



iCAN Digital Precision
Cancer Medicine

FIMM
Institute for Molecular Medicine Finland
Nordic EMBL Partnership for Molecular Medicine
HiLIFE UNIT



Novel combinations to target *NPM1* mutated acute myeloid leukemia

Daniela Mendoza-Ortiz

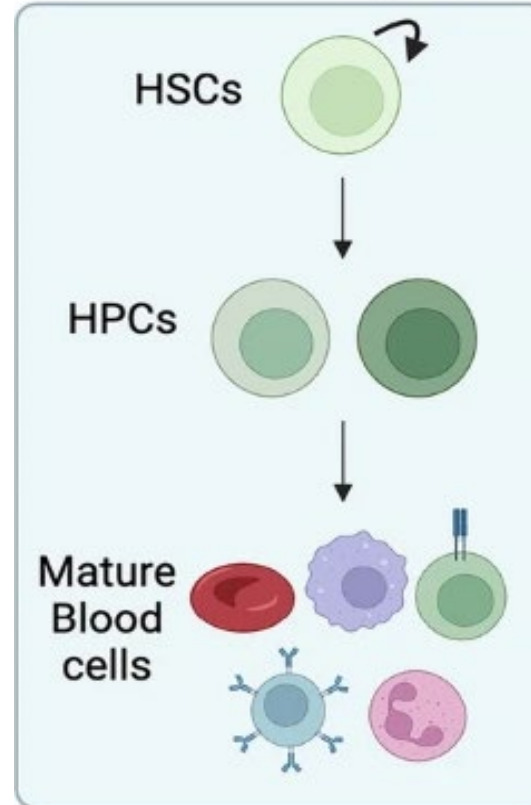
26.09.2024

Disclosures

Nothing to disclose

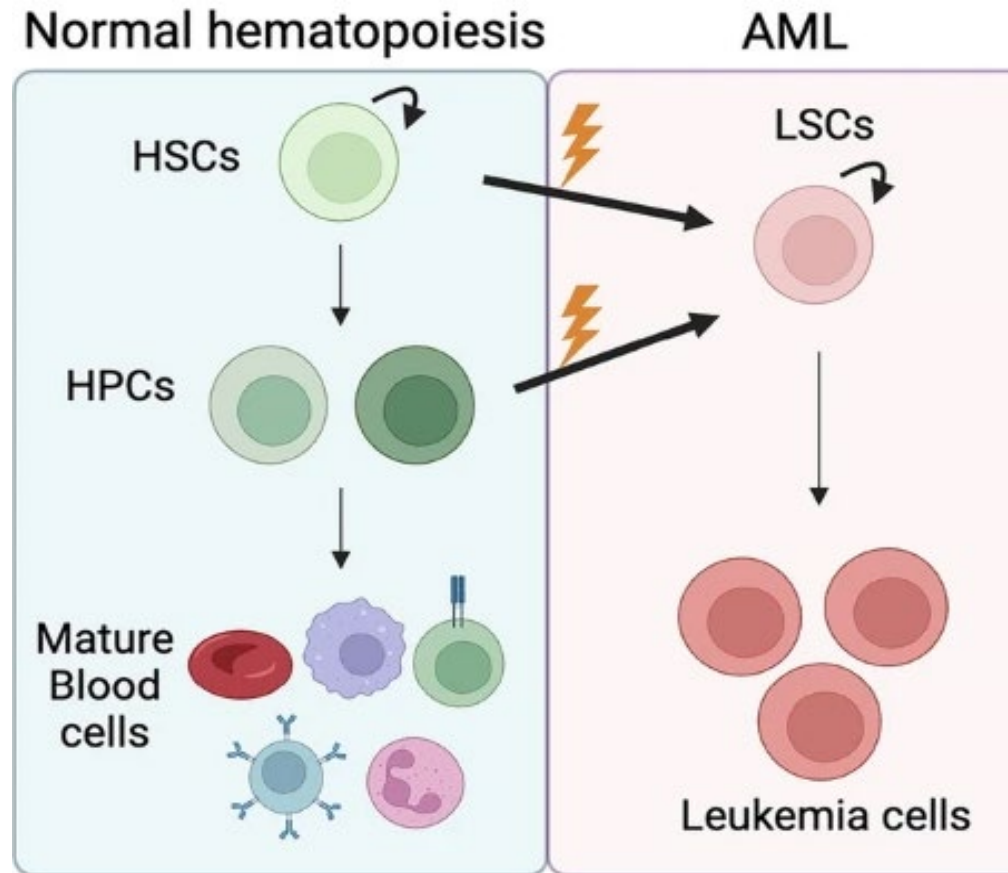
Acute myeloid leukemia (AML)

Normal hematopoiesis



Modified from Shi et al. 2024.

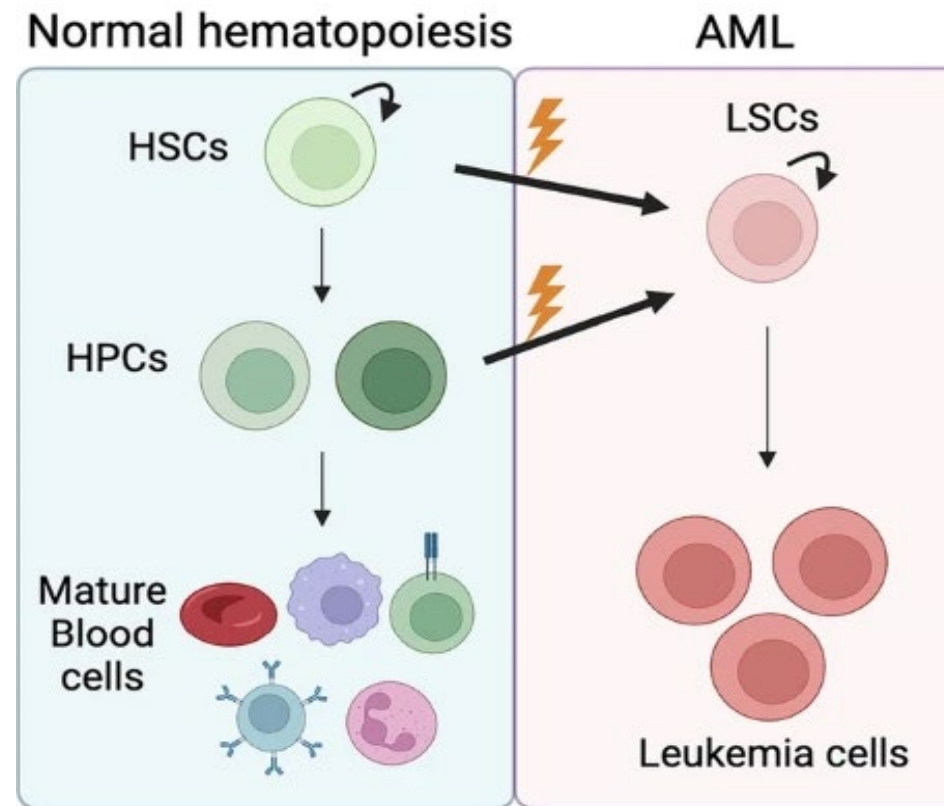
Acute myeloid leukemia (AML)



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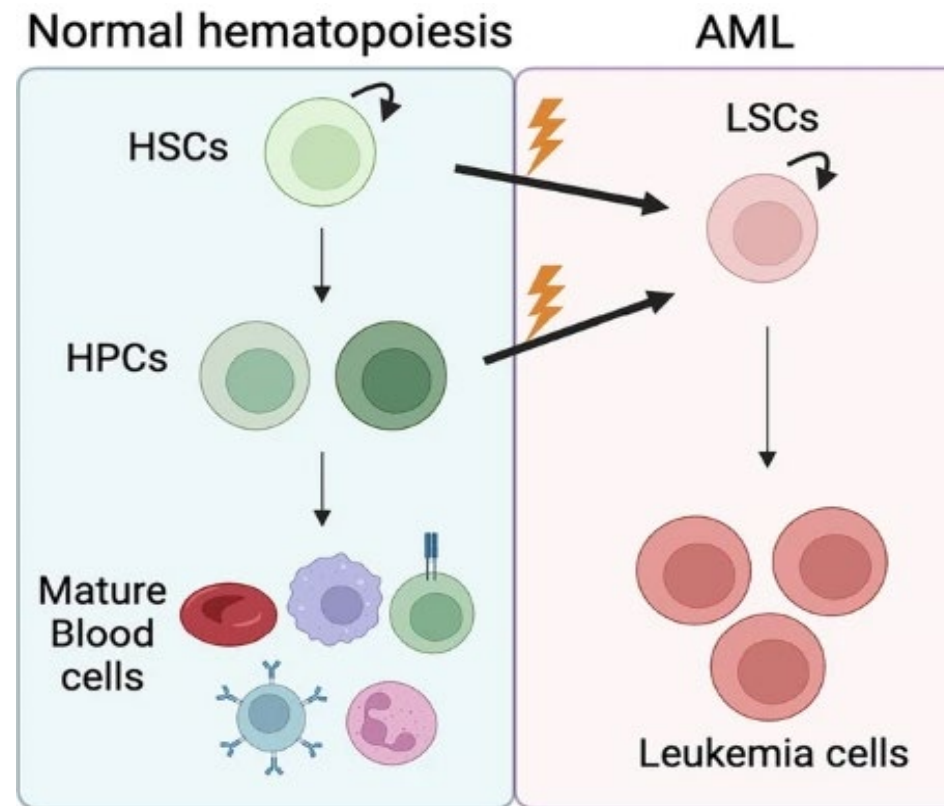
- Cancer of the blood and bone marrow (BM)



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Acute myeloid leukemia (AML)

