



Project BASTION „From Basic to Translational Research in Oncology”

Kick-off Meeting International Advisory Board Meeting

Warsaw

November 27th 2012

Tomasz Stokłosa



Philadelphia chromosome, chronic myelogenous leukemia and imatinib...



CD34+

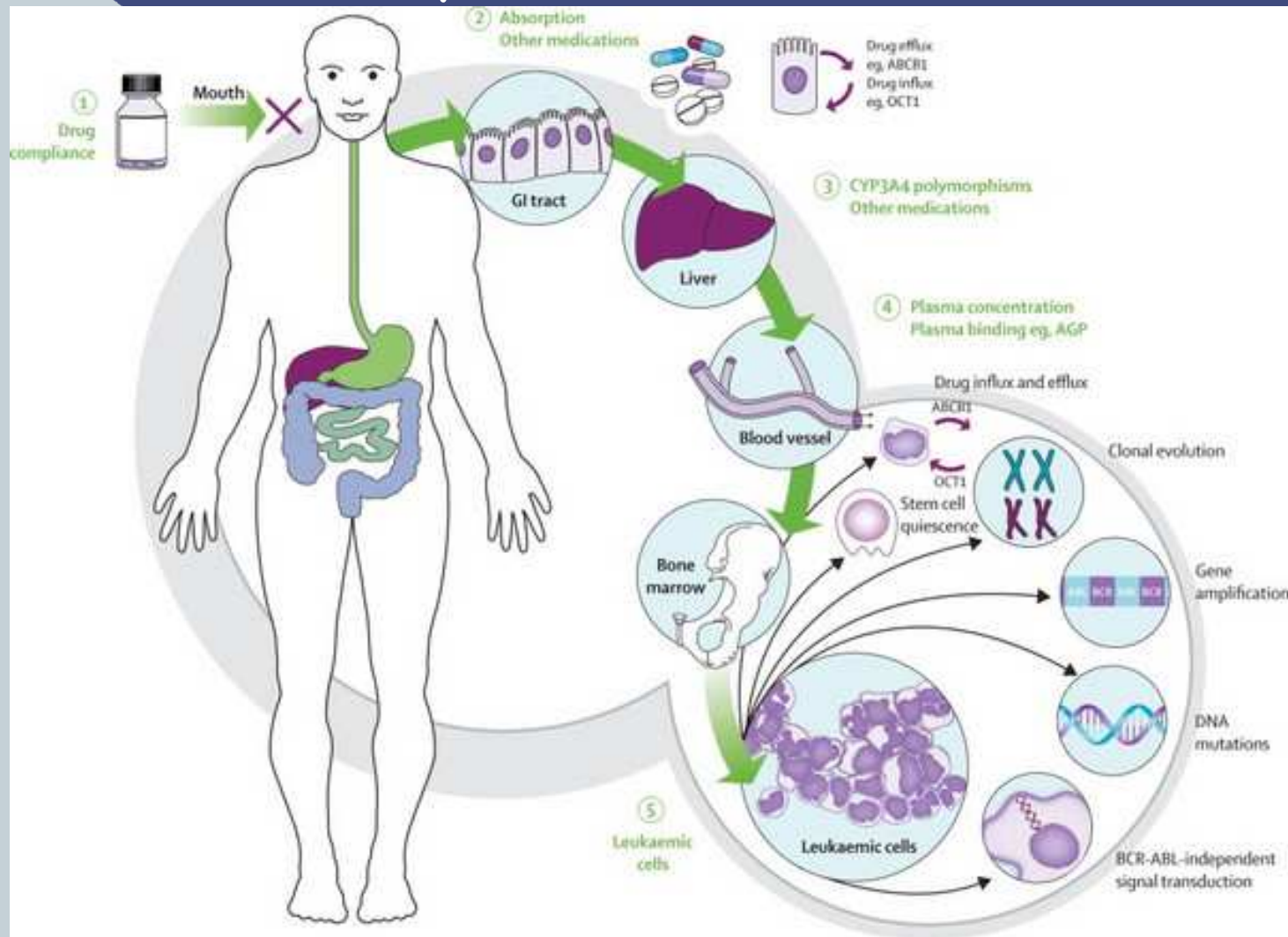
CD34+
Ph+



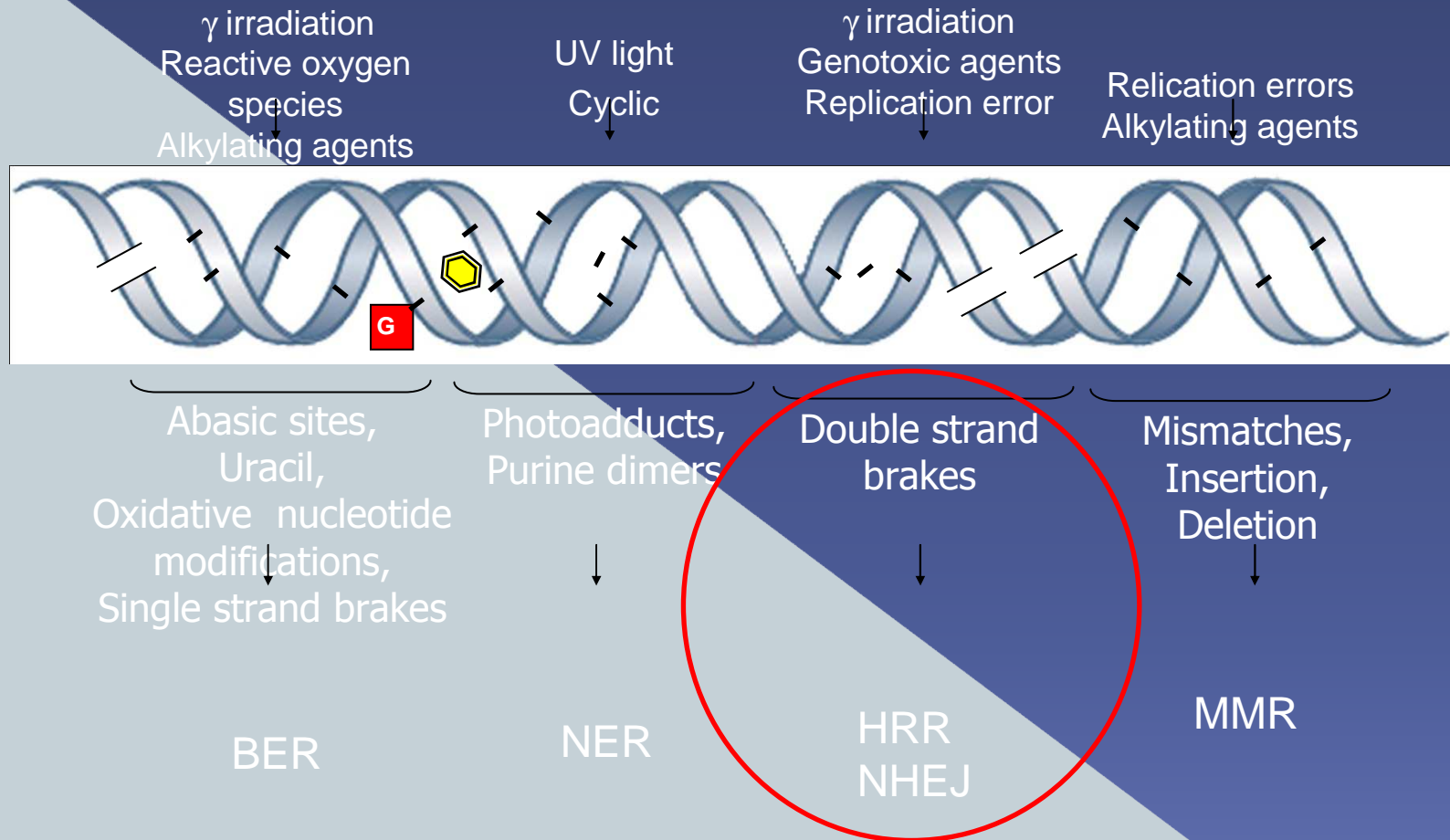
Chronic phase

>10 years

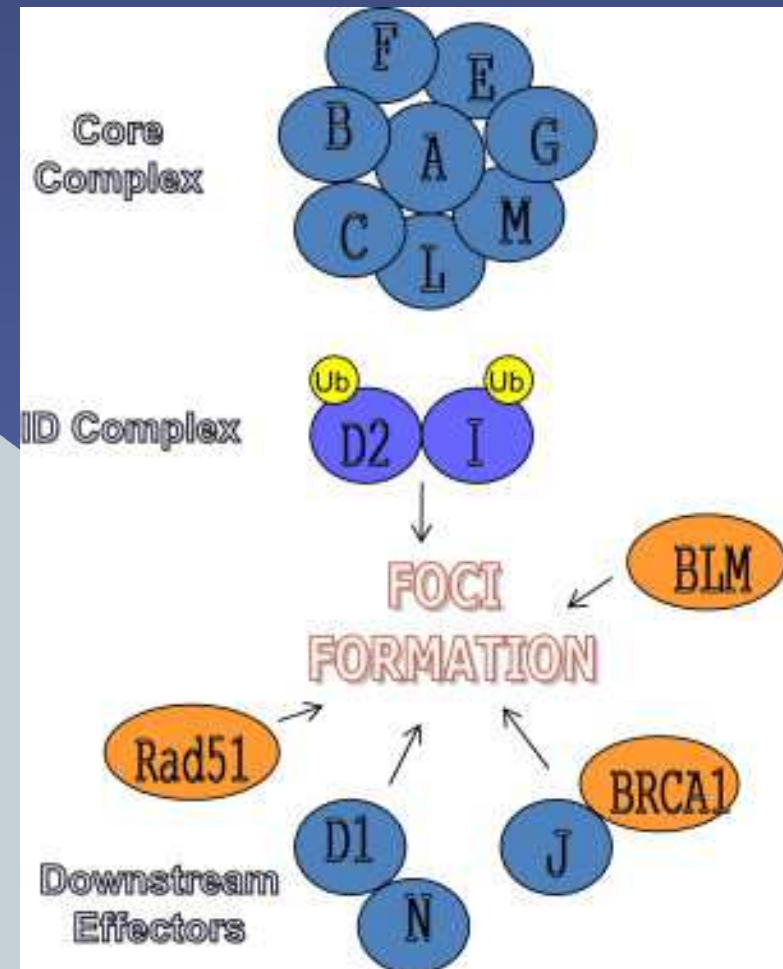
Molecular mechanisms of drug resistance in chronic myeloid leukemia



