



VII SIMPOSIO NACIONAL  
de ONCOLOGÍA de PRECISIÓN

Vigo, 20 y 21 de febrero de 2025

# Oncología de precisión: problemas y soluciones

Federico Rojo





- Nuevas estrategias diagnósticas moleculares en oncología de precisión
- El uso clínico de NGS y la clasificación ESCAT son suficientes para seleccionar pacientes?
- Podemos ser más precisos en la selección de pacientes con la aplicación de la IA en biomarcadores?



# **Nuevas estrategias diagnósticas moleculares en oncología de precisión**

# ESCAT

## ESMO Scale for Clinical Actionability of Molecular Targets

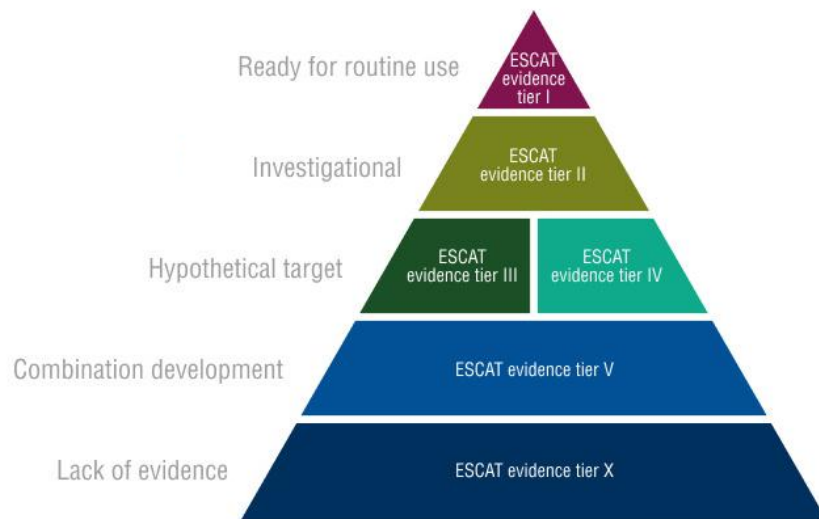
### SPECIAL ARTICLE

## Recommendations for the use of next-generation sequencing (NGS) for patients with advanced cancer in 2024: a report from the ESMO Precision Medicine Working Group

M. F. Mosele<sup>1,2</sup>, C. B. Westphalen<sup>3</sup>, A. Stenzinger<sup>4</sup>, F. Barlesi<sup>1,2,5</sup>, A. Bayle<sup>5,6,7,8</sup>, I. Bièche<sup>9</sup>, J. Bonastre<sup>7,8</sup>, E. Castro<sup>10</sup>, R. Dienstmann<sup>11,12,13</sup>, A. Krämer<sup>14,15</sup>, A. M. Czarnecka<sup>16,17</sup>, F. Meric-Bernstam<sup>18</sup>, S. Michiels<sup>7,8</sup>, R. Miller<sup>19,20</sup>, N. Normanno<sup>21</sup>, J. Reis-Filho<sup>22</sup>, J. Remon<sup>2</sup>, M. Robson<sup>23</sup>, E. Rouleau<sup>24</sup>, A. Scarpa<sup>25</sup>, C. Serrano<sup>11</sup>, J. Mateo<sup>11</sup> & F. André<sup>1,2,5\*</sup>

### Newly added tumour types (2024)

- Advanced non-squamous NSCLC
- Advanced Breast cancer**
- Prostate cancer
- Gastrointestinal stromal tumours (GIST)
- Colorectal cancer
- Sarcoma**
- Ovarian cancers
- Thyroid cancer**
- Cholangiocarcinoma
- Cancer of unknown primary (CUP)**

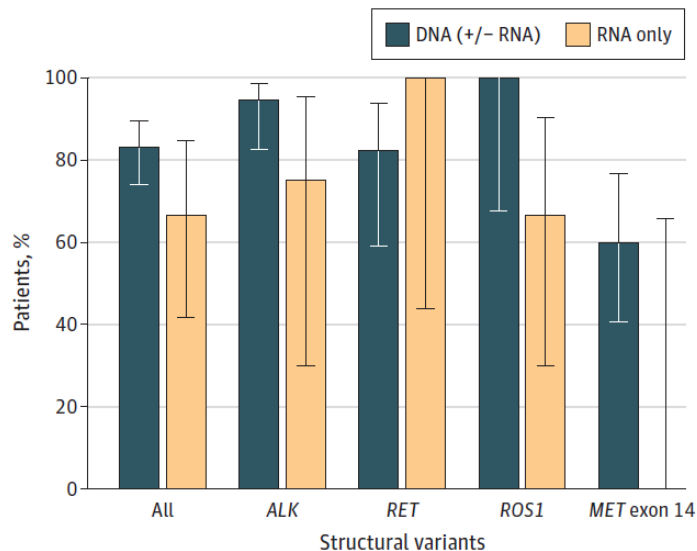




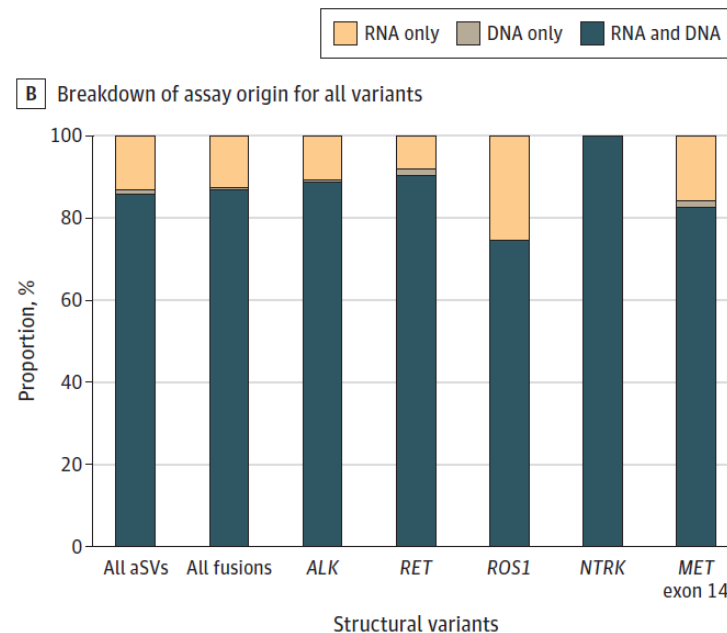
# Actionable Structural Variant Detection via RNA-NGS and DNA-NGS in NSCLC

