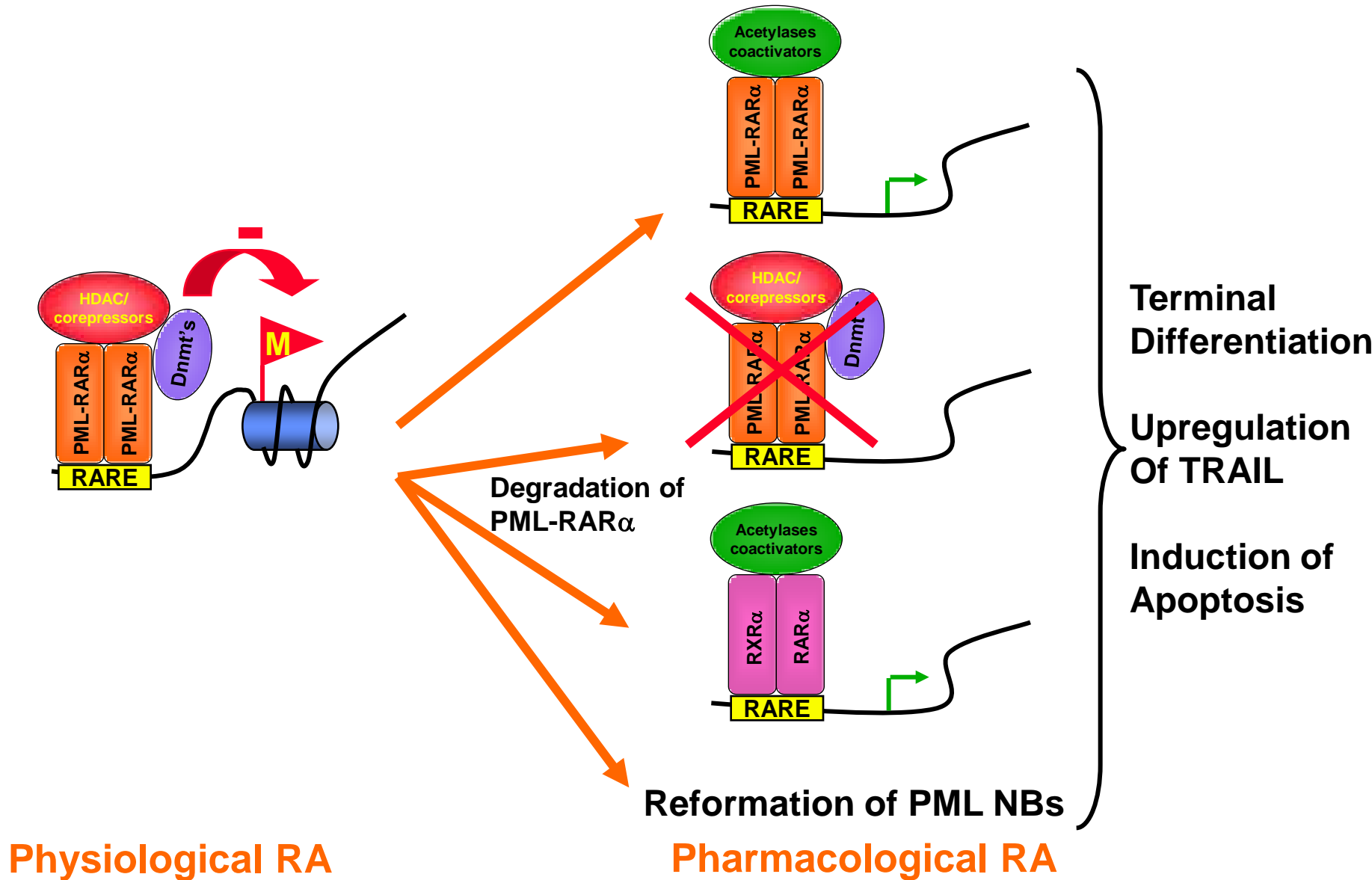


Genes involved in haematopoietic differentiation potentially deregulated by APL-associated fusion proteins

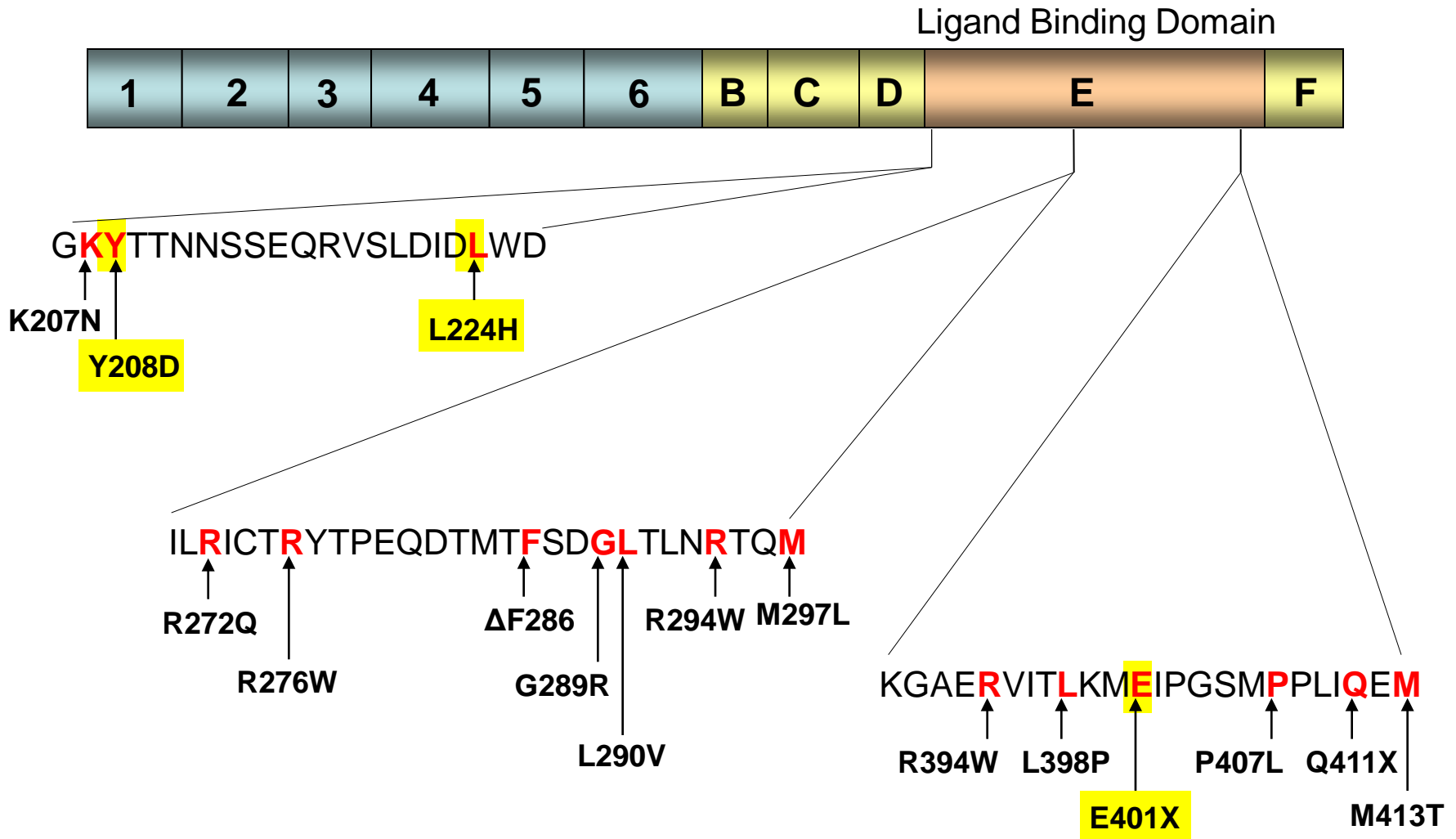
Hox
C/EBP β
RAR β 2
p21^{WAF/CIP}

Target gene regulatory element

Mechanisms of ATRA activity in PML-RAR α associated APL



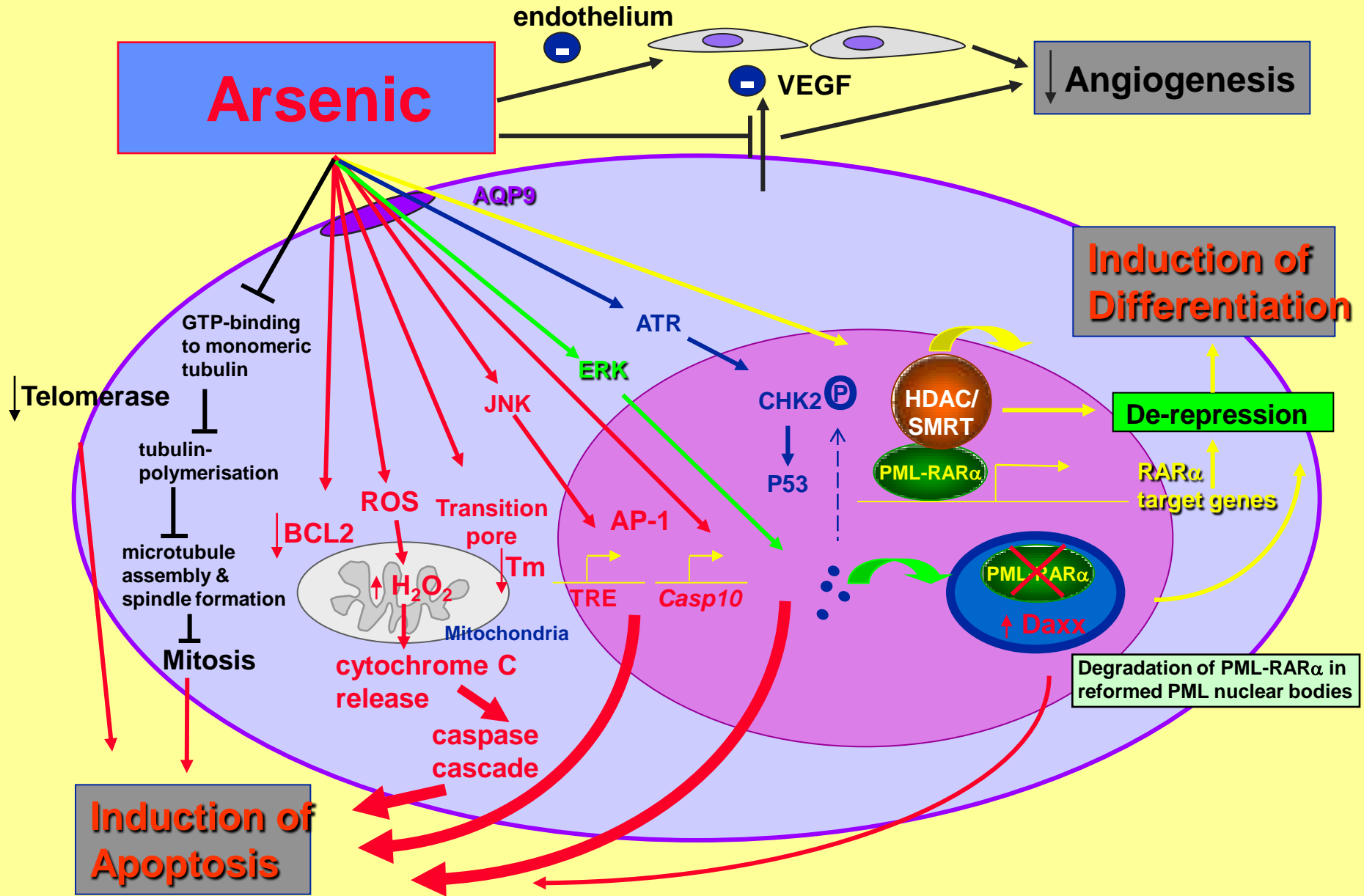
Mutations in PML-RAR α Ligand-Binding Domain (LBD) Found in ATRA-resistant Cell Lines and/or APL Patients



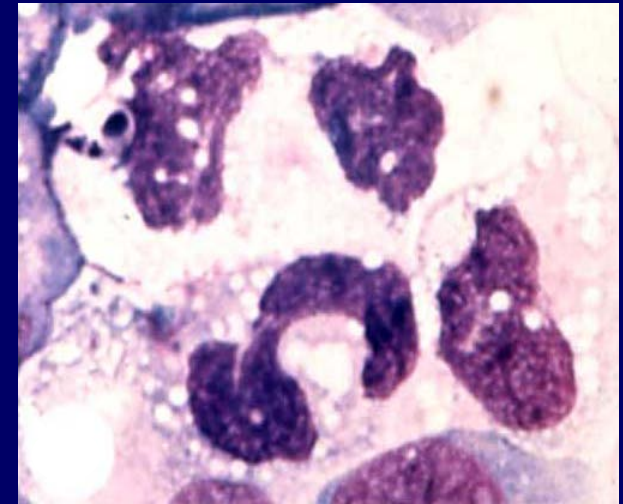
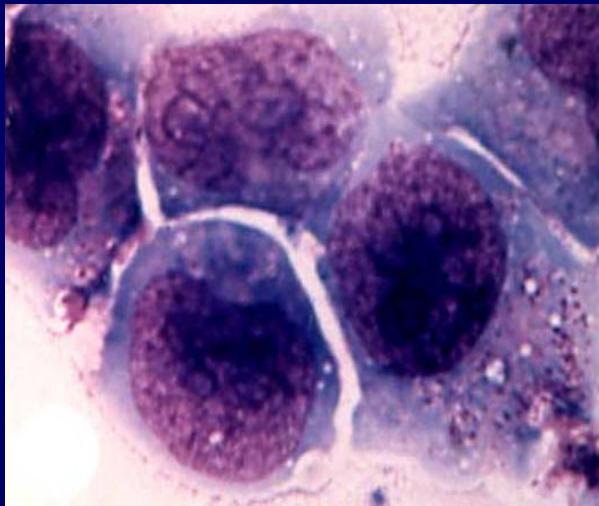
Arsenic activity in APL

