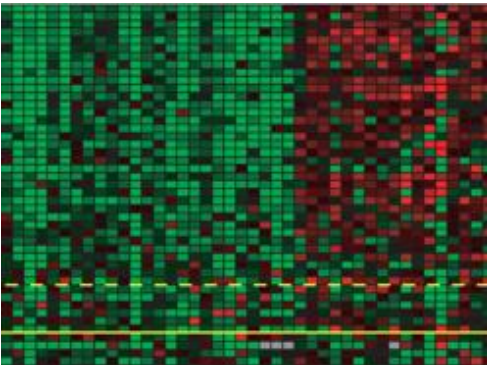
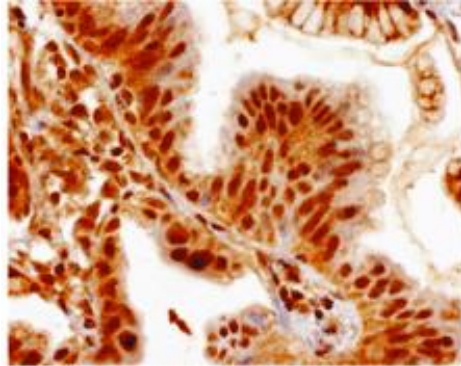
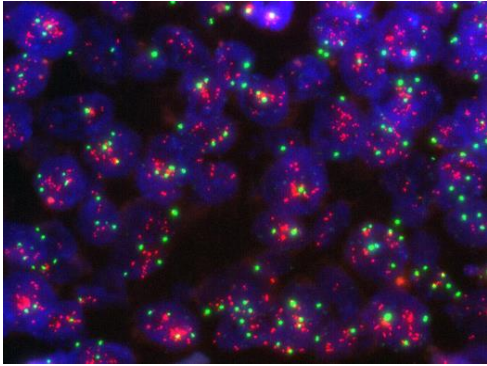


Principles of Laboratory Testing for Treatment Decision-Making in Metastatic Breast Cancer