- Multisite retrospective cohort with 917 patients with advanced NSCLC for both hybrid capture RNA-NGS and DNA-NGS testing
- Concurrent RNA-NGS and DNA-NGS identified 15.3% more patients harboring a SVs compared with DNA-NGS alone, including 14.3% more patients harboring actionable fusions and 18.6% more harboring MET exon 14 skipping alterations

**A** Targeted therapy adherence for eligible patients

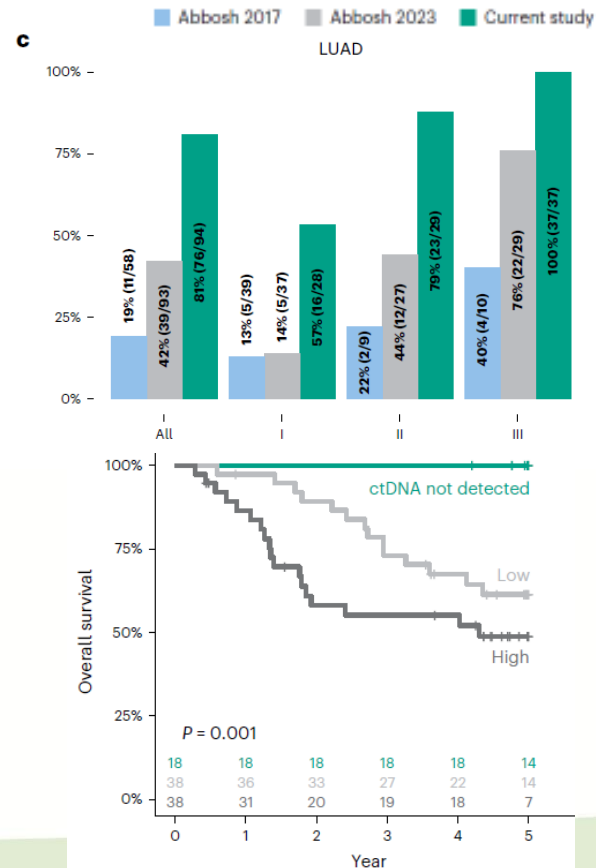
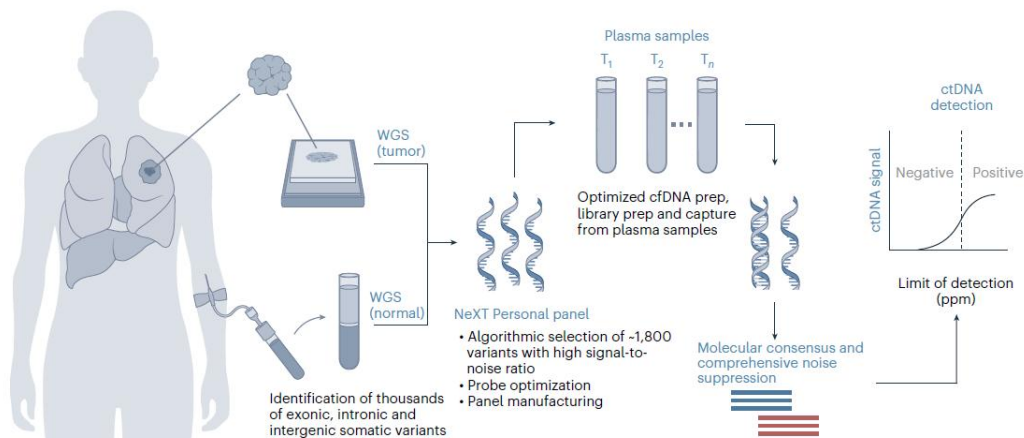


**B** Breakdown of assay origin for all variants



# Ultrasensitive ctDNA NGS for preoperative disease stratification in early-stage lung adenocarcinoma

- NeXT Personal: whole-genome-based, tumor-informed ultrasensitive ctDNA NGS
- Detection at 1–3 ppm of ctDNA with 99.9% specificity
- From 171 patients with early-stage lung cancer from the TRACERx study, 81% of patients with lung adenocarcinoma (LUAD), including 53% with pTNM stage I had ctDNA pre-operatively
- ctDNA predicted worse clinical outcome



# Plasma mutational burden predicts early progression in HR+/HER2- metastatic breast cancer (GEICAM/2014-12 FLIPPER trial)

1. Postmenopausal
2. HR-positive & HER2-negative metastatic breast cancer
3. Endocrine sensitive disease
  - a) relapse after adjuvant endocrine therapy  $\geq 5$  years and disease-free interval  $> 12$  months or
  - b) "de novo" metastatic disease
4. ECOG PS 0-2
5. No prior therapy for MBC

N=189

**Stratification:**

- Visceral vs non visceral metastases
- Disease presentation at study entry: metastatic "de novo" versus recurrent

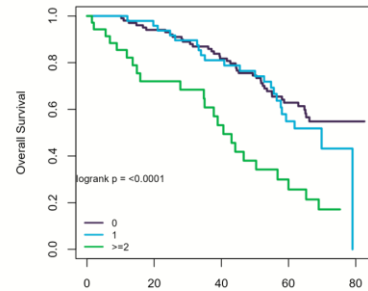
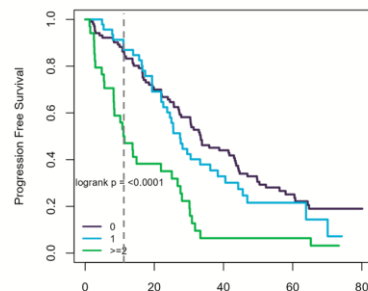
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1:1