- **Effective in PML-RAR α associated APL, including RA resistant cases**
- ***In vivo* response reflects 2 dose dependent activities:**
 - **Apoptosis (0.5-2 μ M)**
 - Triggered by alteration in mitochondrial membrane potentials & transition pore permeability
 - dithiol oxidation or cross-linking
 - Induction of reactive oxygen species
 - **Partial differentiation (0.1-0.5 μ M)**

Potential mechanisms of arsenic activity in APL



Impact of PML-RAR α fusion protein on nuclear architecture and its reversal by targeted therapies



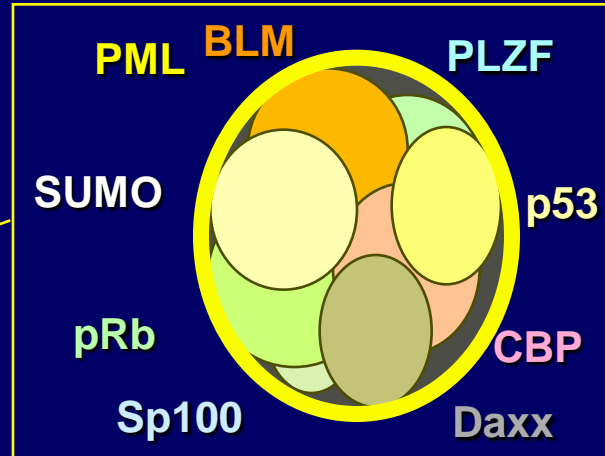
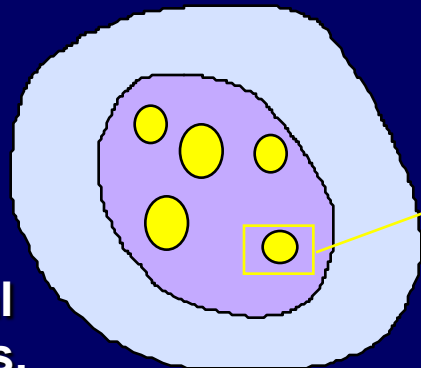
ATRA



Arsenic



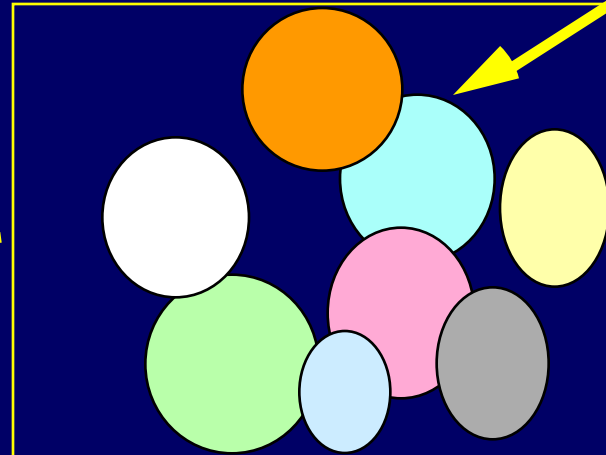
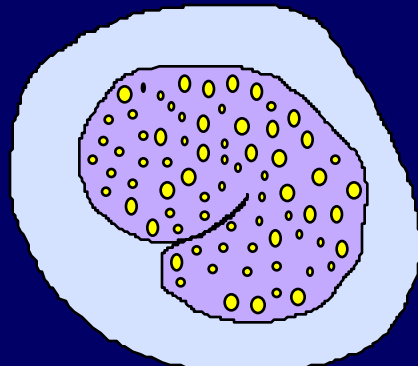
Disruption of PML nuclear bodies in t(15;17) associated APL



NBs implicated in regulation of:

- Apoptosis
- Senescence
- Growth suppression
- Genomic stability

ATRA
Arsenic

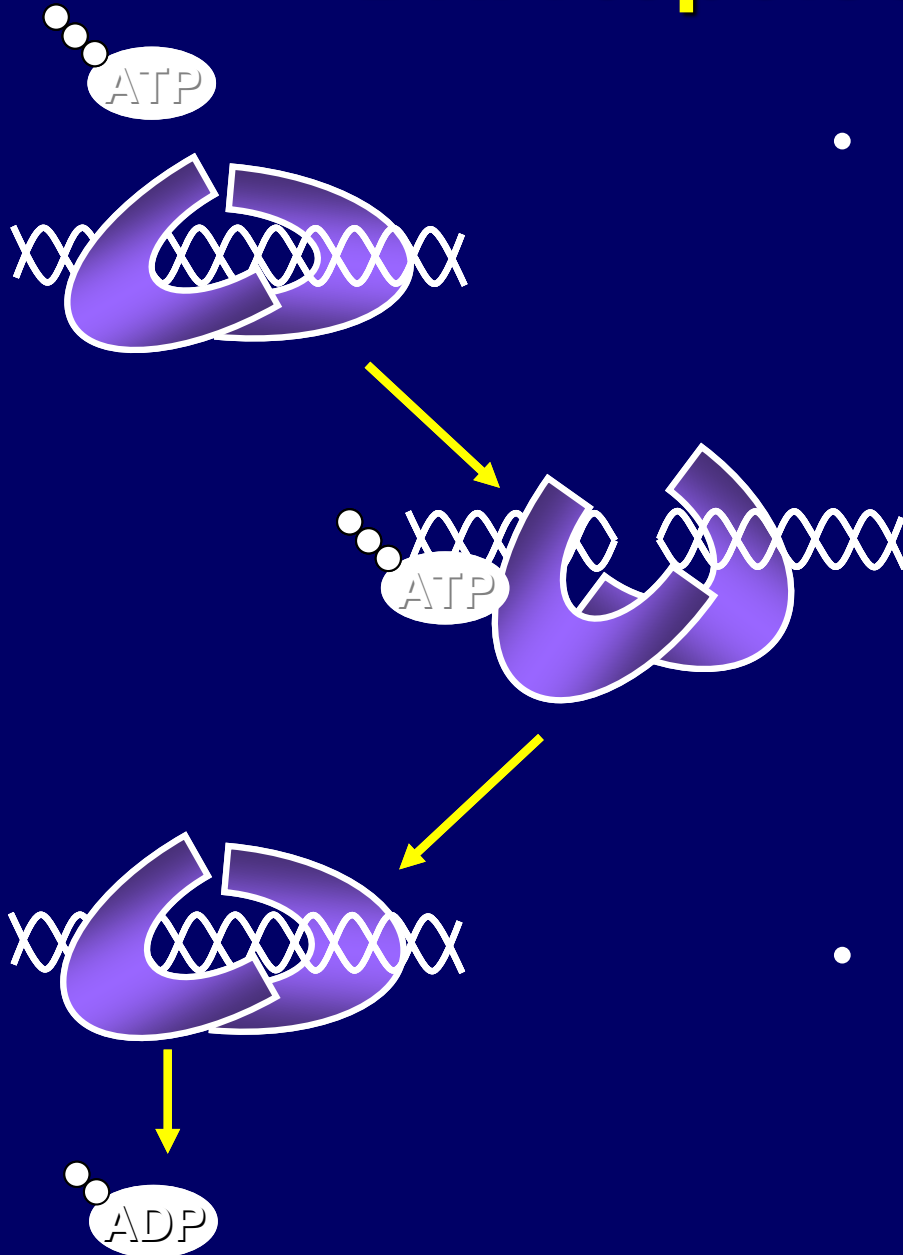


- Transcriptional repression
- Altered retinoid signalling
- Disruption of PML NBs
- Loss of PML growth suppressor & pro-apoptotic effects

Investigating mechanisms underlying formation of leukaemia-associated translocations

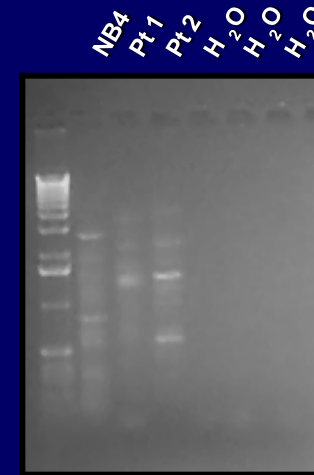
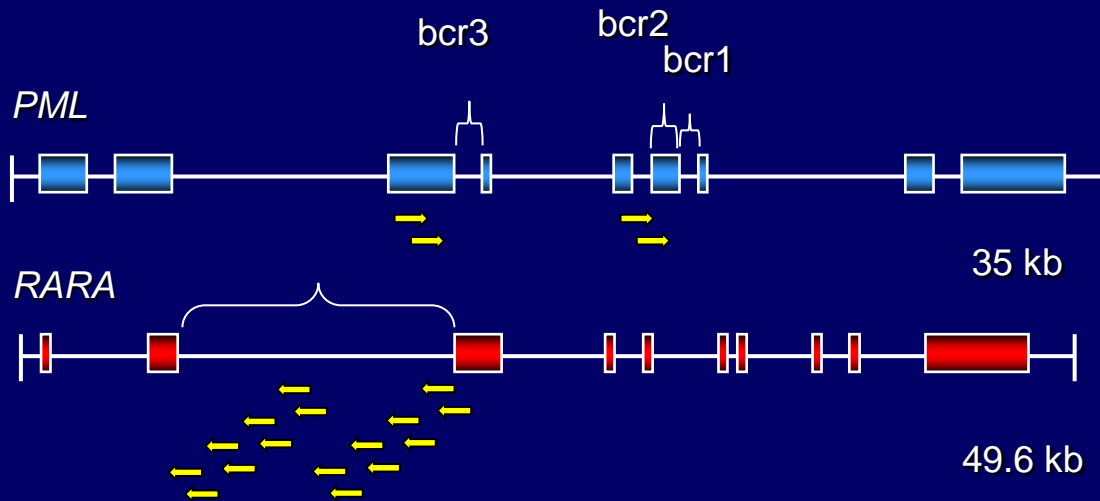
- Insights can potentially be gained from therapy-related leukaemias which have *de novo* counterparts
- Therapy-related leukaemias are becoming an increasing healthcare problem as more patients survive their primary cancers
- Incidence of therapy-related acute promyelocytic leukaemia (t-APL) with t(15;17) rising
- Balanced translocations arise following exposure to chemotherapeutic agents targeting DNA topoisomerase II
- Suggests a role for DNA topoisomerase II in forming translocations, but mechanisms controversial

DNA topoisomerase II

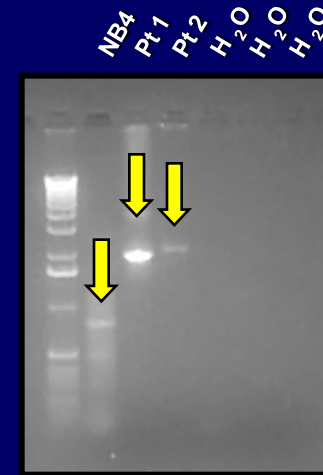


- Catalyses relaxation of supercoiled DNA
 - DNA topoisomerase II homodimer introduces 4-base staggered nicks in DNA as each subunit covalently binds and cleaves one strand
 - A second DNA double helix passes through the cleaved strands
 - Cleaved strands are then re-ligated
- Topoisomerase II-targeted drugs interfere with cleavage-re-ligation equilibrium → net increased cleavage → DNA damage → apoptosis