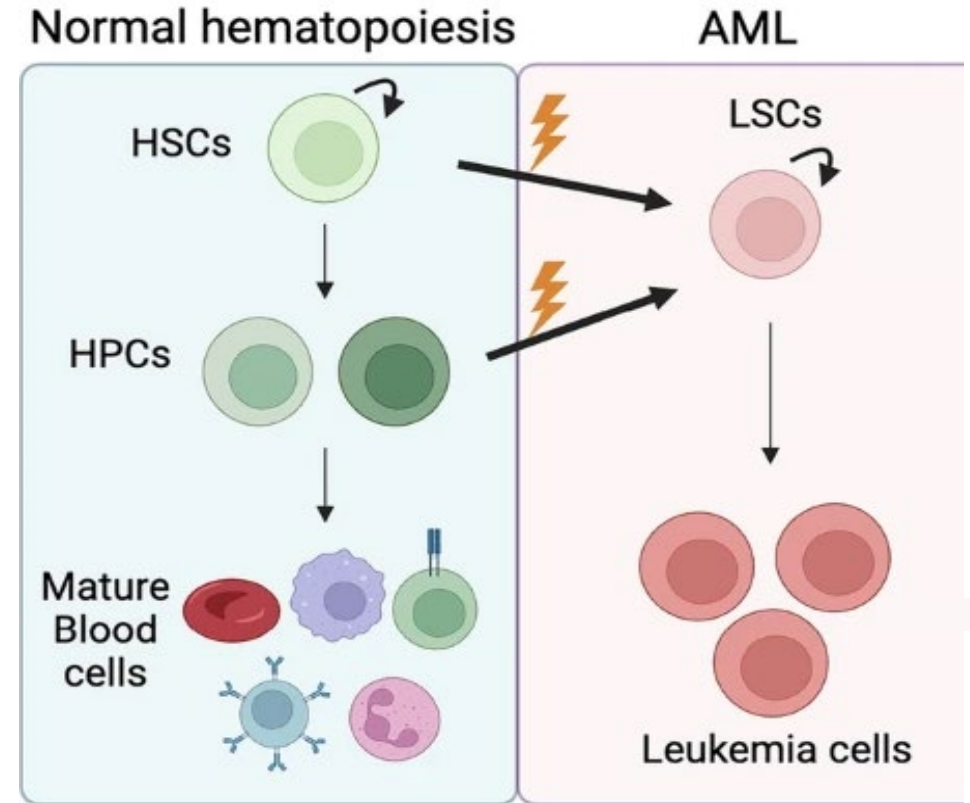
- Cancer of the blood and bone marrow (BM)
- Most common form of leukemia



Modified from Shi et al. 2024.

Acute myeloid leukemia (AML)

- Cancer of the blood and bone marrow (BM)
- Most common form of leukemia
- Treatment options
 - Standard treatment
 - “7 + 3” chemotherapy
 - Hypomethylating agents
 - Targeted treatments → Venetoclax
 - BM transplant



Modified from Shi et al. 2024.

Acute myeloid leukemia (AML)

- Cancer of the blood and bone marrow (BM)
- Most common form of leukemia
- Treatment options
 - Standard treatment
 - “7 + 3” chemotherapy
 - Hypomethylating agents
 - Targeted treatments → Venetoclax
 - BM transplant
- The European Leukemia Net (ELN) 2022

NPM1 mutation as critical marker for risk stratification and guiding treatment decisions.

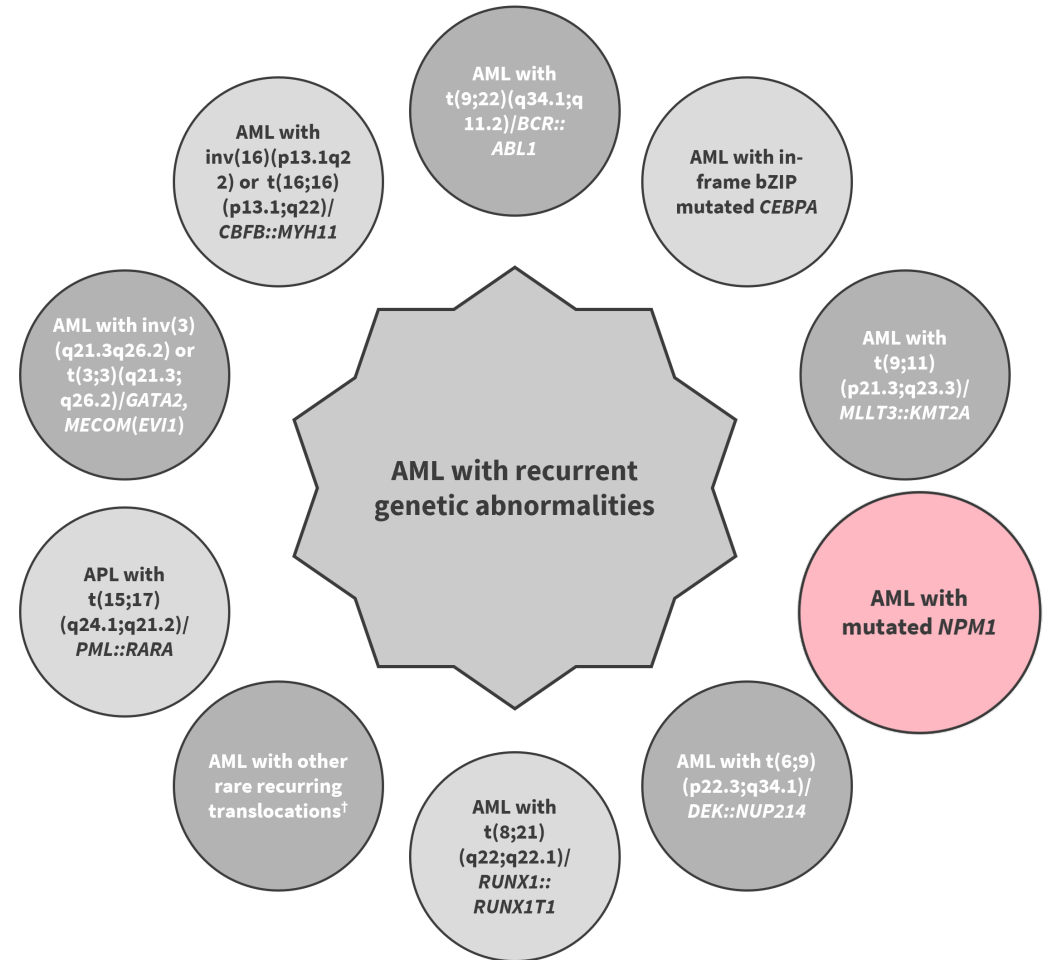
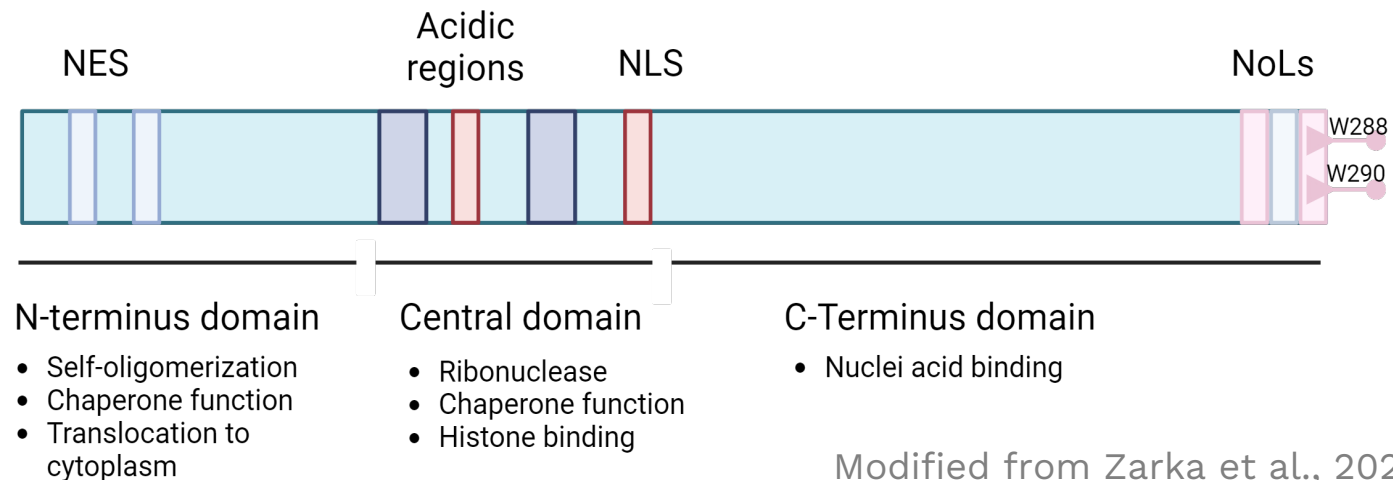


Image Source: AML Hub, 2022.
Retrieved from [AML Hub](#).

