

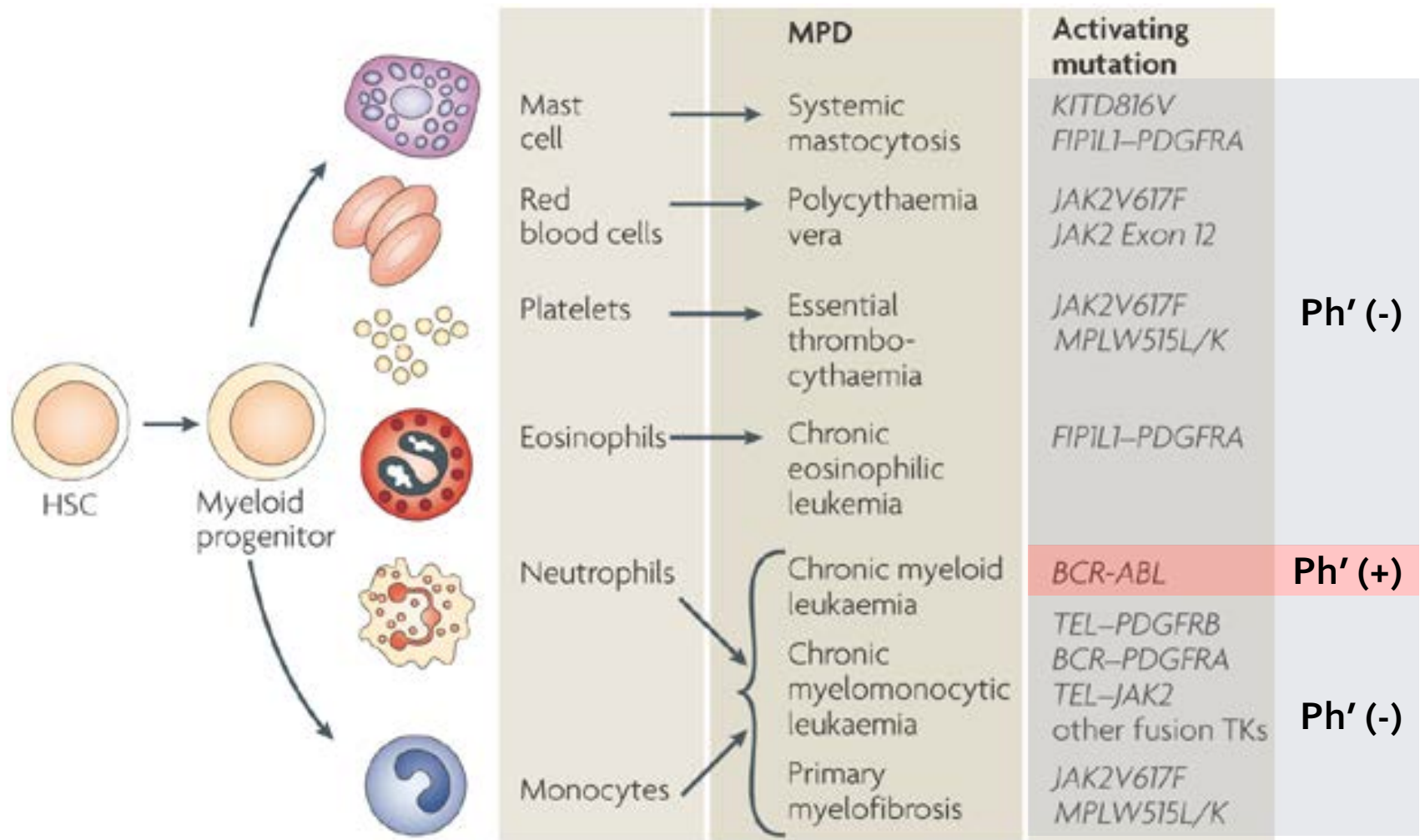
***MULTIPLEX DETECTION OF
JAK₂ AND MPL MUTATIONS
IN MYELOPROLIFERATIVE NEOPLASMS***