Characterisation of t(15;17) genomic breakpoint junctions in t-APL



1st Round of Nested PCR



2nd Round of Nested PCR

Patient Breakpoint Sequence

PML 1886

GGATTCCCATAGGTGCACACCCACACCCCTCCCAGCATGCATCCTAGGCAGTTCATAATG

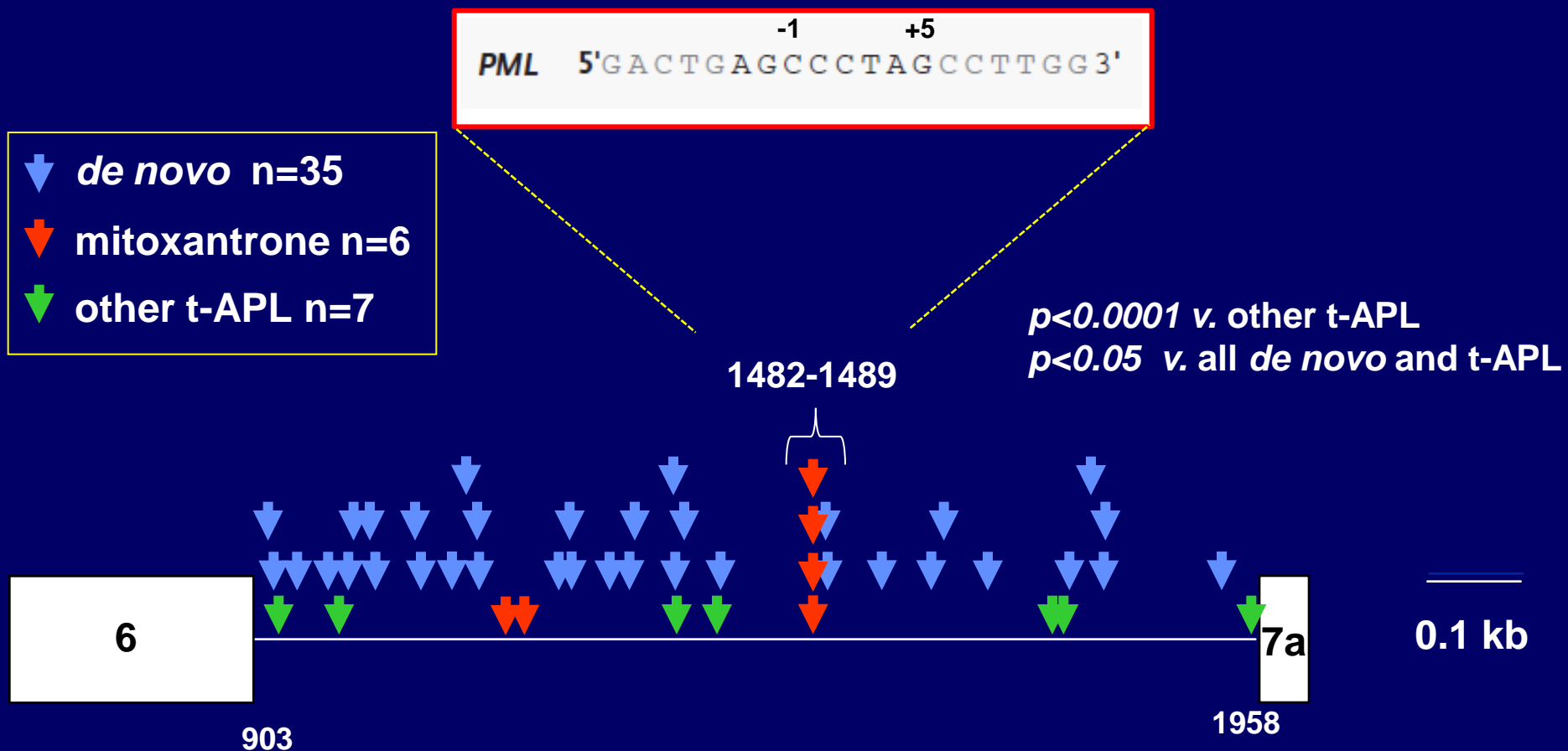
Breakpoint Sequence

GGATTCCCATAGGTGCACACCCACACCCCTGATGGGGGGTACCCAGAATAATGGGCTTTT

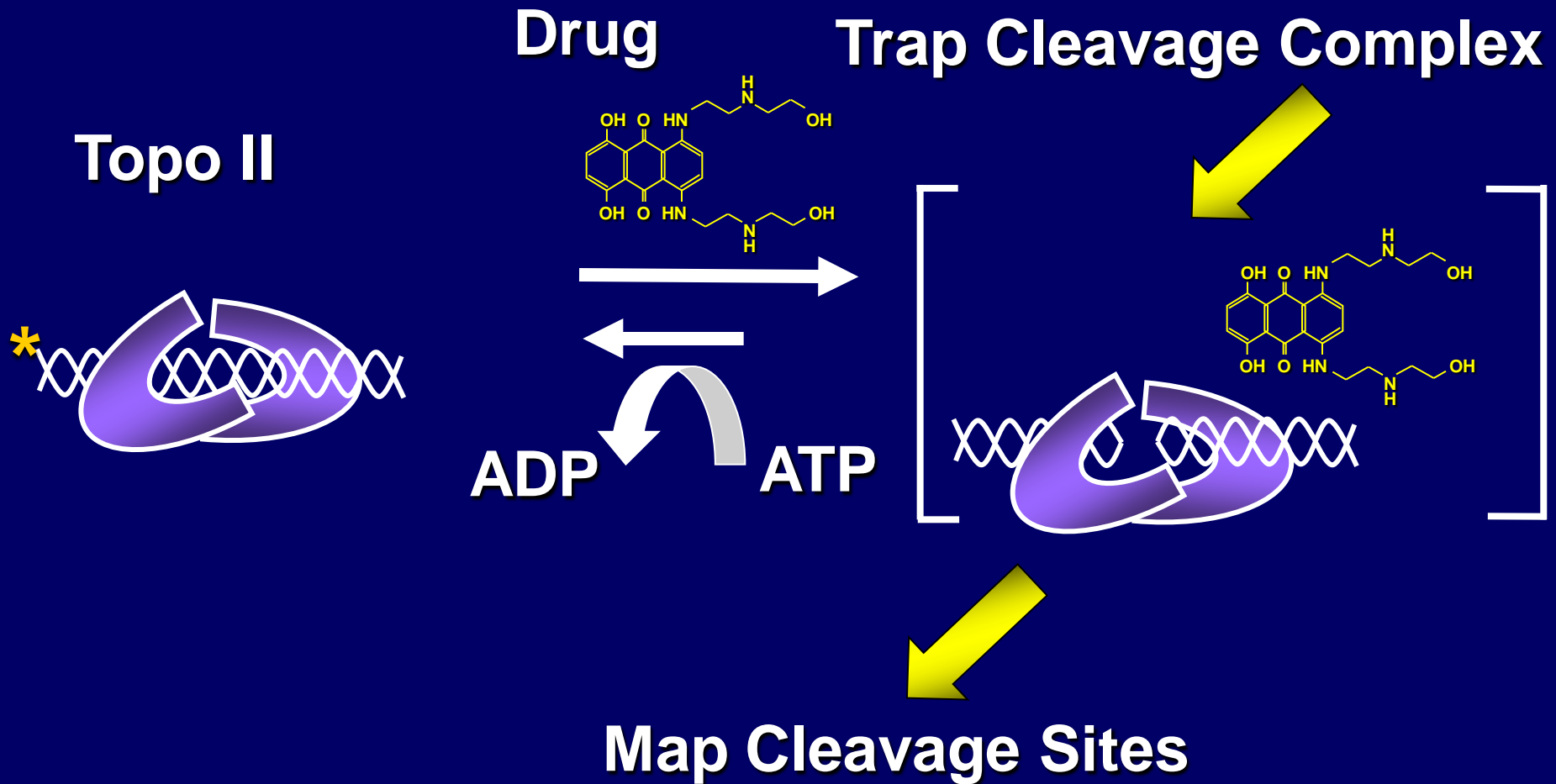
RARA

AGACCCCTTTGTCATGCCATCTCTCCCAGATGGGGGGTACCCAGAATAATGGGCTTTT

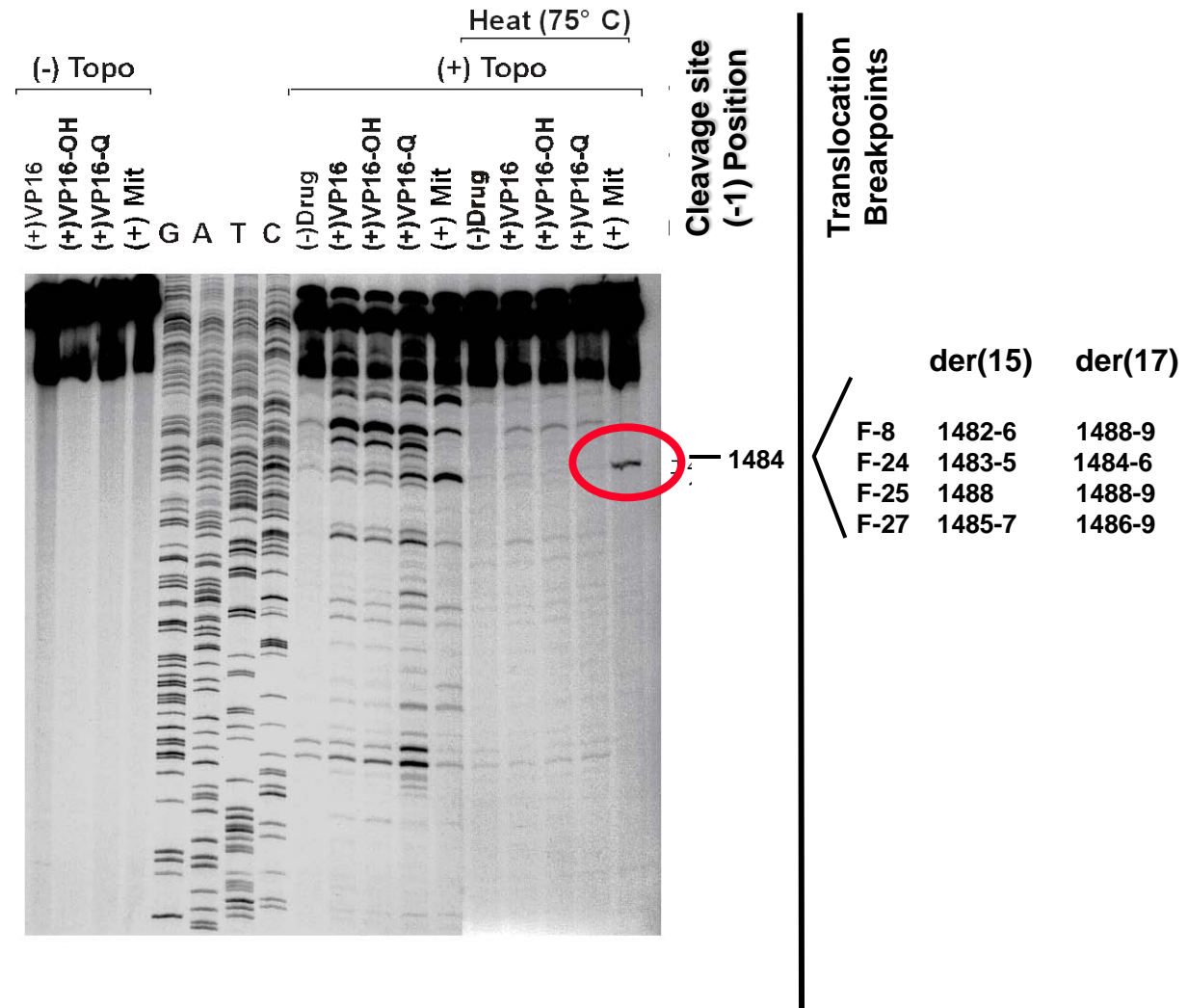
Defining mechanisms underlying AML as a complication of cancer therapy: An increasing healthcare problem



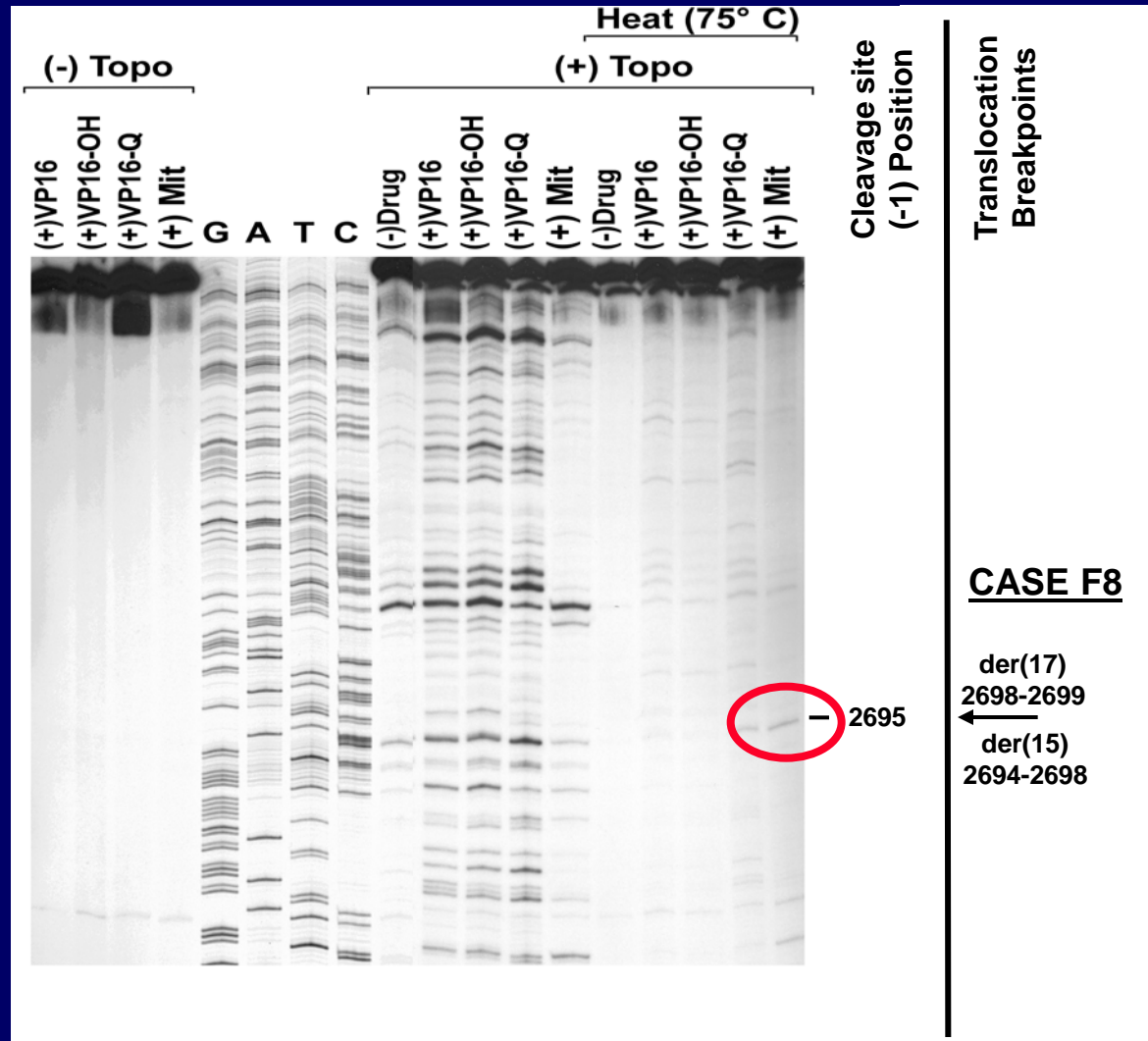
DNA topoisomerase II *in vitro* cleavage assay