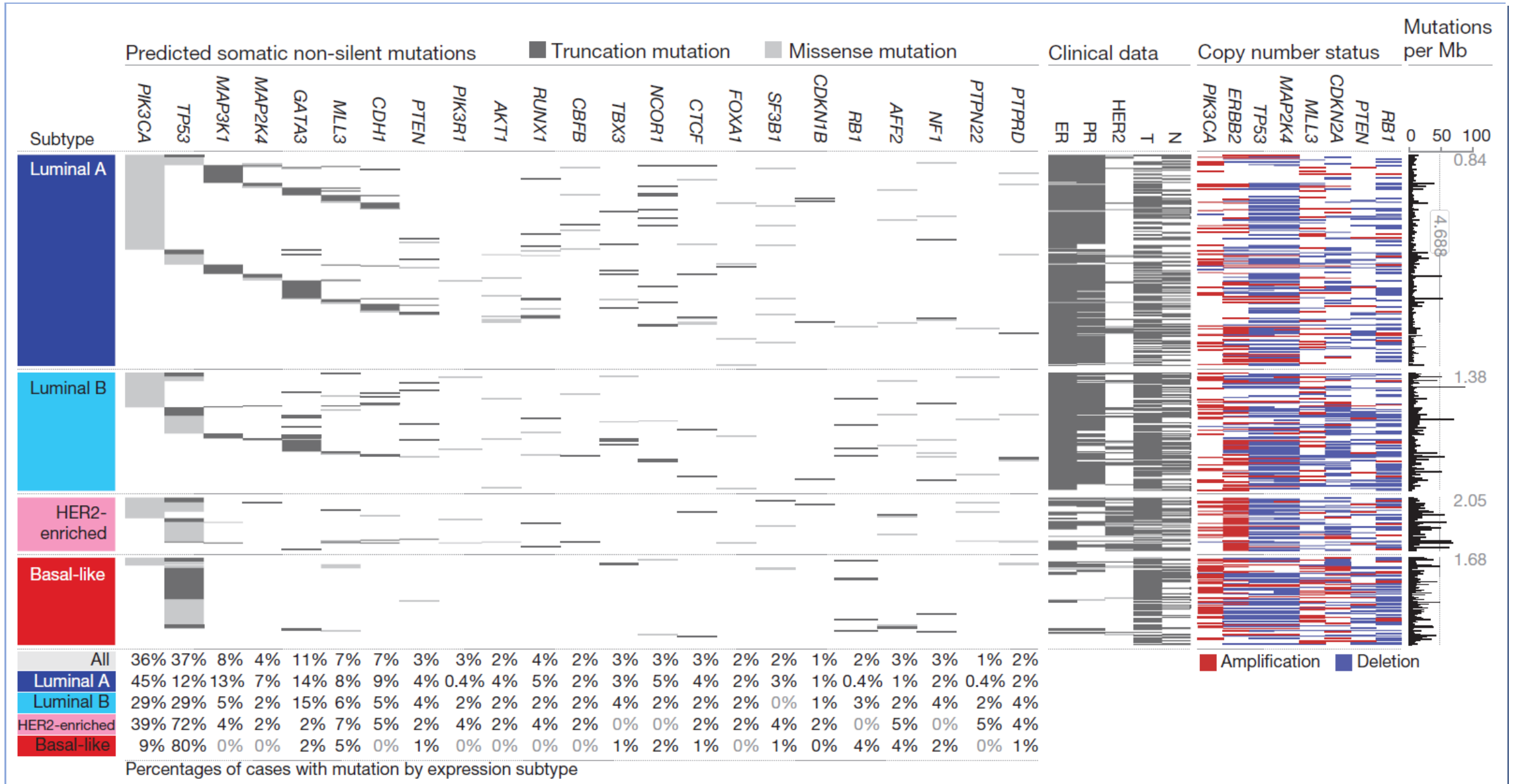


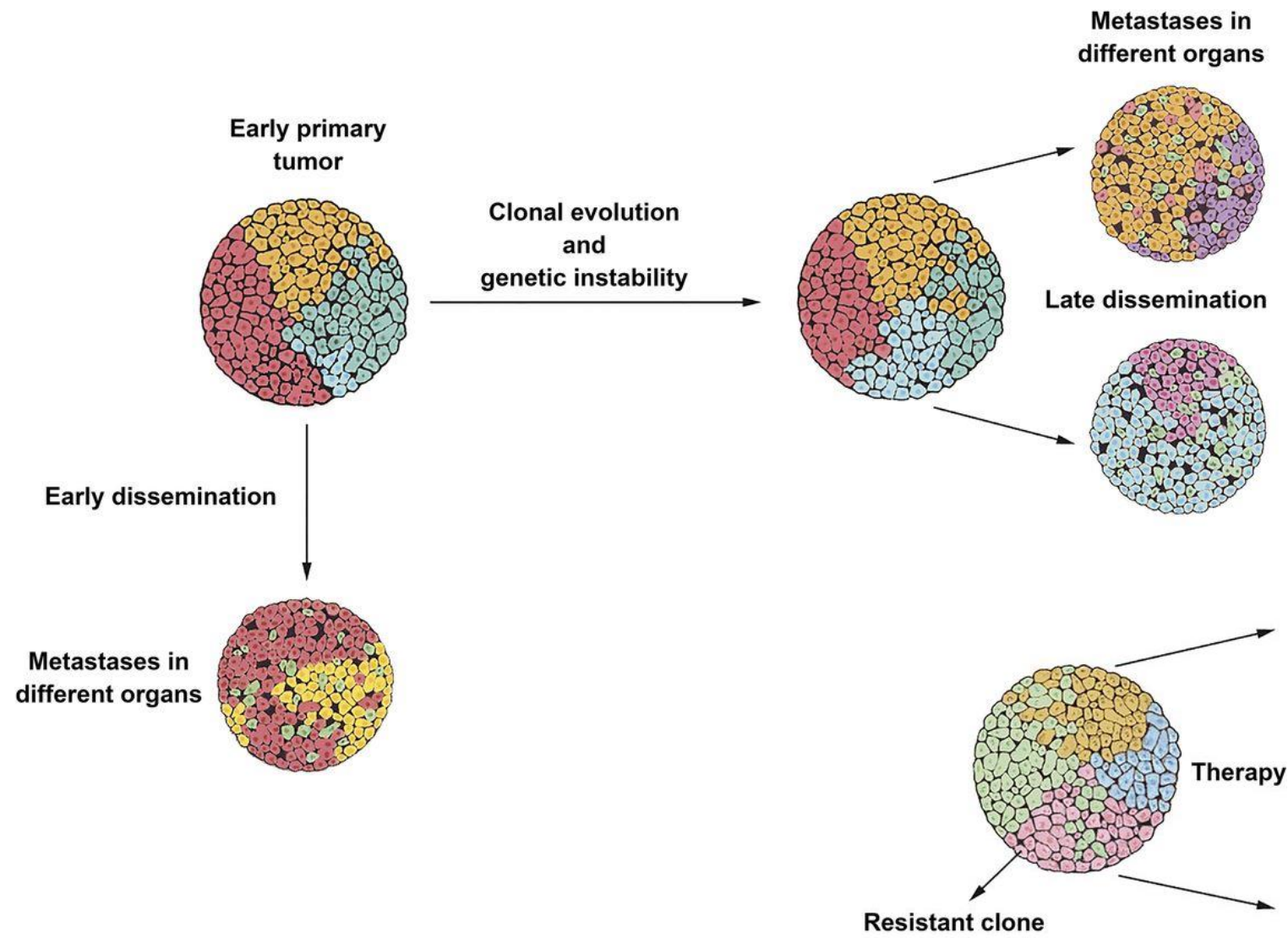
Debu Tripathy, MD

Department of Breast Medical Oncology

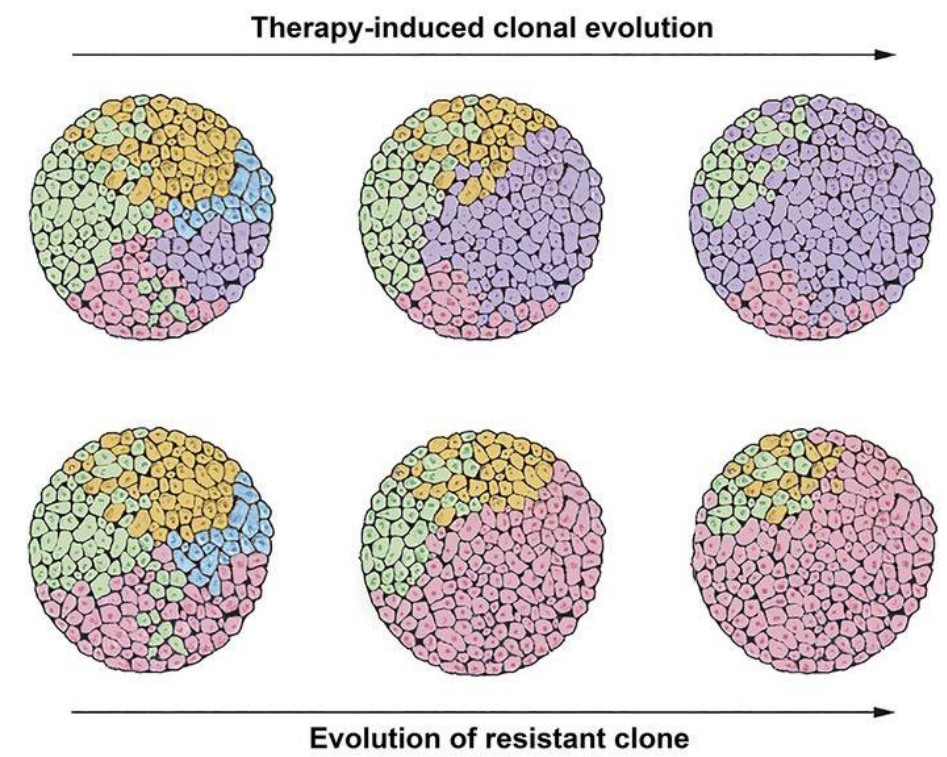
**The University of Texas MD Anderson Cancer Center,
Houston, TX**