MILENA IVANOVA, PhD

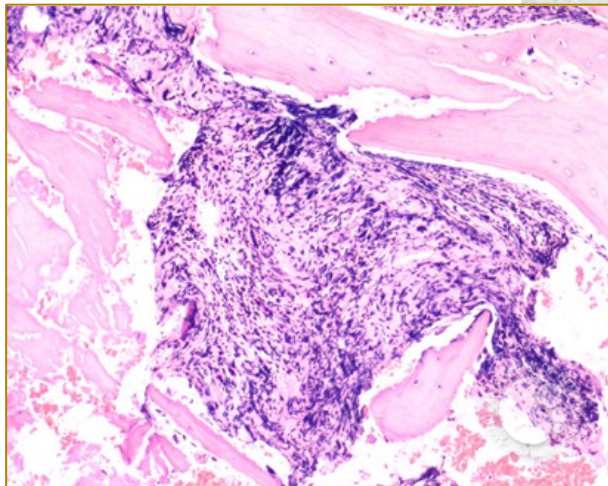
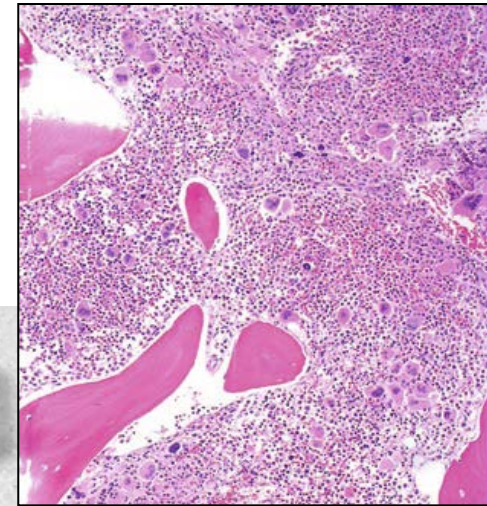
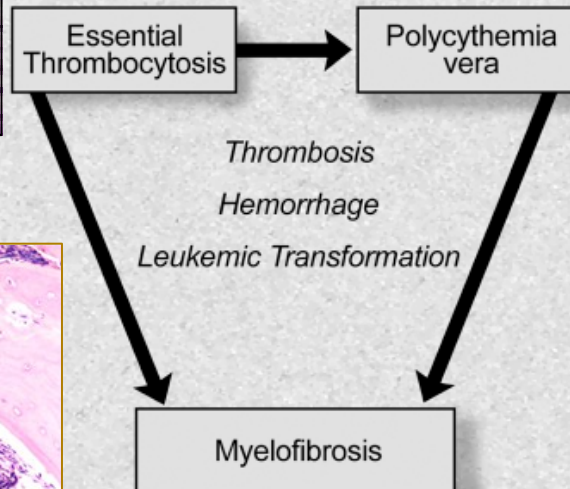
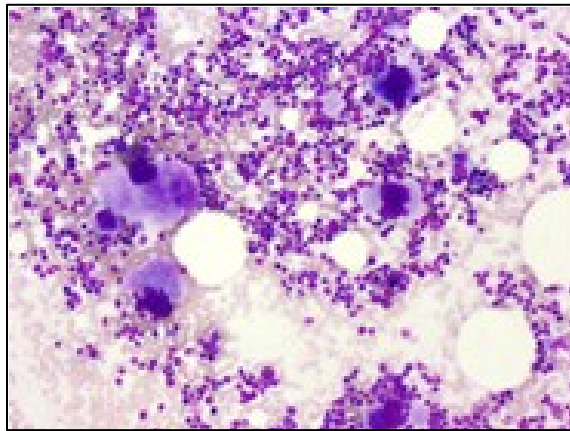
Department of Clinical Immunology

University Hospital Alexandrovska, Sofia, Bulgaria

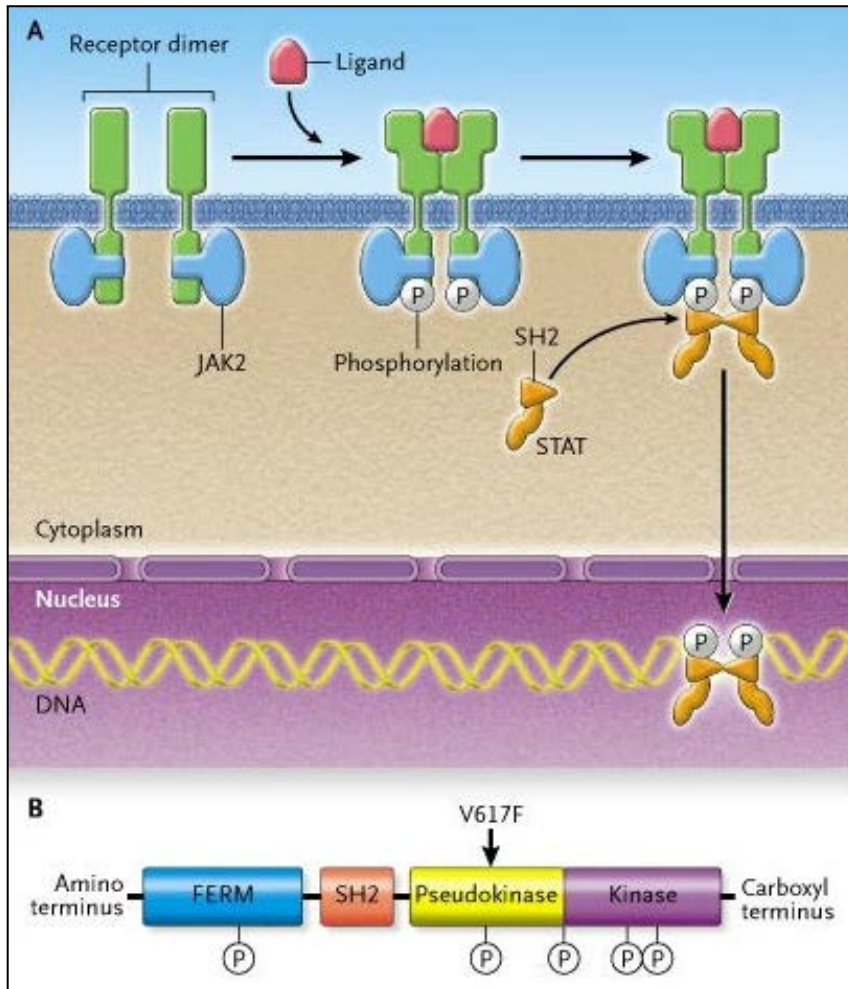
MYELOPROLIFERATIVE NEOPLASMS (MPNs) - DEFINITION