Mitoxantrone induces strong, heat-stable topoisomerase II cleavage at *PML* translocation breakpoint hotspot



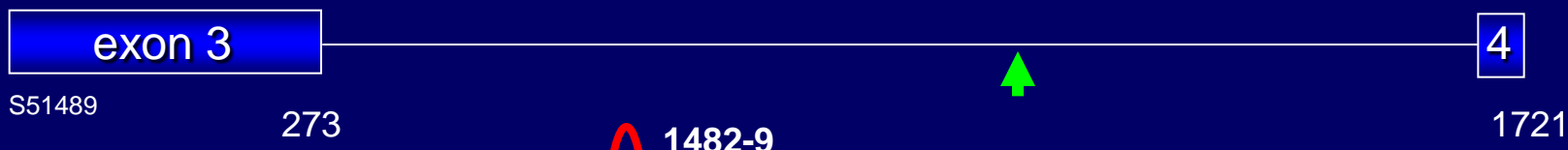
Mitoxantrone induces DNA topoisomerase II cleavage at *RARA* translocation breakpoints in t-APL



PML breakpoint clustering in t-APL arising after mitoxantrone treatment of malignant and benign conditions

PML

bcr3

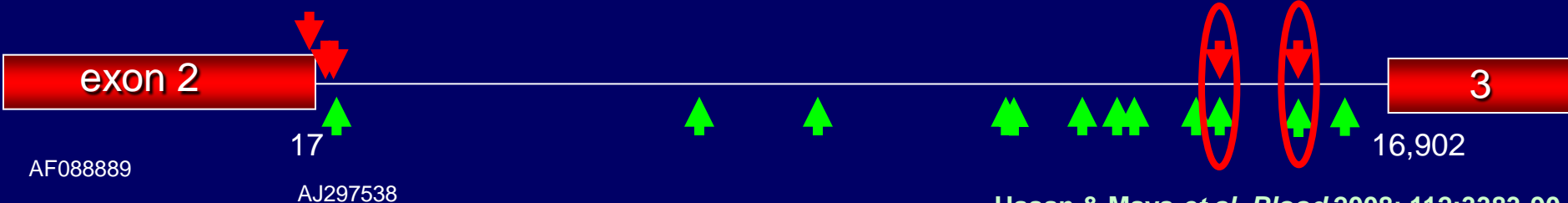


bcr1

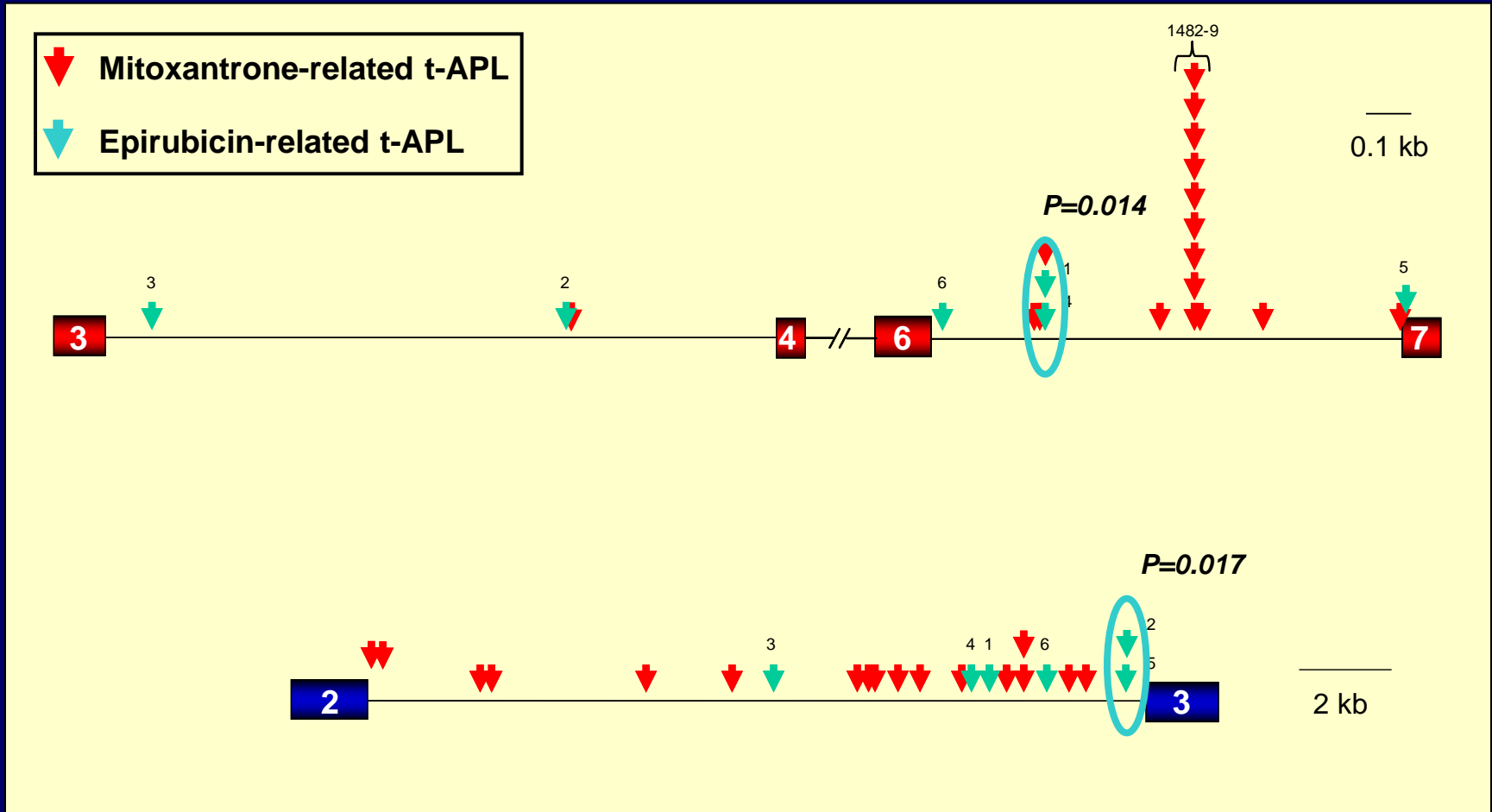


↓ Mitoxantrone for breast cancer
↑ Mitoxantrone for multiple sclerosis

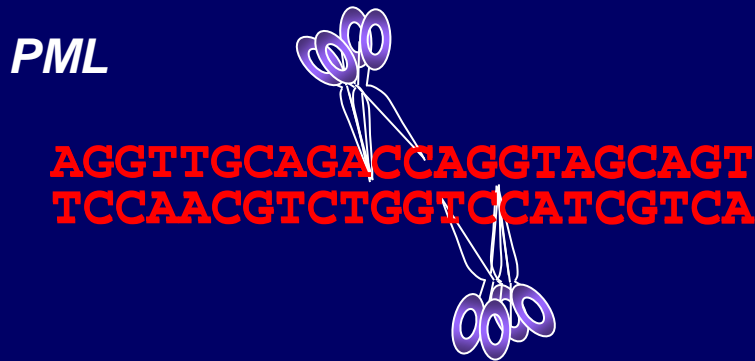
RARA



Genomic breakpoints in epirubicin-related t-APL cluster outside mitoxantrone hotspot



Model for formation of the t(15;17) chromosomal translocation in mitoxantrone-related t-APL



