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Via VIII Febbraio, 2

**05 Aprile**

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Auditorium  
Via Altinate, 71



# **Serum thymidine kinase activity in patients with HR-positive/HER2-negative advanced breast cancer treated with ribociclib plus letrozole: Results from the prospective BioltaLEE trial**

**Luca Malorni, Giampaolo Bianchini, Roberta Caputo, Alberto Zambelli, Fabio Puglisi, Giulia V Bianchi, Lucia Del Mastro, Ida Paris, Filippo Montemurro, Giacomo Allegrini, Marco Colleoni, Stefano Tamberi, Claudio Zamagni, Marina E Cazzaniga, Michele Orditura, Valentina Guarneri, Daniela Castelletti, Matteo Benelli, Mariacristina Di Marino, Grazia Arpino, Michelino De Laurentiis**

**Luca Malorni**

SOS Ricerca Traslazionale  
SOC Oncologia  
Ospedale di Prato  
Azienda USL Toscana Centro

# Disclosures

- **Research support: Pfizer, Novartis**
- **Advisory role: Pfizer, Novartis, Lilly, Roche, Menarini Stemline**
- **Travel arrangements: Celgene, Menarini Stemline**



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Original Research

## Serum thymidine kinase activity in patients with HR-positive/HER2-negative advanced breast cancer treated with ribociclib plus letrozole: Results from the prospective BioItaLEE trial



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### AWARDS:





**Monaleesa-2**

January 2014- March 2015

**Monaleesa-3**

June 2015- June 2016

**Monaleesa-7**

December 2014- August 2016

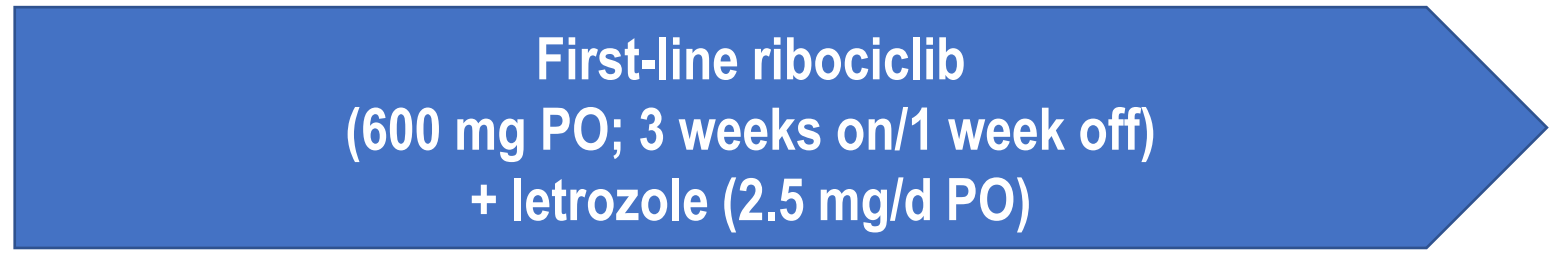
# BioltaLee- Study Design



N=287

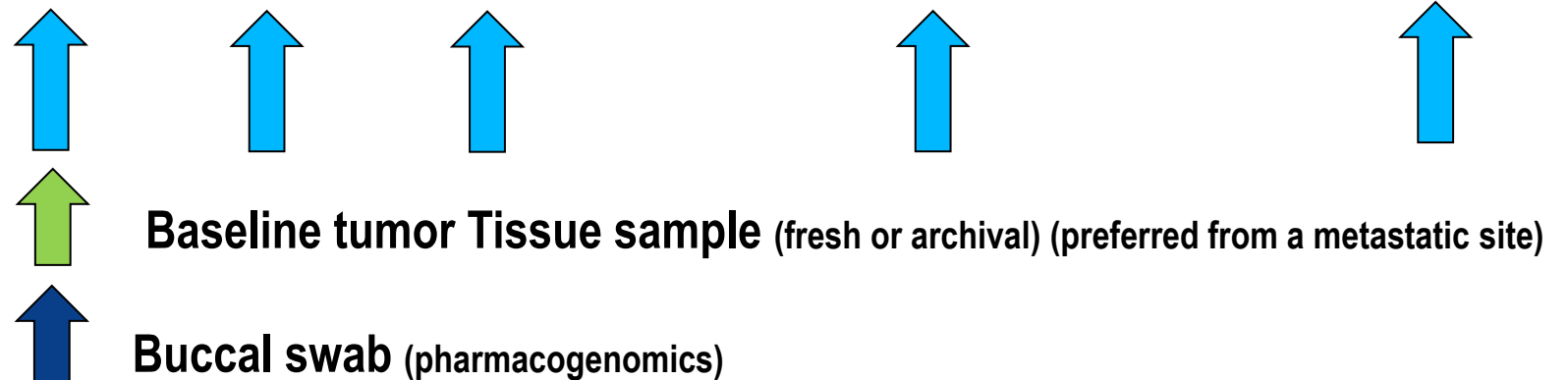
- Postmenopausal women with HR+, HER2- aBC (locoregionally recurrent not amenable to surgery or metastatic)
- No prior systemic hormonal therapy or chemotherapy for aBC
- TFI >12 months\*
- Patients willing to undergo blood and tumor sample collection at baseline and at a scheduled timeframe

Enrollement from 02 February to 28 November 2018  
Across 47 Italian centers



Baseline (D0)      Day 15 Cycle 1 (D15)      Day 1 Cycle 2 (C2D1)      First Imaging (~ 3 months) (FI)      Disease progression Or end of treatment (EOT)

- Liquid biopsy for:**
- ctDNA analysis
  - Serum thymidine kinase activity

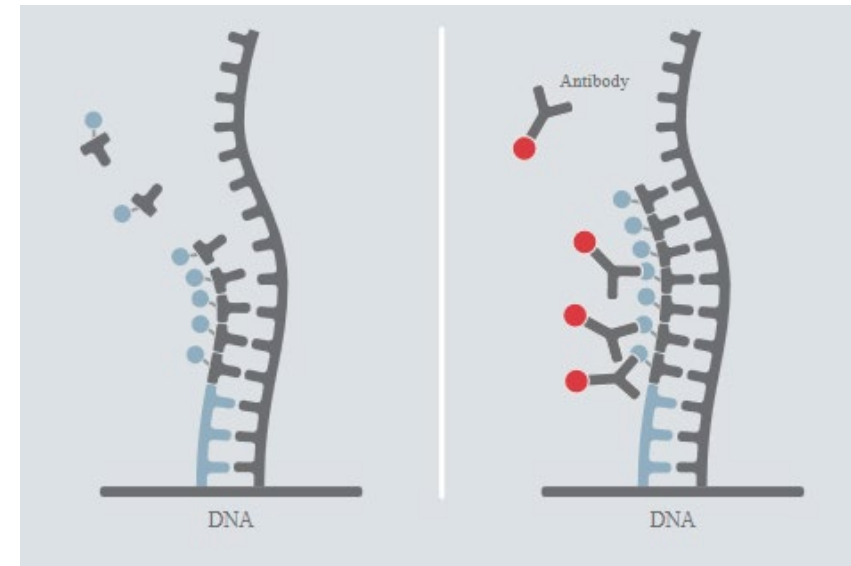
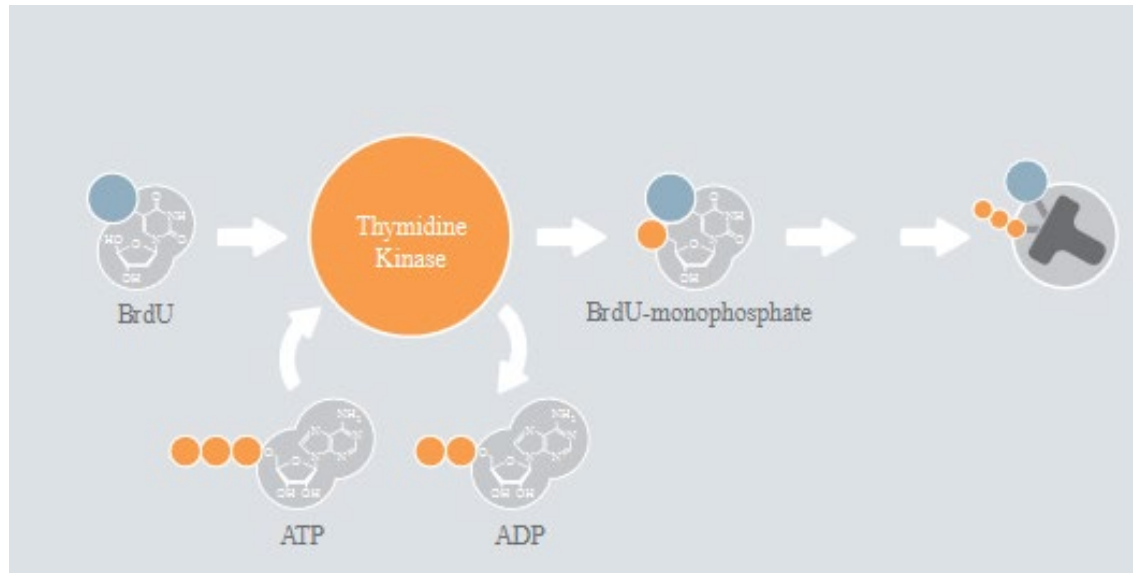


# THYMIDINE KINASE ACTIVITY

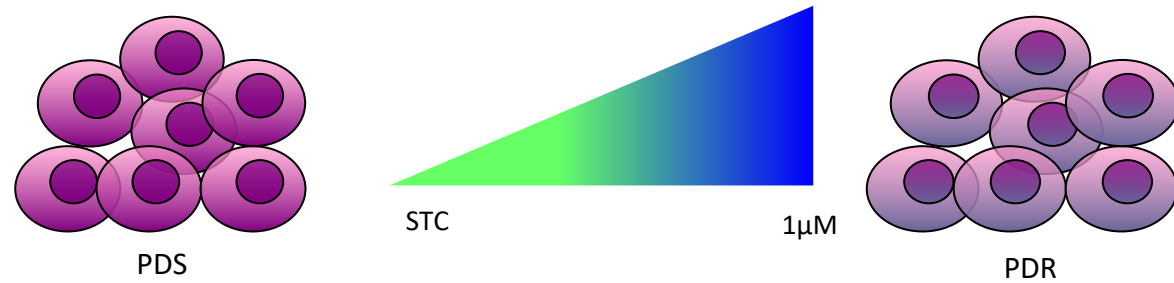
- TK1 is a cell cycle dependent enzyme playing a critical role in cell proliferation
- TK1 activity rapidly increases after the G1-S transition and then declines
- Cancer cells can secrete pathological levels of TK1 detectable in blood

## *“Liquid Ki67”*

The ELISA based DiviTum™ assay (Biovica International, Uppsala, Sweden) determines the enzymatic activity of TK1 in blood serum/plasma or cell cultures.

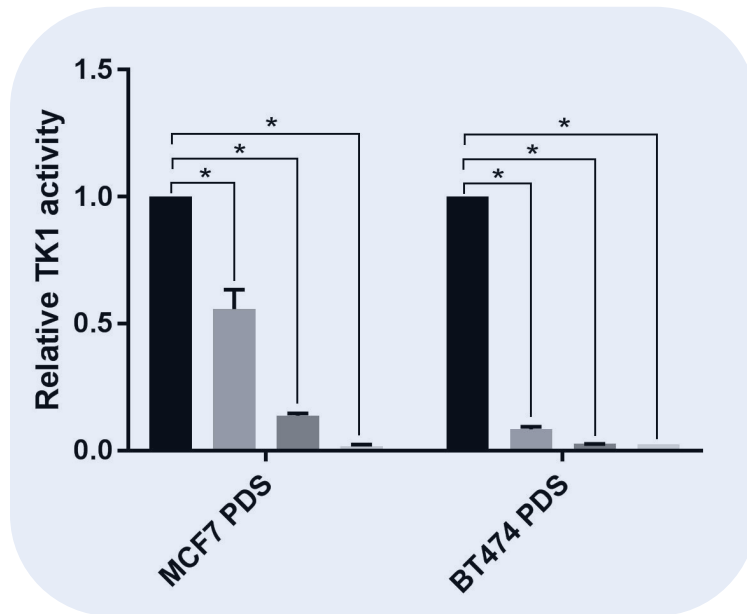


# TK1 ACTIVITY AND CDK4/6 INHIBITION

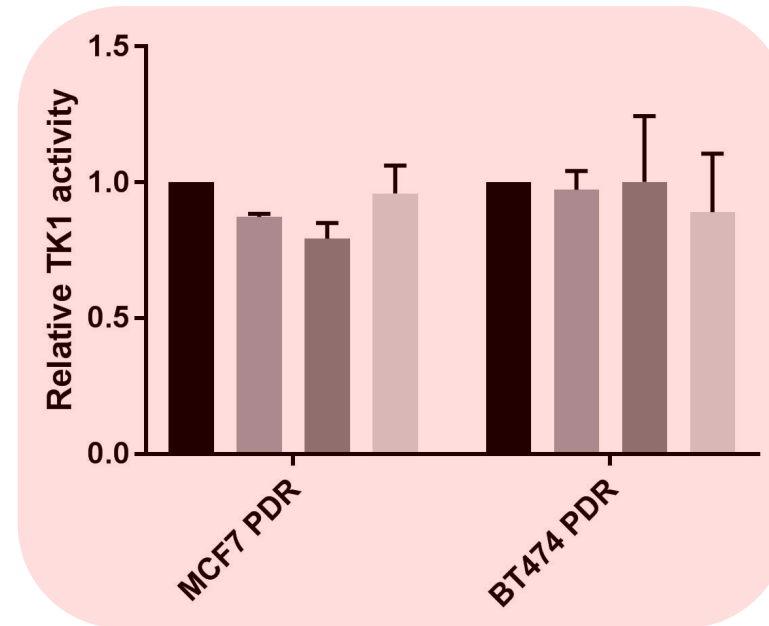


TK1 activity (assessed at day 3)