THE CLASSICAL VIEW OF Ph'(-) MPNS

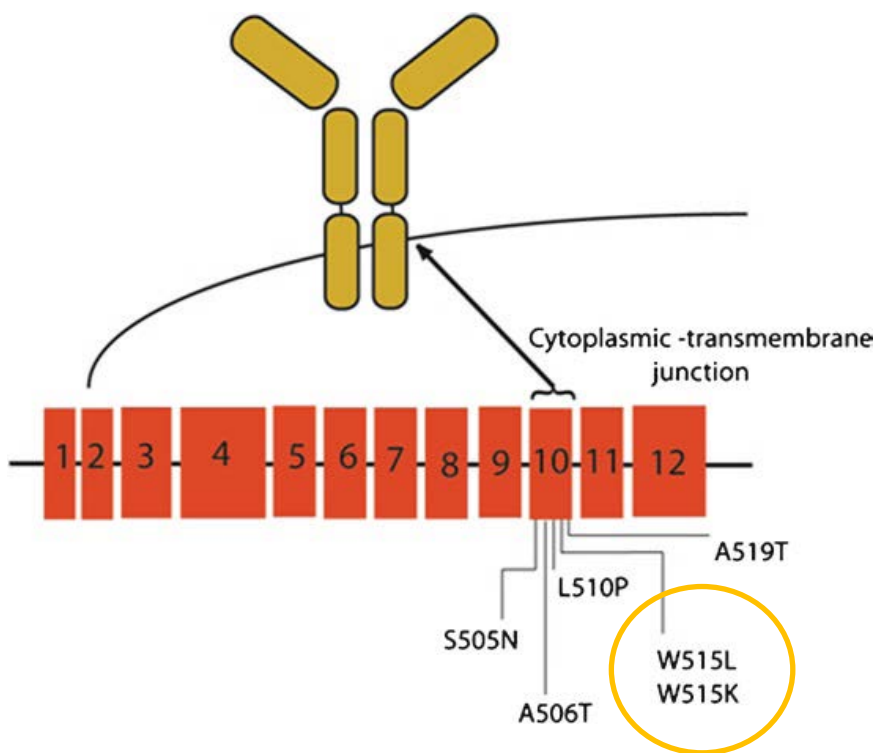


JAK2 V617F MUTATION FOR ALL Ph' (-) MPNS



<i>Disease entity</i>	<i>Frequency (%)</i>
Polycythemia vera	81–99
Essential thrombocythemia	41–72
Primary myelofibrosis	39–57

MPL EXON 10 MUTATIONS IN CONGENITAL THROMBOCYTOSIS, ET AND PMF



<i>Disease entity</i>	<i>Frequency (%)</i>
Polycythemia vera	0
Essential thrombocythemia	3–5
Primary myelofibrosis	8–10

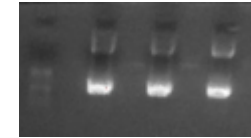
THE MISSION IMPOSSIBLE?

The MUSTs for the best method to detect the most common MPN mutations:

- ✓ Diagnose the vast majority of Ph⁺(-) MPN cases
- ✓ Be easy, rapid and reproducible
- ✓ Be sensitive and quantitative
- ✓ Allows multiplexing
- ✓ Be open for further development
- ✓ Be affordable

JAK2 V617F ANALYSIS BY PCR-SSOP USING LUMINEX® xMAP™ TECHNOLOGY

PCR of target sequences (345bp of exon 14) using 5' biotinylated primers



Hybridization with specific LNA modified capture probes conjugated via carbodiimide coupling to carboxylated beads