**Experimental Arm:**  
Fulvestrant +  
Palbociclib  
n = 94

**Control Arm:**  
Fulvestrant +  
Placebo  
n = 95

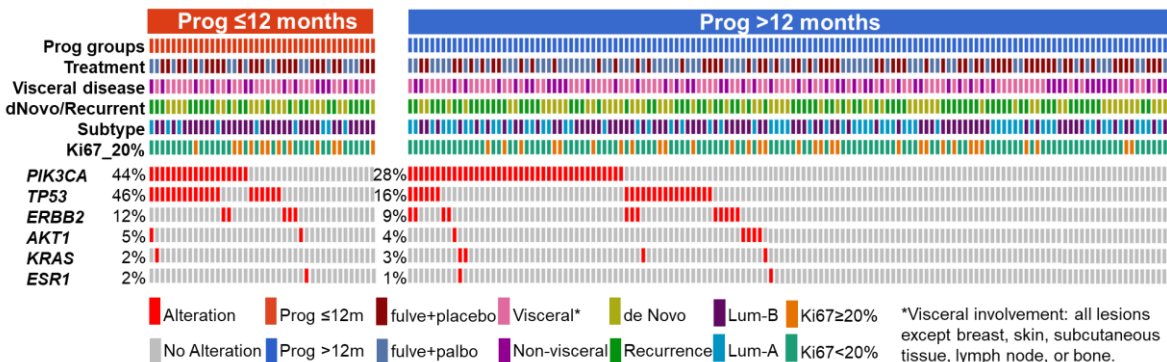
PIK3CA/TP53 # of baseline mutations by PFS/OS



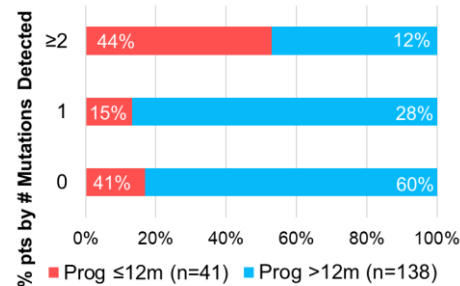
| # mut | N   | Events     | mPFS  | HR (CI95%)       | p-value |
|-------|-----|------------|-------|------------------|---------|
| 0     | 103 | 74 (71.8%) | 33.38 | 1                |         |
| 1     | 49  | 36 (73.5%) | 27.43 | 1.47 (0.97-2.21) | 0.0662  |
| ≥2    | 35  | 32 (91.4%) | 11.22 | 3.6 (2.25-5.76)  | <0.0001 |

| # mut | N   | Events     | mOS   | HR (CI95%)       | p-value |
|-------|-----|------------|-------|------------------|---------|
| 0     | 103 | 40 (38.8%) | NR    | 1                |         |
| 1     | 49  | 22 (44.9%) | 69.85 | 1.34 (0.78-2.27) | 0.2860  |
| ≥2    | 35  | 23 (65.7%) | 40.61 | 2.92 (1.66-5.12) | 0.0002  |

Baseline mutational profiles by progression groups (PFS-12m)



PIK3CA/TP53 # of baseline mutations by PFS-12m



| # mut | N   | Prog ≤12m   | Prog >12m   | OR (CI95%)        | p-value |
|-------|-----|-------------|-------------|-------------------|---------|
| 0     | 100 | 17 (17%)    | 83 (83%)    | 1                 |         |
| 1     | 45  | 6 (13.33%)  | 39 (86.67%) | 0.89 (0.31-2.55)  | 0.8338  |
| ≥2    | 45  | 18 (52.94%) | 16 (47.06%) | 5.84 (2.29-14.87) | 0.0002  |

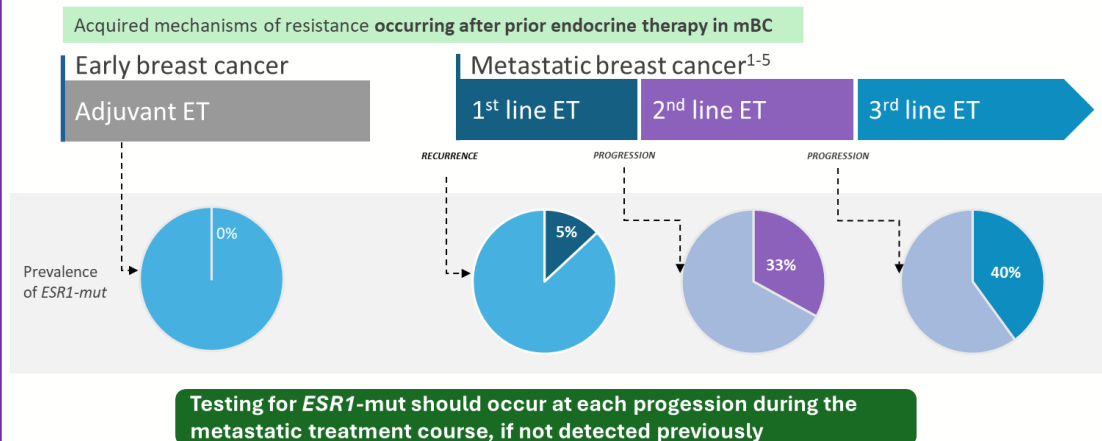
# ESR1 mutations emerge in ~40% metastatic breast cancer after endocrine therapy

## International guidelines for *ESR1* mutation testing in ER+/HER2- metastatic breast cancer



National Comprehensive Cancer Network®

- **Routine testing for *ESR1*-mut at recurrence or PD on ET** (with or without CDK4/6i)
- Use a CLIA-certified assay **on blood or tissue obtained at PD**
- **Blood-based ctDNA testing is preferred**
- Consider **retesting** if *ESR1*-mut is not detected.
- Test for *ESR1*-mut at relapse or progression after ET plus CDK4/6i on plasma or tissue
- Assess for *ESR1*-mut at progression following prior lines of ET with NGS or PCR (tumor tissue or blood)