Palbociclib sensitive



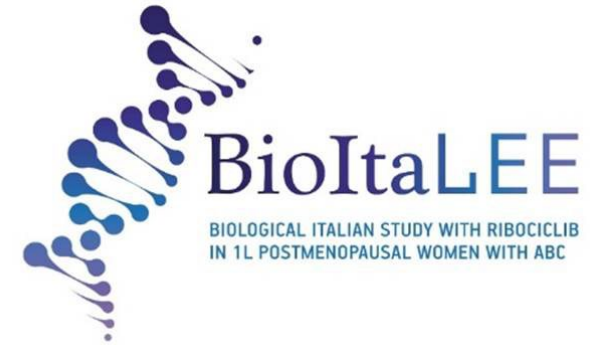
Palbociclib resistant



TK1 is strongly modulated by treatment in sensitive cells but not in resistant cells

- DMSO, D3
- Palbociclib 50nM, D3
- Palbociclib 350nM, D3
- Palbociclib 1µM, D3

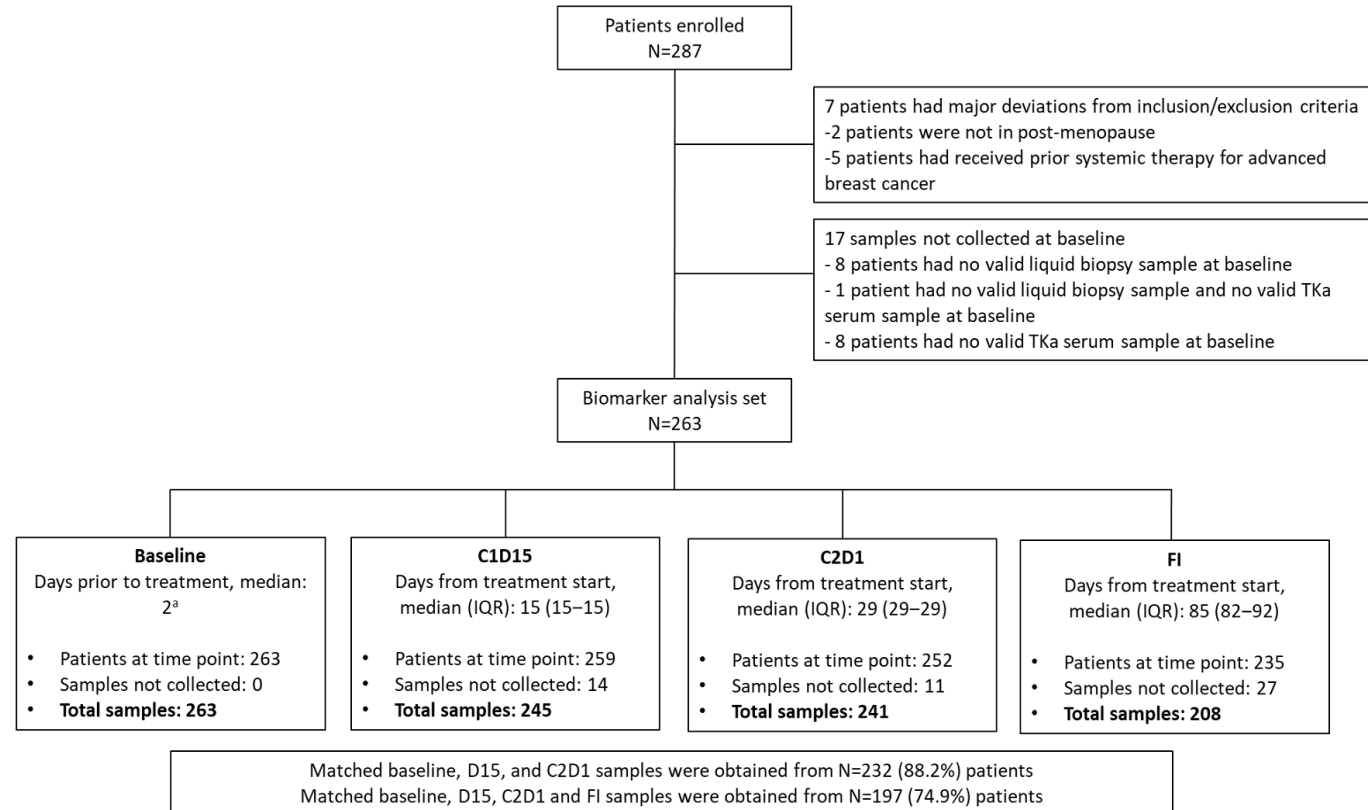
# BioltaLee- Objectives and endpoints



- The primary objective of the BioItaLEE study was to identify ctDNA alterations, their changes during treatment and possible association with clinical outcomes;
- Evaluation of **sTKa levels over time** during treatment with ribociclib plus letrozole and their association with clinical outcomes was a **key, pre-specified secondary objective** of the study;
- time to progression and progression-free survival (PFS) were secondary end-points.



# BioltaLee- CONSORT diagram

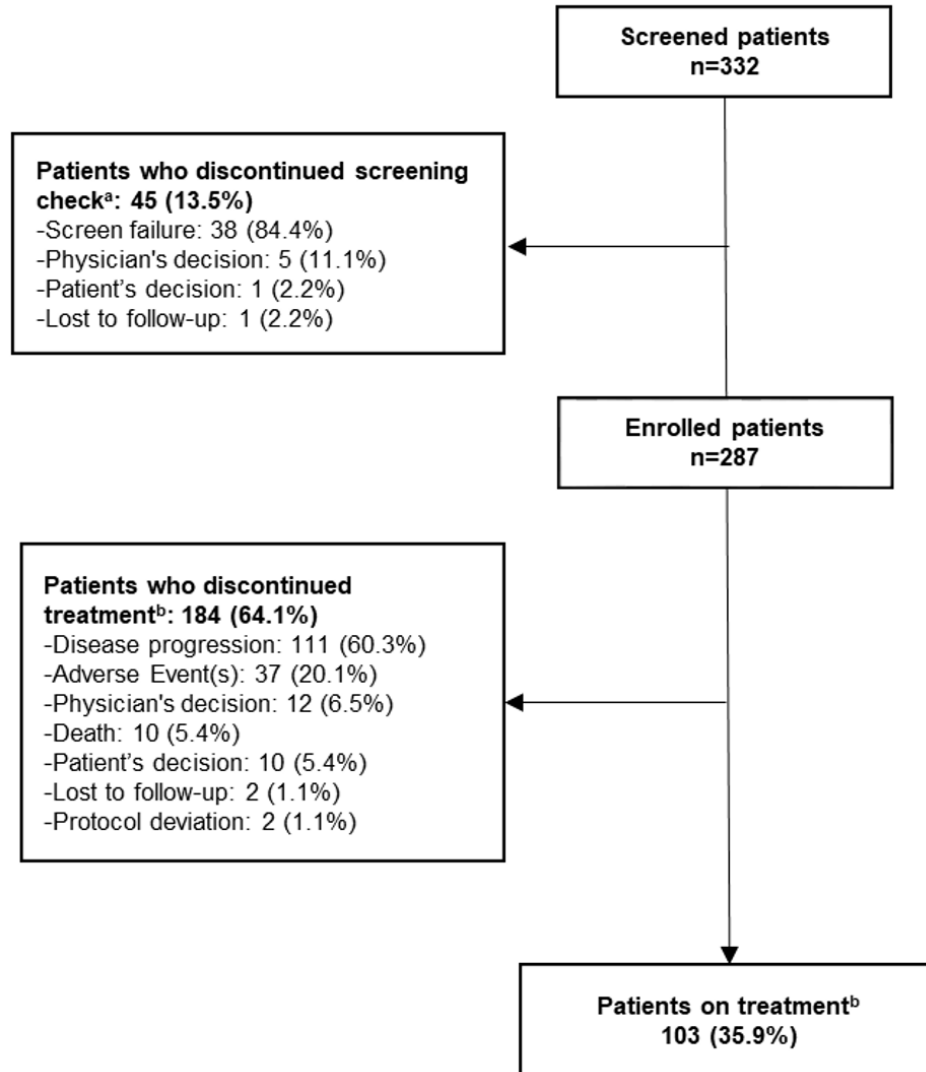


Out of 287 pts enrolled, 263 (92%) compose the Biomarker analysis set

matched Baseline, D15 and C2D1  
232 (88%)

matched Baseline, D15, C2D1 and First Imaging  
197 (75%)

# BioltaLee- CONSORT diagram



at time of analysis  
64.1% (n = 184) had discontinued treatment  
35.9% (n = 103) were still on ribociclib plus  
letrozole

median follow-up was 26.9 months  
(range, 22.3–32.3)

# BioltaLee- Patients' characteristics

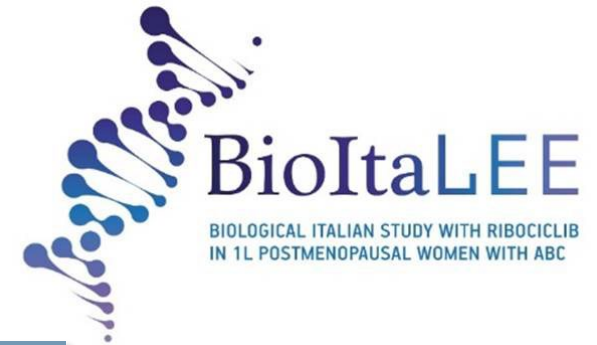


Table 1  
Patient demographics and disease characteristics.

Patient characteristic	Enrolled patients N = 287	Biomarker population N = 263
Age (years), median (IQR)	65.5 (59.0–71.0)	65.7 (60.0–72.0)
≥70 years, n (%)	98 (34.2)	93 (35.4)
ECOG PS, n (%)		
0	205 (71.4)	191 (72.6)
1	77 (26.8)	68 (25.9)
2	5 (1.7)	4 (1.5)
Disease characteristics, n (%)		
Tumour subtype		
Luminal A <sup>a</sup>	83 (28.9)	74 (28.1)
Luminal B	185 (64.5)	173 (65.8)
Unknown	19 (6.6)	16 (6.1)
Disease status		
<i>De novo</i> metastatic <sup>b</sup>	114 (39.7)	105 (39.9)
Recurrent	173 (60.3)	158 (60.1)
Disease-free interval period, n (%) <sup>c</sup>		
≤2 years	19 (11.0)	18 (11.4)
>2 years and ≤5 years	10 (5.8)	8 (5.1)
>5 years and ≤7 years	18 (10.4)	14 (8.9)
>7 years	118 (68.2)	111 (70.3)
Missing	8 (4.6)	7 (4.4)
Metastatic sites, n (%) <sup>d</sup>		
Bone	206 (71.8)	193 (73.4)
Bone only	64 (22.3)	62 (23.6)
Visceral	127 (44.3)	114 (43.3)
Liver	41 (14.3)	36 (13.7)
Lung	96 (33.5)	87 (33.1)
Other visceral	18 (6.3)	17 (6.5)
CNS	0	0
Lymph nodes	159 (55.4)	142 (54.0)
Skin	8 (2.8)	8 (3.0)
Breast	21 (7.3)	21 (8.0)
Other	28 (9.8)	26 (9.9)
Number of organs of interest involved, n (%)		
0	2 (0.7)	1 (0.4)
1	107 (37.3)	99 (37.6)
2	124 (43.2)	113 (43.0)
≥3	54 (18.8)	50 (19.0)

median age 65y (35%>70 y)  
PS ECOG 0 72%

Luminal B 65%

*de novo* M+ 40%  
long DFI (>7y) 70%

visceral M+ 43%  
bone-only 23%

# BioltaLee- efficacy



Of the 166 patients with measurable disease at baseline, 149 patients performed at least one post-baseline imaging evaluation up to the cut-off date.