Genomic Landscape in Receptor Subtypes of Breast Cancer

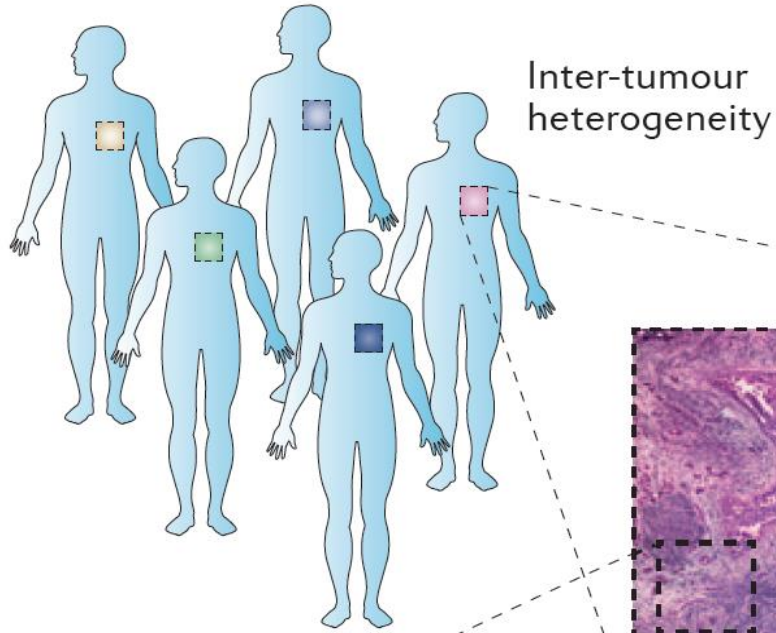




The Fluid State of the Tumor Micro-Ecosystem: Genomic Evolution and Adaptive Resistance

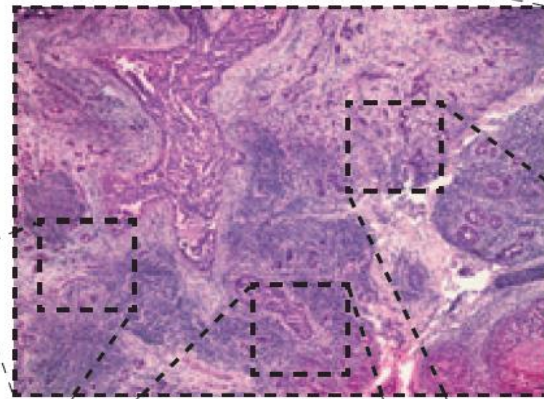


Breast Cancer Heterogeneity Patients and Tumors



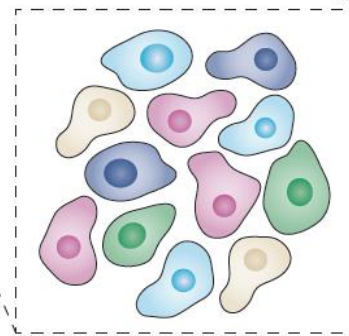
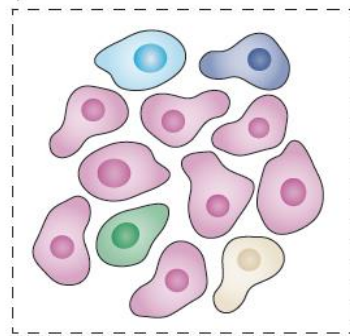
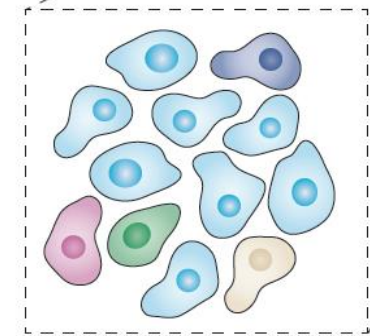
Between patients: **Subtypes and clinical courses**

Within a patient: **Evolution, divergent paths**



Intra-tumour heterogeneity

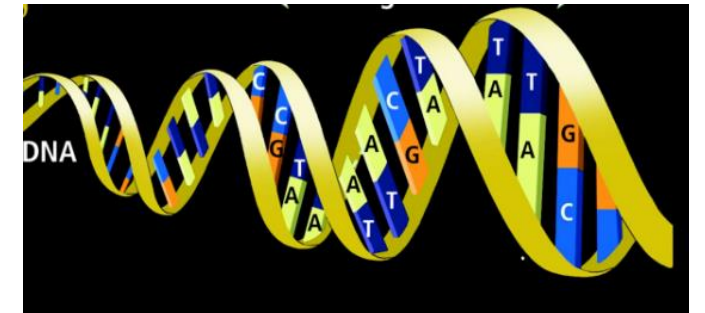
Within Tumours:
subclones



Dominance of clone 1

Dominance of clone 2

Mixed dominance



What, How, and Why Do We Test in Metastatic Breast Cancer?

Test	Material Measured	Purpose
Estrogen/Progesterone receptor	Protein	Effectiveness of hormonal (endocrine) therapies
HER2 receptor	Protein and/or DNA	Effectiveness of HER2-targeted therapies
PD-L1	Protein	Effectiveness of immunotherapy
Germline (Hereditary) gene mutations	Germline DNA	Genetic counseling, cancer prevention, Effectiveness of PARP inhibitors therapy
Tumor Mutations	Tumor DNA	Tumor gene mutations that show potential activity of specific targeted therapies
Tumor Gene Expression (RNA)	Tumor RNA	Expression profiles to optimize and personalize treatment (experimental)

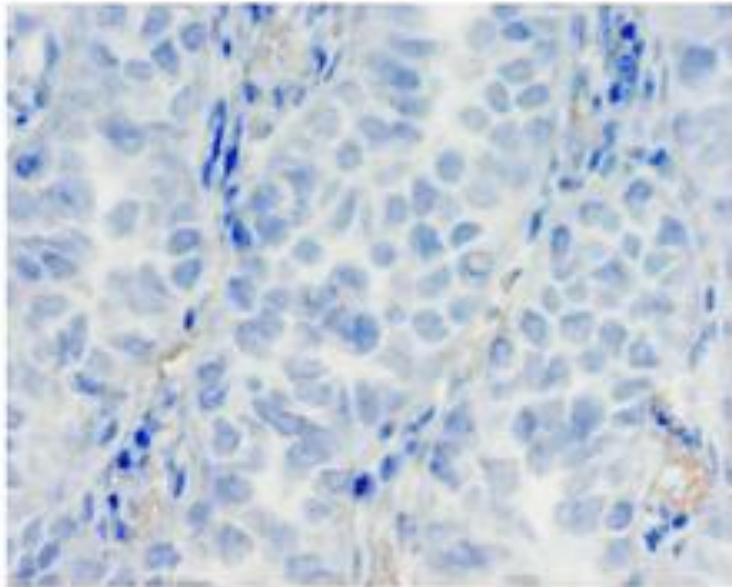
Recommendations for Testing in Metastatic Breast Cancer