Nucleophosmin 1 (NPM1)

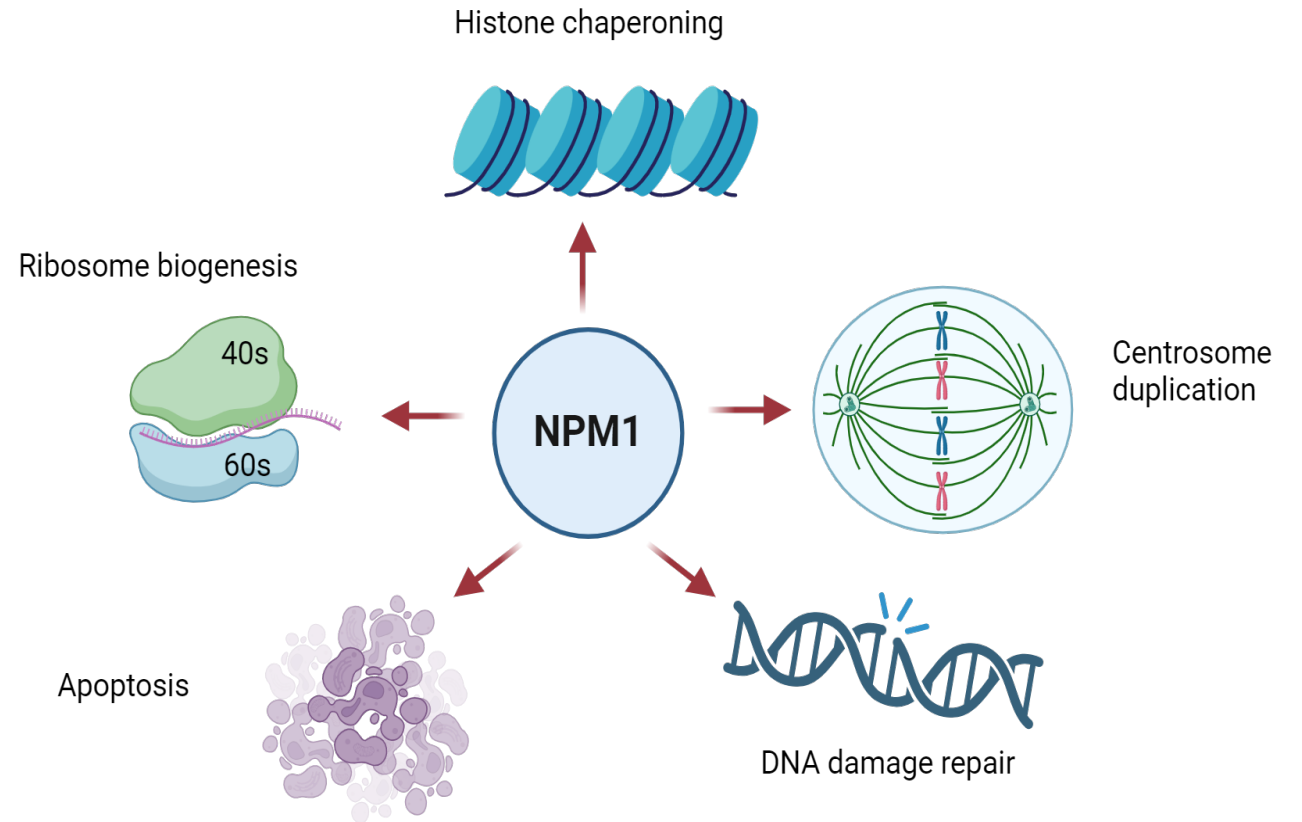
- Nucleus-cytoplasmic shuttling protein
 - Nucleolar localization



Modified from Zarka et al., 2020.
Created with BioRender

Nucleophosmin 1 (NPM1)

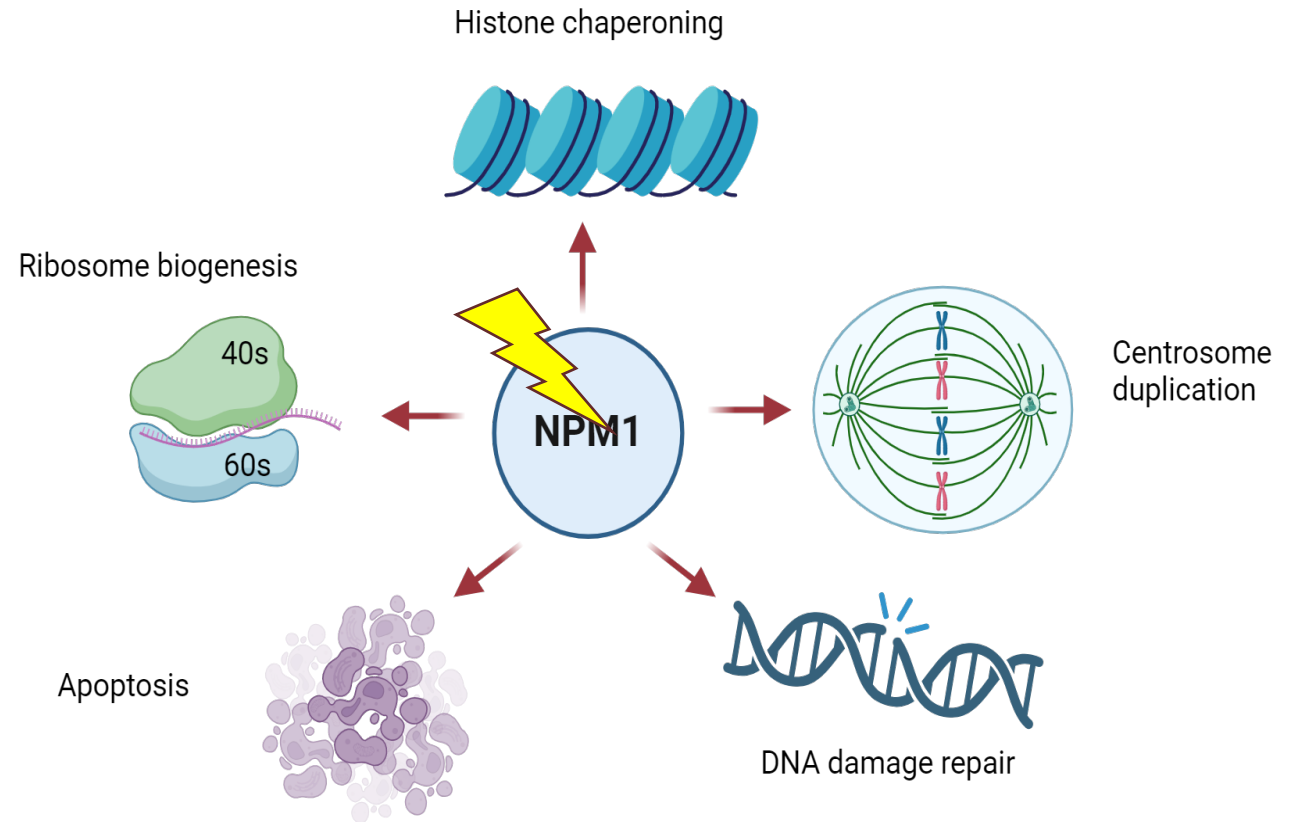
- Nucleus-cytoplasmic shuttling protein
 - Nucleolar localization
- Multifunctional protein with critical roles for normal cell proliferation and survival



Modified from Brunetti et al., 2019.
Created with BioRender

Nucleophosmin 1 (NPM1)

- Nucleus-cytoplasmic shuttling protein
 - Nucleolar localization
- Multifunctional protein with critical roles for normal cell proliferation and survival
- Dysregulation can contribute to cancer development



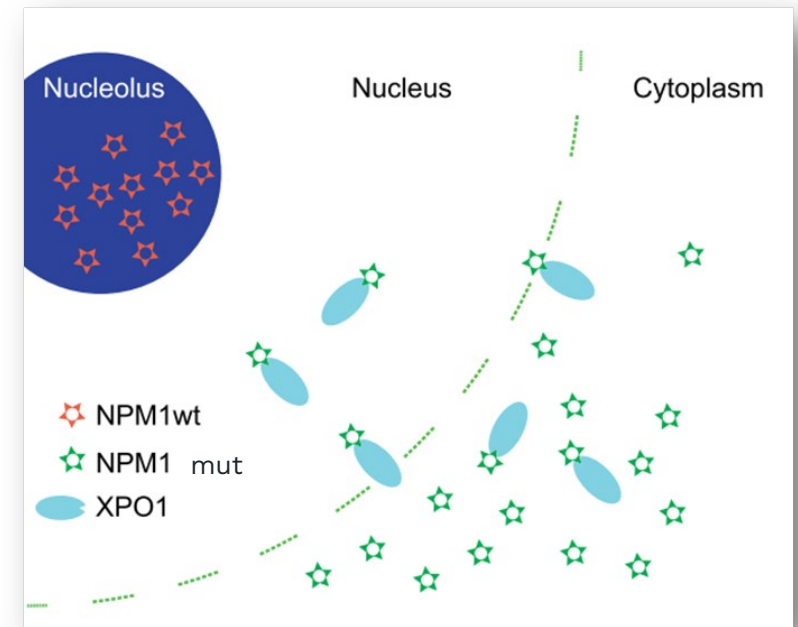
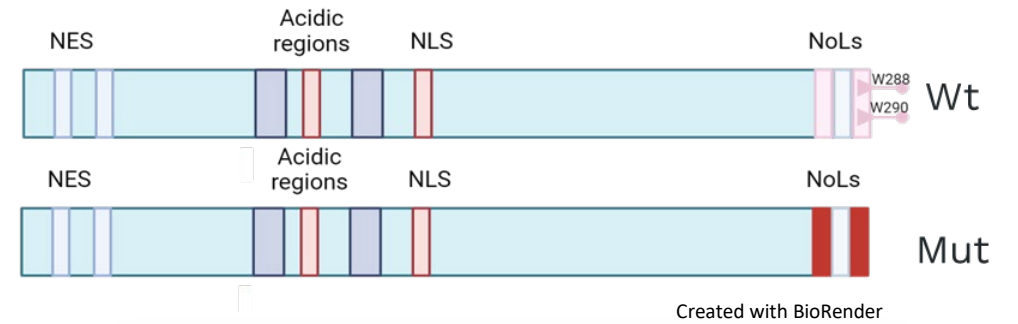
Modified from Brunetti et al., 2019.
Created with BioRender

NPM1 in AML (*NPM1* mut)

- Most common AML genetic alteration

NPM1 in AML (*NPM1* mut)