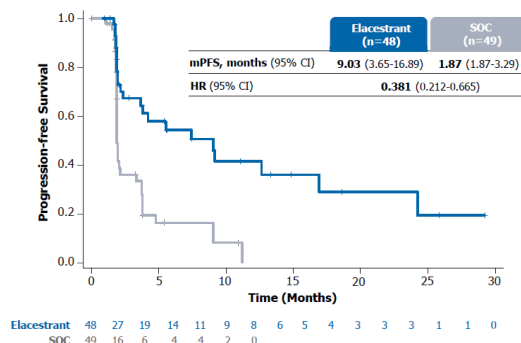


# ESR1 mutations emerge in ~40% metastatic breast cancer after endocrine therapy

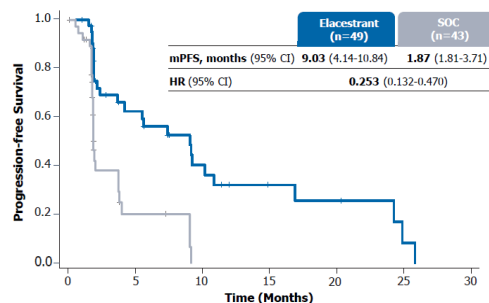
Elacestrant was associated with a PFS benefit across ESR1 mutation variants (D538G, Y537S, and Y537N)

Elacestrant was associated with a clinically meaningful improvement in mPFS independently of ESR1 VAF

ESR1-mut, D538G variant\*

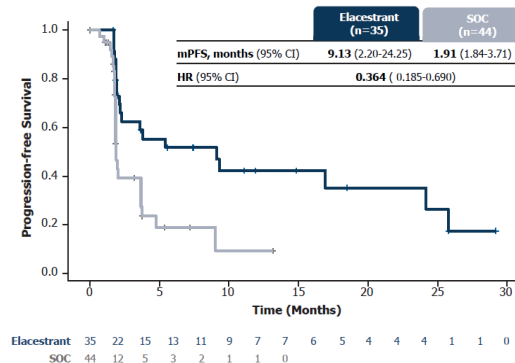


ESR1-mut, Y537S/N variants\*

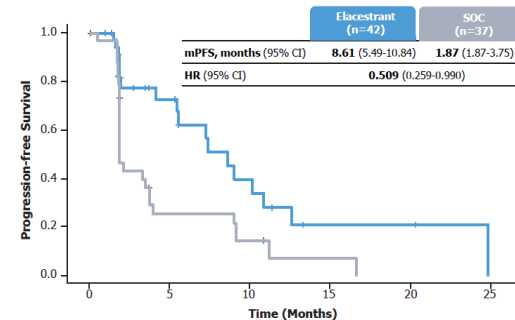


D538G, Y537S and Y537N represent nearly 90% of ESR1 mutations detected

High VAF (≥1.2%)



Low VAF (<1.2%)





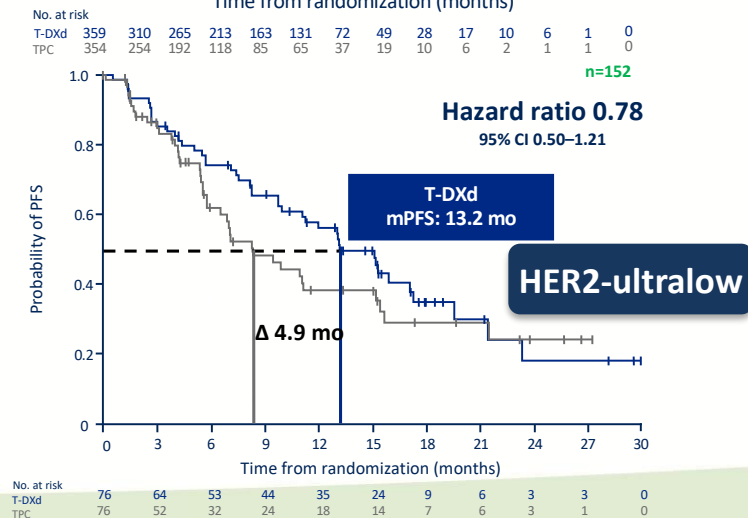
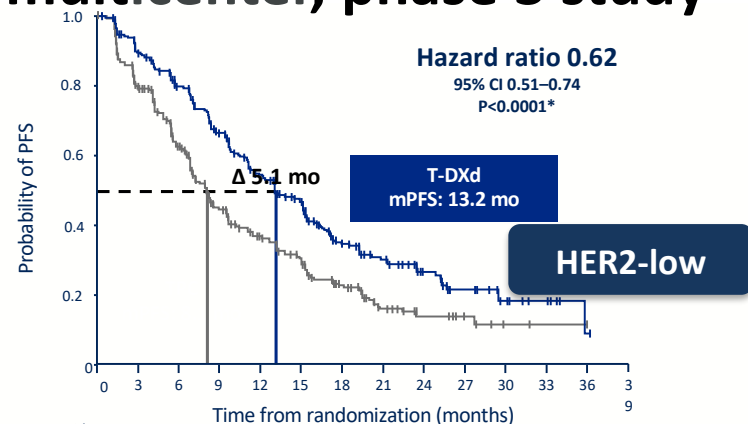
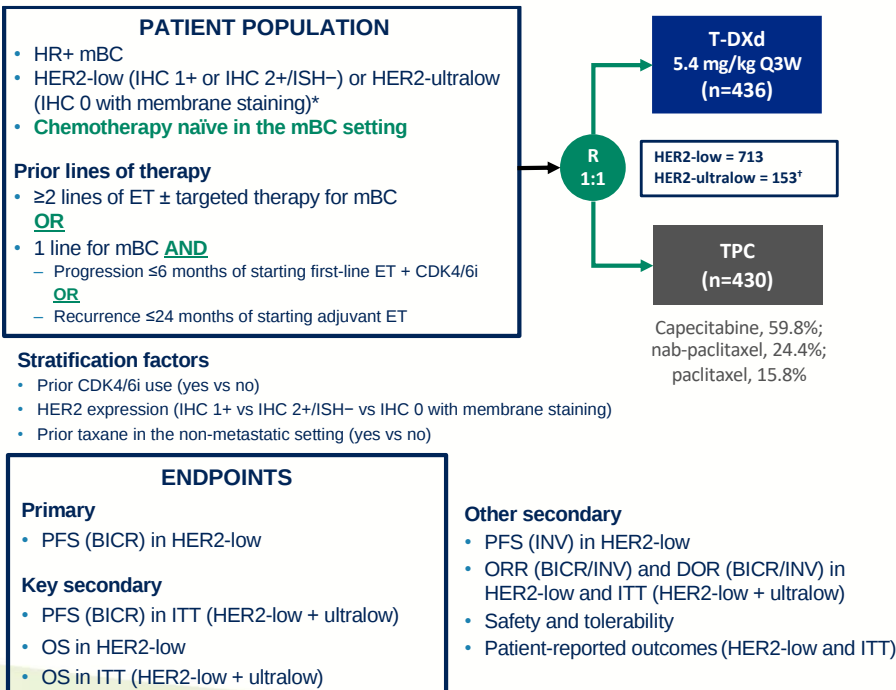
**El uso clínico de NGS y la clasificación ESCAT son suficientes para seleccionar pacientes?**