AGGTTGCAGA CCCCACACCTCCGG
TCCAACGTCTGGTC TGTGGAGGCC

CCGGAGTT CCAGGTAGCAGT
GGCCTCAAGGGG CATCGTCA

Exonucleolytic Digestion of Bases

AGGTTGCAGA CCCCACACCTCCGG
TCCAACGTCTGGTC TGTGGAGGCC

Nonhomologous End Joining

AGGTTGCAGACACACCTCCGG
TCCAACGTCTGGTGTGGAGGCC

Nonhomologous End Joining

CCGGAGTTCACAGGTAGCAGT
GGCCTCAAGGGGTCATCGTCA

Ligation

AGGTTGCAGACACACCTCCGG
TCCAACGTCTGGTGTGGAGGCC

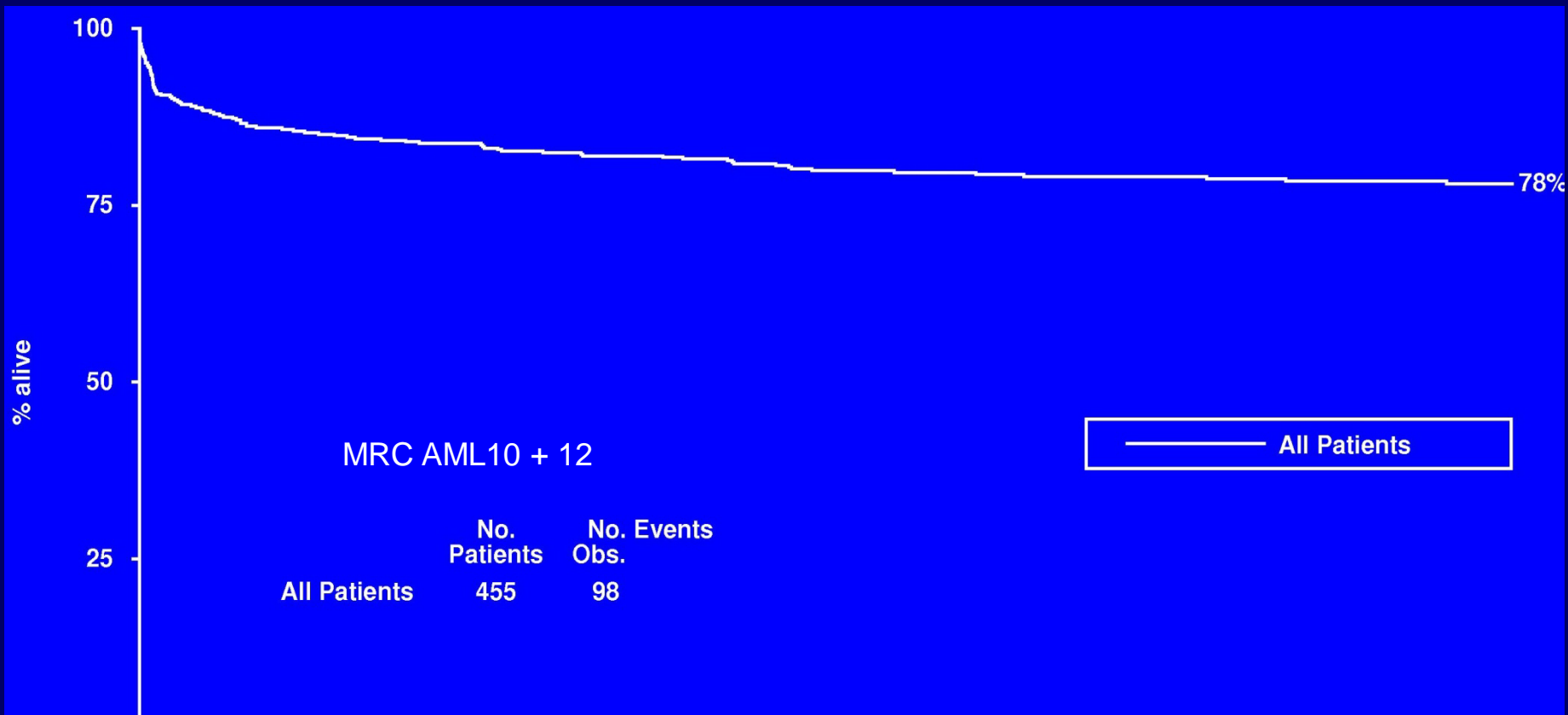
Ligation, Gap Fill In

CCGGAGTTCACAGGTAGCAGT
GGCCTCAAGGGTCCATCGTCA

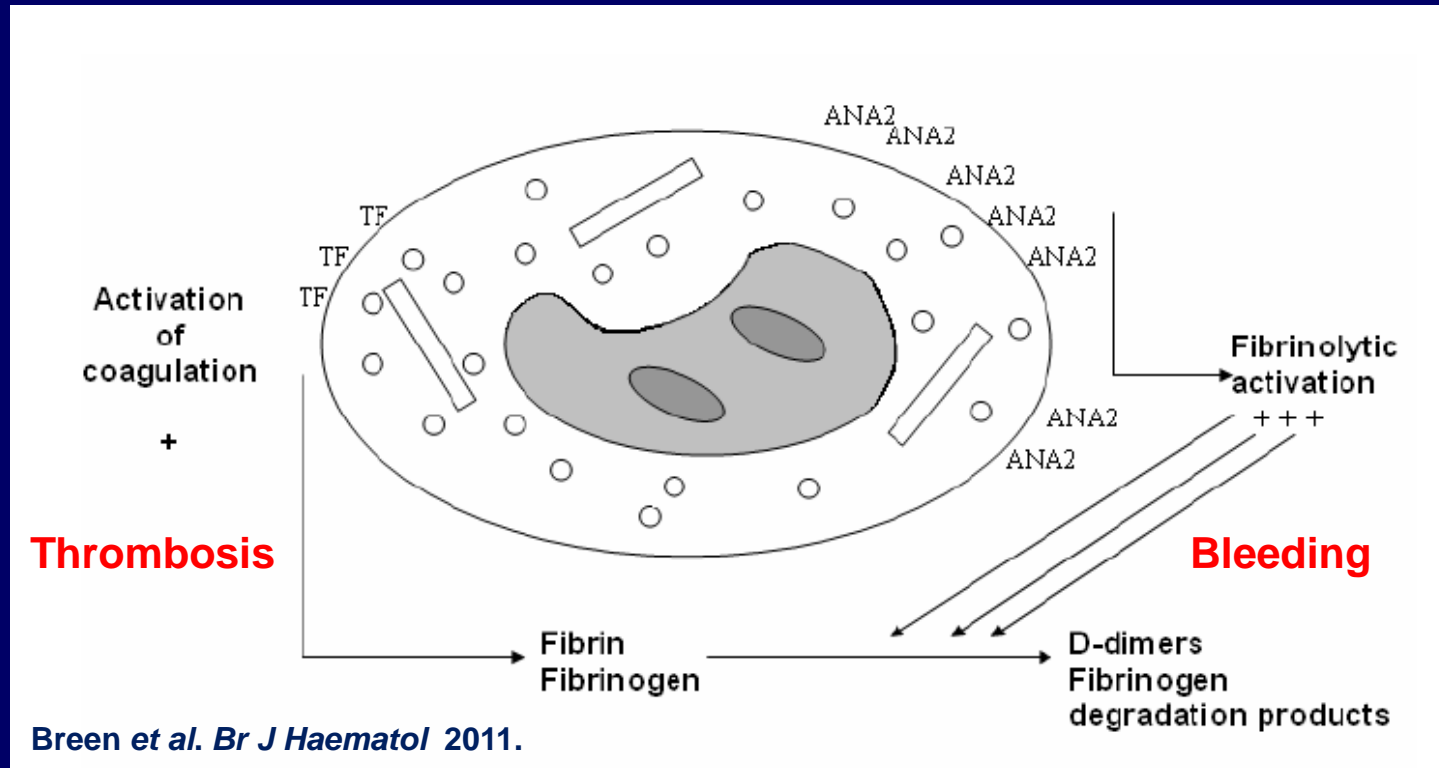
PML-RARA

RARA-PML

The challenge: How to improve on outcome already achieved with ATRA & anthracycline-based chemotherapy?



APL coagulopathy



- **Early death remains a significant problem:**

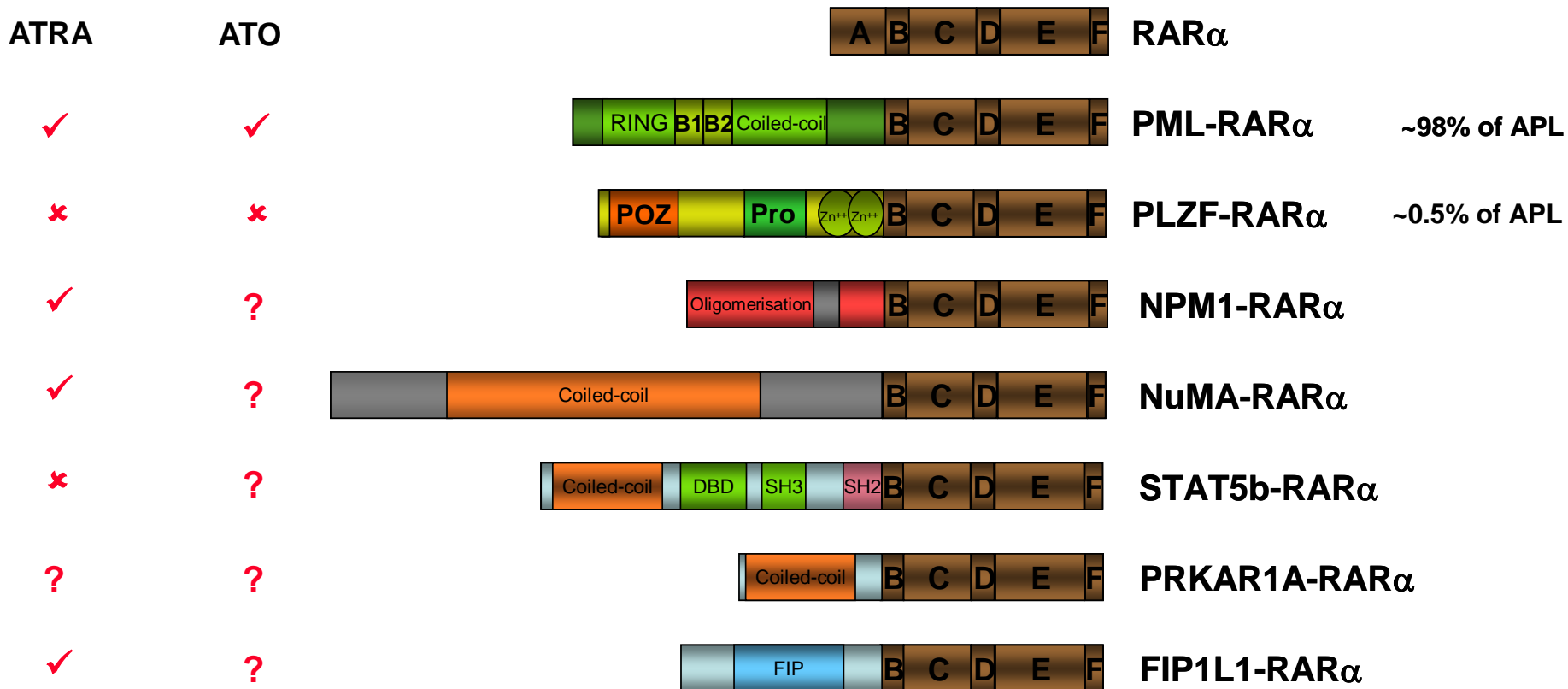
- 17% in SEER registry data (Park et al, *Blood* 2011; 118: 1248-54)
- 30 of 105 APL cases arising between 1997-2006 identified in Swedish population died within 30d (29%) (Lehmann et al, *Leukemia* 2011; 25: 1128-34)

Initial management of APL

- Commence ATRA as soon as diagnosis suspected
- Implementation of supportive care for coagulopathy
- Important to establish presence of PML-RAR α fusion, which predicts sensitivity to ATRA & arsenic:
 - PML immunofluorescence
 - Cytogenetics
 - PCR

Molecular basis of APL: Implications for targeted therapy

Sensitivity to
targeted therapy



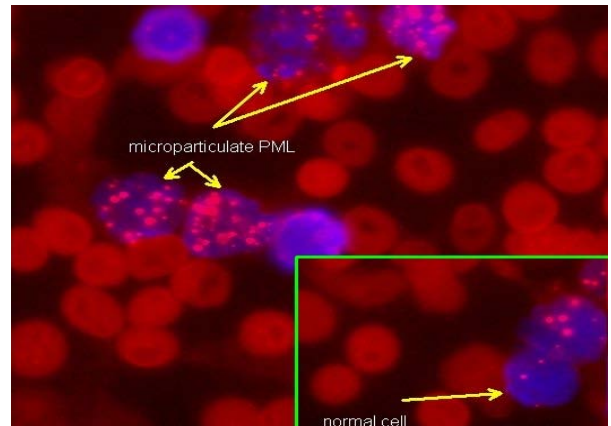
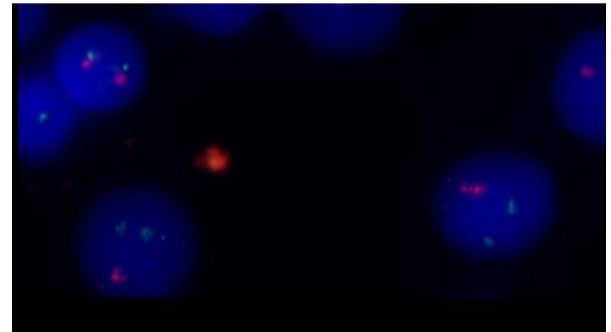
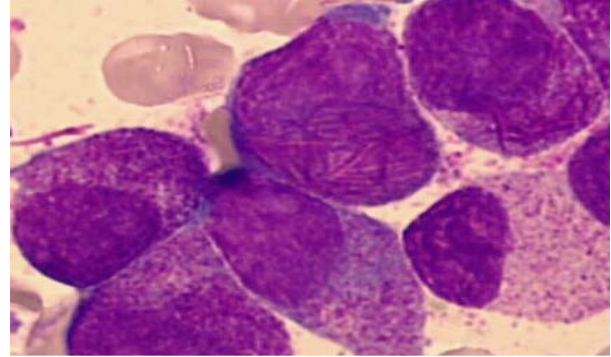
Is it APL?

- **Case History**

- 25yr old female
- Presented – PE (23.12.09)
- FBC: Hb 10g/dl, WBC 0.5, Plt 86
- Marrow – APL?
- Cytogenetics: Normal
- FISH: No *PML-RARA* fusion signal
- PML antibody test +ve
- RT-PCR: *PML-RARA* +ve
(reciprocal *RARA-PML* neg)

Diagnosis: APL secondary to
PML-RARA insertion

➔ ATRA + Idarubicin

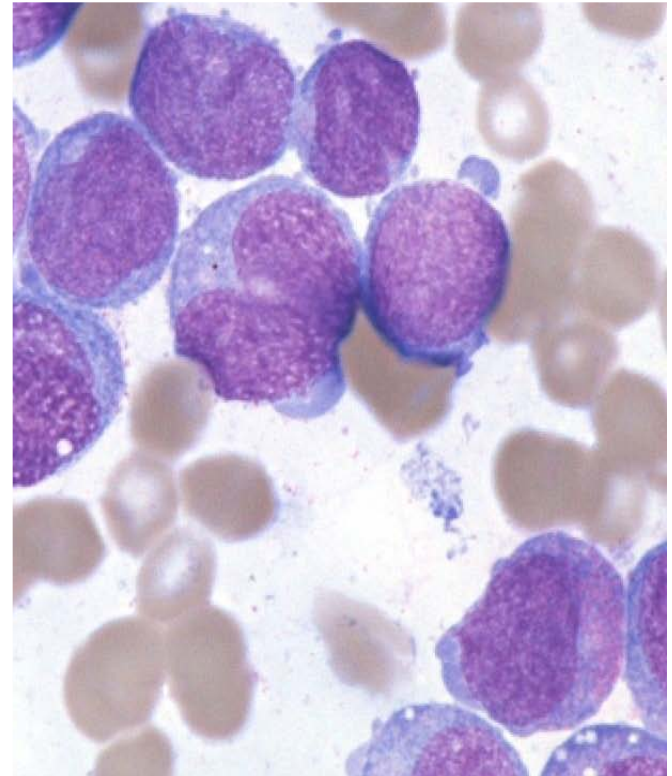


Importance of molecular diagnostics to guide appropriate therapy

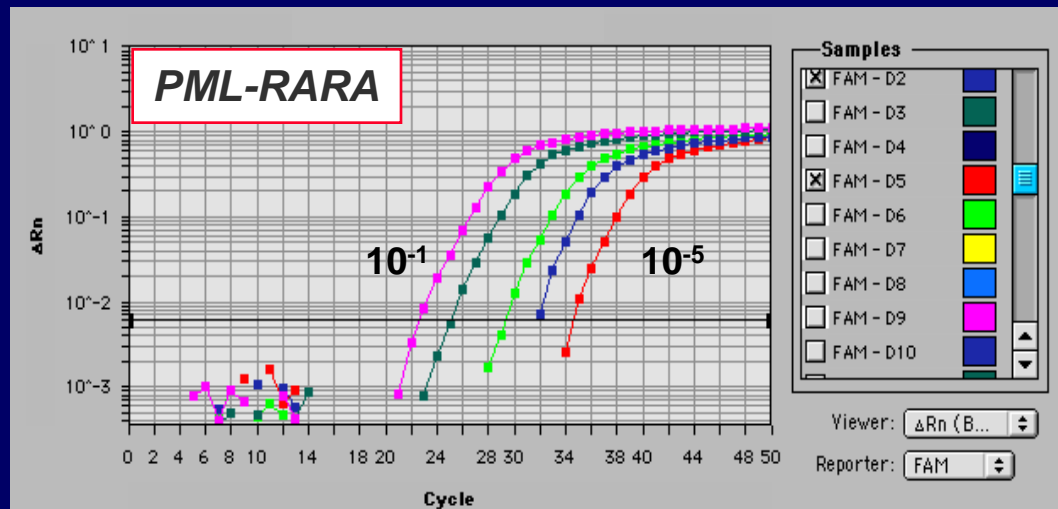
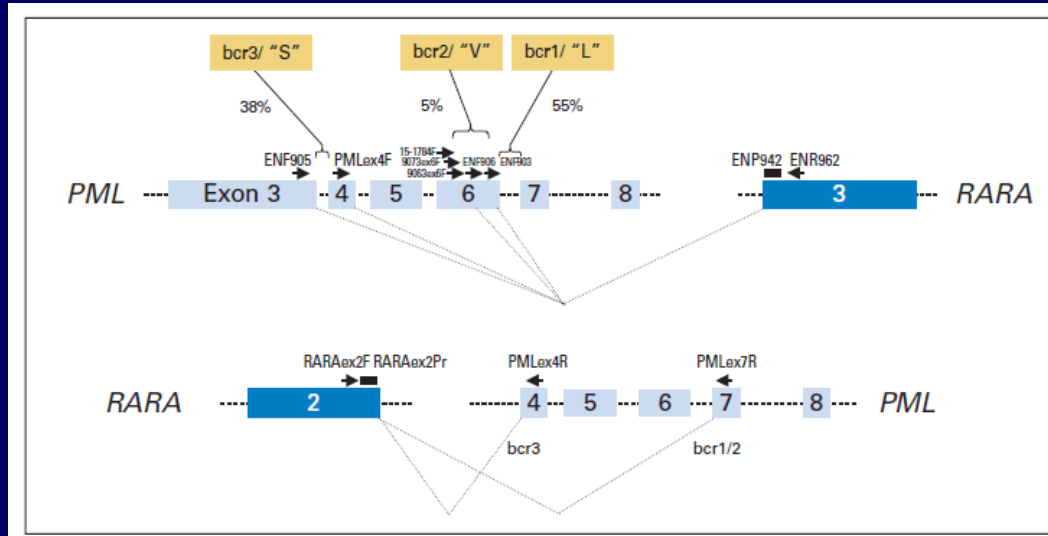
• Case History