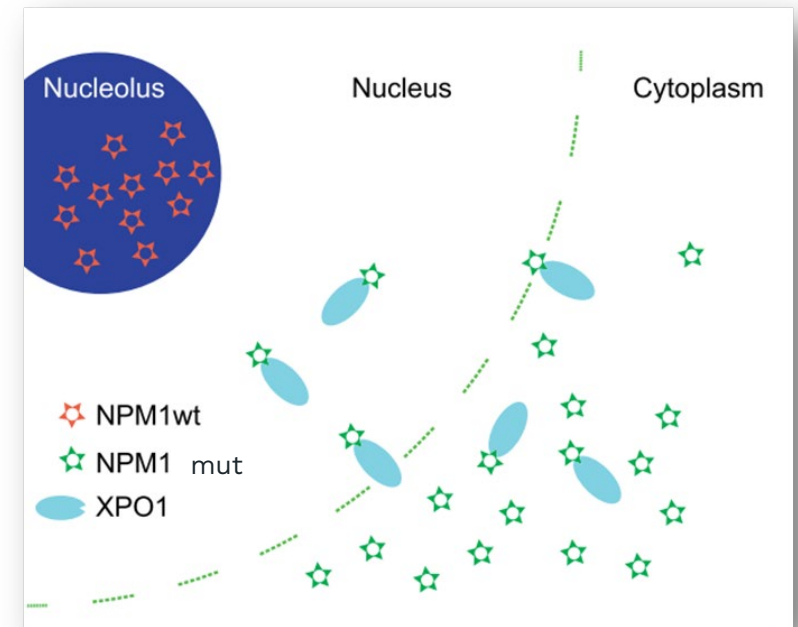
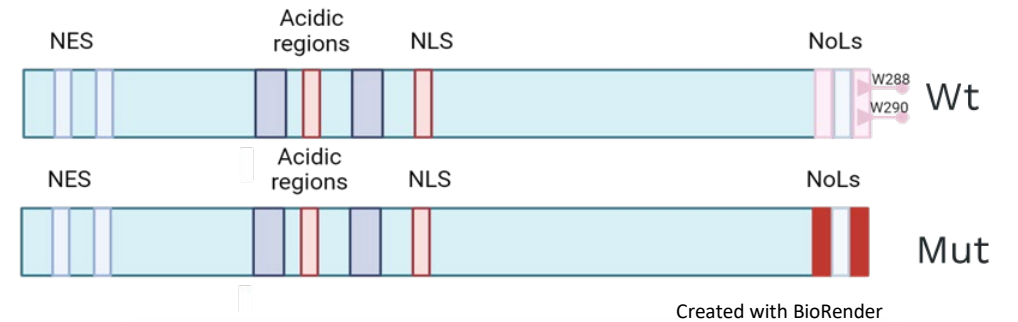
- Most common AML genetic alteration
- Frameshift mutation at the C-terminus domain



Modified from Brunetti et al., 2019.

NPM1 in AML (*NPM1* mut)

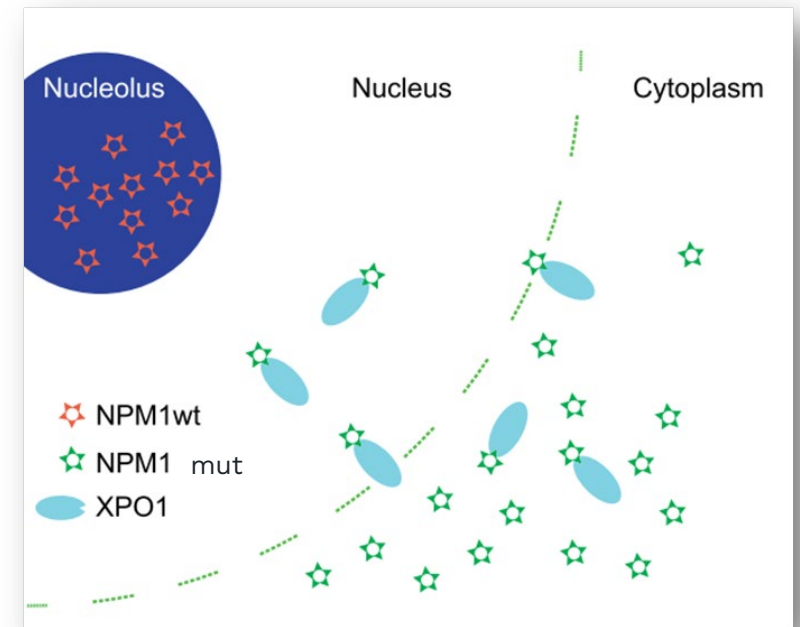
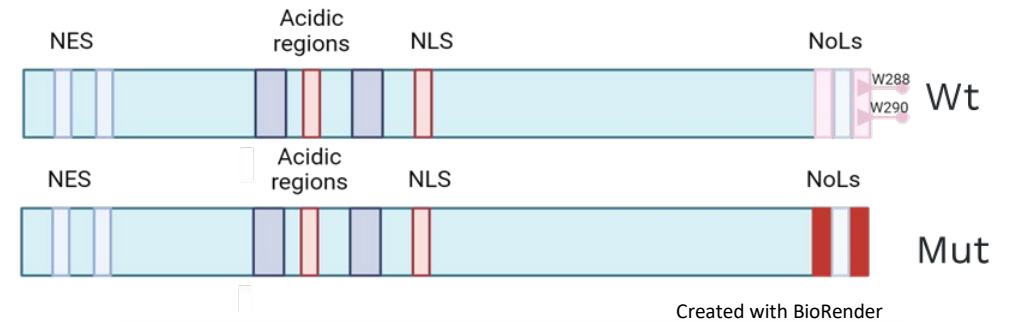
- Most common AML genetic alteration
- Frameshift mutation at the C-terminus domain
 - Aberrant cytoplasmic localization



Modified from Brunetti et al., 2019.

NPM1 in AML (*NPM1* mut)

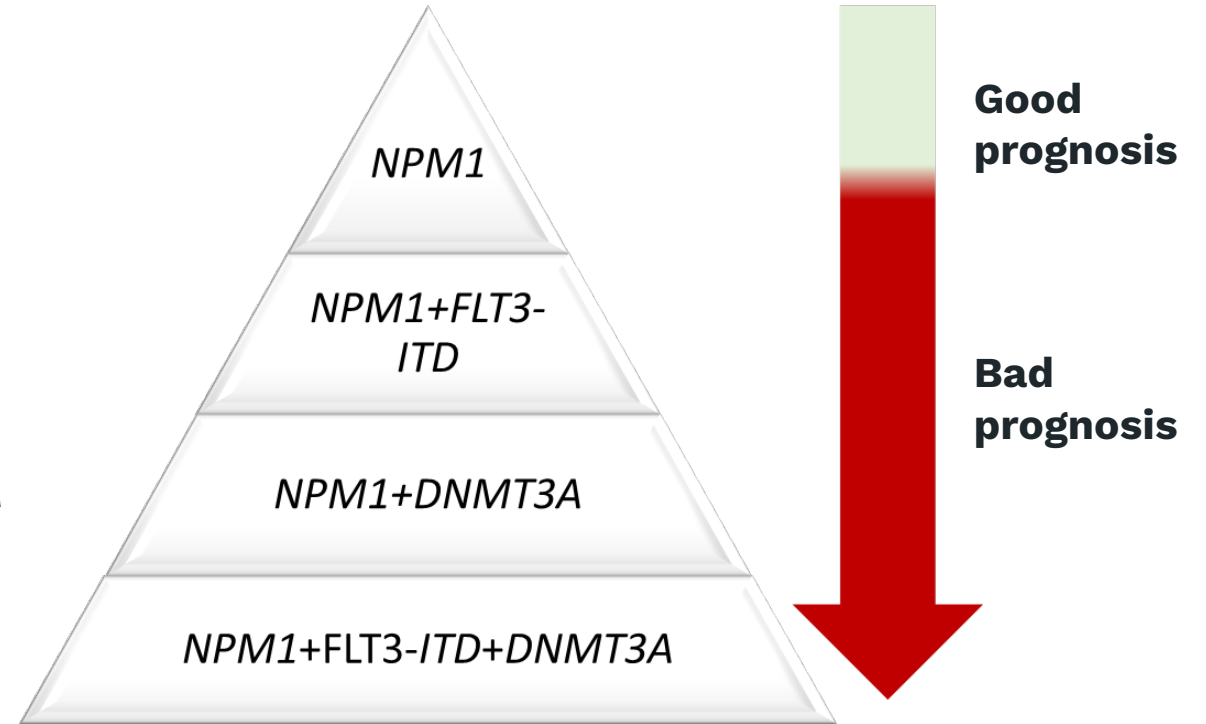
- Most common AML genetic alteration
- Frameshift mutation at the C-terminus domain
 - Aberrant cytoplasmic localization
- Associated with increased *HOX* genes expression



Modified from Brunetti et al., 2019.