# DESTINY-Breast06:

## Open-label, randomized, multicenter, phase 3 study

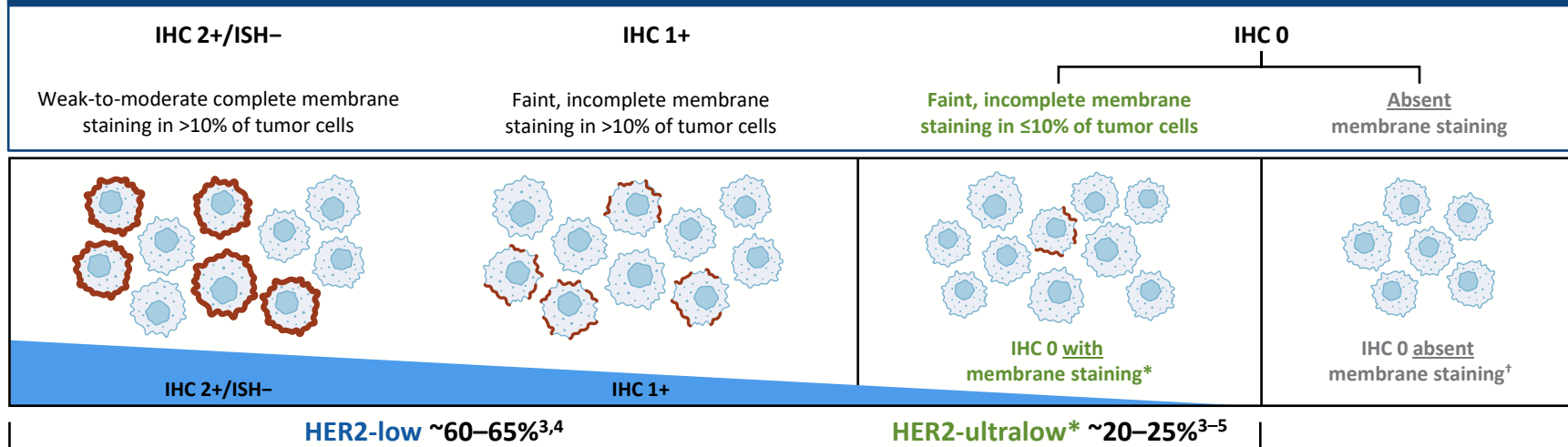
T-DXd demonstrated a statistically significant and clinically meaningful improvement in PFS compared with standard-of-care chemotherapy in **HER2-low**  
PFS improvement with T-DXd vs TPC in **HER2-ultralow** was consistent with results in **HER2-low**



# Targeting 'low' and 'ultralow' HER2-expressing tumors in mBC<sup>1</sup>

- In the DESTINY-Breast06 trial, T-DXd's efficacy is studied in HR+, HER2-low and HER2-ultralow,
- Moves T-DXd to an earlier line of treatment in HER2-low mBC (vs DESTINY-Breast04)
- Expands the proportion of patients who may benefit from T-DXd to ~85% of HR+, HER2- mBC

## HER2 IHC categories within HR+, HER2-negative mBC (per ASCO/CAP<sup>2</sup>):



DESTINY-Breast06 patient population: ~85% of HR+, HER2-negative mBC

\*HER2-ultralow (HER2 IHC 0 with membrane staining of any intensity in ≤10% of tumor cells) was referred to as HER2 IHC >0 to <1+ in the DESTINY-Breast06 protocol; †no membrane staining is observed

1. Curigliano G et al. Presented at: ASCO Annual Meeting; May 31 – June 4, 2024; Chicago, IL. Presentation LBA1000; 2. Wolff AC, et al. *J Clin Oncol.* 2023;41:3867–3872; 3. Denkert C, et al. *Lancet Oncol.* 2021;22:1151–1161; 4. Chen Z, et al. *Breast Cancer Res Treat.* 2023;202:313–323; 5. Mehta S, et al. *J Clin Oncol.* 2024;42(Suppl. 16):e13156 (Abstract)

# DESTINY-PanTumor02: Phase 2 study of T-DXd for HER2-expressing solid tumors

## Key eligibility criteria

- Advanced solid tumors not eligible for curative therapy
- 2L+ patient population
- HER2 expression (IHC 3+ or 2+)
  - Local test or central test by HercepTest if local test not feasible (ASCO/CAP gastric cancer scoring<sup>1</sup>)<sup>a</sup>
- Prior HER2-targeting therapy allowed
- ECOG/WHO PS 0–1

## Baseline characteristics

- 267 patients received treatment; 202 (75.7%) based on local HER2 testing
  - 111 (41.6%) patients were IHC 3+ based on HER2 test (local or central) at enrollment, primary efficacy analysis (all patients)
  - **75 (28.1%) patients were IHC 3+ on central testing**, sensitivity analysis on efficacy endpoints (subgroup analyses)
- Median age was 62 years (23–85) and **109 (40.8%) patients had received ≥3 lines of therapy**