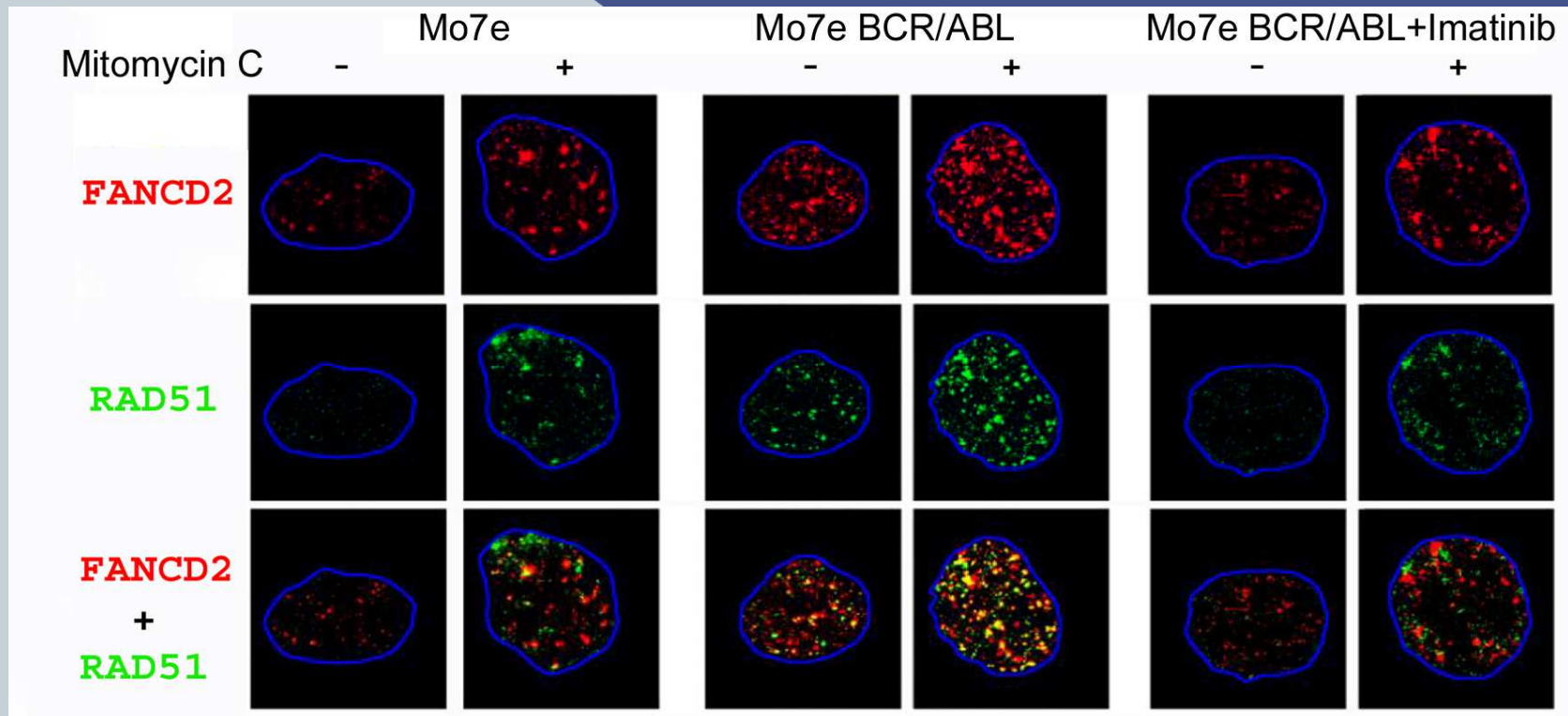
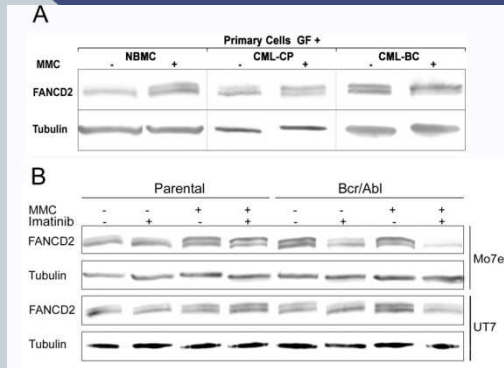
Major DNA repair mechanisms



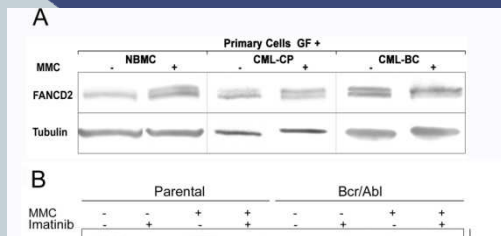
Role of Fanconi Anemia proteins in DNA damage response