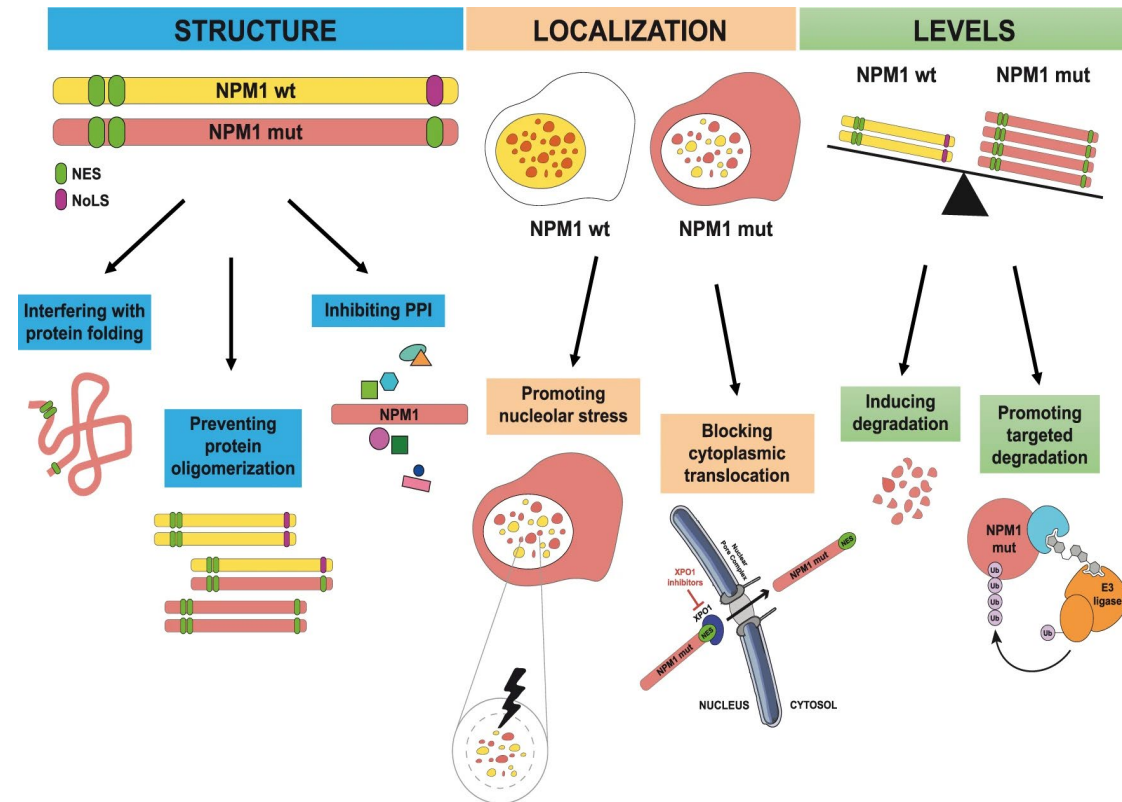
NPM1 in AML (*NPM1* mut)

- Most common AML genetic alteration
- Frameshift mutation at the C-terminus domain
 - Aberrant cytoplasmatic localization
- Associated with increased *HOX* genes expression
- Often co-mutated with *FLT3-ITD* and/or *DNMT3A*



NPM1 in AML (*NPM1* mut)

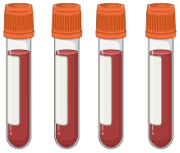
- Most common AML genetic alteration
- Frameshift mutation at the C-terminus domain:
 - Aberrant cytoplasmic localization
- Associated with increased *HOX* genes expression
- Often co-mutated with *FLT3-ITD* and/or *DNMT3A*
- Treatment:
 - Standard therapy → High relapse rates
 - The heterogeneity of *NPM1* mut confers different treatment strategies



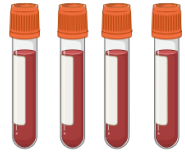
Ranieri et al., 2020.