- 75yr old female
- WBC $15 \times 10^9/l$
- Suspected M3v
- Randomised to AIDA in AML17 trial
- Cytogenetics: Normal
- FISH: No *PML-RARA* fusion signal
- RT-PCR: *PML-RARA* neg

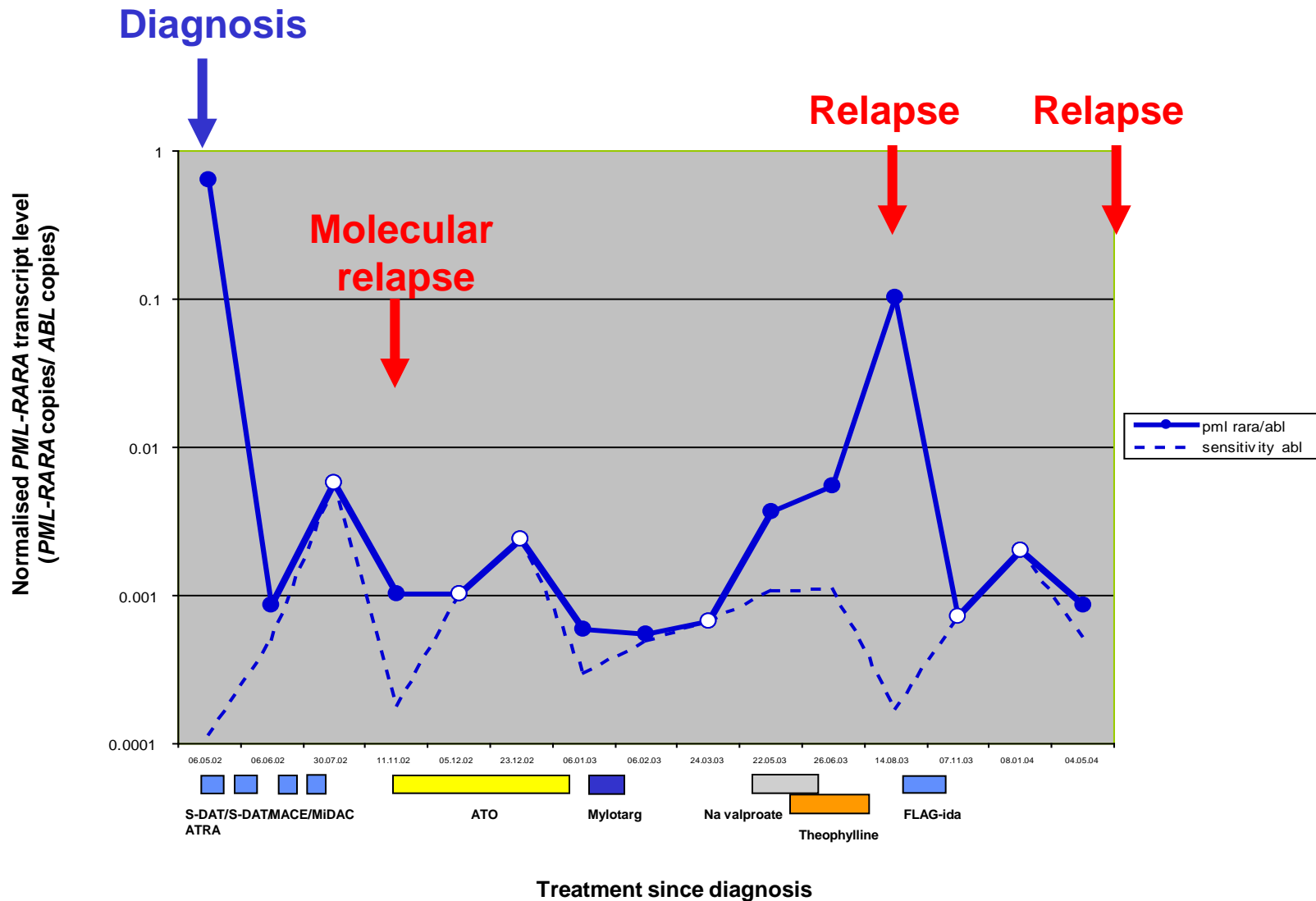
Diagnosis: NPM1 mutant AML



Real-time quantitative PCR (RQ-PCR) amplification of *PML-RARA* transcripts can detect submicroscopic levels of leukaemia

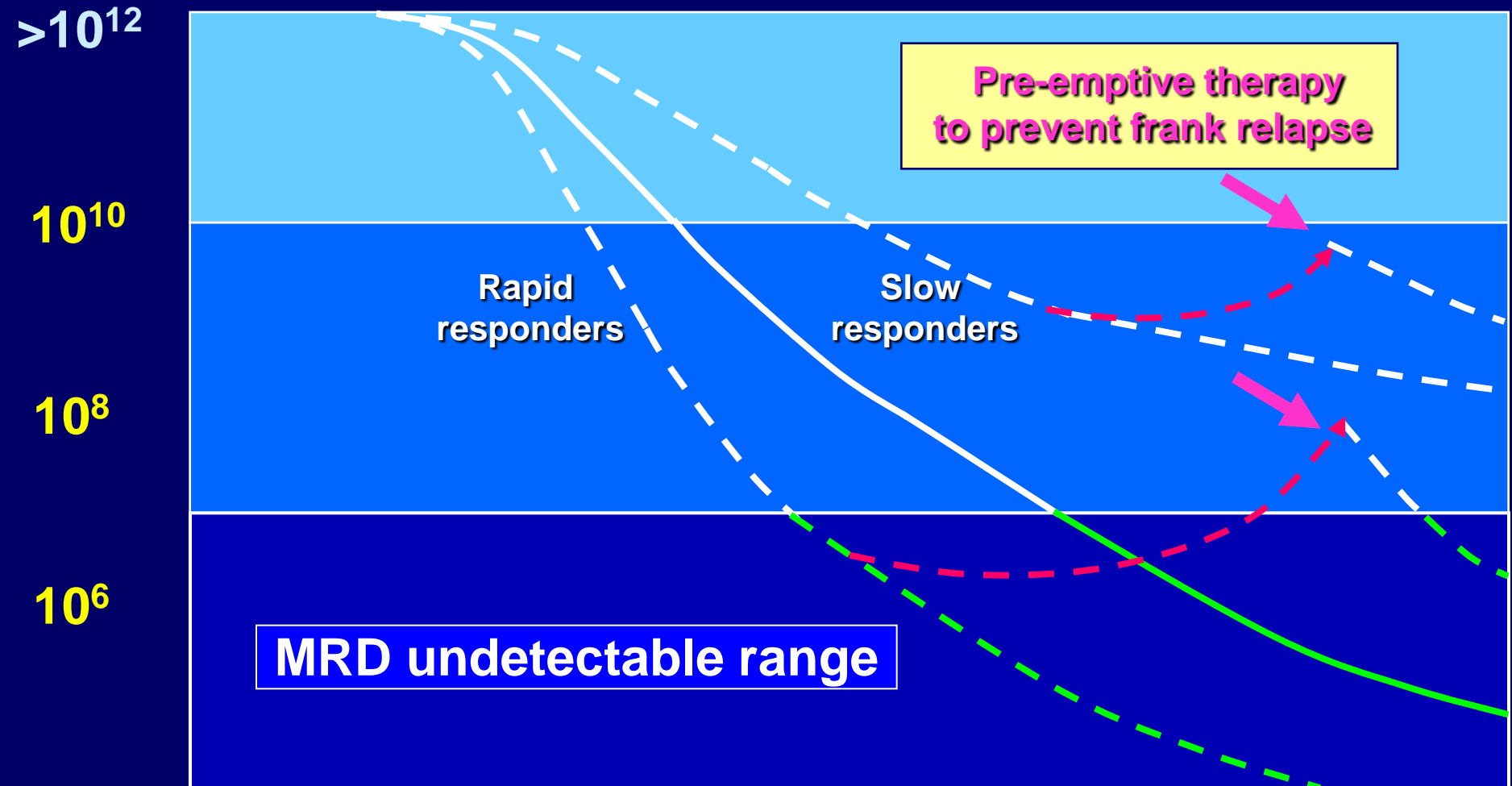


MRD monitoring using RQ-PCR to predict relapse in APL



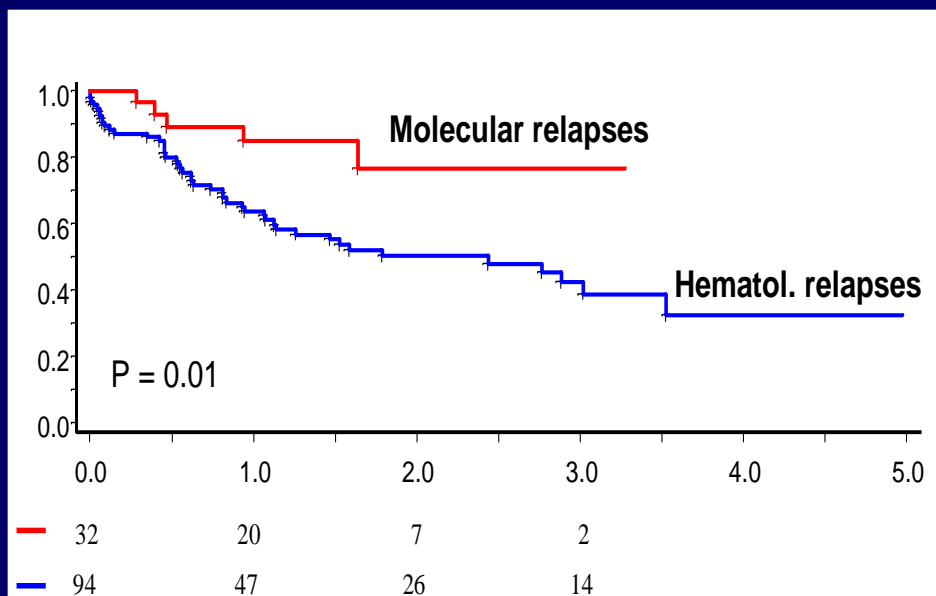
Use of sequential MRD monitoring to direct pre-emptive therapy to prevent impending relapse

Leukaemic cell burden

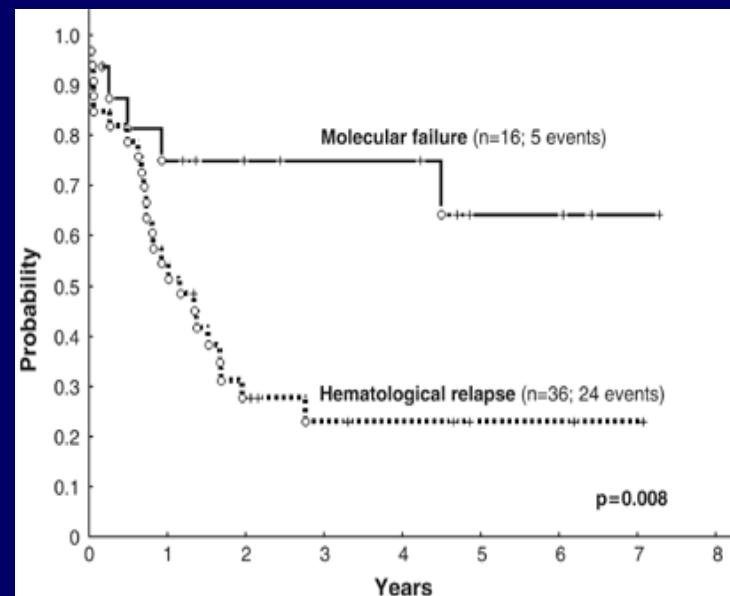


Defining role of MRD monitoring in APL: Outcome according to disease burden at time of relapse