FANCD2 pathway is hyperactivated in CML cells



FANCD2 pathway is hyperactivated in CML cells



Leukemia (2011), 1–9

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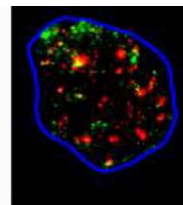
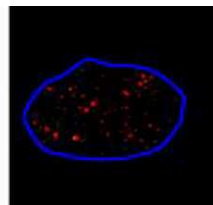
ORIGINAL ARTICLE

Monoubiquitinated Fanconi anemia D2 (FANCD2-Ub) is required for BCR-ABL1 kinase-induced leukemogenesis

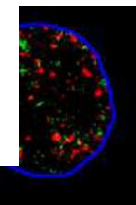
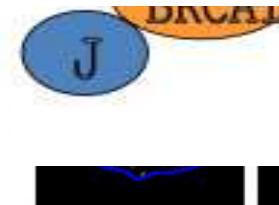
M Koptyra^{1,5}, T Stoklosa², G Hoser³, E Glodkowska-Mrowka², I Seferynska⁴, A Klejman⁵, J Blasiak⁶ and T Skorski^{1,5}

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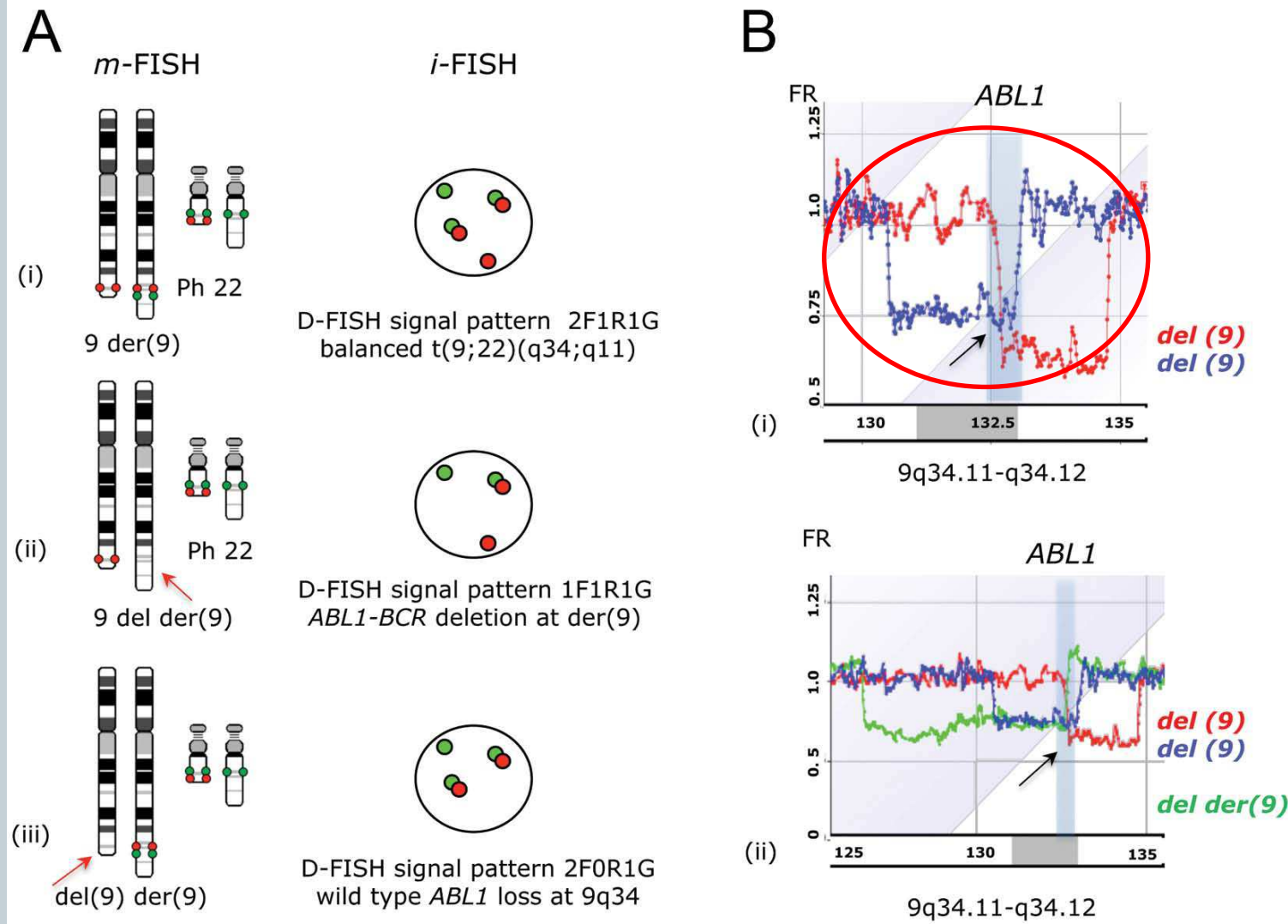
FANCD2
+
RAD51



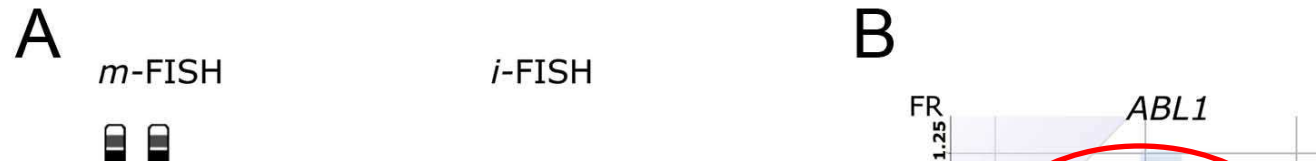
Downstream
Effectors



Microdeletions in normal *ABL* allele who didn't respond to TKI (lack of CCyR in 12 m).