Project background

226 *NPM1*
wildtype
samples

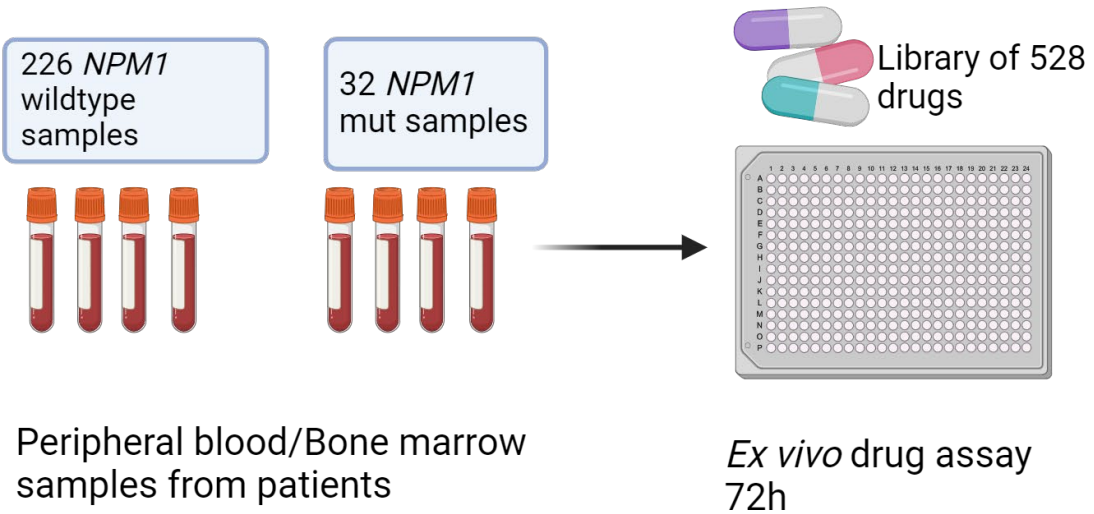


32 *NPM1*
mut samples

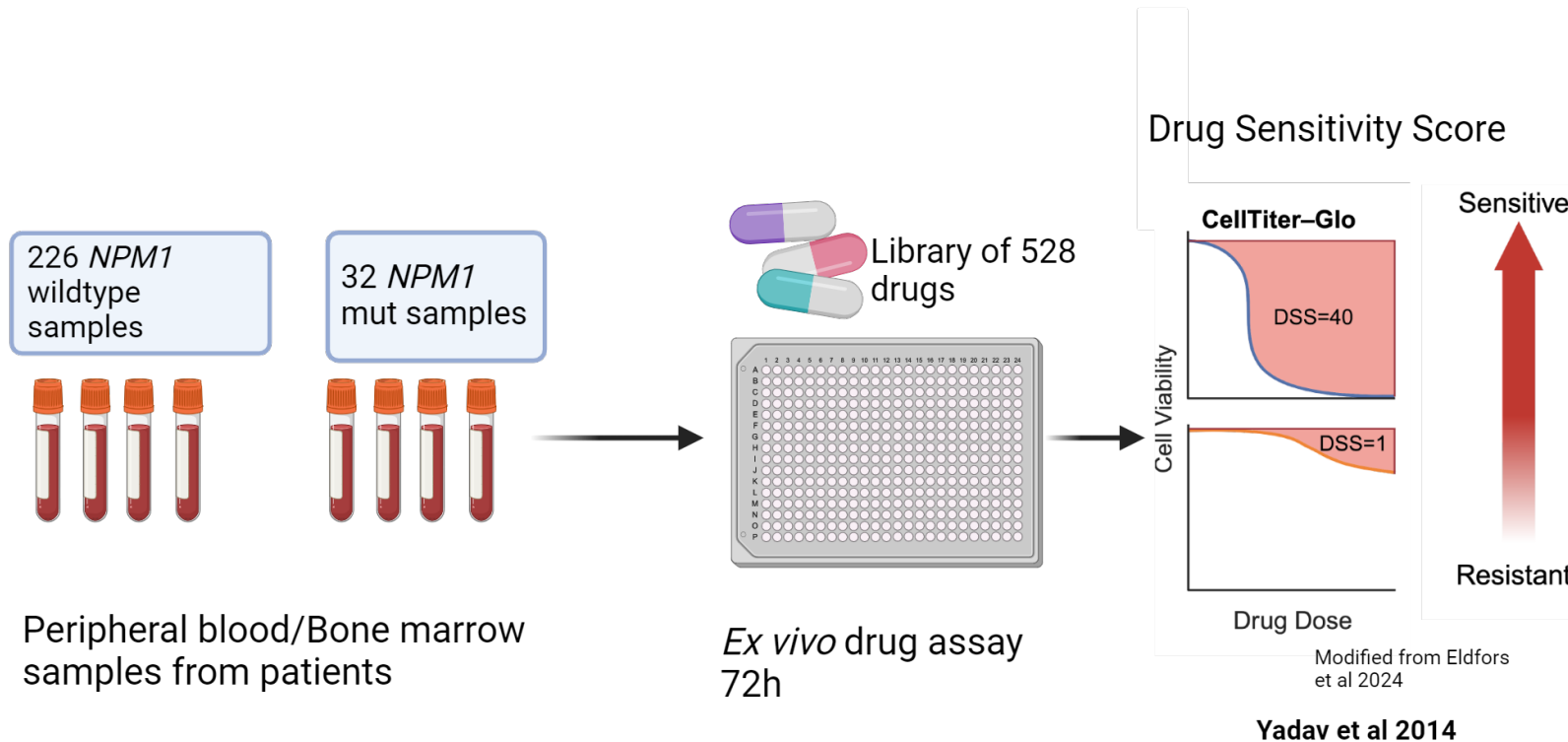


Peripheral blood/Bone marrow
samples from patients

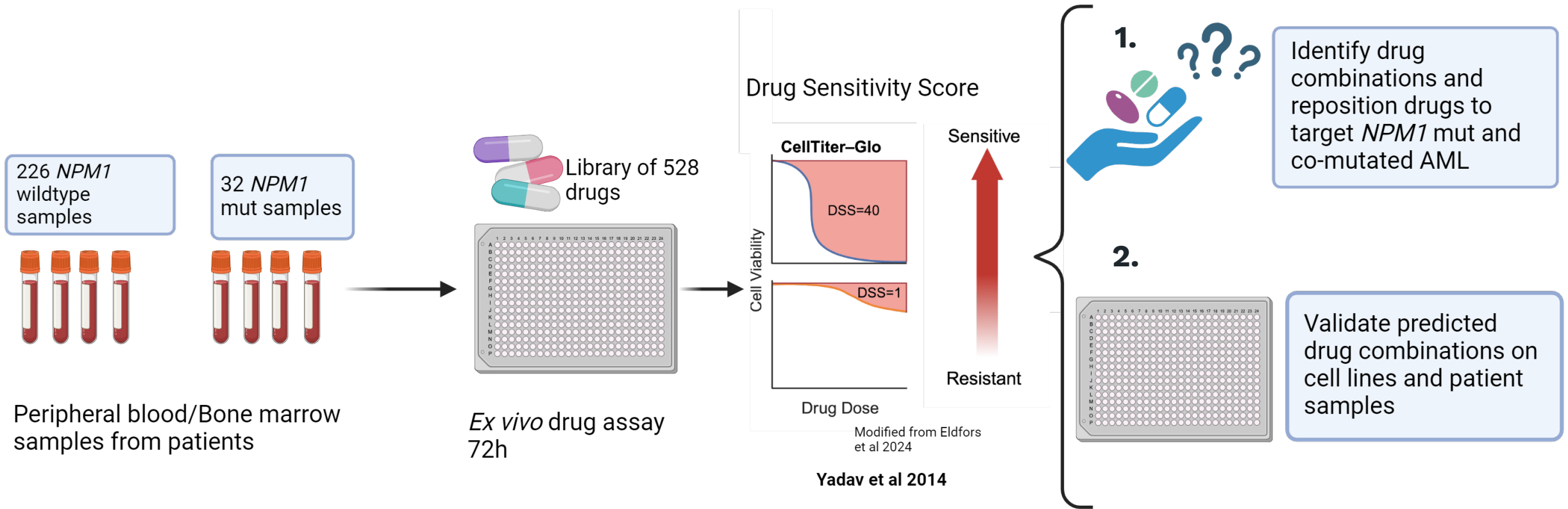
Project background



Project background

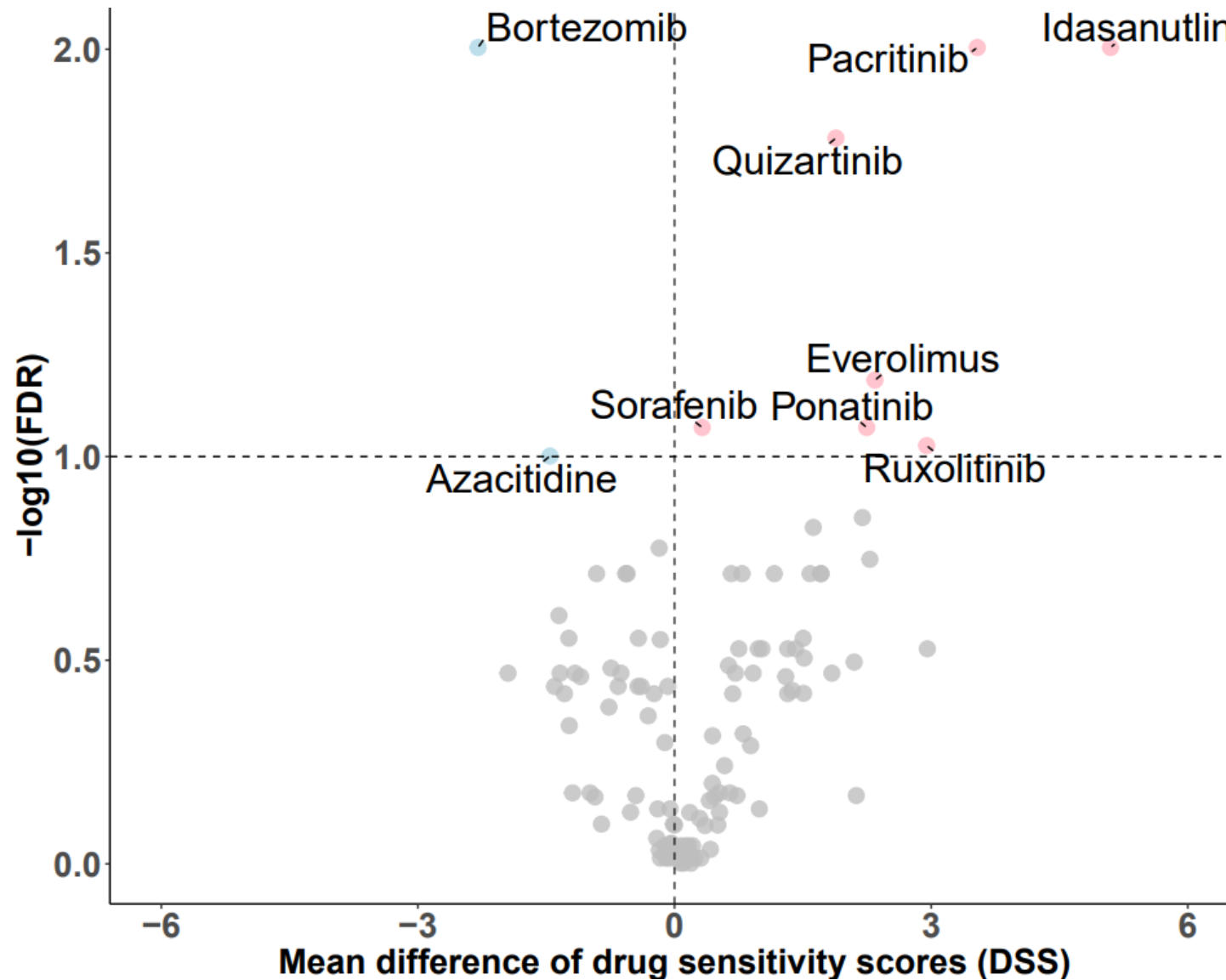


Project background



NPM1 mut samples exhibited increased sensitivity to pacritinib

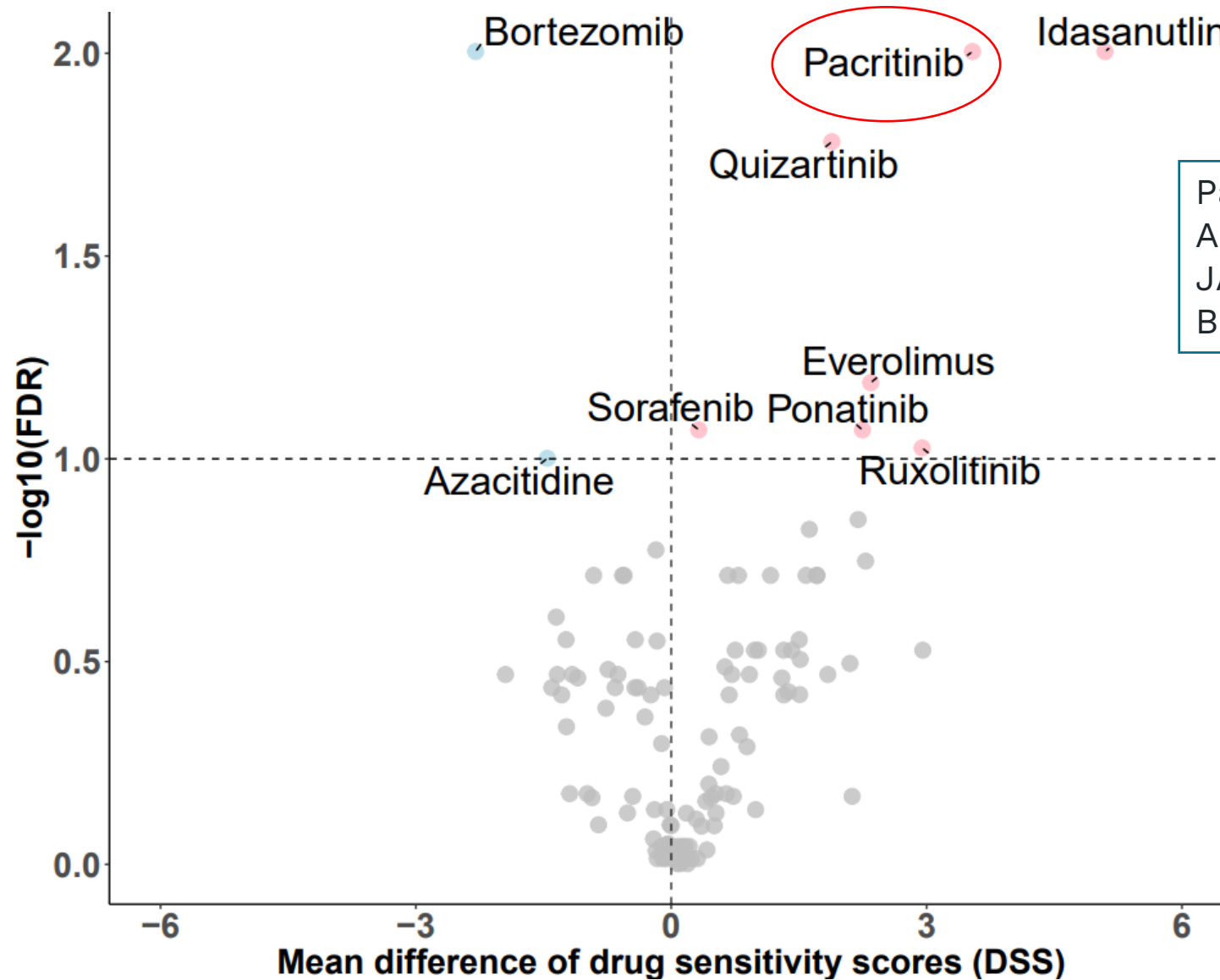
Less sensitive in *NPM1* mut samples ← → More sensitive in *NPM1* mut samples