## Best overall response:

- PR 52.3%
- CR 0.7%  
(ORR 53%)
- SD 35.6%
- PD 10.7%
- unkn 0.7%

Median PFS was 23.4 months (95% CI, 20.8– NE)

# BioltaLee- efficacy

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# Monaleesa 2

**Table 2.** Best Overall Response, According to Local Assessment.

Response	Ribociclib Group
Patients with measurable disease at baseline — no.	256
Confirmed best overall response — no. (%)	
Complete response	8 (3.1)
Partial response	127 (49.6)
Stable disease	95 (37.1)
Progressive disease	13 (5.1)
Unknown	13 (5.1)
Overall response†	
No. of patients	135
Percentage of patients (95% CI)	52.7 (46.6–58.9)

Median PFS was 25.3 months (95% CI 23.0–30.3)

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## BASELINE

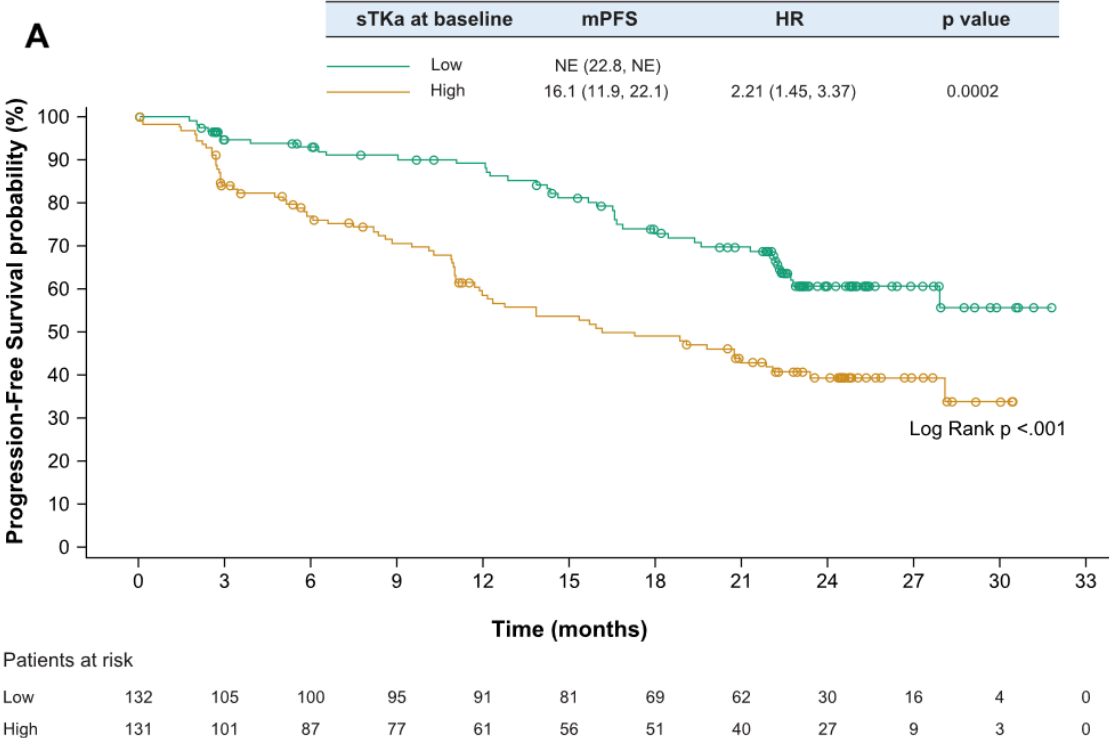
**From 263 available samples at baseline,  
median sTKa was 74.8 Du/L (19–9412)**

**11,8% of patients had sTKa levels below  
LOD (20Du/L)**

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## BASELINE

### Baseline (median cut-off)



From 263 available samples at baseline, median sTKa was 74.8 Du/L (19–9412)

11,8% of patients had sTKa levels below LOD (20Du/L)

patients with high sTKa had a worse PFS compared to those with low sTKa  
HR 2.21; 95% CI, 1.45, 3.37; P = 0.0002

High sTKa at baseline was associated with worse prognosis

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## BASELINE

Characteristic	sTKa levels at baseline		p value
	Low N=132	High N=131	
<b>Age (years), median (range)</b>	67 (47–86)	64 (47–81)	0.0214 <sup>a</sup>
<b>ECOG PS at baseline, n (%)</b>			
Grade 0	101 (76.5)	90 (68.7)	0.0725 <sup>b</sup>
Grade 1	31 (23.5)	37 (28.2)	
Grade 2	0	4 (3.05)	
<b>Time from first diagnosis to study treatment start (years), median (range)</b>	7.43 (0.04–31.88)	2.97 (0.04–26.73)	0.0687 <sup>a</sup>
<b>Patient disease status, n (%)</b>			
<i>De novo</i>	49 (37.1)	56 (42.75)	0.3515 <sup>b</sup>
Recurrent	83 (62.9)	75 (57.25)	
<b>Ki67 categories, n (%)</b>			
<20%	62 (47.0)	45 (34.35)	0.0278 <sup>b</sup>
20-35%	43 (32.6)	45 (34.35)	
>35%	16 (12.1)	33 (25.2)	
Missing	11 (8.3)	8 (6.1)	
<b>Tumor type</b>			
Luminal A	45 (34.1)	29 (22.1)	0.038 <sup>b</sup>
Luminal B	77 (58.3)	96 (73.3)	
Unknown	10 (7.6)	6 (4.6)	

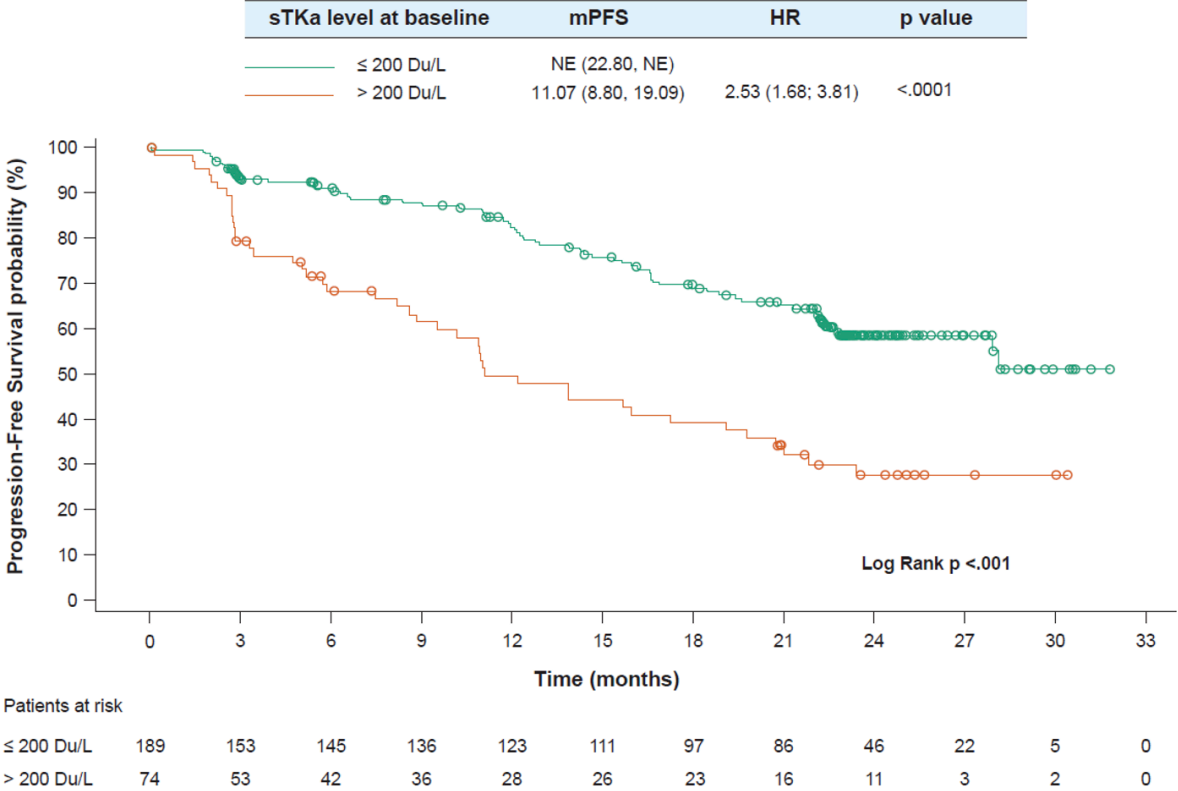
High sTKa at baseline was associated with younger age and LumB status



# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## BASELINE

### Baseline (200 Du/L cut-off)

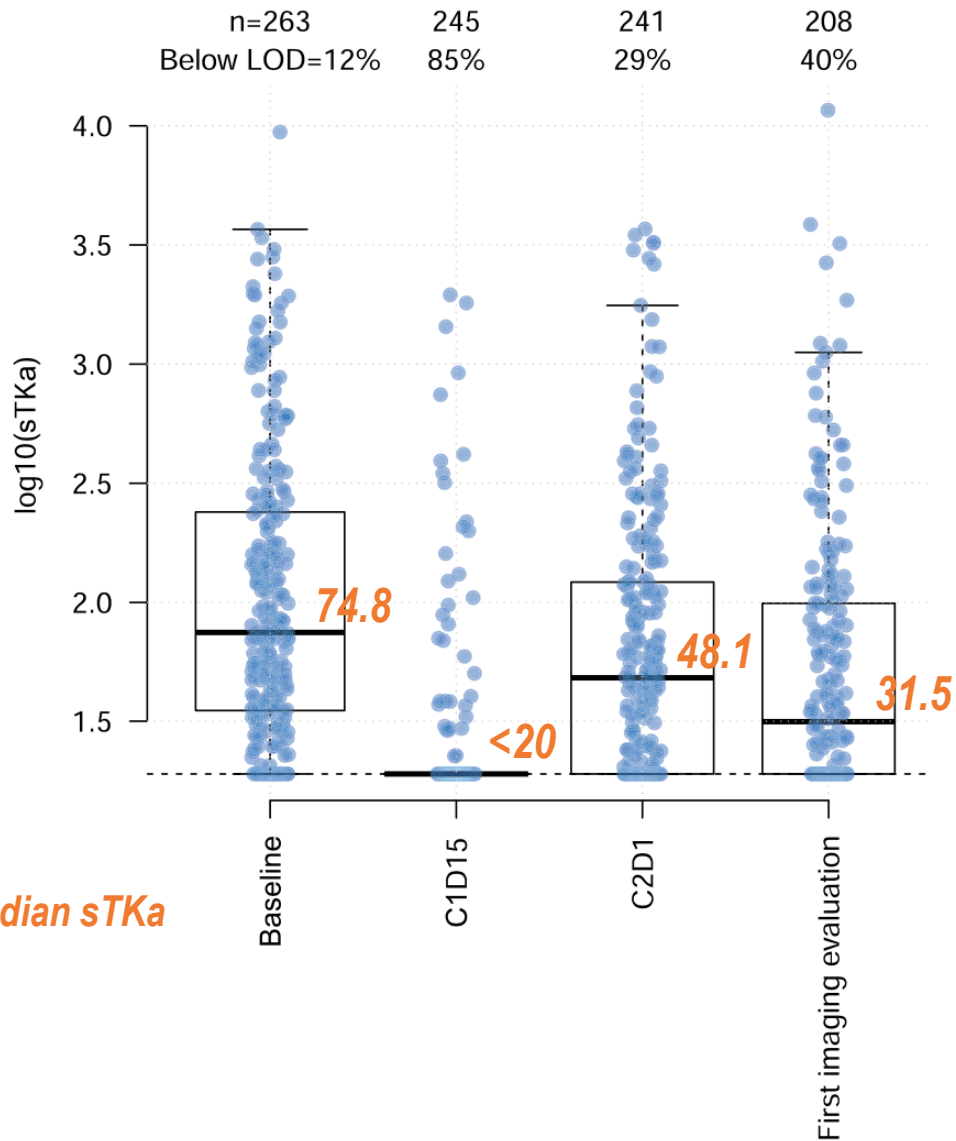


When using a cut-off of 200 Du/L, 74 (28.1%) patients had high and 189 (71.9%) had low sTKa levels at baseline.

patients with high sTKa had a worse PFS compared to those with low sTKa  
 HR, 2.53; 95% CI, 1.68, 3.81; P < 0.0001

# Serum Thymidine Kinase 1 in the BIOITALee trial- key findings

## Early changes during C1



\*median sTKa

Early sTKa clearance (levels below LOD) was observed in:

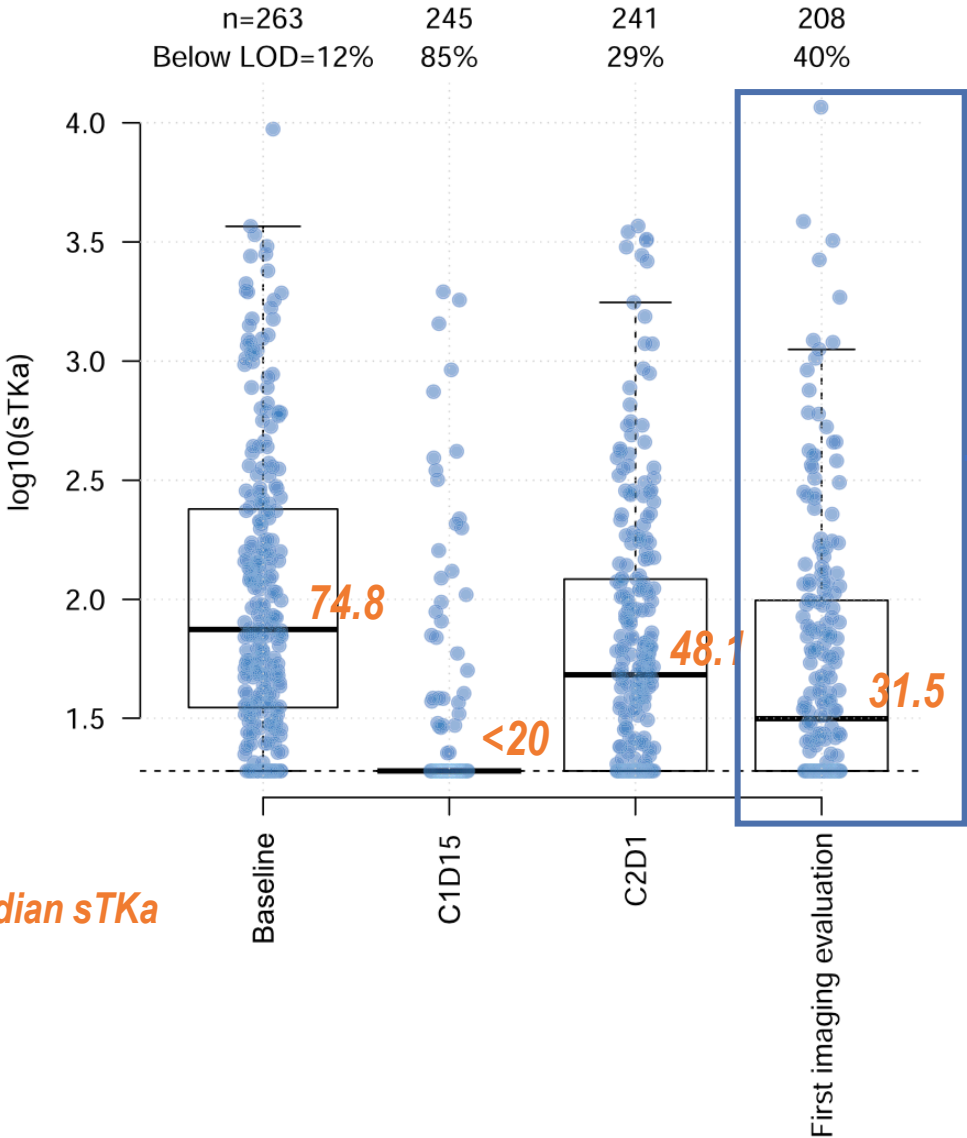
- 85% of patients at D15 and
- 29% of patients at C2D1

A rebound at C2D1 was seen in 68.5% of patients

A significant reduction in sTKa was observed upon ribociclib + letrozole treatment at D15 and C2D1 but rebound is common

# Serum Thymidine Kinase 1 in the BIOITALee trial- key findings

## Early changes during C1



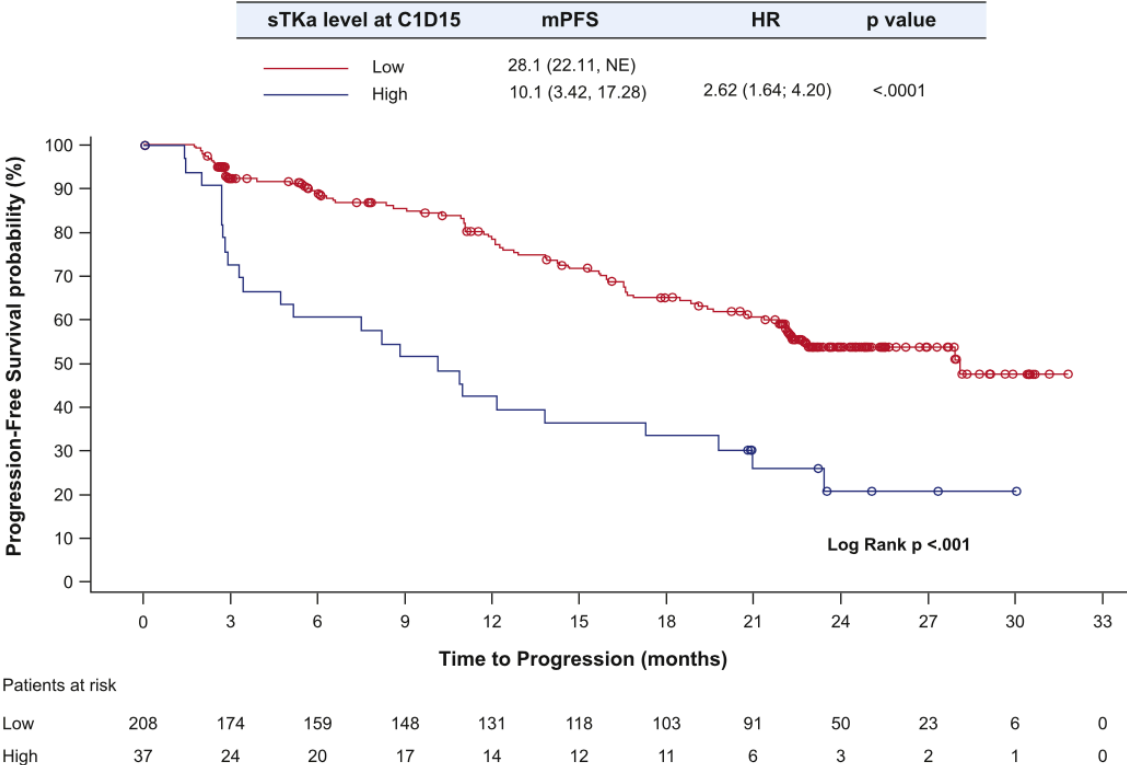
\*median sTKa

median sTKa at First Imaging was 31.5 Du/L (IQR, 19.0–99.2)

- median sTKa at first imaging according to treatment interval from RIBO:
- ongoing treatment (n = 113; 54.3%), median sTKa 26.9 Du/L (IQR, 19.0–95.1)
  - off-treatment ≤7 days (n=67; 32.2%) median sTKa 19.0 Du/L (IQR, 19.0–56.9)
  - off-treatment > 7 days (n = 28; 13.5%) median sTKa 116.0 Du/L (IQR, 63.8–429.1)

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings D15

## D15 (LOD cut-off)



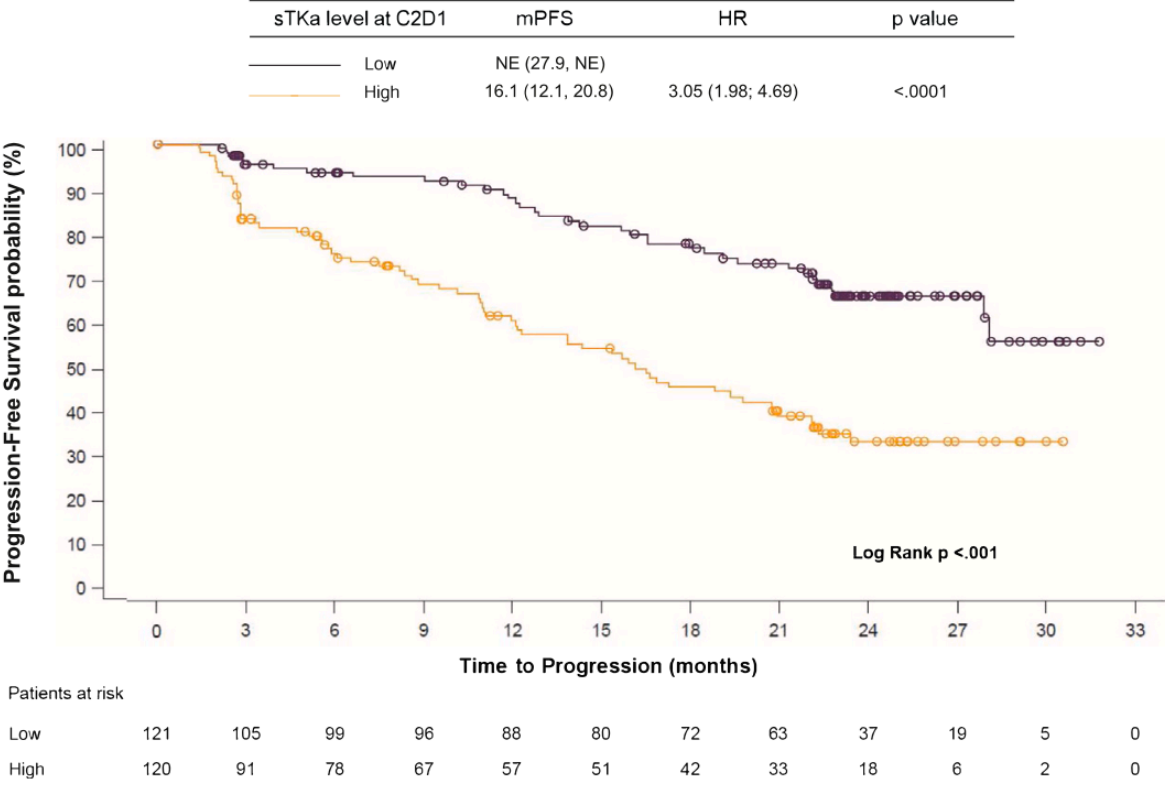
**sTKa clearance at D15 was not observed in 37 pts (15%)**

**these pts had worse outcome compared to those with sTKa clearance  
HR, 2.62; 95% CI, 1.64, 4.20; P < 0.0001**

**Lack of sTKa clearance (<LOD) at D15 was associated with poor prognosis**

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings C2D1

## C2D1 (median cut-off)



**median sTKa at C2D1 was 48.1 Du/L (IQR, 19.0–121.7)**

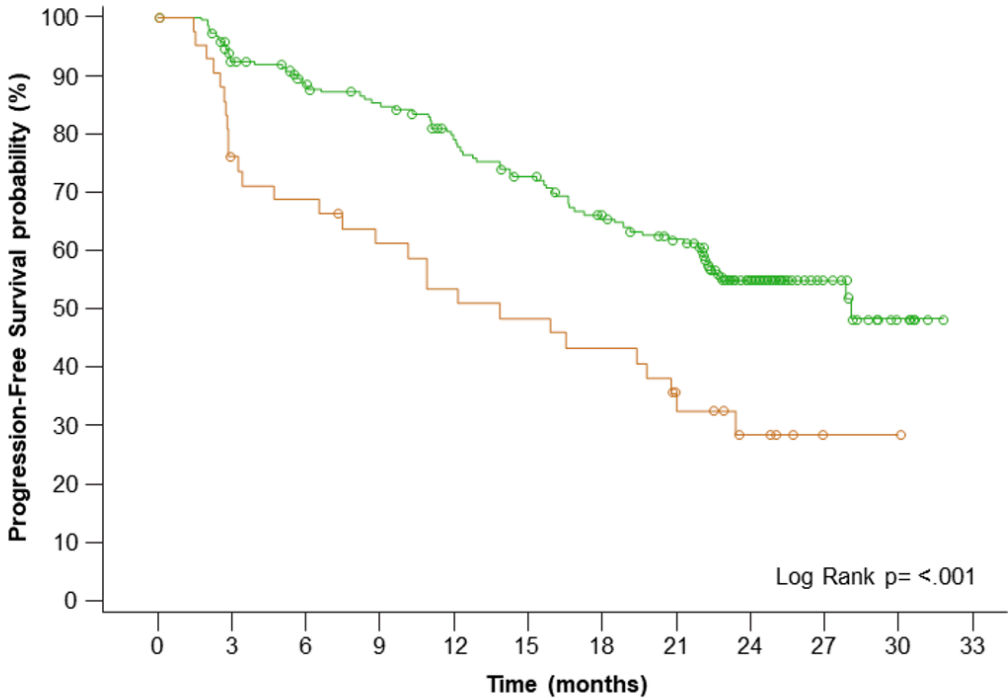
**pts with high sTKa at C2D1 had a worse PFS compared with pts with low sTKa HR, 3.05; 95% CI, 1.98, 4.69; P < 0.0001**

**High sTKa (>median) at C2D1 was associated with poor prognosis**

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings C2D1

## C2D1 (200 Du/L cut-off)

sTKa level at C2D1	mPFS	HR	p value
≤ 200 Du/L	28.1 (22.2, NE)		
> 200 Du/L	13.8 (6.5, 21.0)	2.31 (1.48; 3.62)	0.0003



Patients at risk	0	3	6	9	12	15	18	21	24	27	30	33
≤ 200 Du/L	198	165	149	139	124	112	97	86	49	24	6	0
> 200 Du/L	43	31	28	24	21	19	17	10	6	1	1	0

When using a cut-off of 200 Du/L, 43 patients had high and 198 had low sTKa levels at C2D1

pts with high sTKa at C2D1 had a worse PFS compared with pts with low sTKa  
HR, 2.31; 95% CI, 1.48, 3.62; P 0.0003

High sTKa (>200 Du/L) at C2D1 was associated with poor prognosis

# Multivariate analysis

Table 2  
Multivariate analysis of PFS for sTKa levels at different time points in the study.

Characteristic	Baseline HR (95% CI)	<i>P</i> value	C1D15 HR (95% CI)	<i>P</i> value	C2D1 HR (95% CI)	<i>P</i> value
sTKa levels at time point (high versus low)	2.21 (1.45, 3.37)	0.0002	2.62 (1.64, 4.20)	<0.0001	3.05 (1.98, 4.69)	<0.0001
Recurrent versus <i>de novo</i> disease	0.98 (0.66, 1.45)	0.9147	0.98 (0.65, 1.46)	0.9037	1.06 (0.70, 1.59)	0.7877
Tumour type (luminal B versus luminal A)	1.38 (0.83, 2.27)	0.2097	1.69 (1.02, 2.80)	0.0431	1.63 (0.97, 2.74)	0.0631
No visceral metastases versus visceral metastases	0.65 (0.42, 1.02)	0.0612	0.75 (0.48, 1.17)	0.2096	0.66 (0.41, 1.05)	0.0783
≥3 organs involved by metastases versus <3 organs involved by metastases	0.9 (0.53, 1.51)	0.6860	0.94 (0.55, 1.59)	0.8085	1.05 (0.61, 1.81)	0.8713

C, cycle; CI, confidence interval; D, day; HR, hazard ratio; PFS, progression-free survival; sTKa, serum thymidine kinase activity.