Microdeletions in normal *ABL* allele who didn't respond to TKI (lack of CCyR in 12 m).



Published OnlineFirst June 21, 2011; DOI:10.1158/0008-5472.CAN-11-0068

Priority Report

Cancer
Research

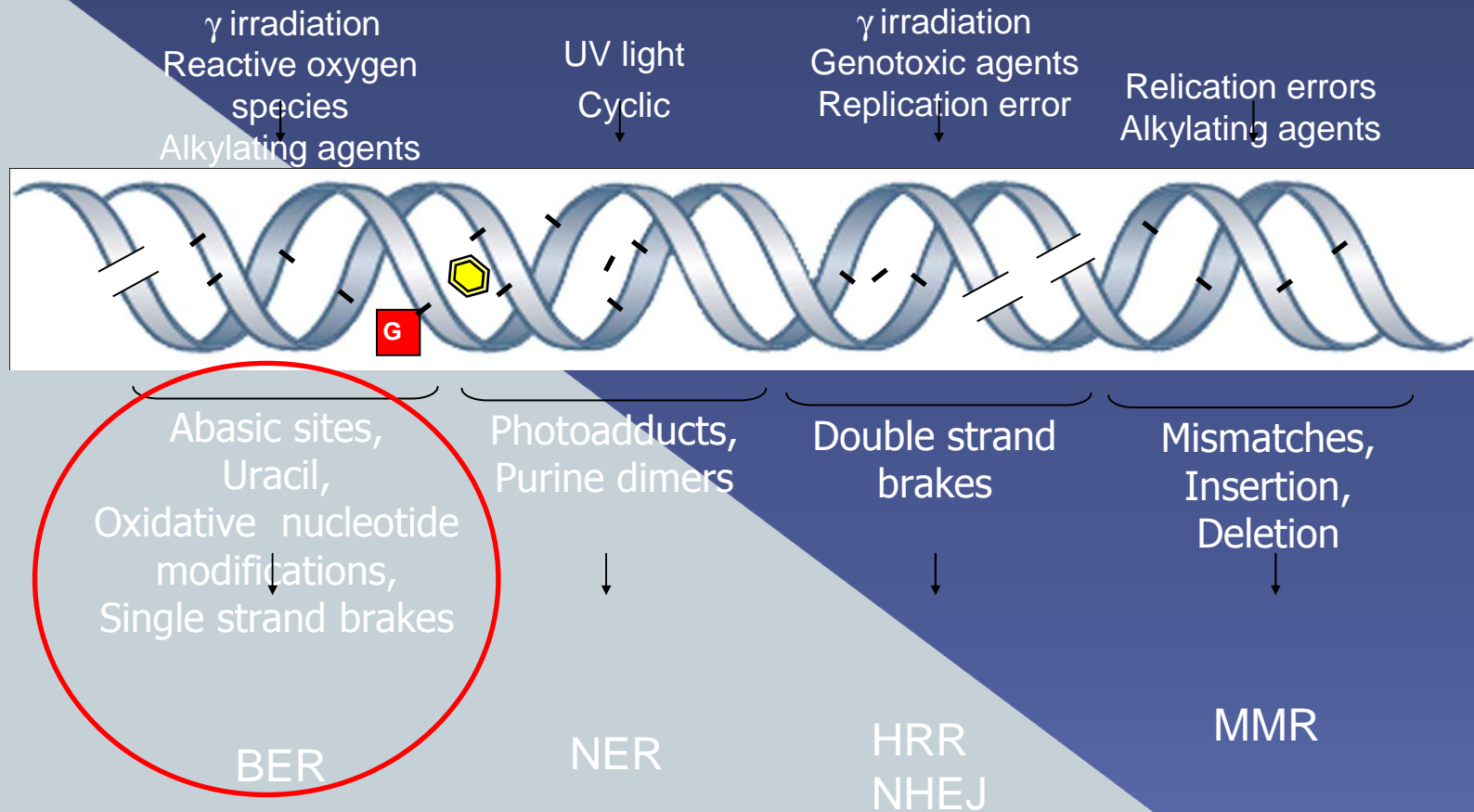
Imatinib Sensitivity in BCR-ABL1-Positive Chronic Myeloid Leukemia Cells Is Regulated by the Remaining Normal *ABL1* Allele

Anna Virgili¹, Mateusz Koptyra², Yashodhara Dasgupta², Eliza Glodkowska-Mrowka³, Tomasz Stoklosa³, Elisabeth P. Nacheva¹, and Tomasz Skorski²