JAK2 wt: 5'-CTCCACAGACACATACTCC
JAK2 V617F 5'-CTCCACAGAAACATACTCC

Washing

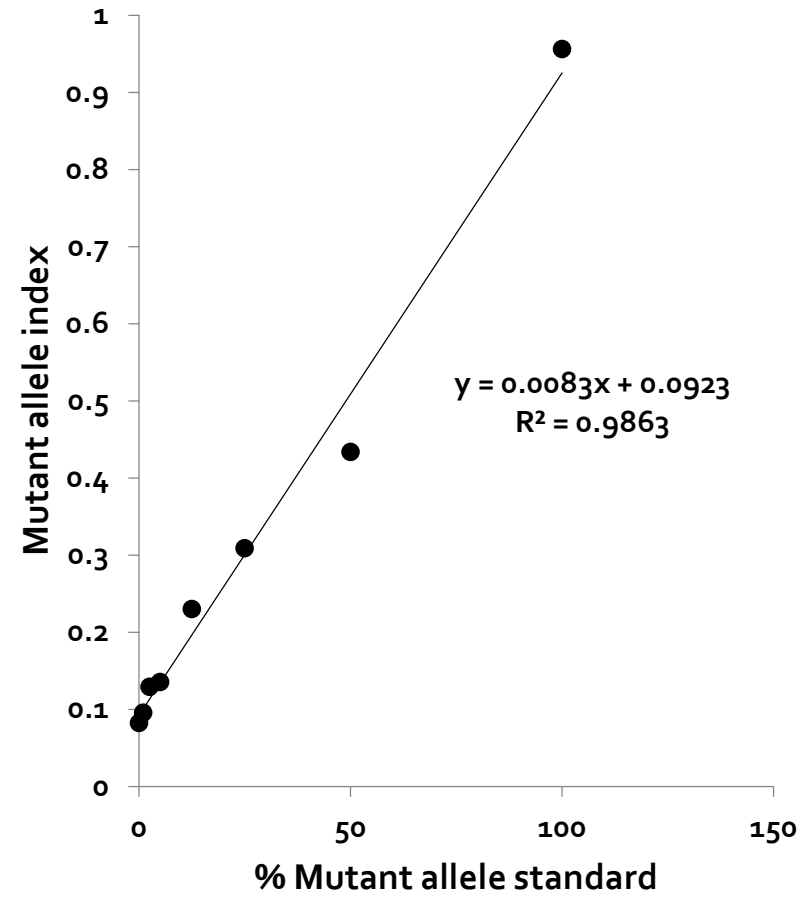
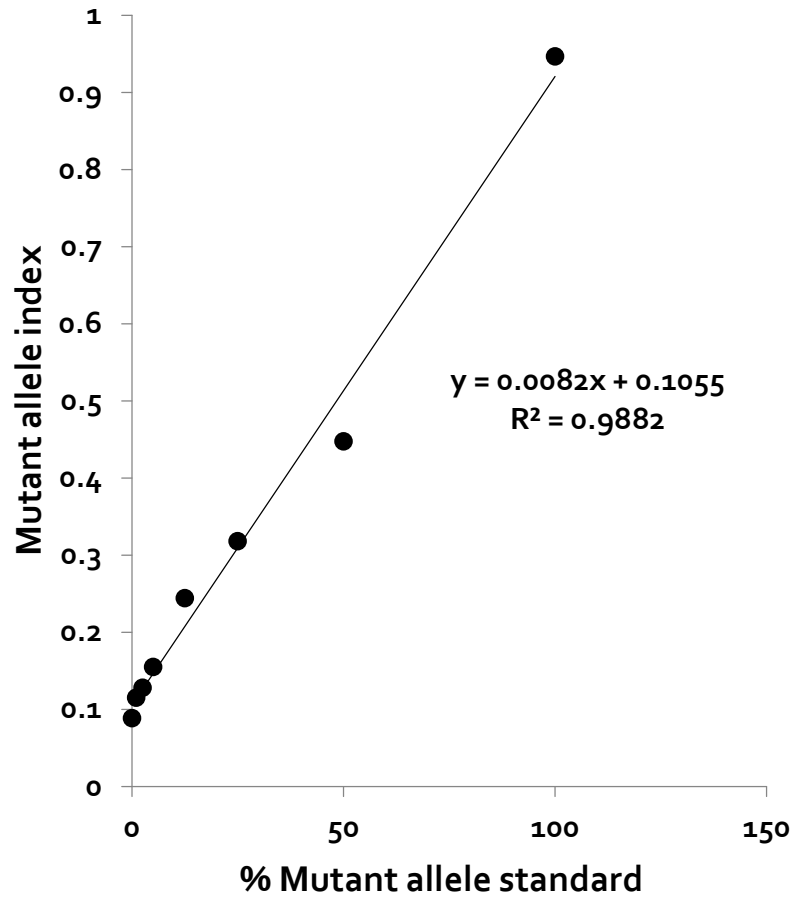


Data acquisition

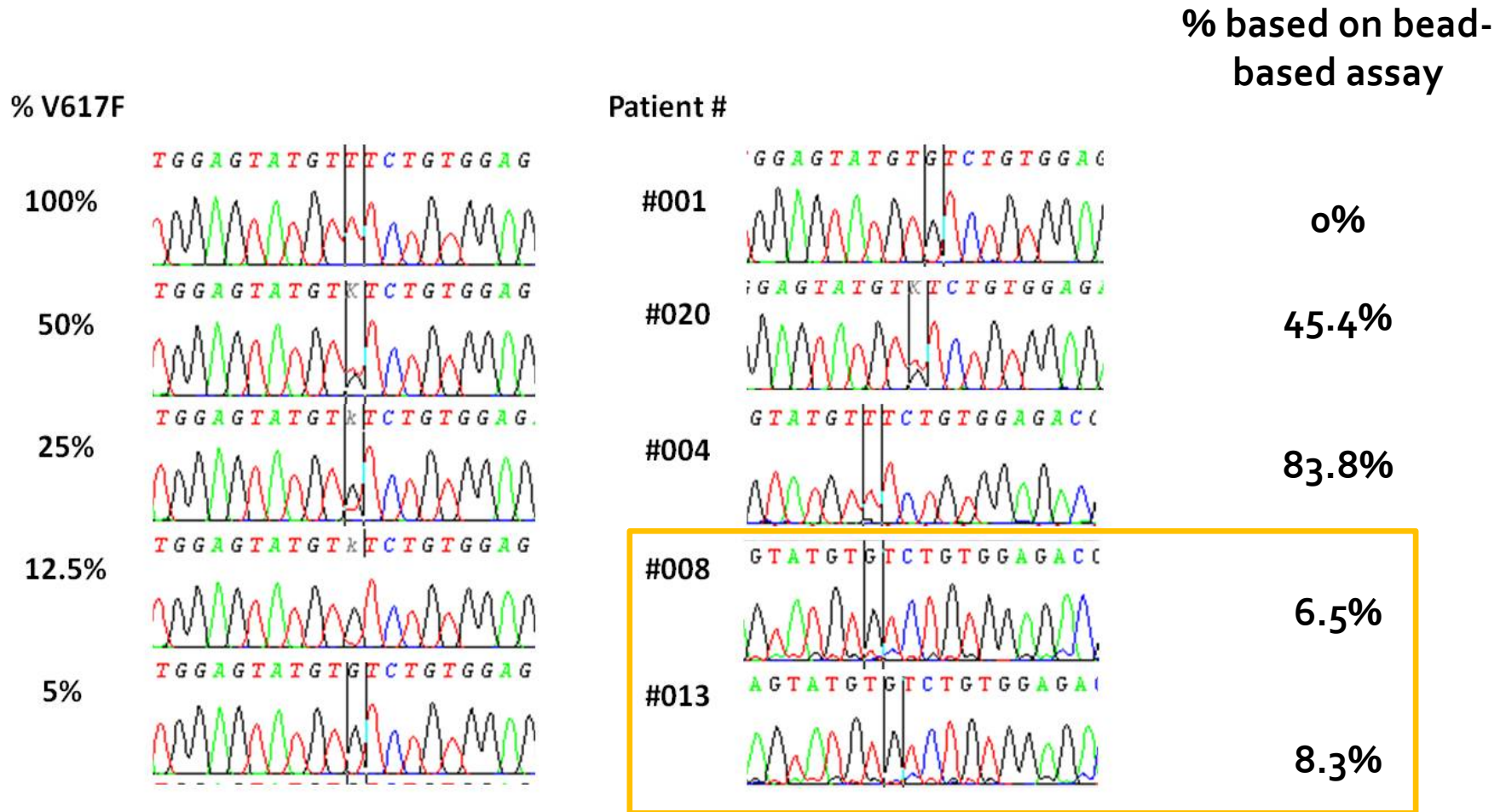
Background correction of MFI
Calculation of mutant/wild type allele index
Index (mutant allele) = [MFI (mutant allele)]/[MFI (mutant allele) + MFI (wt allele)]