GIMEMA



PETHEMA

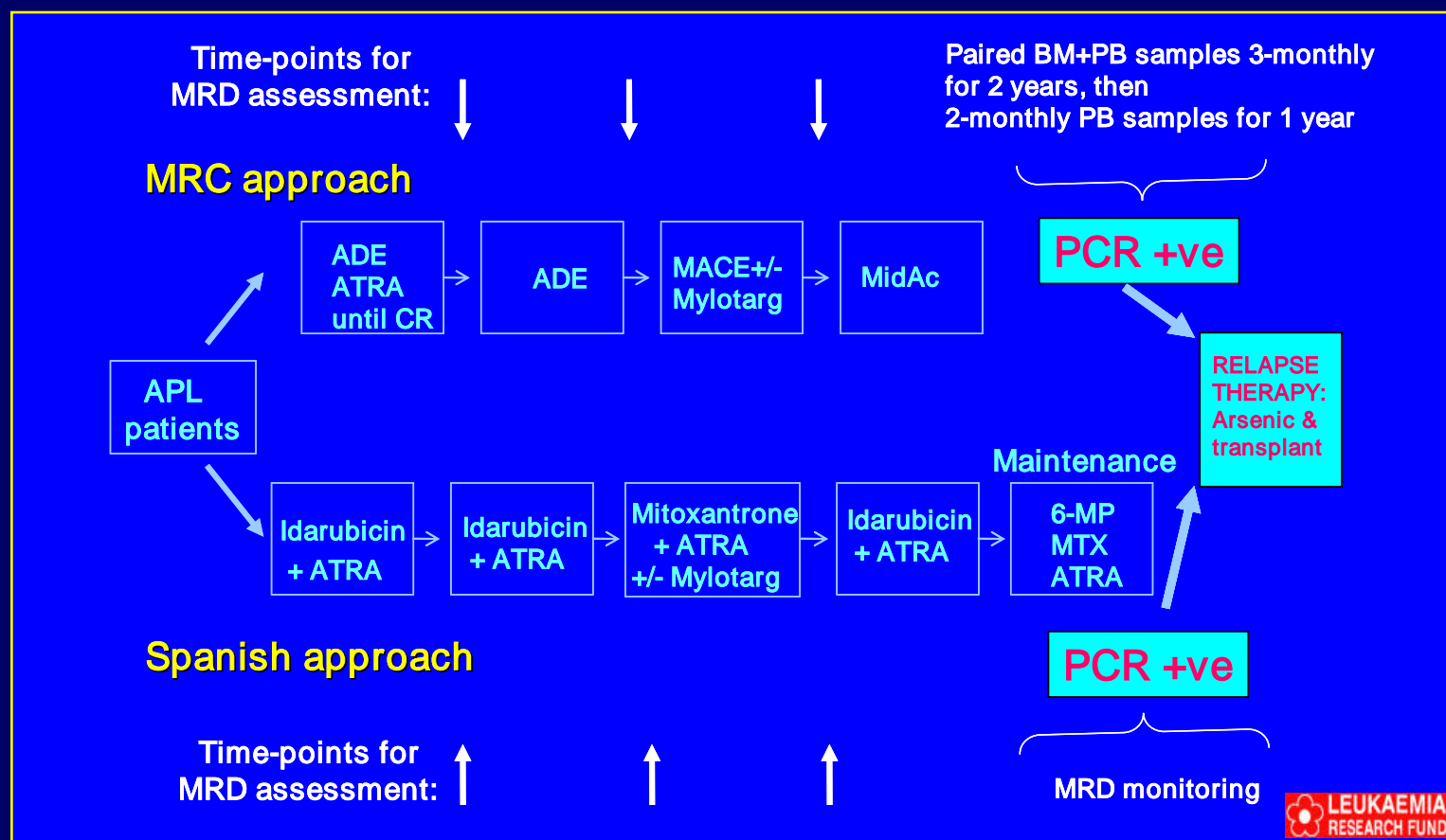


Lo Coco *et al*, *Semin Hematol* 2002; 39 (2 Suppl 1): 14-17.

Esteve *et al*, *Leukemia* 2007; 21: 446-452

Evaluation of MRD monitoring by RQ-PCR to determine treatment approach in patients with APL: MRC AML15 trial

- 281 patients, median age 42 yrs (16-69), median follow-up 26 months (0-56mo)
- 5,207 samples analysed
 - Including 1,832 paired BM+PB samples, median of 18 samples analysed/pt (range, 1-50)



Serial MRD monitoring by RQ-PCR to detect persistent disease/ molecular relapse is strongest independent predictor for clinical relapse

Parameter	Hazard ratio (95% CI)	P-value
Persistent PCR positivity or molecular relapse	33.56 (4.59-245.27)	<0.0001
PCR positivity at post-course 2 timepoint	8.88 (1.25-63.11)	0.0009
Presenting WBC	1.03 (1.01-1.05)	0.004

Multivariable analysis considering also:

- Pre-treatment *PML-RARA* expression level
- *PML* breakpoint
- Kinetics of *PML-RARA* transcript reduction after induction
- PCR status after any given course of chemotherapy (post #1, #2, #3)

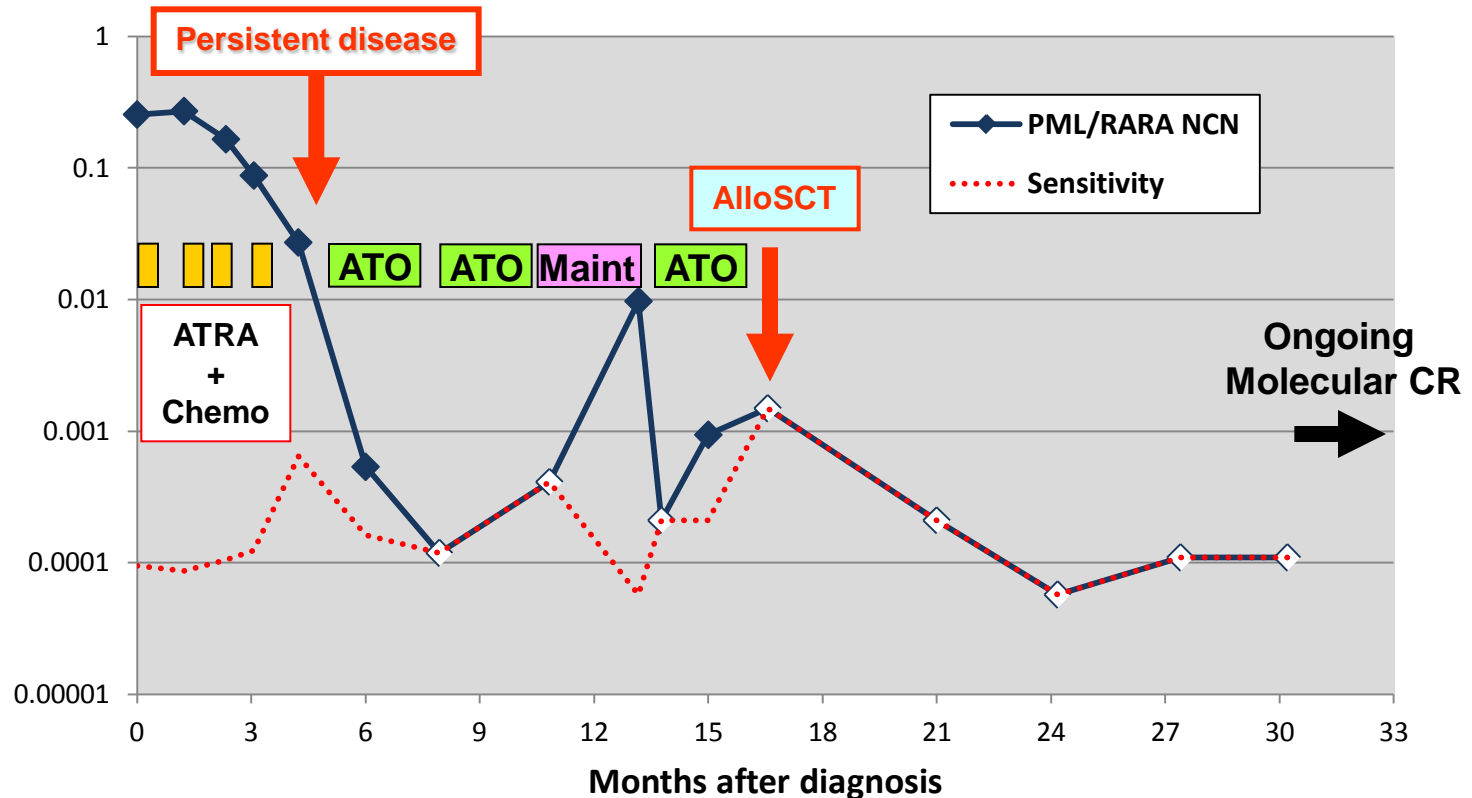
Serial MRD monitoring by standardised RQ-PCR assay to guide patient management in *PML-RARA*+ APL

Case History:

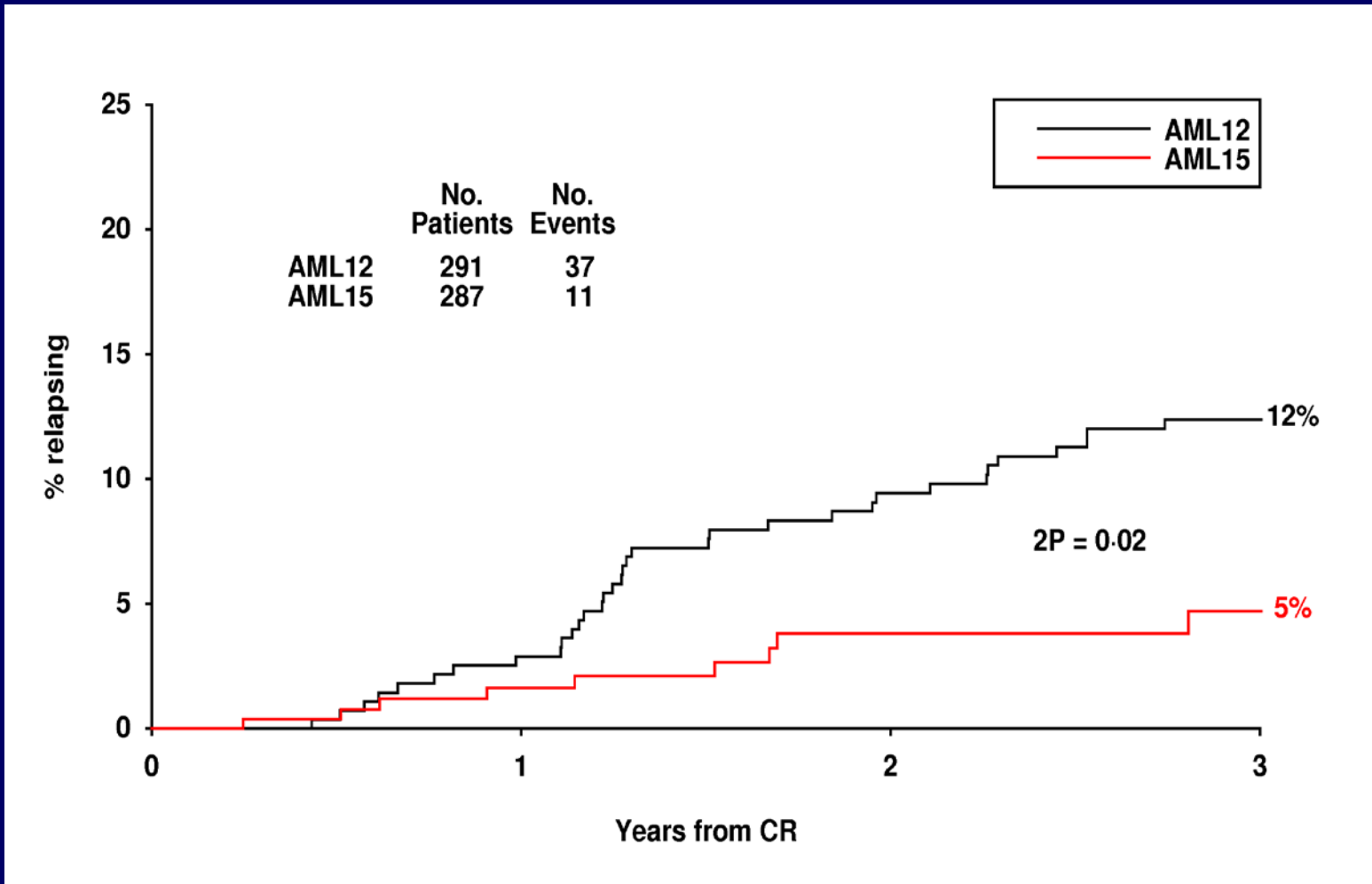
9 month male infant

FBC: Hb 8g/dl, WBC $28.6 \times 10^9/l$, Plt $12 \times 10^9/l$

PML-RARA+ APL

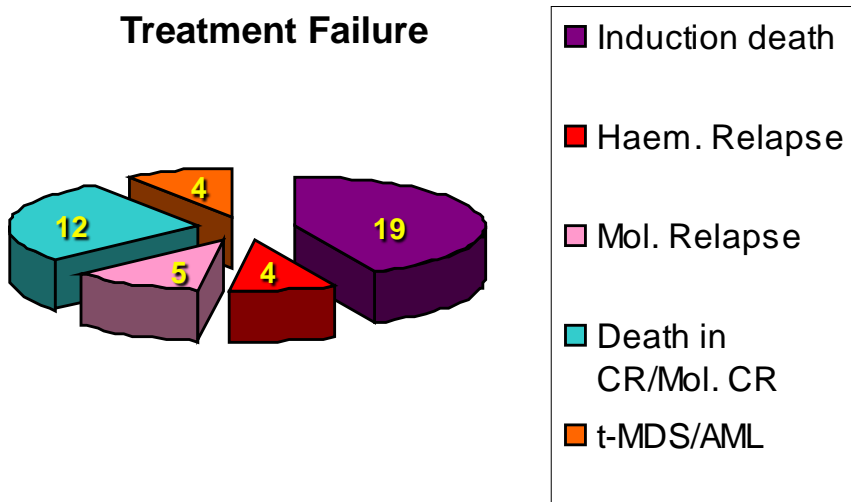


Evaluation of MRD monitoring & pre-emptive therapy to reduce rates of frank relapse in *PML-RARA*+ APL in MRC AML15 trial