**T-DXd**  
5.4 mg/kg Q3W

40 per cohort<sup>b</sup>

|                                                                                     |                                 |
|-------------------------------------------------------------------------------------|---------------------------------|
|  | <b>Endometrial cancer</b>       |
|  | <b>Cervical cancer</b>          |
|  | <b>Ovarian cancer</b>           |
|  | <b>Bladder cancer</b>           |
|  | <b>Other tumors<sup>c</sup></b> |
|  | <b>Biliary tract cancer</b>     |
|  | <b>Pancreatic cancer</b>        |

## Primary endpoint

- Confirmed ORR (investigator)

## Secondary endpoints

- DOR, DCR, PFS, OS
- Safety

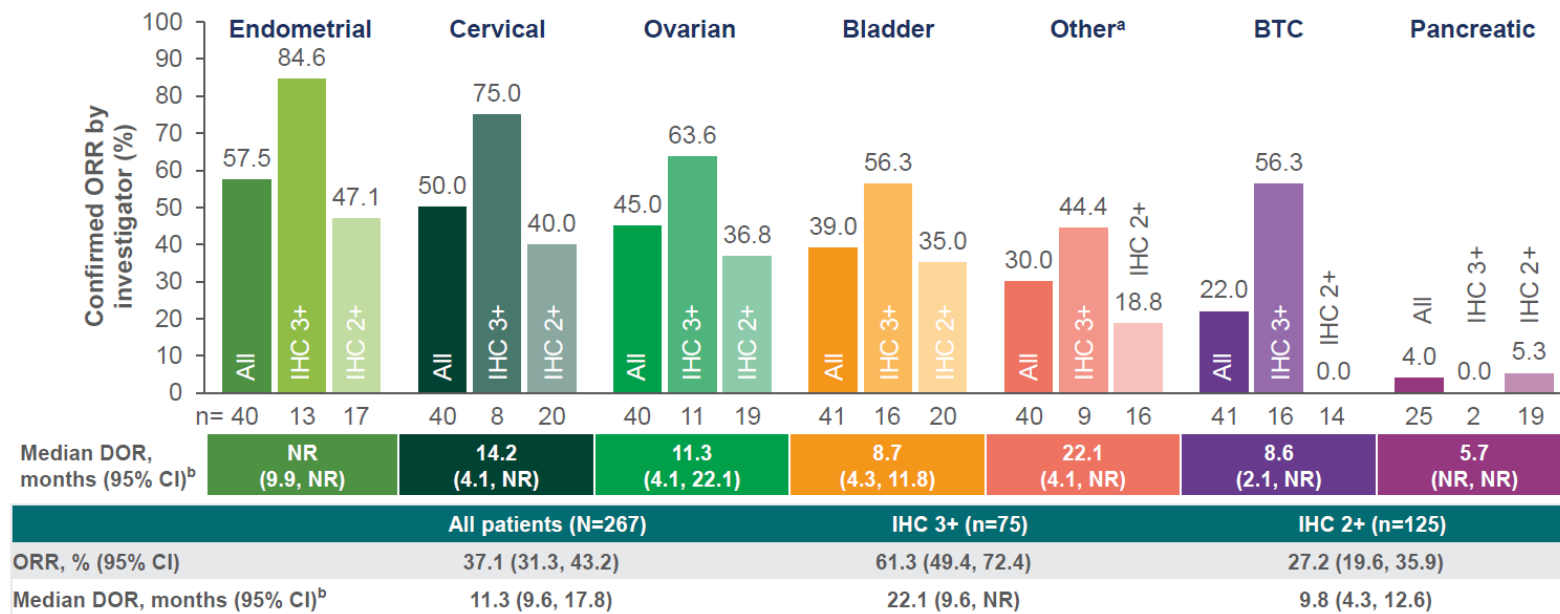
## Exploratory analysis

- Subgroup analyses by HER2 status

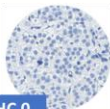
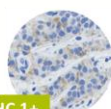
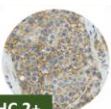
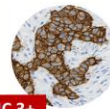
Primary analysis  
data cutoff: Jun 8, 2023  
Median follow up: 12.75 mo