- **Imaging** (CT chest, abdomen, pelvis and bone scan or whole body PET/CT scan, if indicated brain MRI)
- **Genetic counseling and germline DNA testing**
- **Biopsy of metastatic lesion for histological confirmation of cancer, ER, PR and HER2 testing**
- **After 1st line therapy (or sooner), tumor gene next generation DNA and/or RNA sequencing**

Estrogen and Progesterone Receptor Immuno-Staining: Scoring by % Staining and Intensity

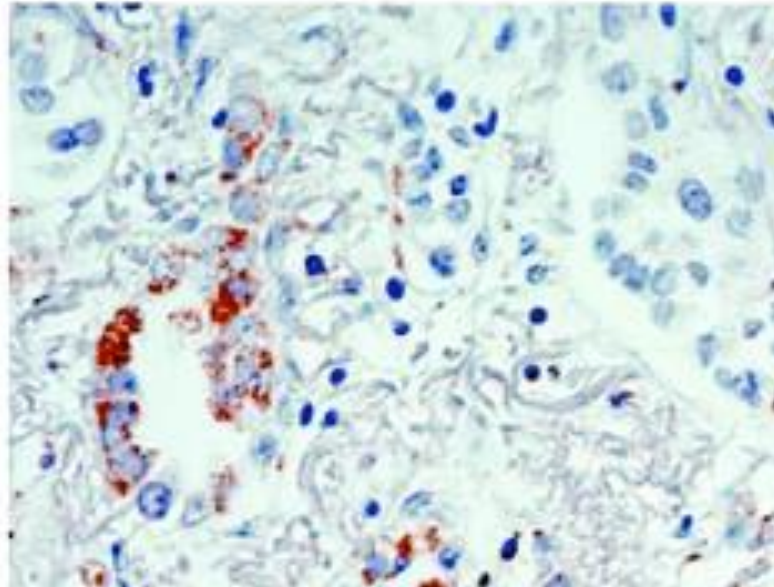
Estrogen Receptor Assay Examples

Negative



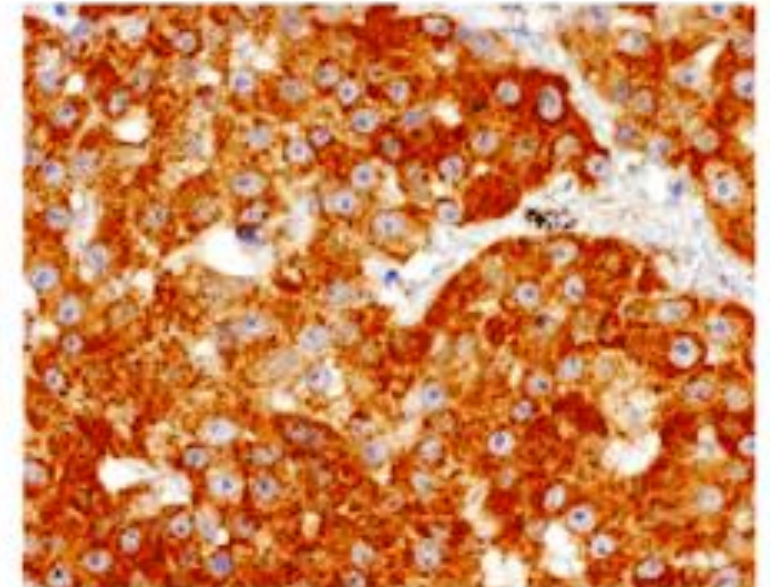
$$0 + 0 = 0$$

Weak



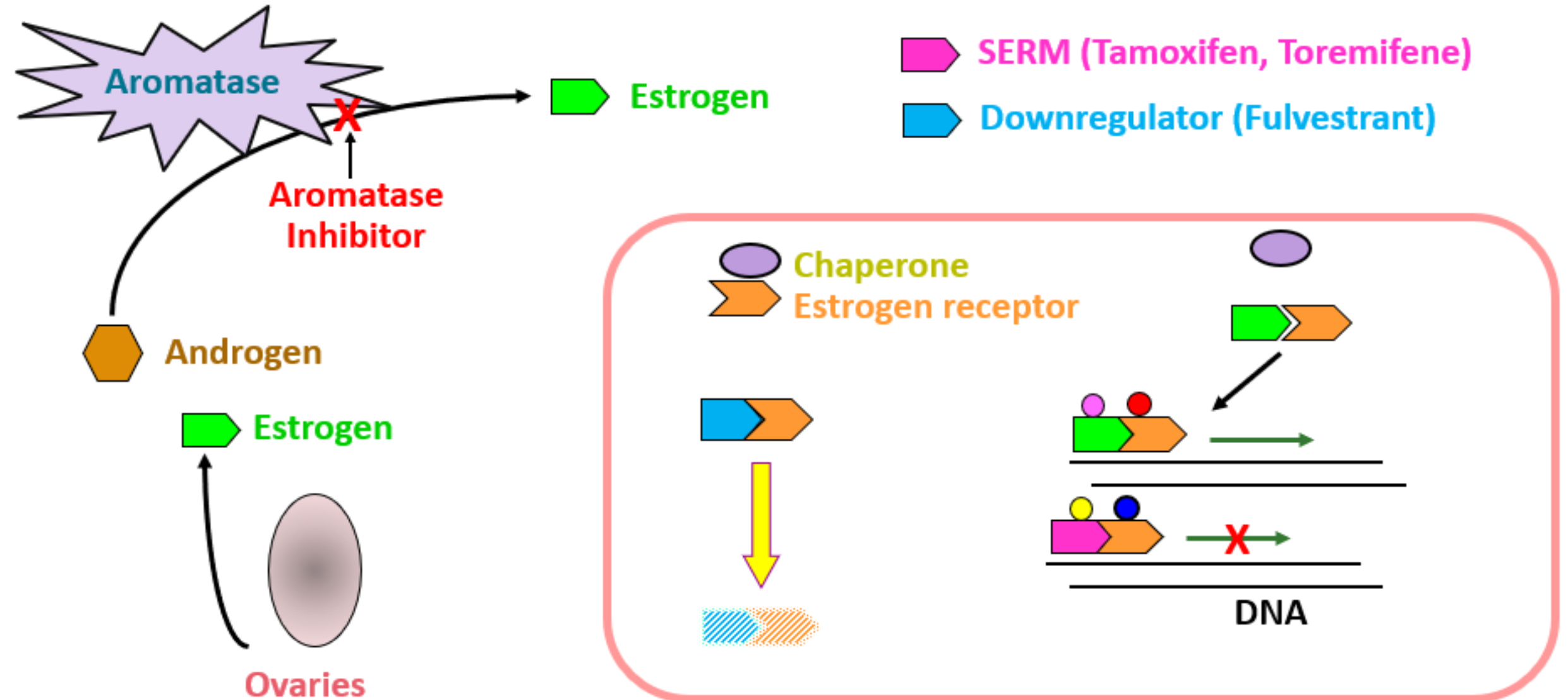
$$2 + 1 = 3$$

Strong

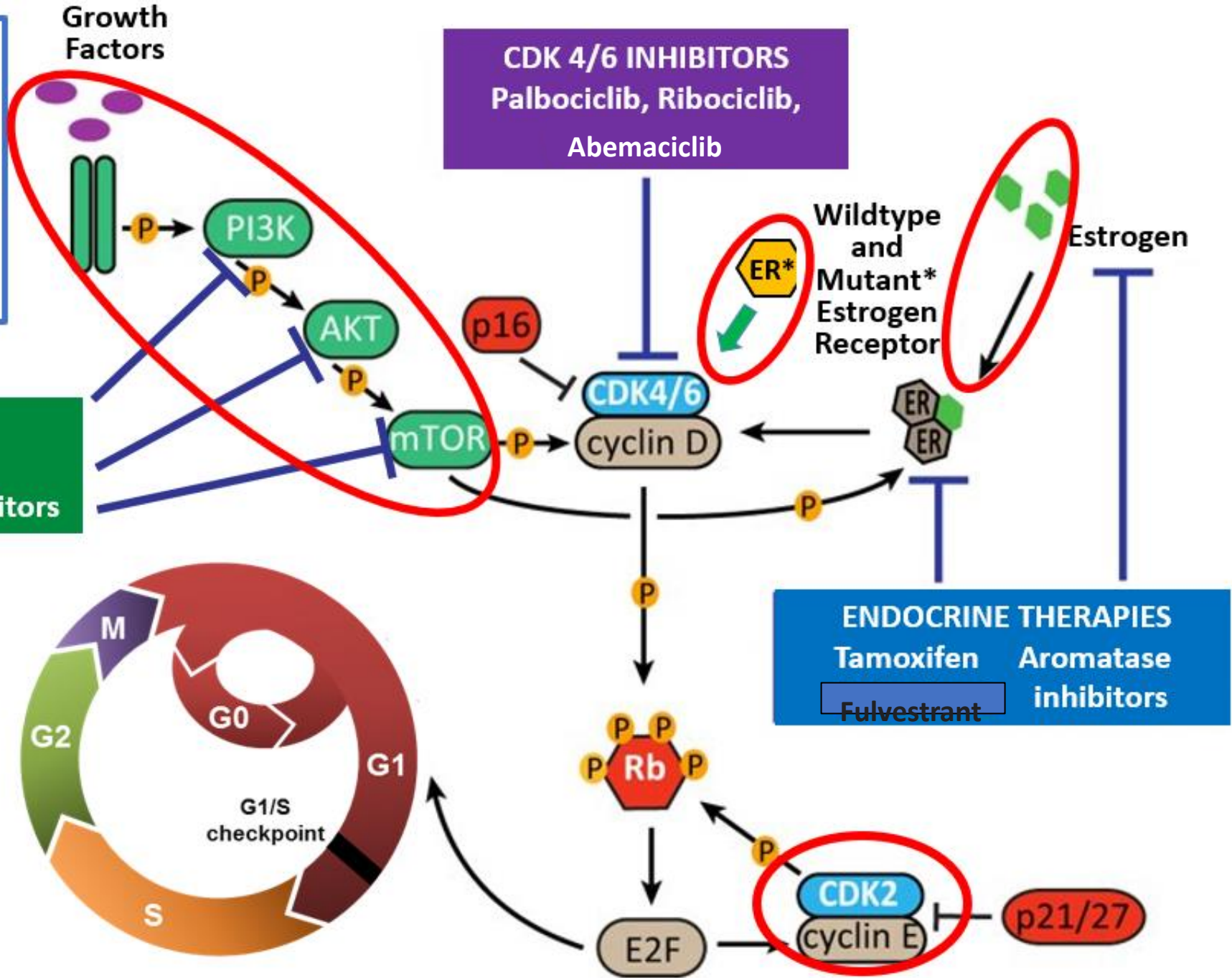


$$3 + 5 = 8$$

Estrogen, Estrogen Receptor and Different Endocrine Therapies Activities in HR+ Breast Cancer