Data analysis

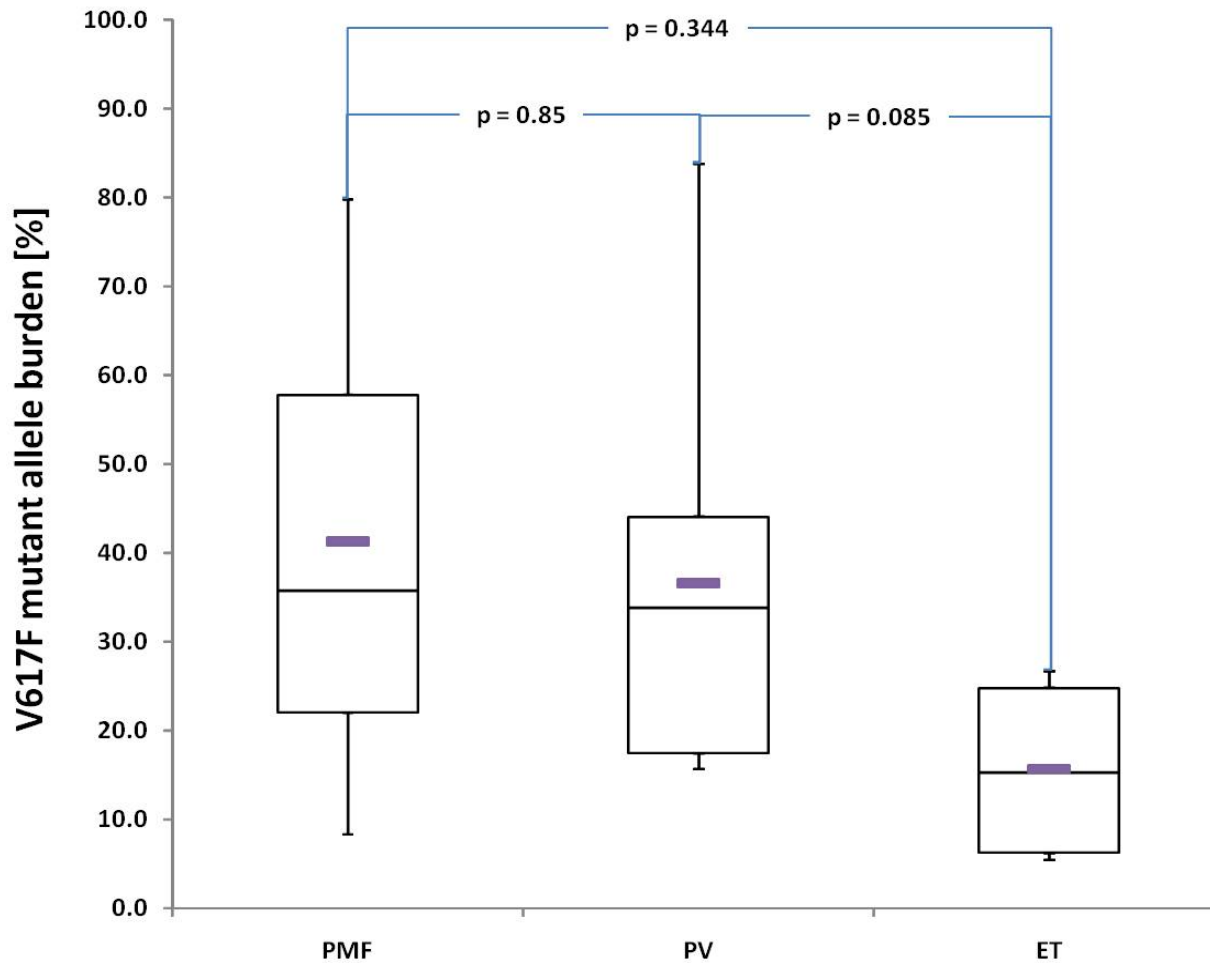
REPRODUCIBLE QUANTITATION OF V6₁₇F ALLELE ON PLASMID STANDARDS AT 1% SENSITIVITY LEVEL