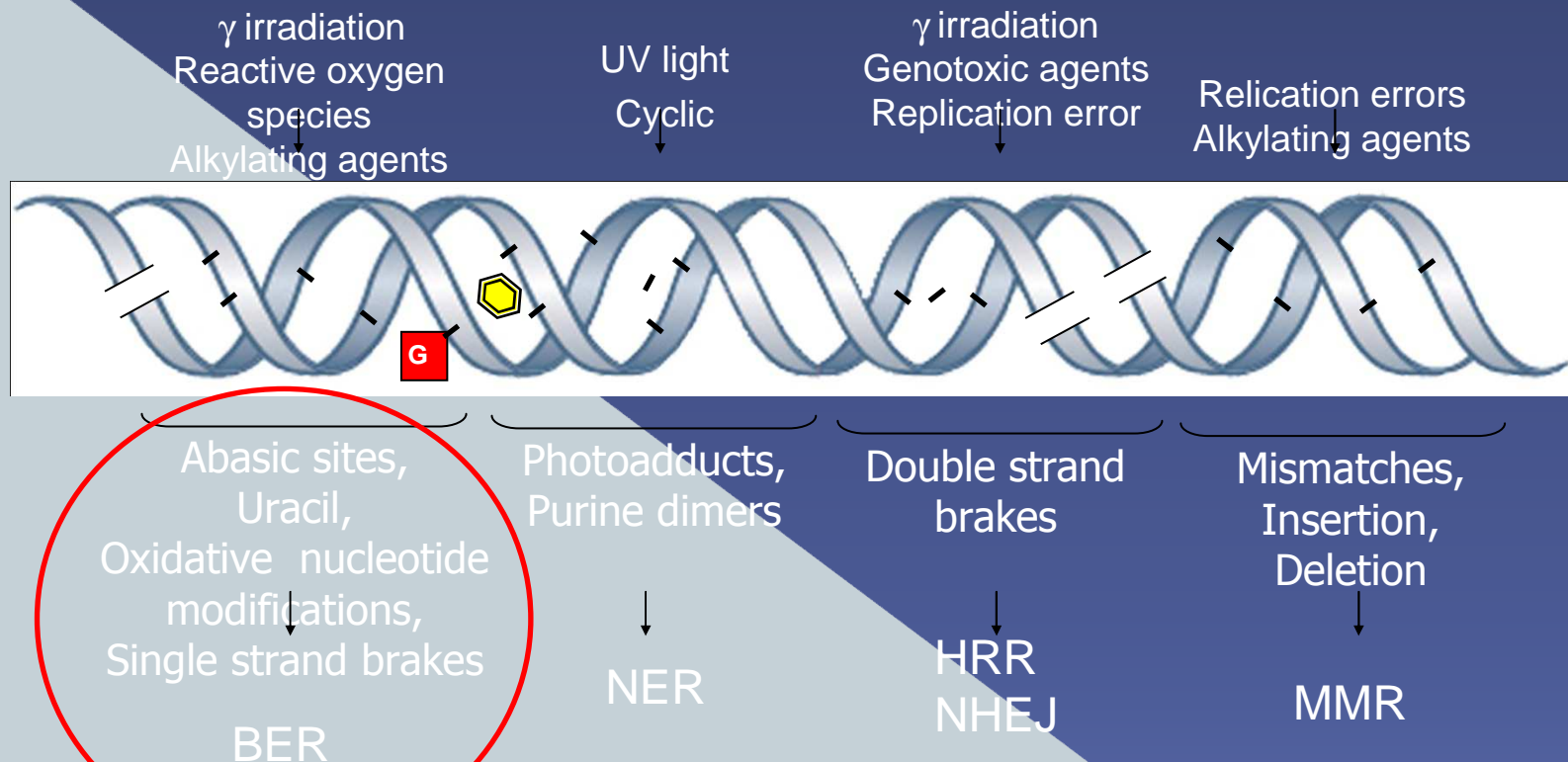
Abstract

Chronic myeloid leukemia in chronic phase (CML-CP) cells that harbor oncogenic *BCR-ABL1* and normal *ABL1* allele often become resistant to the ABL1 kinase inhibitor imatinib. Here, we report that loss of the remaining normal *ABL1* allele in these tumors, which results from cryptic interstitial deletion in 9q34 in patients who did not achieve a complete cytogenetic remission (CCyR) during treatment, engenders a novel unexpected mechanism of imatinib resistance. BCR-ABL1-positive *Abli*^{-/-} leukemia cells were refractory to imatinib as indicated by persistent BCR-ABL1-mediated tyrosine phosphorylation, lack of BCR-ABL1 protein degradation, increased cell survival, and clonogenic activity. Expression of ABL1 kinase, but not a kinase-dead mutant, restored the antileukemic effects of imatinib in ABL1-negative chronic myelogenous leukemia (CML) cells and in BCR-ABL1-positive *Abli*^{+/+} murine leukemia cells. The intracellular concentration of imatinib and expression of its transporters were not affected, although proteins involved in BCR-ABL1 degradation were down-regulated in *Abli*^{-/-} cells. Furthermore, 12 genes associated with imatinib resistance were favorably deregulated in *Abli*^{-/-} leukemia. Taken together, our results indicate that loss of the normal ABL1 kinase may serve as a key prognostic factor that exerts major impact on CML treatment outcomes. *Cancer Res*; 71(16): 5381–6. ©2011 AACR.