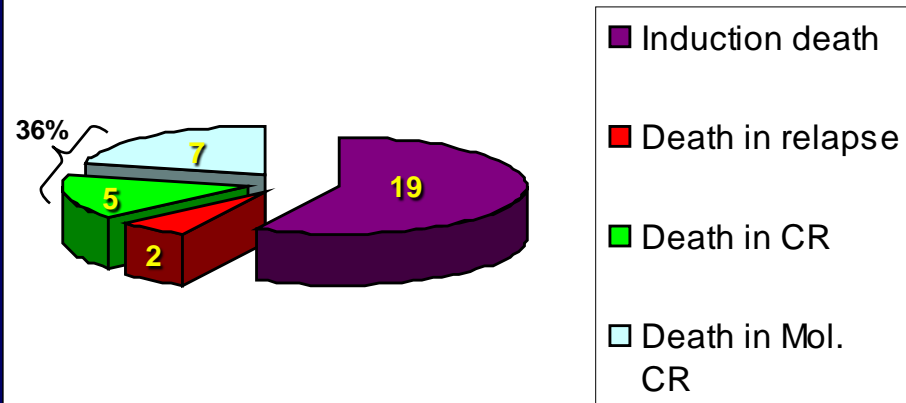


Causes of treatment failure & death in *PML-RARA+* APL in MRC AML15

Treatment Failure



Causes of death



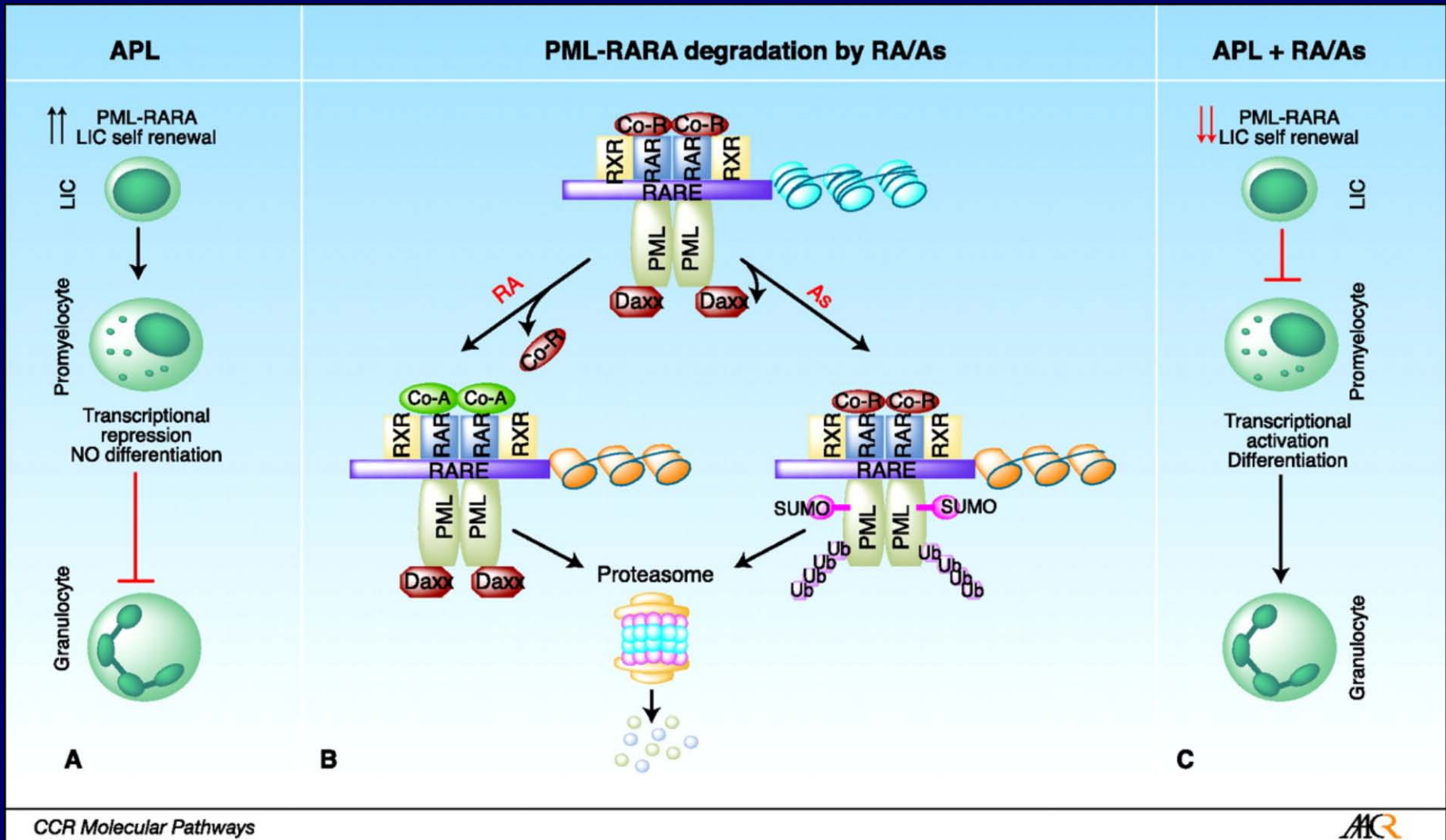
Treatment-related toxicity

~5% Death in CR/Molecular CR

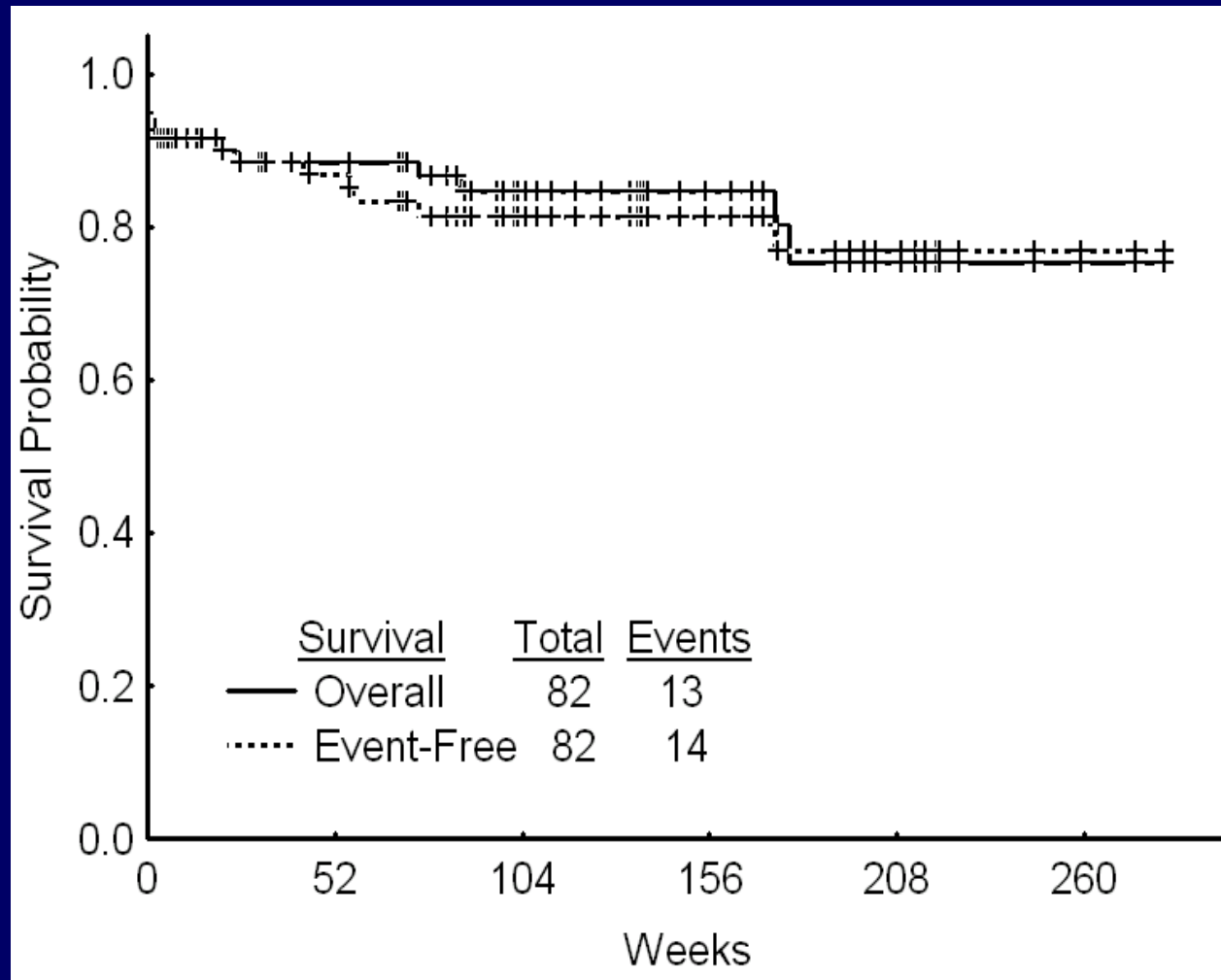
~5% Treatment curtailment/amendment (~3% with significant cardiac toxicity)

~2% t-MDS/ t-AML

ATO and ATRA exert synergistic activity *in vivo*

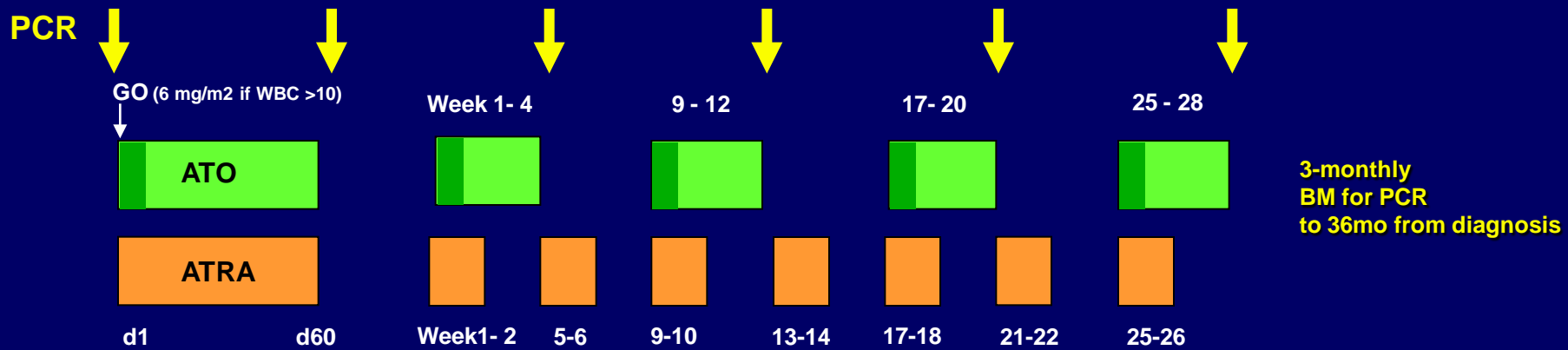


Outcome of *PML-RARA*+ APL following ATRA+ATO based therapy: Updated MD Anderson experience

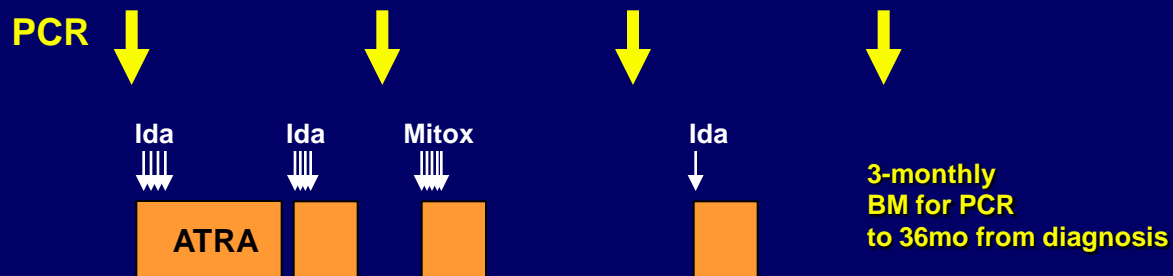


MRD monitoring to guide de-intensified treatment for *PML-RARA*+ APL in AML17 trial

CHEMO-FREE



AIDA



Acknowledgements

MRC/NCRI trials

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Elizabeth MacIntyre

Rayne Institute, King's College London

Bernd Zeisig
Eric So

Cancer Genetics Lab, King's College London

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