BETTER SENSITIVITY ON CLINICAL SAMPLES THAN DIRECT DNA SEQUENCING

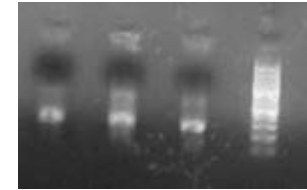


QUANTIFICATION OF THE V617F ALLELE BURDEN IN MPN PATIENTS



MPL EXON 10 MUTATIONS ANALYSIS BY PCR-SSOP USING LUMINEX® XMAP™ TECHNOLOGY

PCR of target sequences (exon 10)
using 5' biotinylated primers



Hybridization with specific LNA modified
capture probes conjugated via carbodiimide
coupling to carboxylated beads

MPL wt: 5'-GAAACTG**CC**ACTCAGCA
W₅₁₅A 5'-GAAACTG**CG**CCTCAGCA
W₅₁₅K 5'-GAAACTG**CTT**CCTCAGCAG
W₅₁₅L 5'-GAAACTG**CA**ACTCAGCAG
W₅₁₅R 5'-GAAACTG**CCT**CCTCAGCAG

Washing

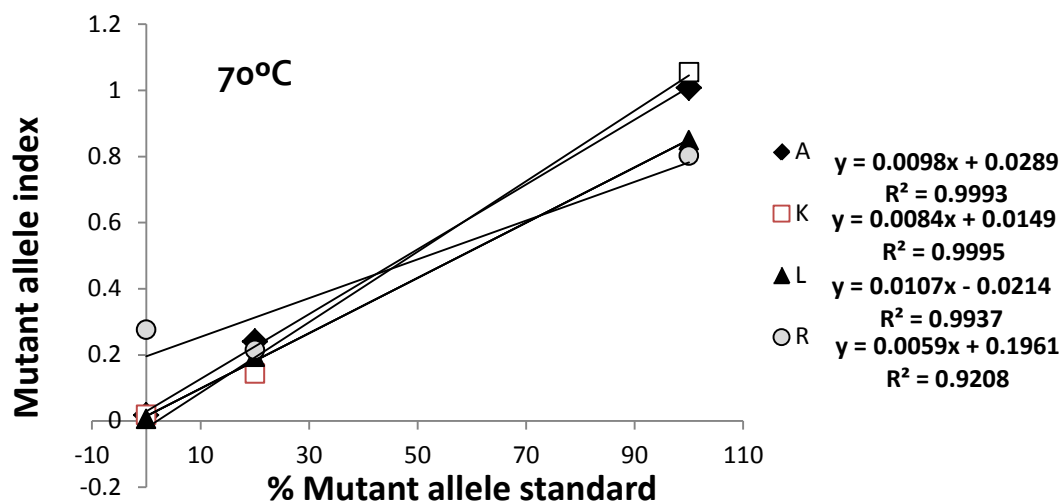
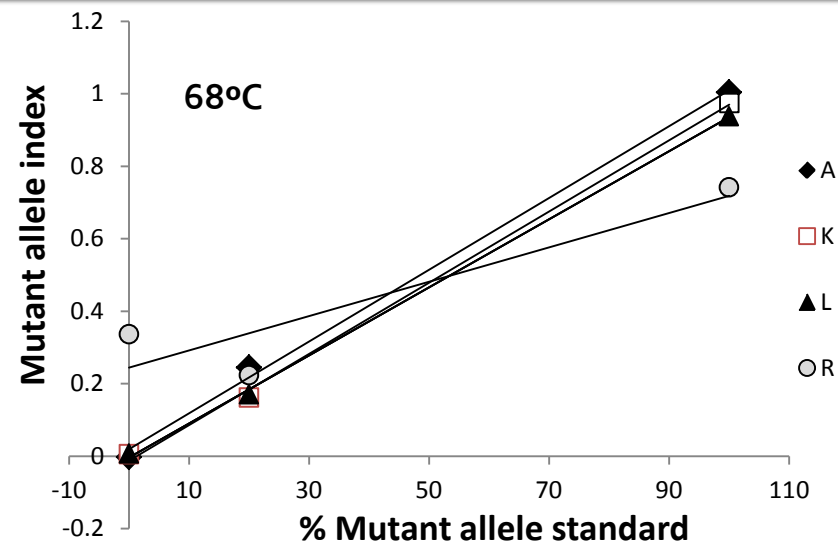
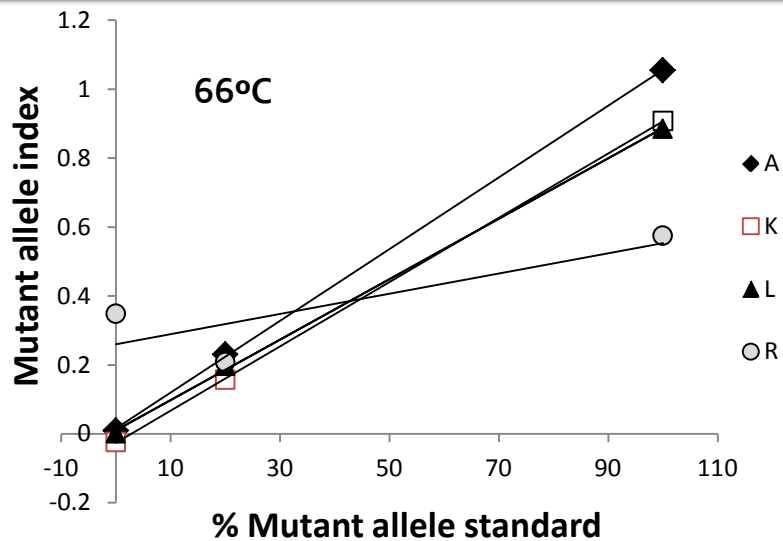