**sTKa at all timepoints is independently prognostic**

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## Dynamic patterns

**Pattern 1**  
TKa <LOD at D15  
and C2D1

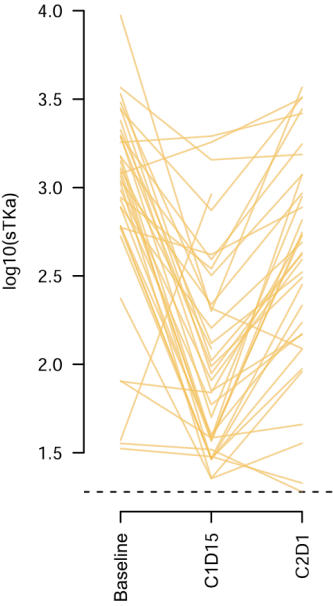
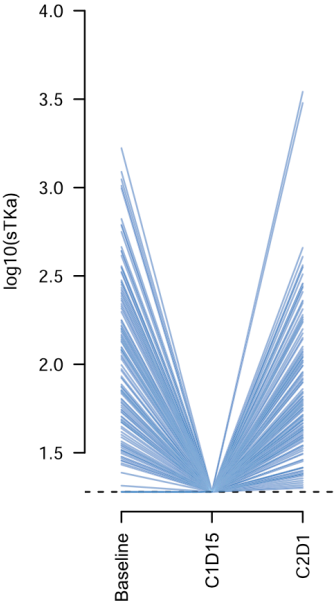
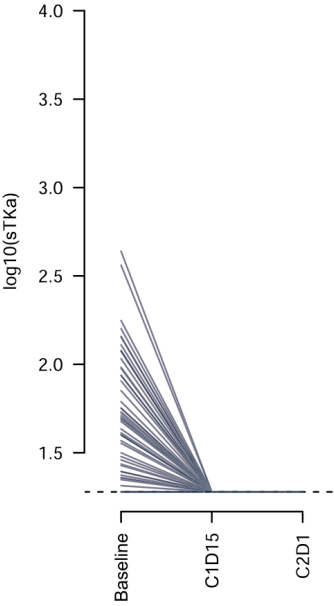
**Pattern 2:**  
TKa <LOD at D15 and  
>LOD at C2D1

**Pattern 3:**  
TKa >LOD at D15

**A** PATTERN 1 (n=62)

PATTERN 2 (n=135)

PATTERN 3 (n=37)





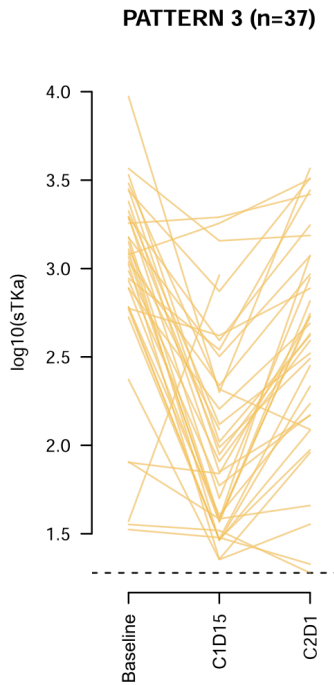
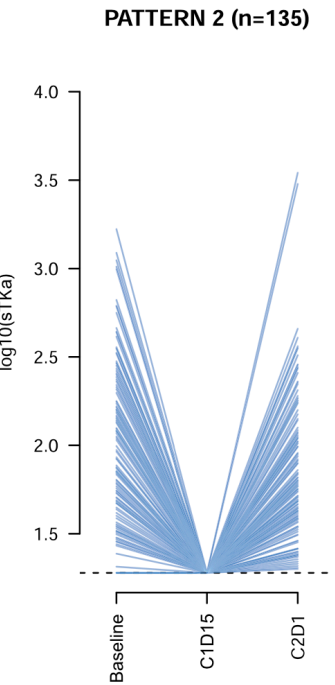
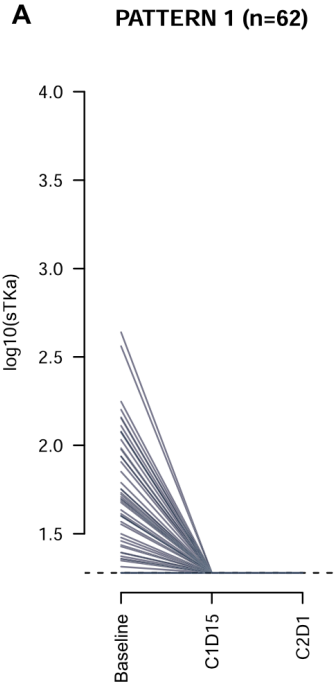
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## Dynamic patterns

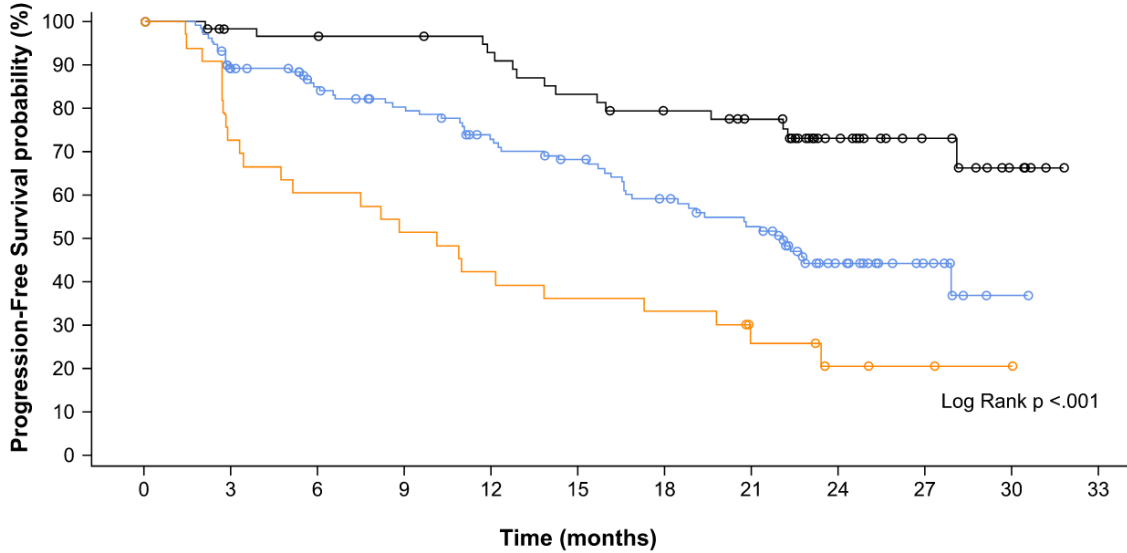
**Pattern 1**  
TKa <LOD at D15 and C2D1

**Pattern 2:**  
TKa <LOD at D15 and >LOD at C2D1

**Pattern 3:**  
TKa >LOD at D15



sTKa pattern	mPFS	HR	p value
Pattern 1	NE (28.1, NE)		
Pattern 2	22.1 (16.8, NE)	2.89 (1.57, 5.31)	0.0006
Pattern 3	10.1 (3.1, 17.3)	5.65 (2.84, 11.23)	<0.0001



Patients at risk

	0	3	6	9	12	15	18	21	24	27	30	33
Pattern 1	62	54	53	52	49	44	40	36	22	12	5	0
Pattern 2	135	111	97	88	76	69	58	50	27	11	1	0
Pattern 3	37	24	20	17	14	12	11	6	3	2	1	0

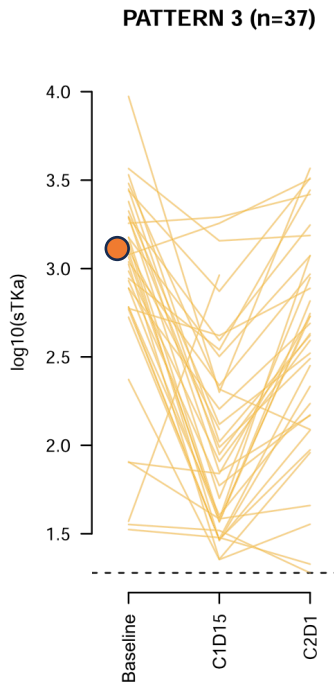
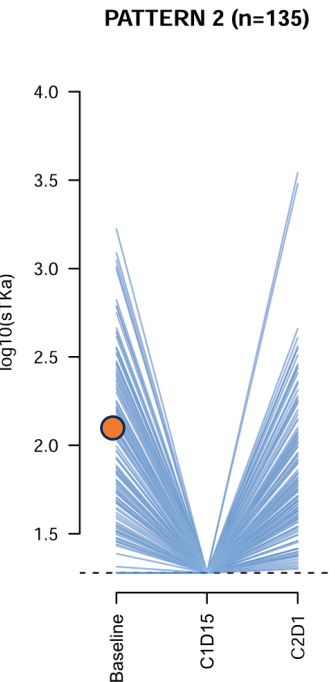
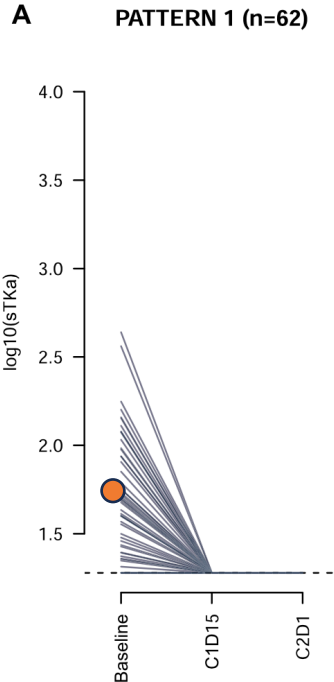
# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## Dynamic patterns

**Pattern 1**  
TKa <LOD at D15 and C2D1

**Pattern 2:**  
TKa <LOD at D15 and >LOD at C2D1

**Pattern 3:**  
TKa >LOD at D15

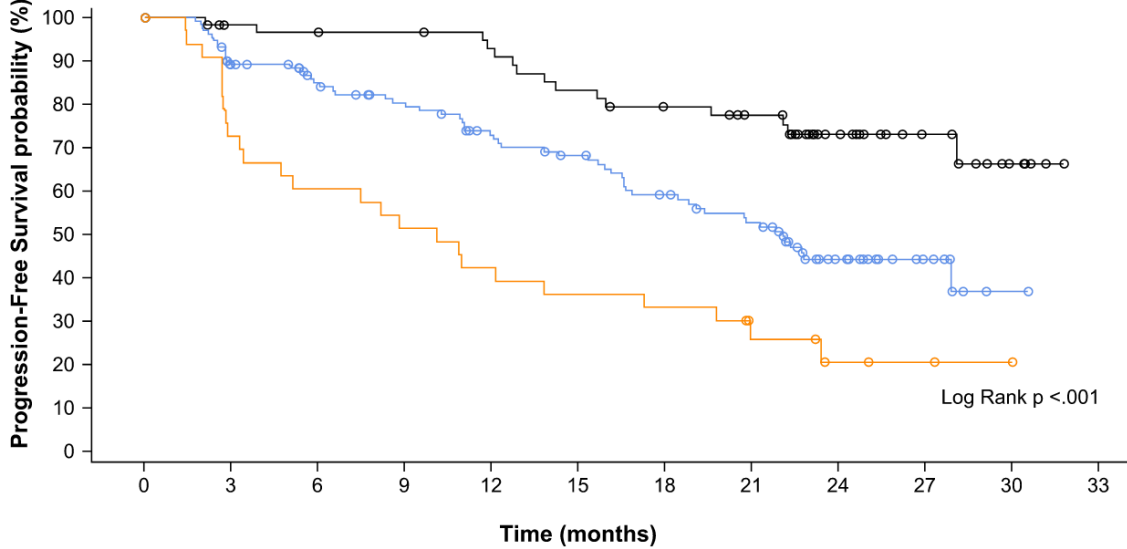


**median sTKa**  
**39.8 Du/L**  
**(19.0–80.6)**

**83.8Du/L**  
**(43.5–178.1)**

**1163.0Du/L**  
**(606.8–1950.8)**

sTKa pattern	mPFS	HR	p value
Pattern 1	NE (28.1, NE)		
Pattern 2	22.1 (16.8, NE)	2.89 (1.57, 5.31)	0.0006
Pattern 3	10.1 (3.1, 17.3)	5.65 (2.84, 11.23)	<0.0001