Major DNA repair mechanisms



Major DNA repair mechanisms



Leukemia (2012), 1–6

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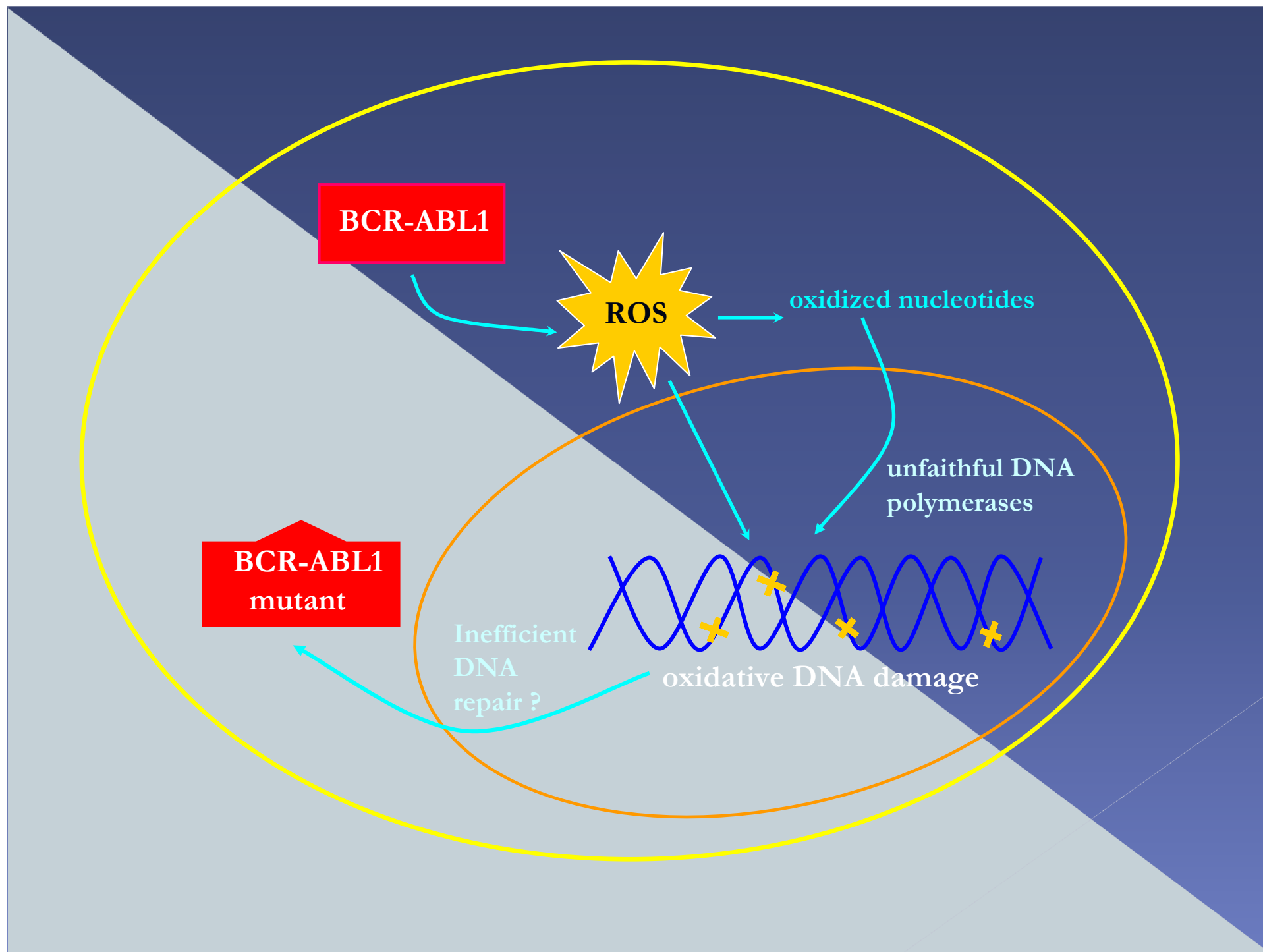
www.nature.com/leu



ORIGINAL ARTICLE

BCR-ABL1 kinase inhibits uracil DNA glycosylase UNG2 to enhance oxidative DNA damage and stimulate genomic instability

A Slupianek¹, R Falinski¹, P Znojek², T Stoklosa^{1,2}, S Flis¹, V Doneddu³, D Pytel^{1,5}, E Synowiec⁴, J Blasiak⁴, A Bellacosa³ and T Skorski¹



The diagram features a red rectangular box labeled 'BCR-ABL1' on the left. A curved cyan arrow points from this box to a yellow starburst shape in the center labeled 'ROS'. From the 'ROS' starburst, a straight cyan arrow points to the text 'oxidized nucleotides' on the right. The entire diagram is set against a dark blue background with a large yellow arc and a smaller orange arc.