Data acquisition

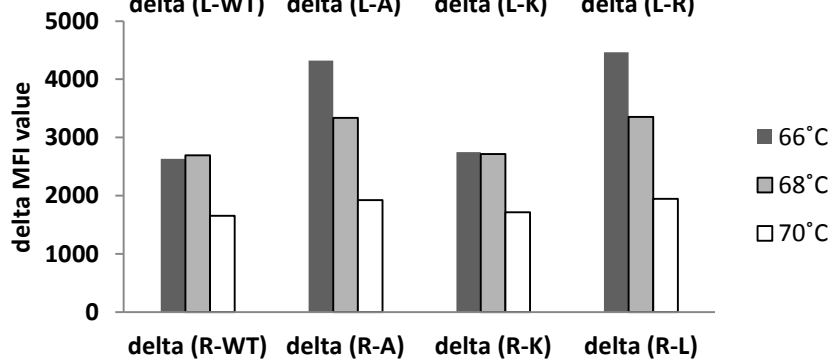
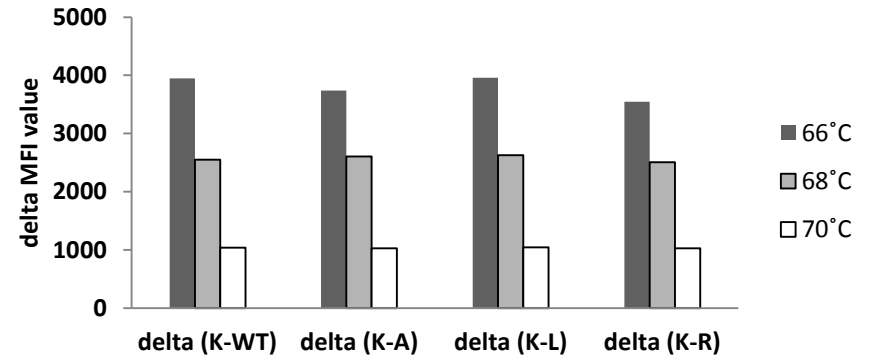
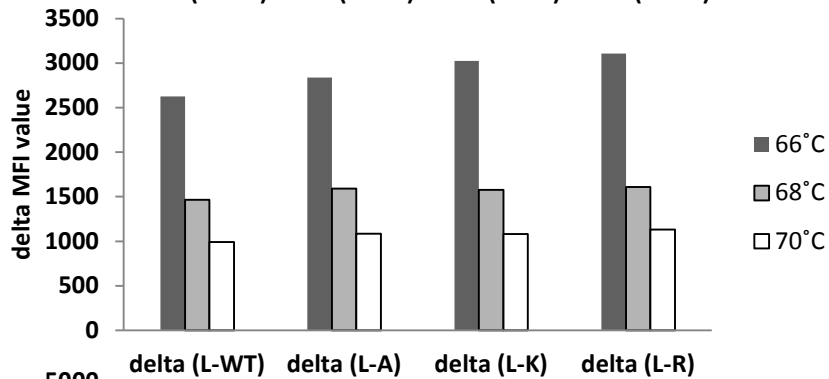
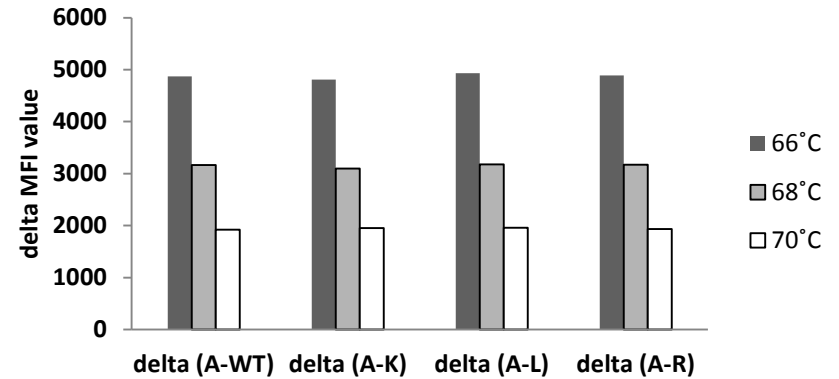
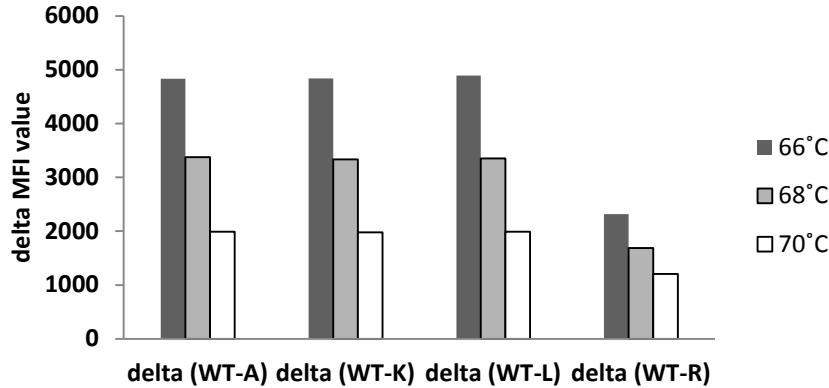
Data analysis