Primary Drivers of Growth in HR+/HER2- Breast Cancer

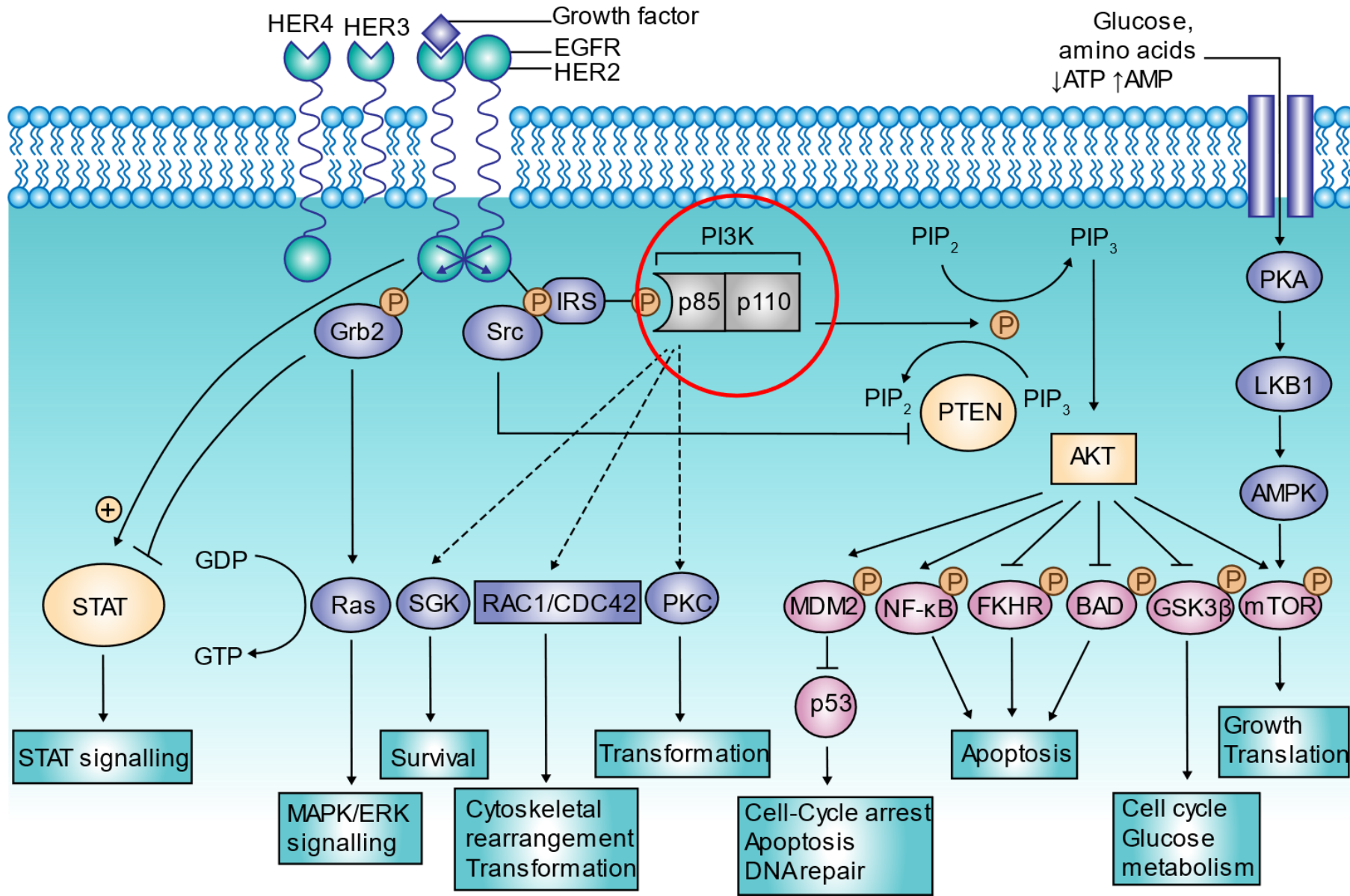


SIGNAL TRANSDUCTION INHIBITORS
PI3 kinase, AKT and mTOR inhibitors

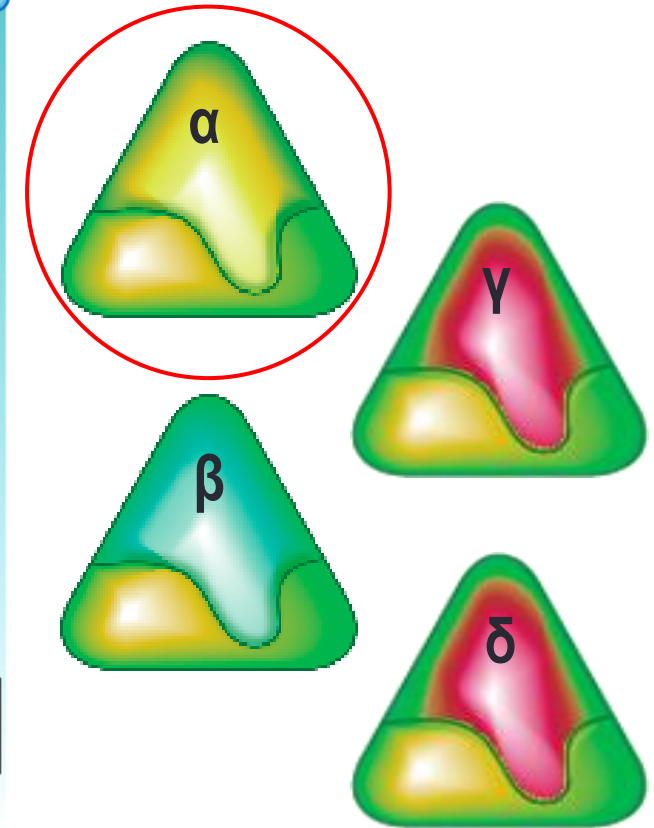
These also mediate resistance to endocrine therapy