Patients at risk

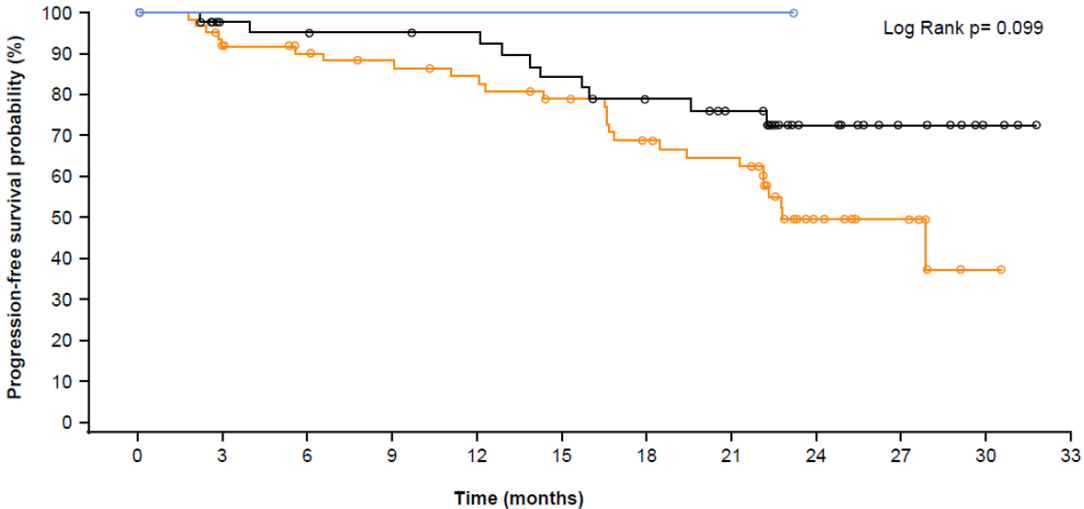
	0	3	6	9	12	15	18	21	24	27	30	33
Pattern 1	62	54	53	52	49	44	40	36	22	12	5	0
Pattern 2	135	111	97	88	76	69	58	50	27	11	1	0
Pattern 3	37	24	20	17	14	12	11	6	3	2	1	0

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## Dynamic patterns

Low baseline sTKa (<median)

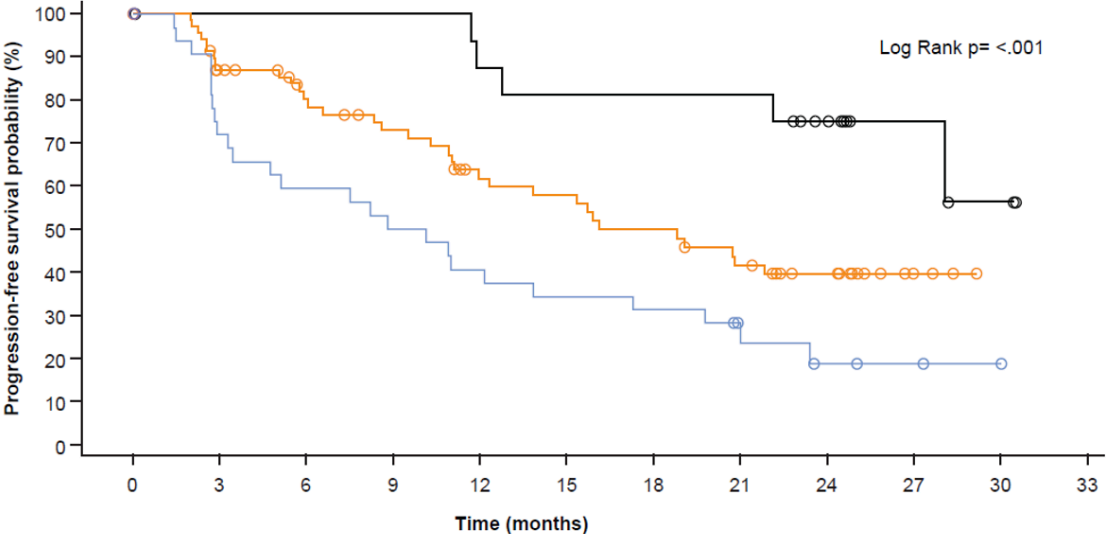
sTKa pattern	mPFS	HR	p value
Pattern 1	NE (NE, NE)		
Pattern 2	22.8 (19.4, NE)	2.55 (1.2; 5.6)	0.0184
Pattern 3	NE (NE, NE)	<0.01 (<0.01; NE)	0.9904



Patients at risk	0	3	6	9	12	15	18	21	24	27	30	33
Pattern 1	46	38	37	36	35	31	27	23	14	8	3	0
Pattern 2	65	55	51	48	45	40	33	30	12	8	1	0
Pattern 3	3	1	1	1	1	1	1	1	0			

High baseline sTKa (>median)

sTKa pattern	mPFS	HR	p value
Pattern 1	NE (22.1, NE)		
Pattern 2	16.1 (12.0, NE)	3.3 (1.2; 9.5)	0.0243
Pattern 3	9.5 (3.3, 17.3)	6.2 (2.1; 18.2)	0.001



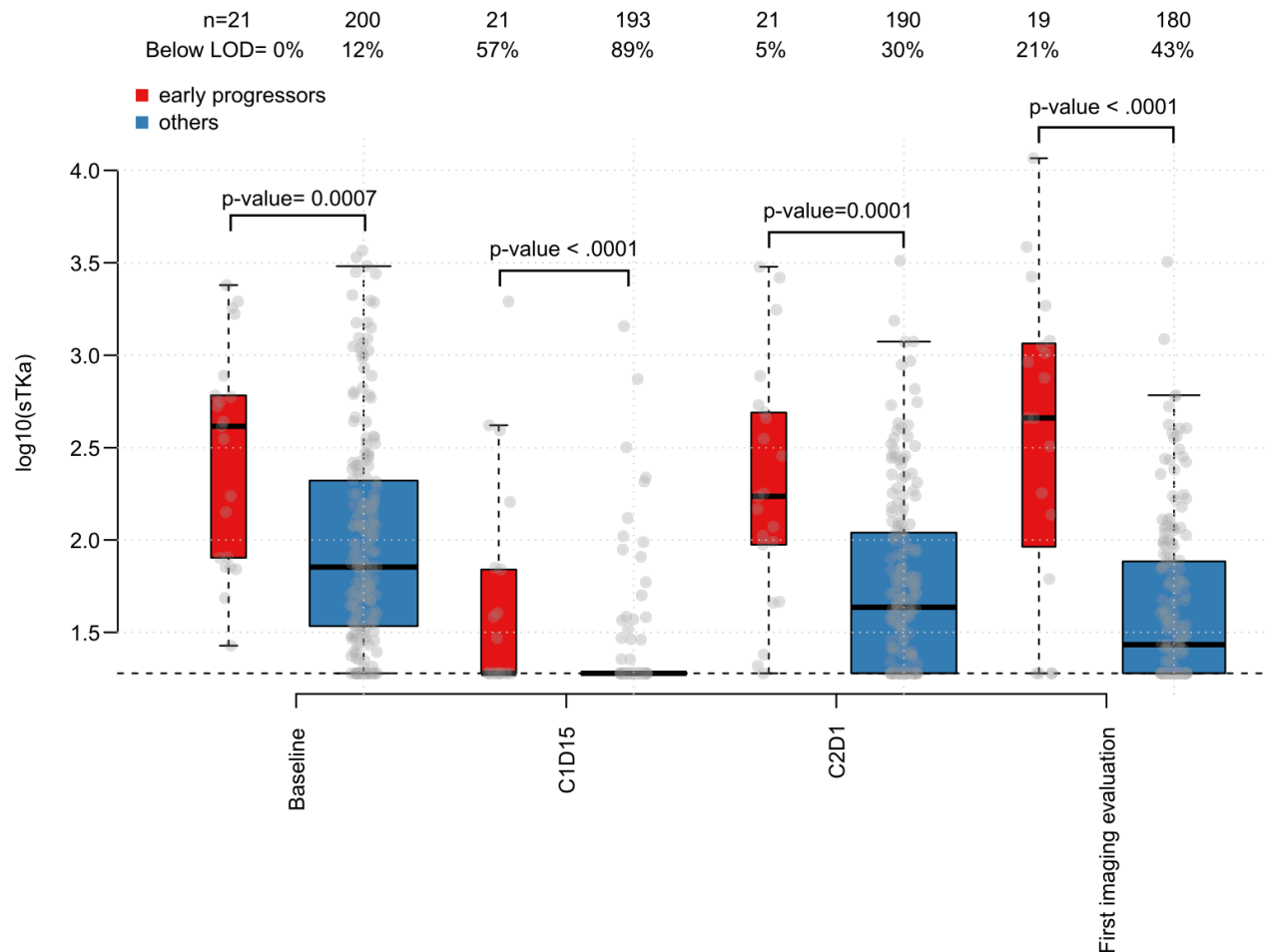
Patients at risk	0	3	6	9	12	15	18	21	24	27	30	33
Pattern 1	16	16	16	16	14	13	13	13	8	4	2	0
Pattern 2	70	56	46	40	31	29	25	20	15	3	0	0
Pattern 3	34	23	19	16	13	11	10	5	3	2	1	0

**Pts with High baseline sTKa with optimal sTKa response (pattern 1) have excellent prognosis**

# Serum Thymidine Kinase 1 (TKa) in the BIOITALee trial- key findings

## First Imaging

Out of 208 patients with evaluable disease at FI and a valid sTKa value, 20 (9.6%) had progressive disease as assessed by imaging and clinical evaluation at this time point (**early progressors**)



- **Early progressors** had higher sTKa levels at all time-points
- Only 4 out of 101 (4.0%) patients with sTKa < median at FI had disease progression at this timepoint

Can we omit radiological evaluation at first imaging (3 months) if sTKa is low?

# TKa DATA IN PATIENTS TREATED WITH ET+CDK4/6i: changes during treatment

- First/second line tx with PALBO+FUL
- endocrine resistant MBC

- <LLOD D15 83%
- rebound at D28 54%

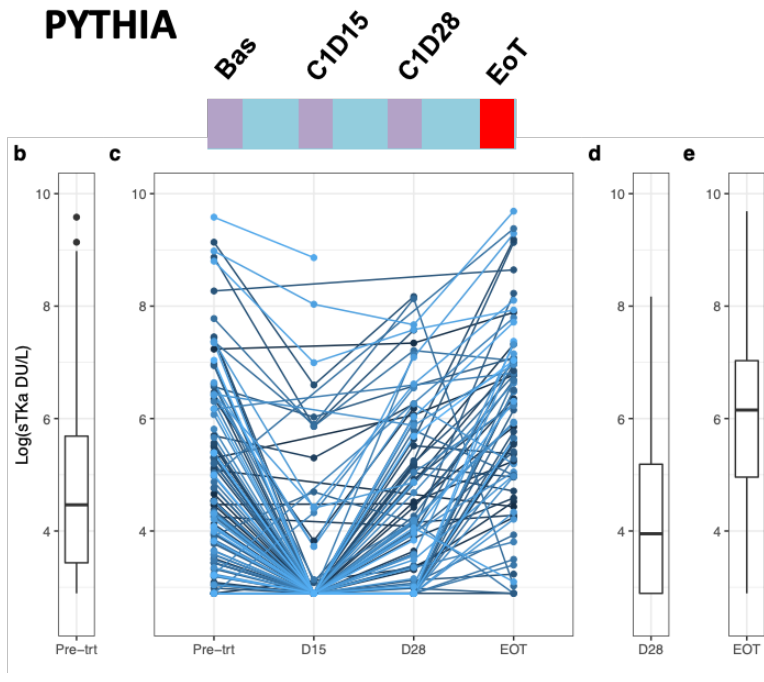
- First line tx with RIBO+LET
- endocrine sensitive MBC