Background correction of MFI
Calculation of mutant/wild type allele index
Index (mutant allele) = [MFI (mutant allele)]/[MFI (mut allele 1) + ... + MFI (mut allele 4) + MFI (wt allele)]

SELECTION OF THE OPTIMAL HYBRIDIZATION TEMPERATURE BASED ON STANDARD CURVES



DETERMINATION OF THE OPTIMAL HYBRIDIZATION TEMPERATURE



Delta MFI values were above 1000 for temperatures tested

PROCEDURE FOR DETERMINATION OF THE POSITIVE CUT-OFF VALUES



3 repeats
with
all probes mix

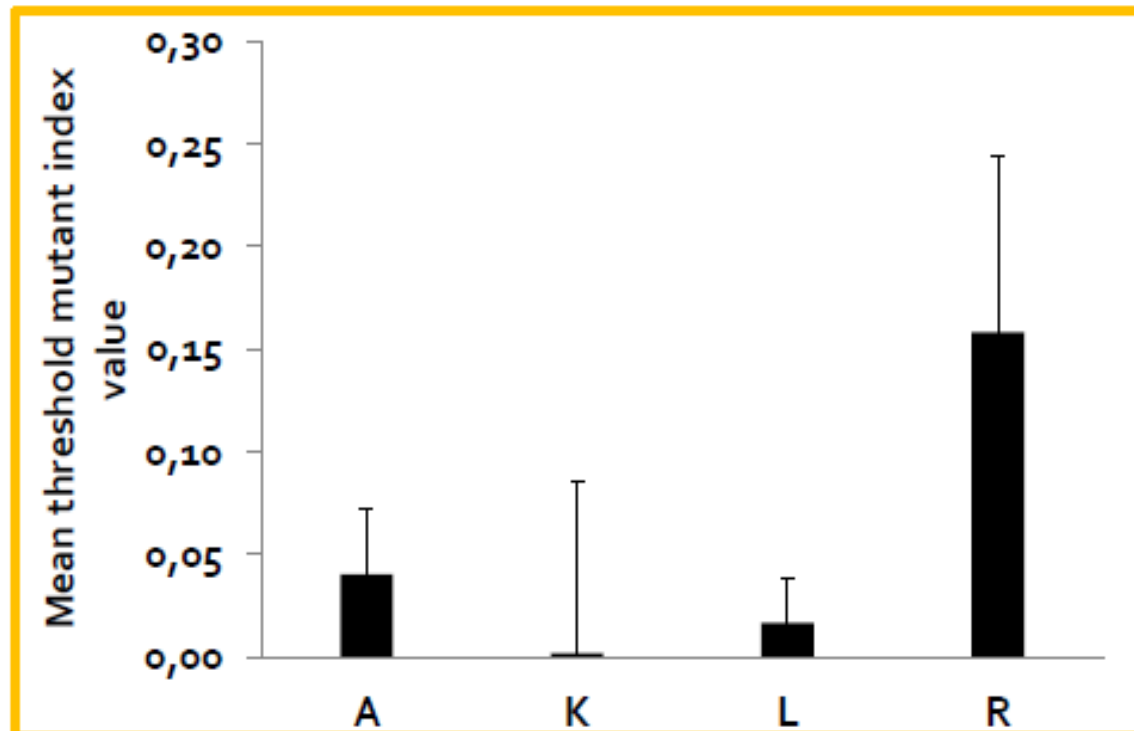


Calculation of the mean
of the maximum values
for each mutant in the
samples where it was
not present



Cut-off value
defined
Mean + SD

100%
WT
W515A
W515L
W515K
W515R



ALL MPL MUTATIONS ARE DETECTED AT 1% SENSITIVITY

%	W515A	W515K	W515L	W515R
0	0.005	0.011	0.008	0.151
1	0.088	0.100	0.094	0.203
2.5	0.145	0.200	0.160	0.247
5	0.300	0.240	0.163	0.264
12.5	0.334	0.444	0.200	0.373
25	0.488	0.545	0.366	0.429

THE MISSION IS POSSIBLE!

Luminex based detection of JAK2 V617F and MPL W515 mutations:

- ✓ Diagnoses the vast majority of Ph'(-) MPN cases
- ✓ It's easy, rapid (4-5h) and reproducible
- ✓ It's sensitive (1%) and quantitative
- ✓ Allows mid-throughput sample processing
- ✓ It's open for further multiplex development
- ✓ It's affordable (Ask the local dealers for the reagents)

ACKNOWLEDGEMENTS

*Prof. Elissaveta Naumova
Dept. of Clinical Immunology*



*Prof. Evguenyi Hadjiev
Dept. of Hematology*



Velizar Shivarov



*Prof. Hiroh Saji,
HLA Laboratory, NPO, Kyoto, Japan*



Фонд

"Научни изследвания"

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