ENDOCRINE THERAPIES
Tamoxifen, Aromatase inhibitors, Fulvestrant

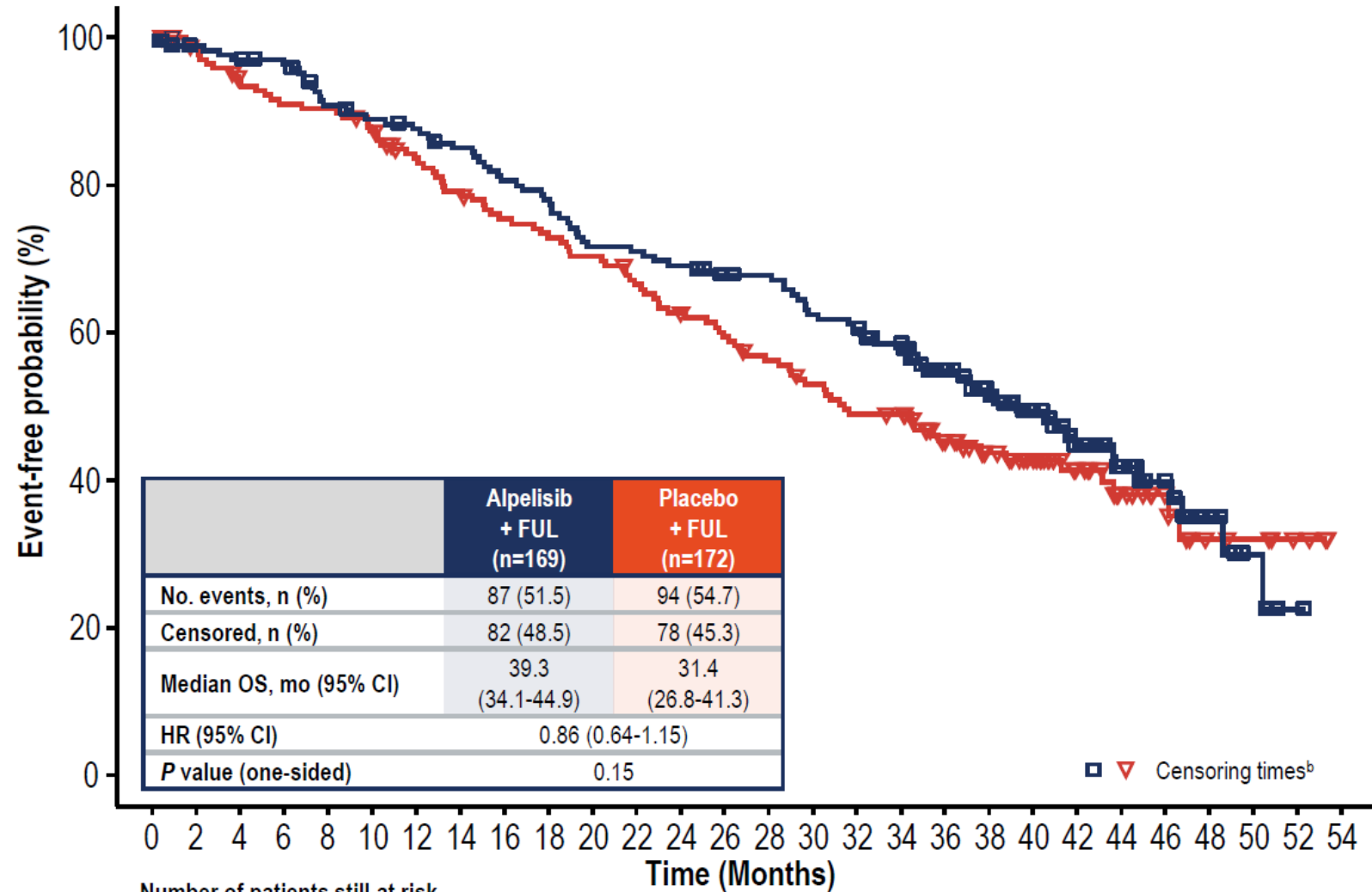
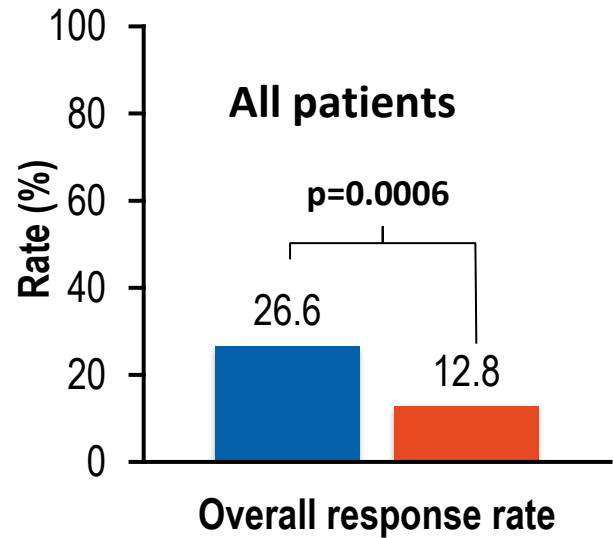
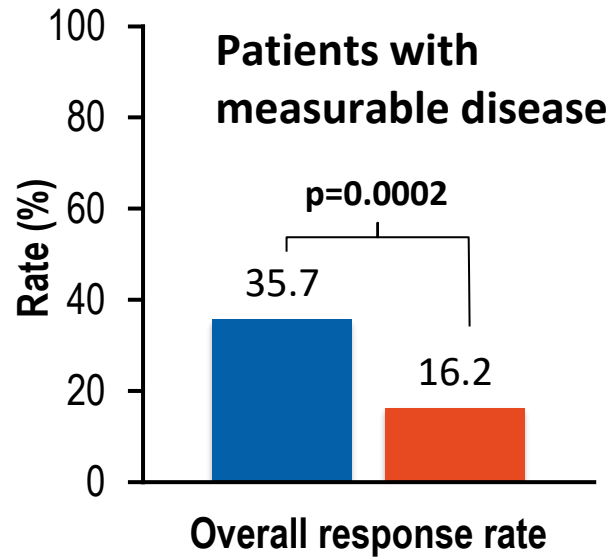
The Importance of the PI3K Pathway in HR+ Breast Cancer



PI3K isoforms



Overall Response Rates and Survival in the *PIK3CA*-Mutant Cohort



Number of patients still at risk

	0	2	4	6	8	10	12	14	16	18	20	22	24	26	28	30	32	34	36	38	40	42	44	46	48	50	52	54
Alpelisib + FUL	169	162	159	156	145	141	138	133	126	122	112	111	108	103	102	94	91	85	68	56	47	35	26	19	9	4	1	0
Placebo + FUL	172	164	155	150	149	143	133	126	119	115	111	104	98	92	86	80	74	73	60	49	42	29	20	13	7	6	3	0