# DESTINY-PanTumor02: Phase 2 study of T-DXd for HER2-expressing solid tumors

## Objective response and duration of response



# ASCO/CAP have released formal HER2 scoring guidelines for breast and gastroesophageal cancers only

|                                      |  IHC 0                                               |  IHC 1+                  |  IHC 2+                                                                                                                                                                                                   |  IHC 3+                                                                                                                                       |                                                                                                                                                                                  |                                                                                                      |
|--------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------|
| <b>Breast<br/>ASCO/CAP2018</b>       | No staining observed or incomplete, faint/barely perceptible membrane staining in $\leq 10\%$ of tumour cells<br><b>HER2 Negative</b> | Incomplete, faint/barely perceptible membrane staining in $>10\%$ of tumour cells<br><b>HER2 Negative</b> | Weak to moderate complete membrane staining in $>10\%$ of tumour cells<br><b>HER2 Equivocal</b><br>ISH positive: HER2/CEP17 ratio $\geq 2$ and HER2 $\geq 4$ , or ratio $< 2$ and HER2 $> 6$ (if IHC 2+/3+)                                                                                  | Complete, intense membrane staining in $>10\%$ of tumour cells<br><b>HER2 Positive</b>                                                                                                                                           |                                                                                                                                                                                  |                                                                                                      |
| <b>GEA<br/>ASCO/CAP2016</b>          | No staining observed or faint membrane staining in $<10\%$ of tumour cells<br><b>HER2 Negative</b>                                    | Faint/barely perceptible membrane staining in $\geq 10\%$ of tumour cells<br><b>HER2 Negative</b>         | Weak to moderate complete, basolateral or lateral membrane in $\geq 10\%$ of tumour cells<br><b>Biopsy:</b> Cluster of at least 5 cells with complete/basolateral/lateral weak/moderate staining<br><b>HER2 Equivocal</b><br>ISH positive: HER2/CEP17 $\geq 2$ or ratio $< 2$ and HER2 $> 6$ | Complete, basolateral or lateral intense membrane in $>10\%$ of tumour cells<br><b>Biopsy:</b> Cluster of at least 5 cells with complete/basolateral/lateral intense staining<br><b>HER2 Positive</b>                            |                                                                                                                                                                                  |                                                                                                      |
| <b>Colorectal<br/>HERACLES</b>       | No staining observed or faint membrane staining in $\leq 10\%$ of tumour cells<br><b>HER2 Negative</b>                                | Faint/barely perceptible membrane staining in $>10\%$ of tumour cells<br><b>HER2 Negative</b>             | Weak to moderate complete, basolateral or lateral membrane in $>10\%$ but $< 50\%$ of cells<br><b>HER2 Negative</b>                                                                                                                                                                          | Weak to moderate complete, basolateral or lateral membrane in $>50\%$ of tumour cells<br><b>HER2 Equivocal</b><br>Mandatory IHC retesting to confirm 50%; ISH required<br>ISH positive: HER2/CEP17 $\geq 2$ in $\geq 50\%$ cells | Complete, basolateral or lateral intense membrane in $>10\%$ but $< 50\%$ of cells<br><b>HER2 Conditionally positive</b><br>Mandatory IHC retesting to confirm 10%; ISH required | Complete, basolateral or lateral intense membrane in $>50\%$ of tumour cells<br><b>HER2 Positive</b> |
| <b>Endometrial<br/>Buza, N et al</b> | No staining observed or faint membrane staining in $\leq 10\%$ of tumour cells<br><b>HER2 Negative</b>                                | Faint/barely perceptible membrane staining in $>10\%$ of tumour cells<br><b>HER2 Negative</b>             | Weak to moderate complete, basolateral or lateral membrane in $>10\%$ but $< 30\%$ of cells<br><b>HER2 Equivocal</b>                                                                                                                                                                         | Weak to moderate complete, basolateral or lateral membrane in $>30\%$ of cells<br><b>HER2 Positive</b>                                                                                                                           | Complete, basolateral or lateral intense membrane in $>10\%$ but $< 30\%$ of cells<br><b>HER2 Positive</b>                                                                       | Complete, basolateral or lateral intense membrane in $>30\%$ of cells<br><b>HER2 Positive</b>        |



# Various HER2 scoring used across endometrial serous, colorectal, urothelial or lung carcinomas

The use of different algorithms for HER2 testing may significantly affect the HER2 positivity rate and the proportion of false positive and false negative results

## Colorectal Carcinoma

| HERACLES trial<br>(Valtorta E. et al.) | ASCO/CAP 2018 Breast | ASCO/CAP 2017<br>Gastric | Other |
|----------------------------------------|----------------------|--------------------------|-------|
| 30%                                    | 16.3%                | 49.4%                    | 4.3%  |

Other responses: original US Food and Drug Administration scoring criteria—HerceptTest (2), positive versus negative (1), CAP 2013 breast guideline (1), based on ordering physician (1), and MyPathway (1).

## Urothelial Carcinoma

| HER2 Score (% of specimens) |     |     |     |     |
|-----------------------------|-----|-----|-----|-----|
| ASCO/CAP<br>Criteria        | 0   | 1+  | 2+  | 3+  |
| Breast 2018                 | 37% | 45% | 5%  | 6%  |
| Gastric 2017                | 37% | 16% | 43% | 11% |

## Endometrial Serous Carcinoma

| Fader et al. | ASCO/CAP 2018<br>Breast | ASCO/CAP 2013<br>Breast | ASCO/CAP 2007<br>Breast | “overall”<br>positive/negative | HerceptTest<br>package insert<br>for Breast | Other |
|--------------|-------------------------|-------------------------|-------------------------|--------------------------------|---------------------------------------------|-------|
| 16%          | 69.4%                   | 1.8%                    | 1.4%                    | 3.6%                           | 5%                                          | 2.9%  |

Other responses: CAP American Society for Clinical Pathology/ASCO gastroesophageal adenocarcinoma (4) and in-house guideline (1)

## Non-Small Cell Lung Cancer

| HER2 Score (% of specimens) |     |     |     |    |
|-----------------------------|-----|-----|-----|----|
| ASCO/CAP<br>Criteria        | 0   | 1+  | 2+  | 3+ |
| Gastric 2017                | 28% | 51% | 17% | 3% |



**Podemos ser más precisos en la selección de pacientes con la aplicación de la IA en biomarcadores?**

# Datopotamab-DXd vs docetaxel for previously treated advanced or metastatic NSCLC: Randomized Phase III TROPION-Lung01 study



2024 World Conference  
on Lung Cancer

SEPTEMBER 7-10, 2024  
SAN DIEGO, CA USA

INTERNATIONAL COLLABORATIVE

IMPACTFUL INSPIRATIONAL INFORMATIVE

#WCLC24

wclc2024.iaslc.org

## TROPION-Lung01

### Study Design (NCT04656652)<sup>1</sup>

#### Key Eligibility Criteria

- NSCLC (stage IIIB, IIIC, or IV)
- ECOG PS of 0 or 1
- No prior docetaxel

#### Without AGA\*

- 1 or 2 prior lines, including platinum CT and anti-PD-(L)1 mAb therapy

#### With AGA

- Positive for *EGFR*, *ALK*, *NTRK*, *BRAF*, *ROS1*, *MET* exon 14 skipping, or *RET*
- 1 or 2 prior approved targeted therapies + platinum-based CT, and ≤1 anti-PD-(L)1 mAb

R 1:1

**Dato-DXd**  
6 mg/kg q3w  
N=299

**Docetaxel**  
75 mg/m<sup>2</sup> q3w  
N=305

#### Stratified by:

Histology<sup>†</sup>, AGA<sup>‡</sup>, anti-PD-(L)1 mAb included in most recent prior therapy, geography<sup>§</sup>

**Dual Primary Endpoints:** PFS by BICR; OS

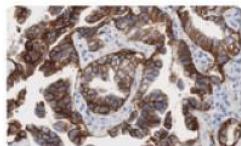
**Secondary Endpoints:** ORR by BICR; DOR by BICR; Safety

Dr Marina Chiara Garassino | Normalized Membrane Ratio of TROP2 by Quantitative Continuous Scoring is Predictive of Clinical Outcomes in TROPION-Lung01