Unpublished data. Do not post

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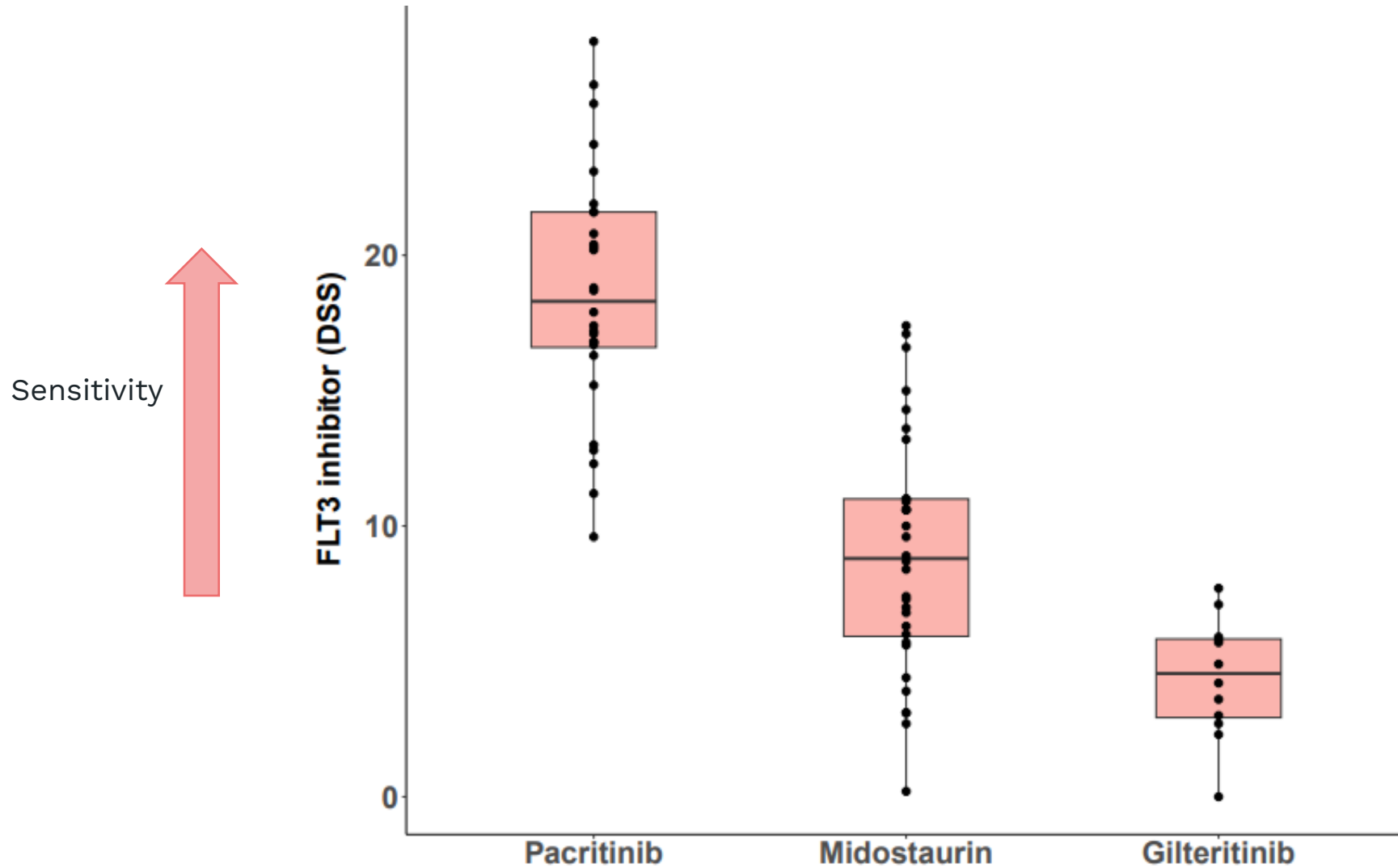
Less sensitive in *NPM1* mut samples ← → More sensitive in *NPM1* mut samples



Pacritinib:
Approved for myelofibrosis
JAK2 and FLT3 inhibitor
Background in *FLT3* mutated AML

Unpublished data. Do not post

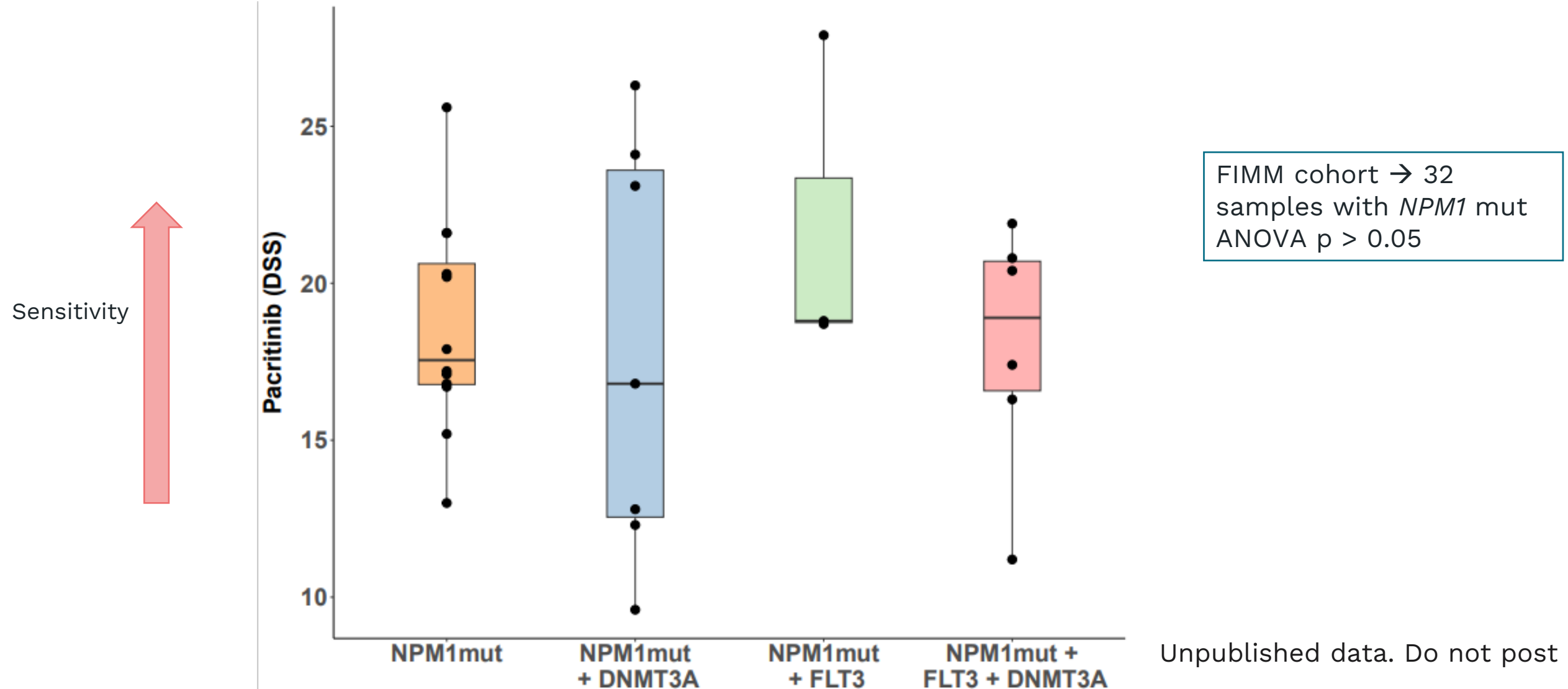
Pacritinib showed higher efficacy in *NPM1* mut samples than conventional FLT3 inhibitors



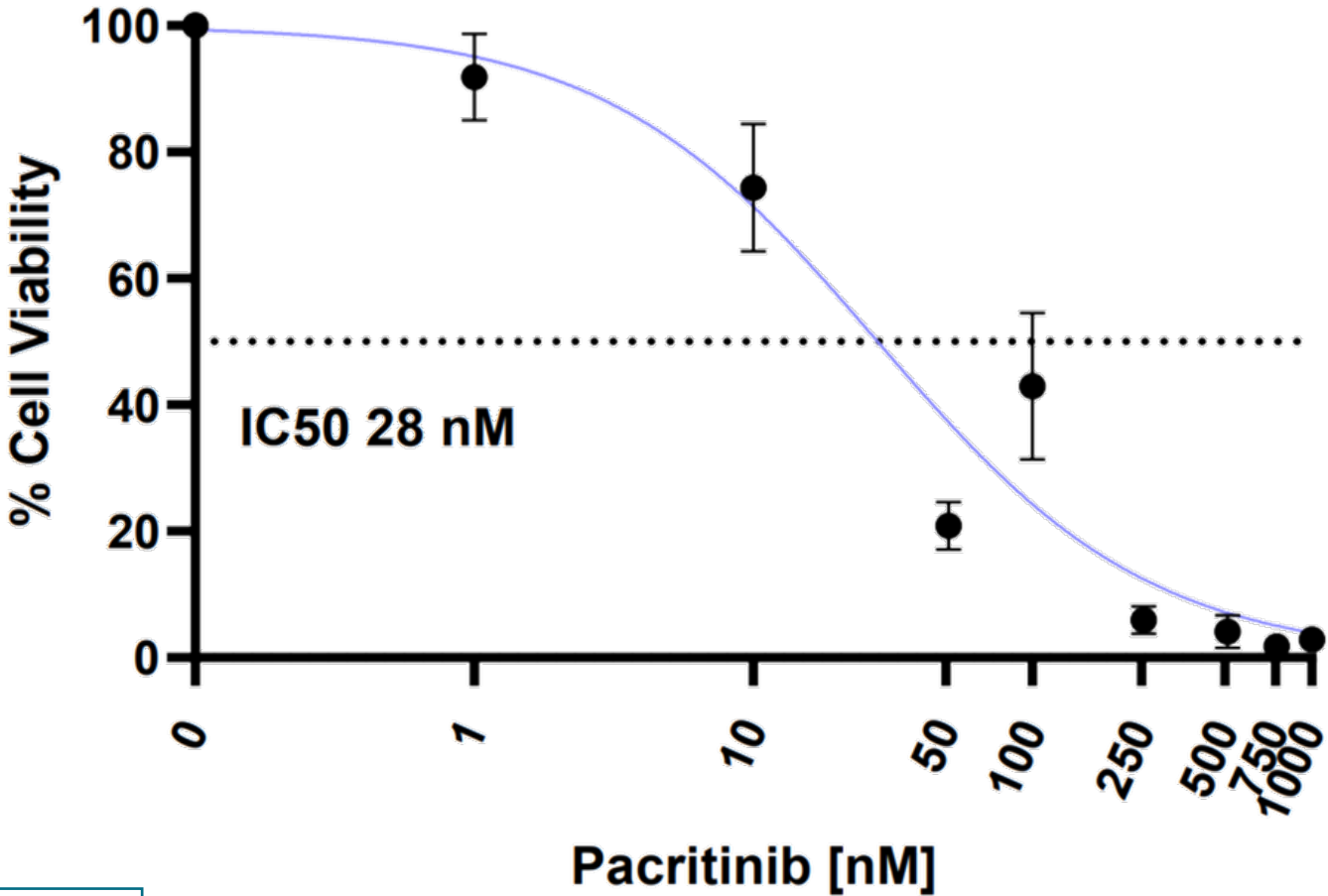
FIMM cohort → 32 samples with *NPM1* mut
ANOVA $p \leq 0.05$

Unpublished data. Do not post

The efficacy of pacritinib did not yield any significance among *NPM1* co-mutated samples.