- <LLOD D15 85%
- rebound at D28 68%

- First/second line tx with PALBO+HT
- endocrine sensitive/resistant MBC

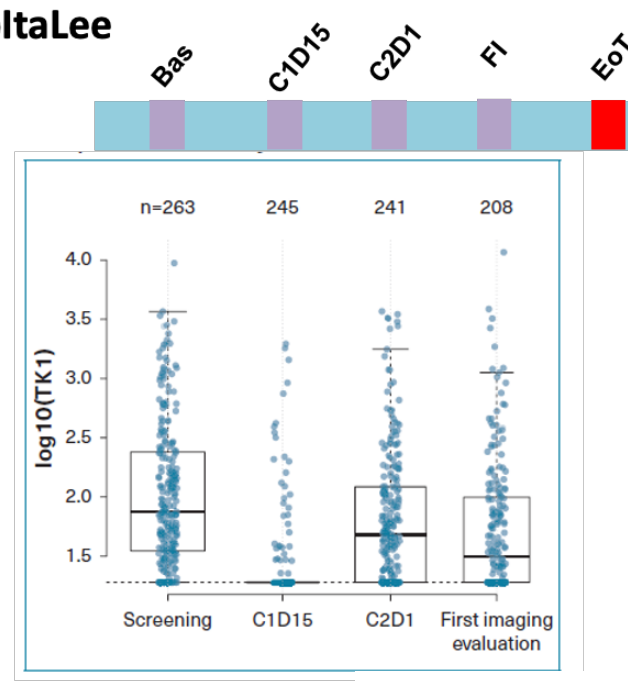
- <LLOD D15 78%
- rebound D28 36%

## PYTHIA



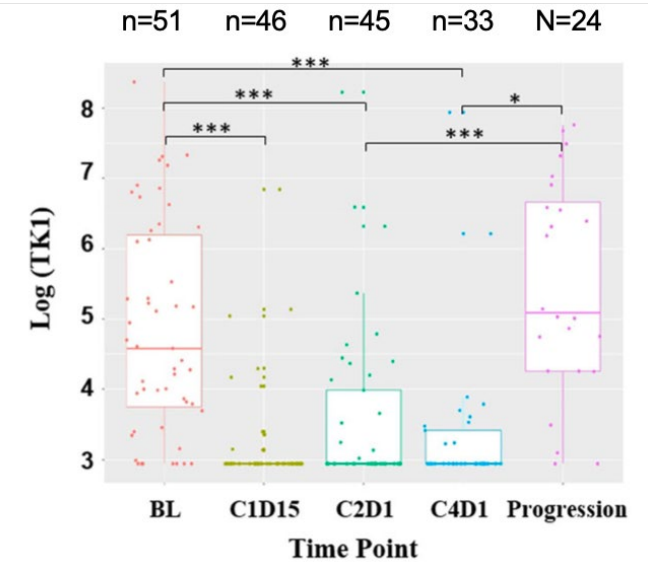
Malorni L. et al, EJC 2022

## BioltaLee



Malorni L. et al, EJC 2023

## WashU palbo dosing trial

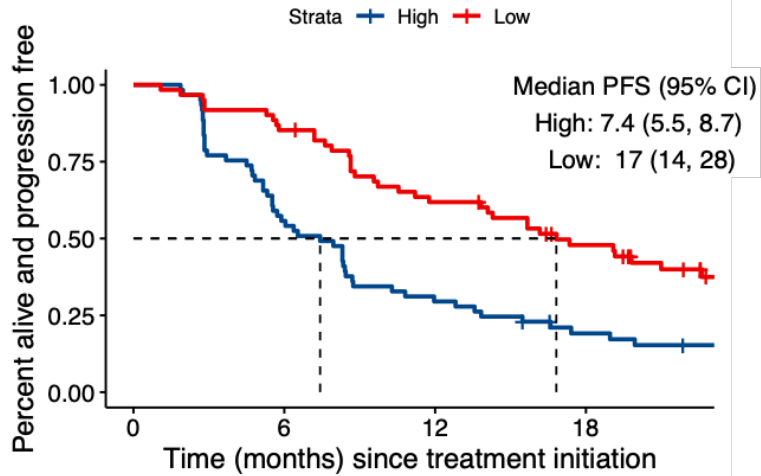


Krishnamurthy, J. npjBC 2022

# TKa DATA IN PATIENTS TREATED WITH ET+CDK4/6i: BASELINE

## PYTHIA

Baseline; median cut-off; 87 Du/L (<20- 14510)



Number at risk

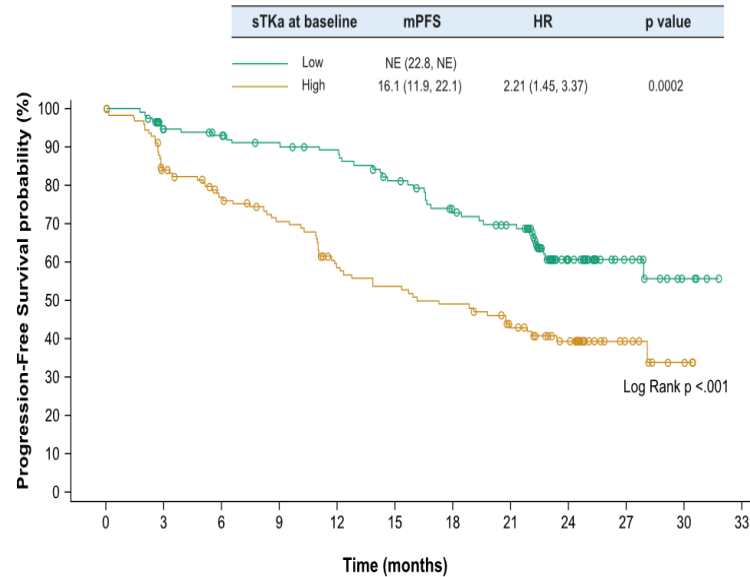
High	61	34	18	10
Low	61	52	37	26

Malorni L. et al, EJC 2022

Multivariable  
HR 1.38; 95% CI: 1.22-1.57  
p<0.001

## BioltaLee

Baseline; median cut-off, 74.8 Du/L (19–9412)



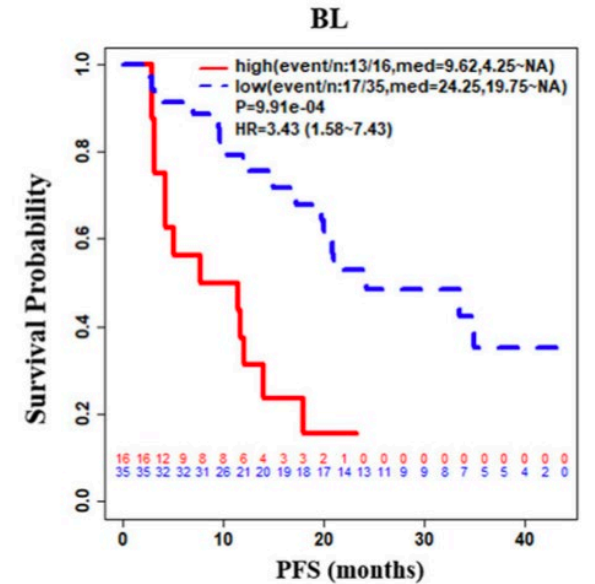
Patients at risk

Low	132	105	100	95	91	81	69	62	30	16	4	0
High	131	101	87	77	61	56	51	40	27	9	3	0

Malorni L. et al, EJC 2023

## WashU palbo dosing trial

Baseline; median cut-off, 97.9 Du/L (42.4–490.3)

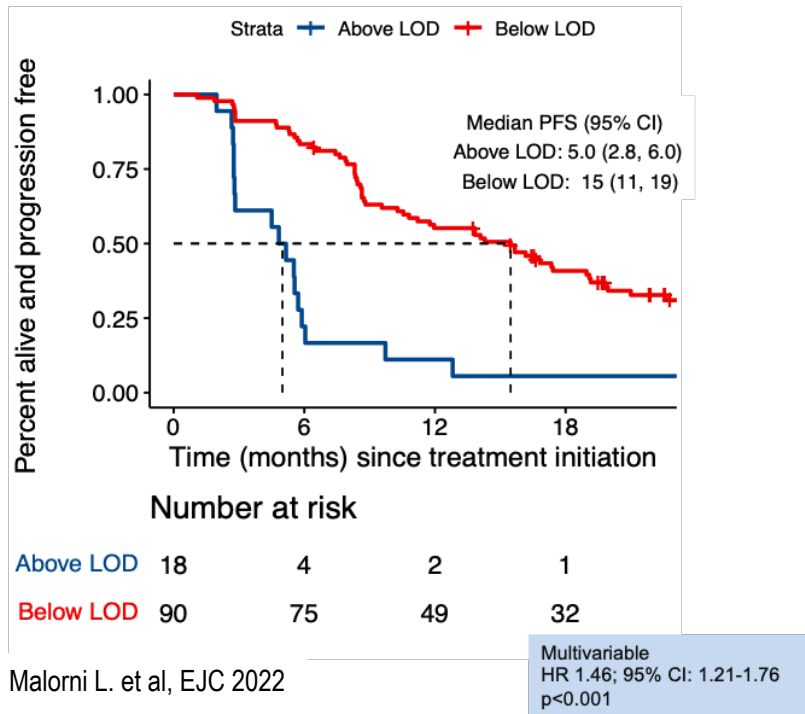


Krishnamurthy, J. npjBC 2022

# TKa DATA IN PATIENTS TREATED WITH ET+CDK4/6i: CYCLE 1 DAY 15

## PYTHIA

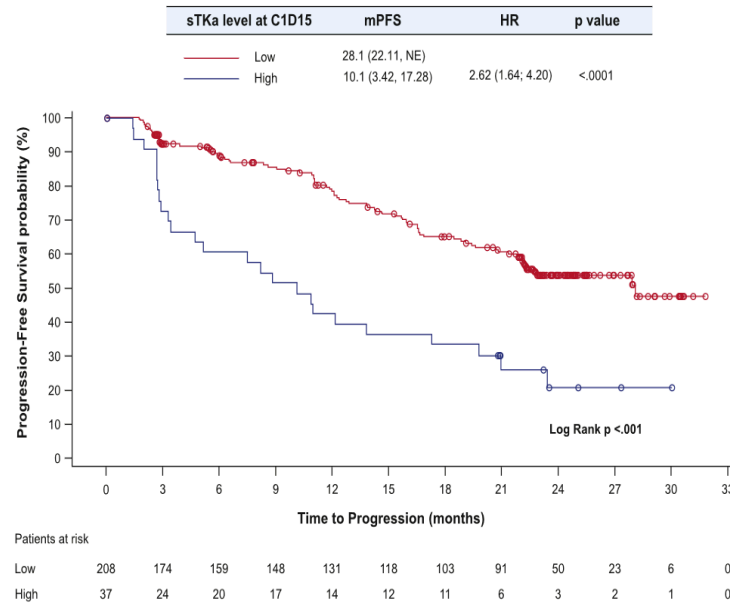
C1D15; LoD cut-off, median <20 Du/L (<20- 7060)



Malorni L. et al, EJC 2022

## BioltaLee

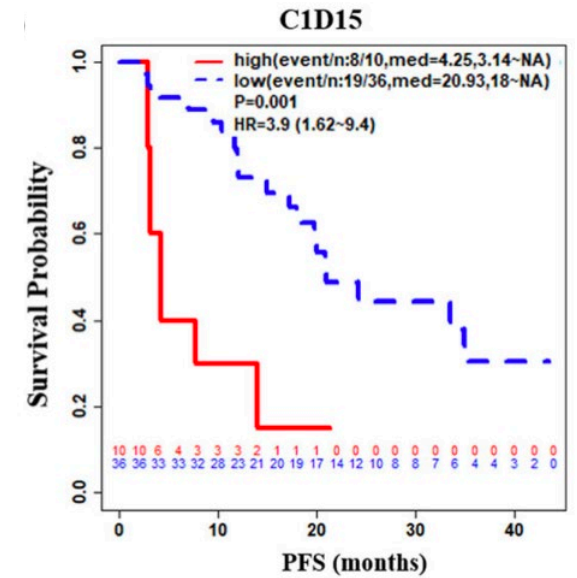
C1D15; LoD cut-off, median 19 Du/L (19–1953)



Malorni L. et al, EJC 2023

## WashU palbo dosing trial

C1D15; LoD cut-off, median <20 Du/L (<20-<20)

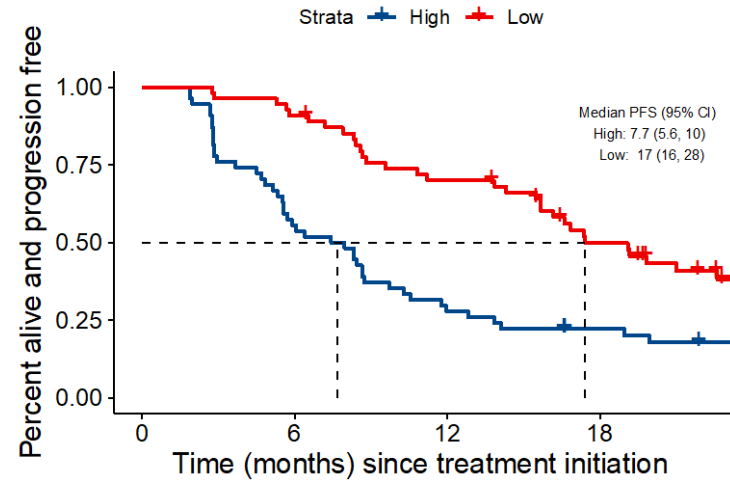


Krishnamurthy, J. npjBC 2022

# TKa DATA IN PATIENTS TREATED WITH ET+CDK4/6i: CYCLE 2 DAY 1

## PYTHIA

C2D1; median cut-off, 52 Du/L (<20, 3533)



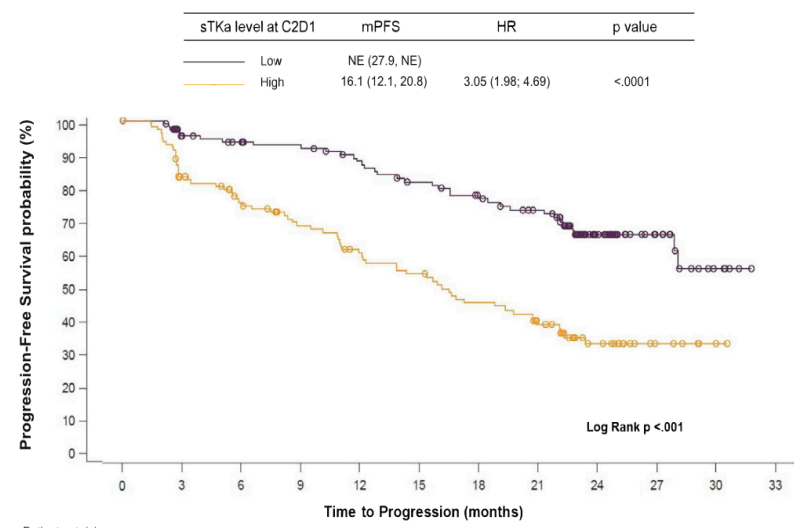
Number at risk

High	54	30	15	10
Low	54	49	37	24

Malorni L. et al, EJC 2022

## BioltaLee

C2D1; median cut-off, 48.1 Du/L (19-3689)