# TROP2 Normalized Membrane Ratio (NMR) measured by Quantitative Continuous Scoring (QCS)

QCS is a novel, fully-supervised computational pathology approach that precisely quantifies and locates targets like TROP2

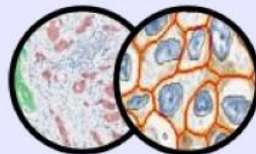
## 1 IHC with TROP2 Assay



## 2 Whole Slide Imaging



## 3 Automated Image Analysis (QCS)



## 4 Patient Biomarker Status Determination

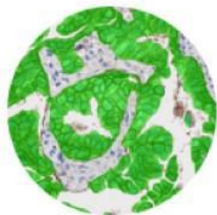


≥75% of tumor cells with  
TROP2 NMR ≤0.56

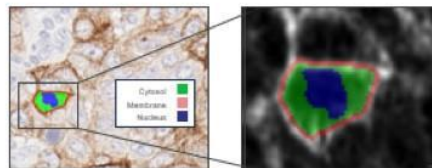


<75% of tumor cells with  
TROP2 NMR ≤0.56\*

### Differentiates tumor from non-tumor



### Measures OD in each tumor cell



Membrane and cytoplasm optical  
density (OD)

### Calculates TROP2 NMR for every tumor cell

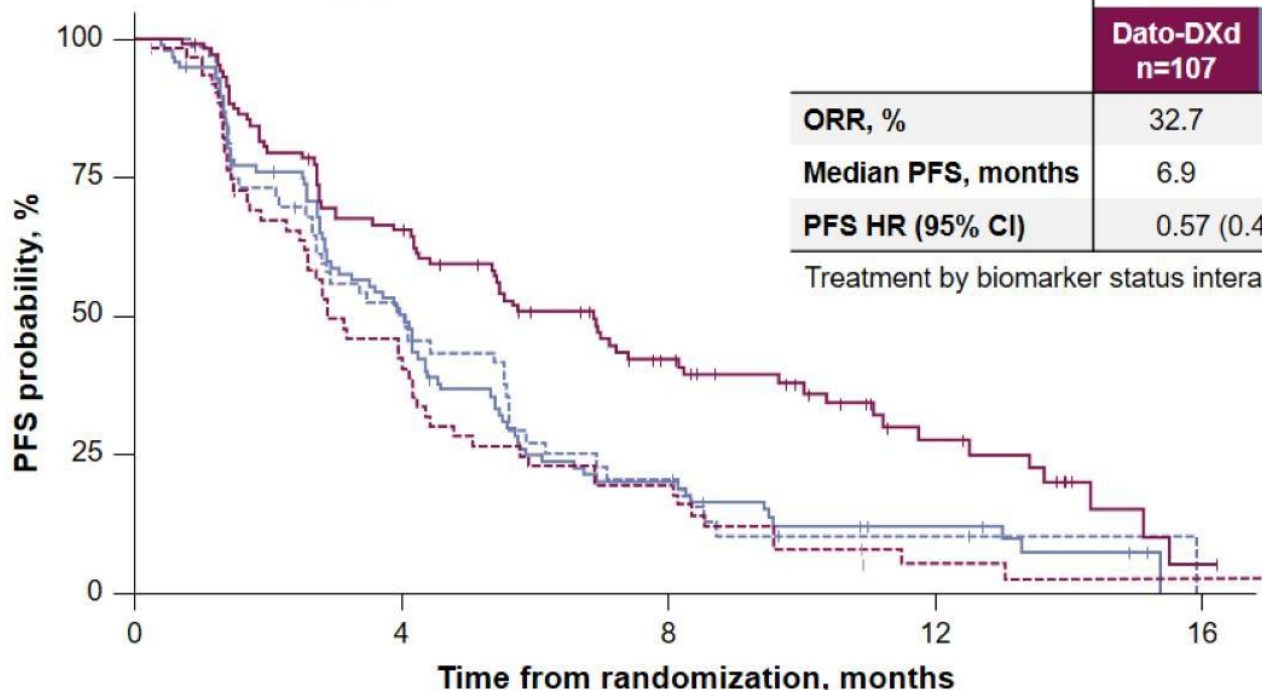
$$\frac{\text{Membrane OD}}{\text{Membrane OD} + \text{Cytoplasm OD}}$$

*Lower NMR → higher cytoplasm proportion*

# Overall BEP: Efficacy by TROP2 QCS-NMR Status

*TROP2 QCS-NMR positivity is predictive for longer PFS with Dato-DXd in the biomarker-evaluable population*

Biomarker-evaluable population, n=352



|                    | TROP2 QCS-NMR+    |                    | TROP2 QCS-NMR-   |                   |
|--------------------|-------------------|--------------------|------------------|-------------------|
|                    | Dato-DXd<br>n=107 | Docetaxel<br>n=107 | Dato-DXd<br>n=65 | Docetaxel<br>n=73 |
| ORR, %             | 32.7              | 10.3               | 16.9             | 15.1              |
| Median PFS, months | 6.9               | 4.1                | 2.9              | 4.0               |
| PFS HR (95% CI)    | 0.57 (0.41–0.79)  |                    | 1.16 (0.79–1.70) |                   |

Treatment by biomarker status interaction: p=0.0063

— Dato-DXd, QCS-NMR+  
- - Dato-DXd, QCS-NMR-  
— Docetaxel, QCS-NMR+  
- - Docetaxel, QCS-NMR-



# Mensajes finales

- La NGS de captura híbrida basada en RNA mejora la capacidad de detectar fusiones génicas
- La biopsia líquida crece como herramienta en medicina de precisión (necesidad de alta sensibilidad!!!)
- Algunas dianas ESCAT I requieren aproximaciones basadas en técnicas básicas como la IHC
- HER2 se extiende como biomarcador pantumor, pero abre interrogantes sobre los criterios de interpretación en algunos tumores y sobre el límite más bajo de expresión (HER2-ultralow)
- La IA aparece por primera vez como herramienta imprescindible para la selección de pacientes en medicina de precisión