AML *NPM1* mut+ *FLT3-ITD*+ *DNMT3A* mouse cell line is sensitive to pacritinib



N = 4



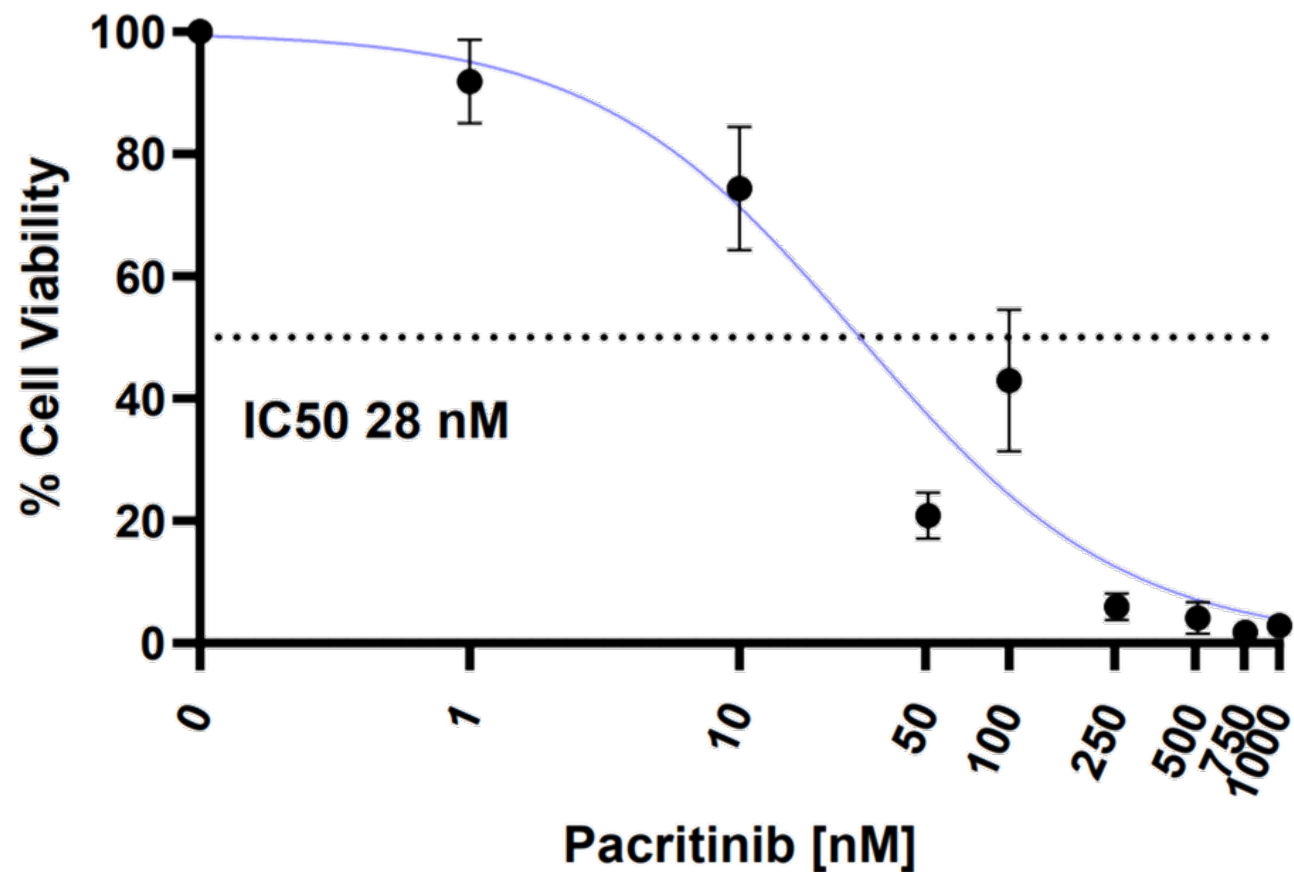
Dr. Kasper Rasmussen

AML mouse model with humanized mutations in *NPM1*, *FLT3* and *DNMT3A*.

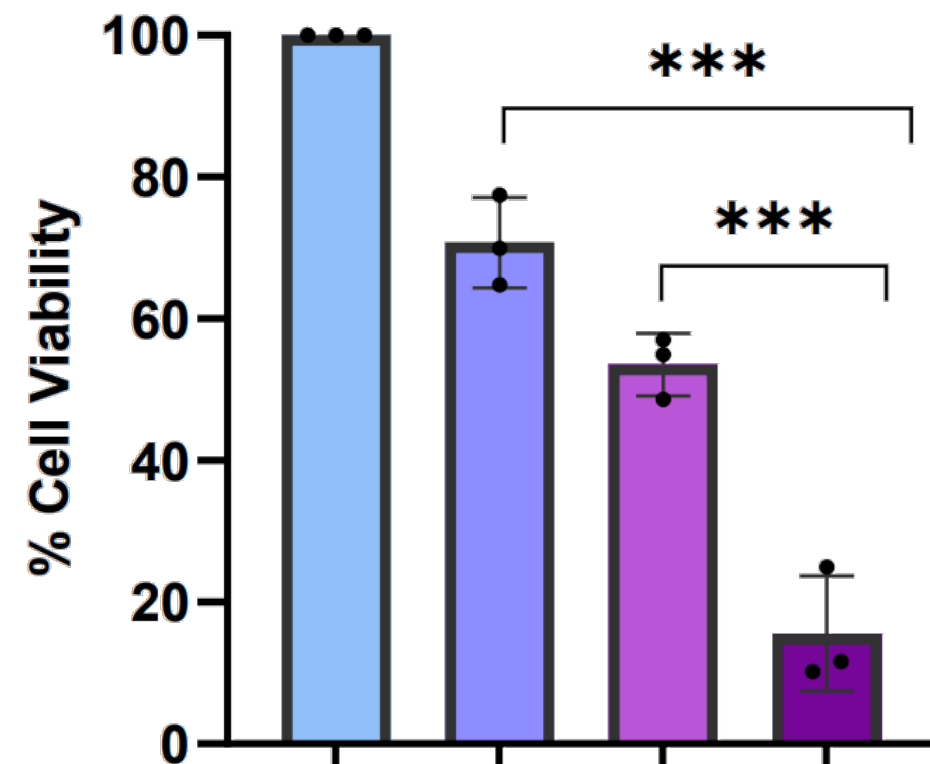
Unpublished data. Do not post

Pacritinib + venetoclax synergizes to induce cell death at low concentrations in AML

NPM1 mut + *FLT3-ITD* + *DNMT3A*



N = 4



Pacritinib [10 nM]

Venetoclax [1nM]

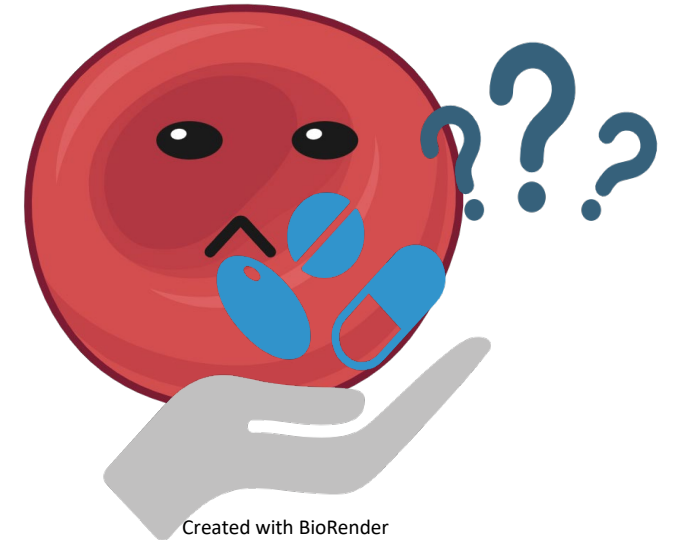
N = 3

ANOVA $p \leq 0.05$

Unpublished data. Do not post

Conclusions and future perspectives

- Pacritinib shows superior activity in *NPM1* mut AML samples compared to other FLT3 inhibitors, with efficacy independent of the co-mutations.
- It effectively induces cell death in *NPM1* triple mutant (*FLT3-ITD* + *DNMT3A*) at a relatively low doses
- The combination of pacritinib with venetoclax enhances its therapeutic effect
- Currently validating the results on *NPM1* mut + *FLT3-ITD*
- Further validation on patients' samples



Acknowledgements

Translational Research and
Personalized Medicine group:
Dr. Caroline Heckman
Dr. Ella Sinervuori
Dr. Mahesh Tambe
Institute for Molecular Medicine
Finland, FIMM. HiLIFE, University
of Helsinki, Finland.

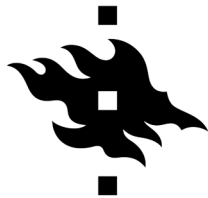


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Research Council of Finland

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