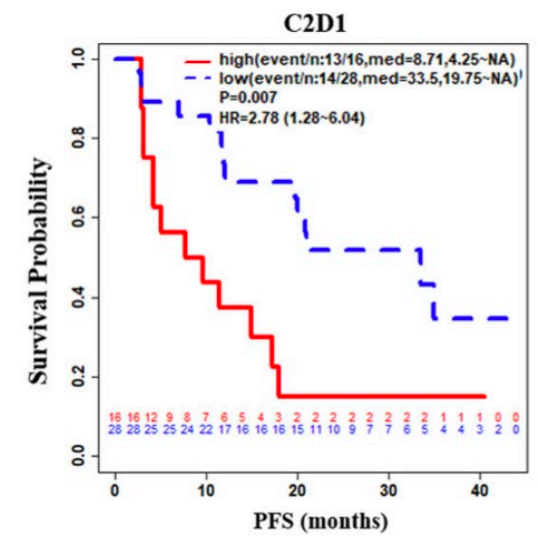
Patients at risk

Low	121	105	99	96	88	80	72	63	37	19	5	0
High	120	91	78	67	57	51	42	33	18	6	2	0

Malorni L. et al, EJC 2023

## WashU palbo dosing trial

C2D1; median cut-off, <20 (<20~54.1)

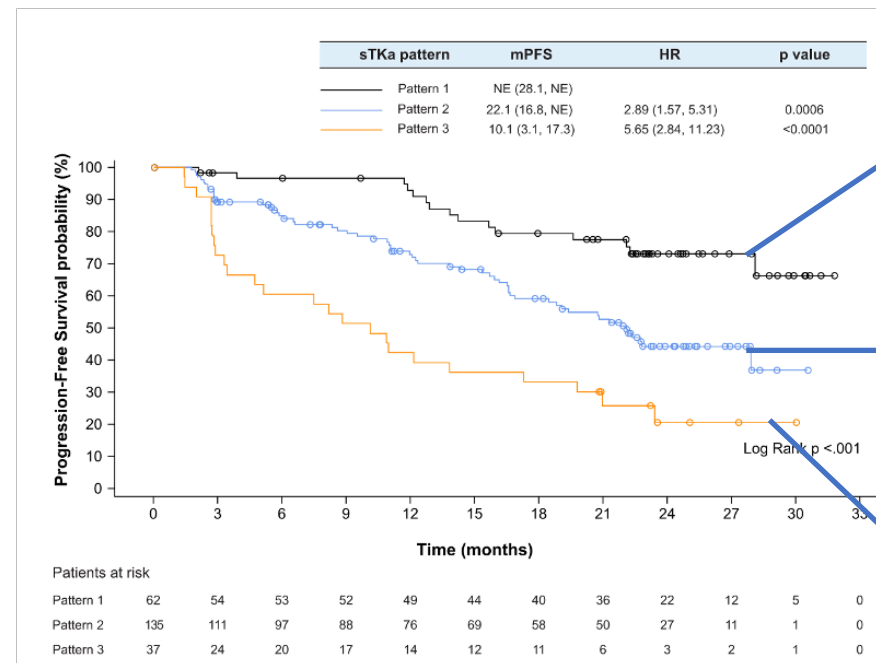
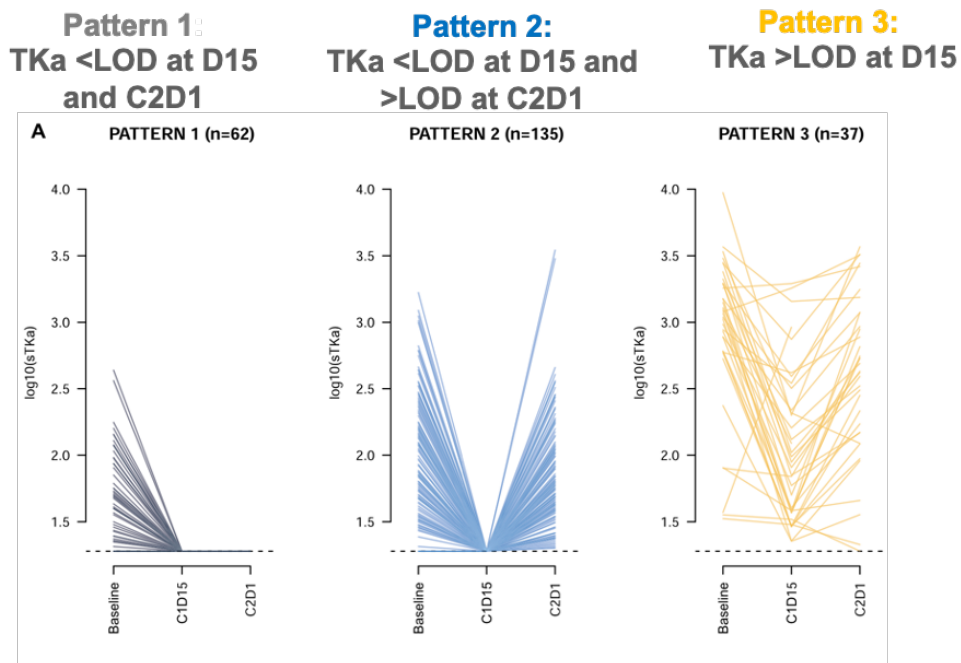


Krishnamurthy, J. npjBC 2022



# Take home messages:

- sTKa appears to be a new promising prognostic and pharmacodynamic biomarker in patients with HR+/HER2- ABC treated with ribociclib plus letrozole as first-line therapy
- Baseline and on-treatment sTKa may give important prognostic and predictive information
- At first imaging, low sTKa is associated with very low risk of clinical/radiological disease progression. Whether sTKa low status can be used to skip radiological examination is currently under investigation (TK IMPACT: NCT04968964)



**26%**  
may forego CDK4/6i in first line?

**58%**  
may need a better HT and/or a continuous CDK4/6i dosing?

**16%**  
may need a better HT+ targeted agent or CHEMO/ADC?

# Take home messages:



Les roses Safrano (Nature morte devant la fenetre ouverte Nice, Place Charles-Flix) (1925)

*Non c'è niente di più difficile per un pittore veramente creativo del dipingere una rosa, perchè prima di tutto deve dimenticare tutte le altre rose che sono state dipinte.*

*Henry Matisse*

# GRAZIE!

**All the patients and their families;  
All the participating centers.**

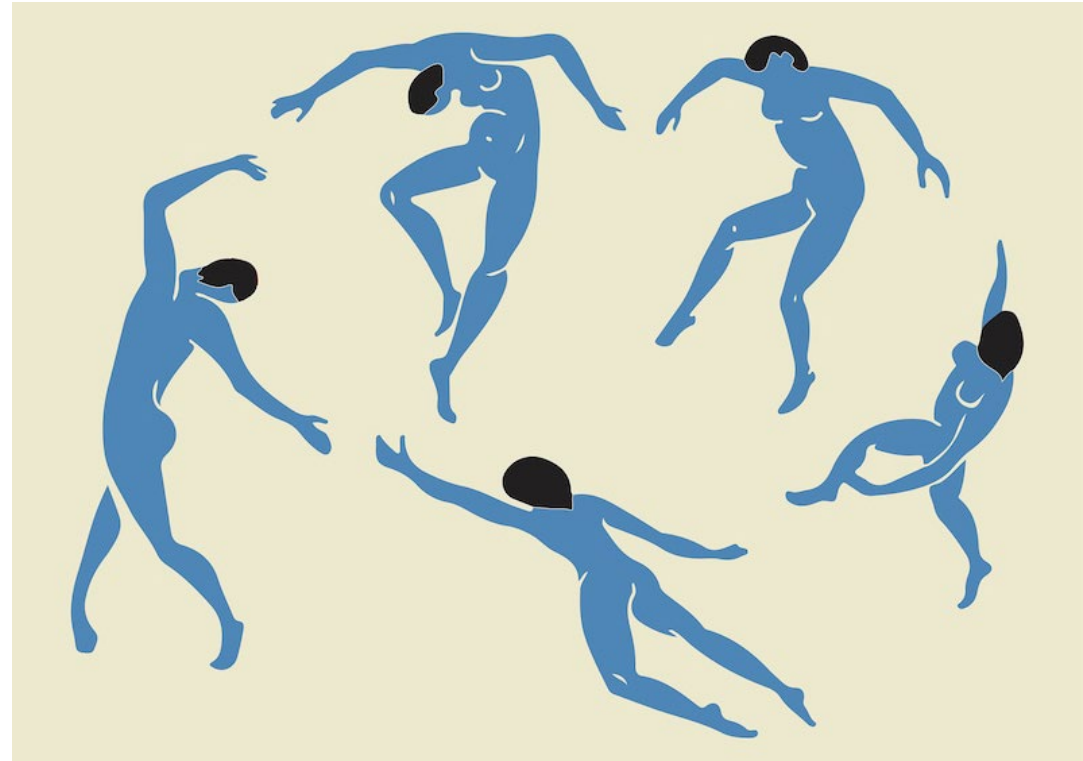
**The BIOItaLee TEAM:**

**Michelino De Laurentiis  
Giampaolo Bianchini  
Grazia Arpino**

**Matteo Benelli  
Maurizio Callari**

**Matteo Suter**

**Donatella Grasso  
Nicola Fenderico  
Daniela Castelletti**



# THANK YOU!

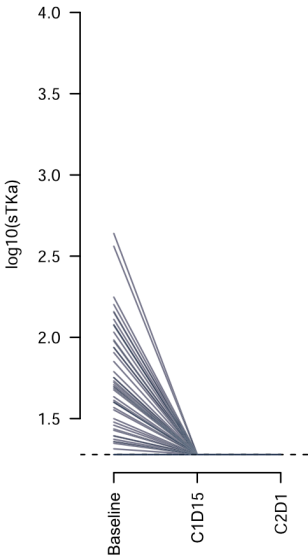


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TKa <LOD at D15 and C2D1

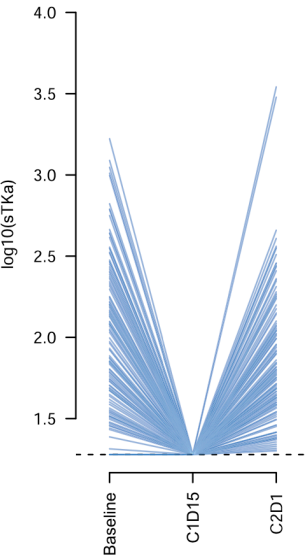
**A** PATTERN 1 (n=62)



**26%**  
may forego CDK4/6i in first line?

**Pattern 2:**  
TKa <LOD at D15 and >LOD at C2D1

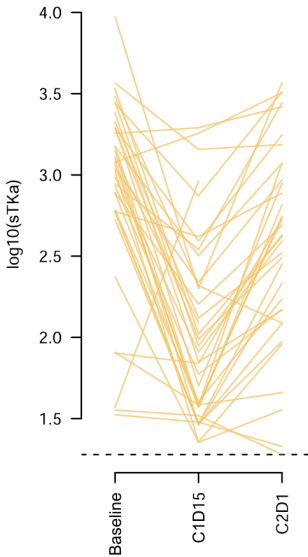
PATTERN 2 (n=135)



**58%**  
may need a better HT and/or a continuous CDK4/6i dosing?

**Pattern 3:**  
TKa >LOD at D15

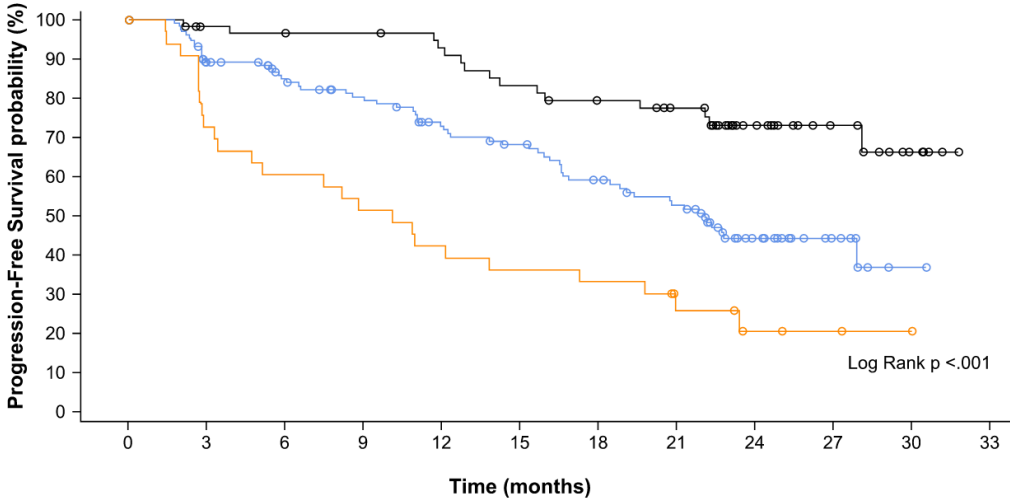
PATTERN 3 (n=37)



**16%**  
may need a better HT+ targeted agent or CHEMO/ADC?

**B**

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Pattern 3	37	24	20	17	14	12	11	6	3	2	1	0