BCR-ABL1

ROS

oxidized nucleotides

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MYELOID NEOPLASIA

Rac2-MRC-cIII-generated ROS cause genomic instability in chronic myeloid leukemia stem cells and primitive progenitors

Margaret Nieborowska-Skorska,¹ Piotr K. Kopinski,¹ Regina Ray,¹ Grazyna Hoser,² Danielle Ngaba,¹ Sylwia Flis,¹ Kimberly Cramer,¹ Mamatha M. Reddy,³ Mateusz Koptyra,¹ Tyrone Penserga,¹ Eliza Glodkowska-Mrowka,⁴ Elisabeth Bolton,¹ Tessa L. Holyoake,⁵ Connie J. Eaves,⁶ Sabine Cerny-Reiterer,⁷ Peter Valent,⁷ Andreas Hochhaus,⁸ Timothy P. Hughes,⁹ Heiko van der Kuip,¹⁰ Martin Sattler,³ Wieslaw Wiktor-Jedrzejczak,¹¹ Christine Richardson,¹² Adrienne Dorrance,¹³ Tomasz Stoklosa,⁴ David A. Williams,¹³ and Tomasz Skorski¹

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- Molecular mechanisms of drug resistance in chronic myeloid leukemia
- *Investigation of the potential targets and markers of sensitivity to tyrosine kinase inhibitors in chronic lymphocytic leukaemia*



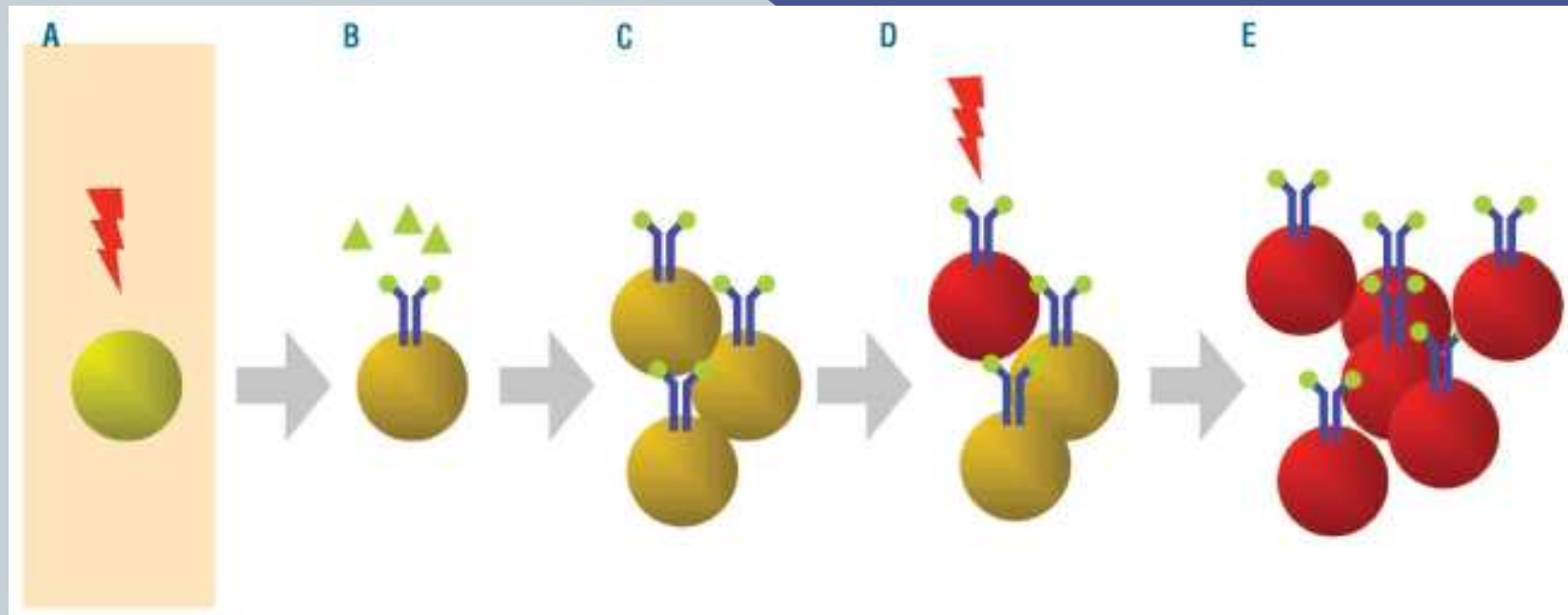
Chronic lymphocytic leukemia (CLL)

Chronic lymphocytic leukemia remains most often diagnosed leukemia in adults and there is no curative treatment for this disease.

Chronic lymphocytic leukemia (CLL)

Chronic lymphocytic leukaemia remains most often diagnosed leukaemia in adults and there is no curative treatment for this disease.

Putative model for the mechanism of
leukemogenesis of CLL.



Chronic lymphocytic leukemia (CLL)

- We plan to investigate potential pathways and targets for TKIs in CLL cells, employing gene expression profiling (GEP) analysis.
- We aim to characterize genes involved in response to these promising novel drugs for CLL and we will try to define group of patients who will mostly benefit from such treatments (based also on cytogenetic as well as molecular characterization of prognostic markers).



Thank you!