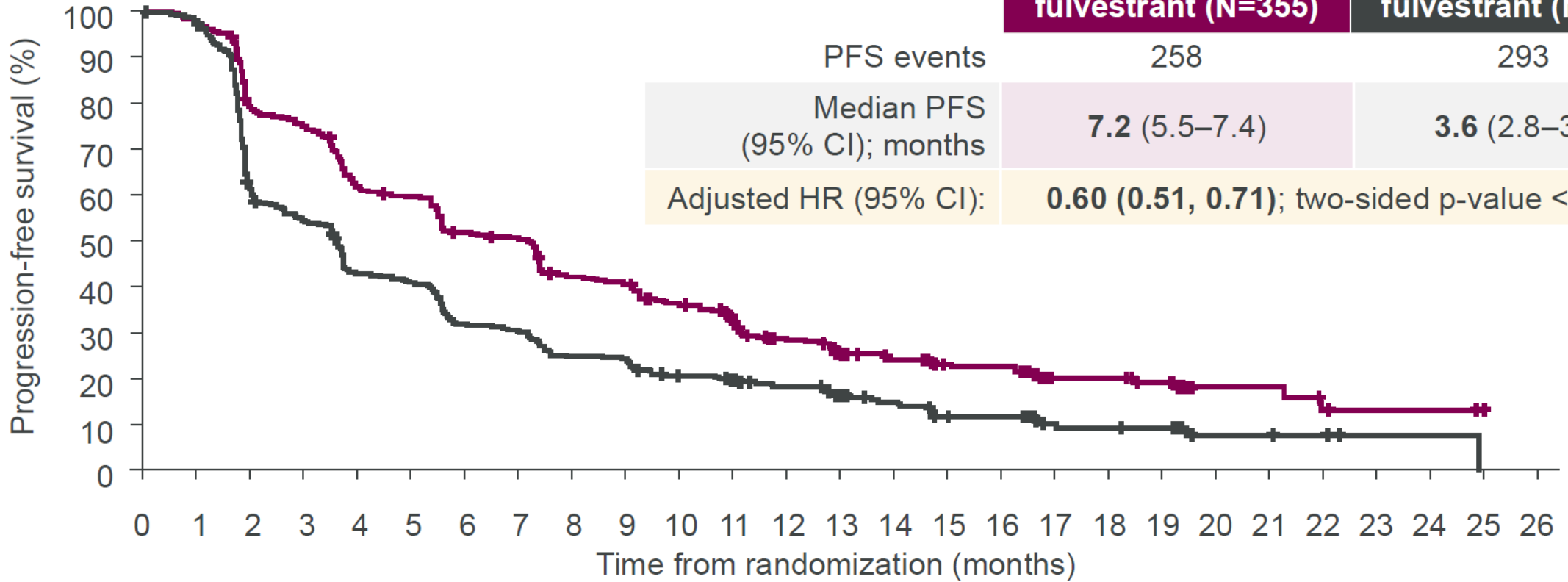
Full Analysis Set, *PIK3CA*-mutant cohort

■ Alpelisib + fulvestrant ■ Placebo + fulvestrant

CAPitello-291 Trial: Fulvestrant +/- Capivasertib (AKT Inhibitor)

- Men and pre/postmenopausal women
- Recurrence while on or <12 months from end of adjuvant AI, or progression while on prior AI for ABC
- ≤2 lines of prior endocrine therapy for ABC
- ≤1 line of chemotherapy for ABC
- Prior CDK4/6 inhibitors allowed (at least 51% required)
- No prior SERD, mTOR inhibitor, PI3K inhibitor, or AKT inhibitor
- HbA1c <8.0% (63.9 mmol/mol) and diabetes not requiring insulin allowed
- FFPE tumor sample from the primary/recurrent cancer available for retrospective central molecular testing

Dual primary endpoint: Investigator-assessed PFS in the overall population

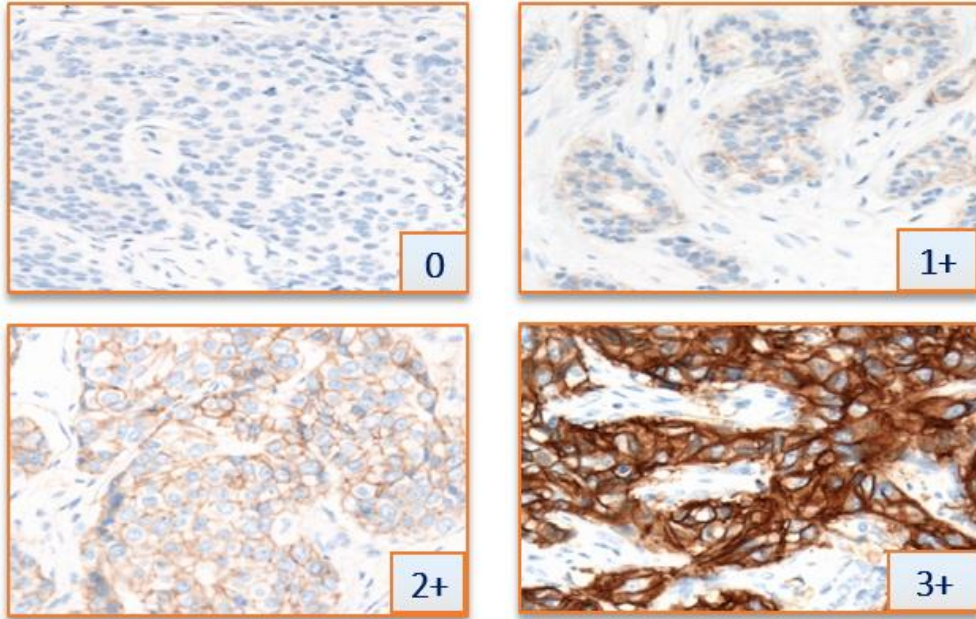


Number of patients at risk

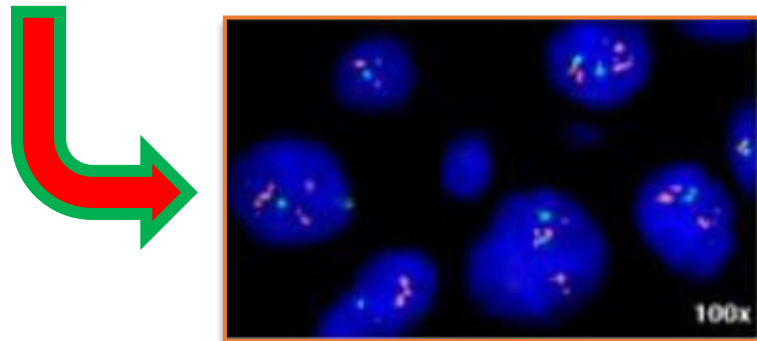
	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26
Capivasertib + fulvestrant	355	330	266	252	207	199	172	166	138	133	115	98	78	64	55	44	43	25	25	21	8	8	5	2	2	1	0
Placebo + fulvestrant	353	329	207	182	142	136	106	100	83	81	66	59	51	41	33	24	23	12	11	10	4	4	3	1	1	0	0

HER2 Testing: Technique and Interpretation are Critical

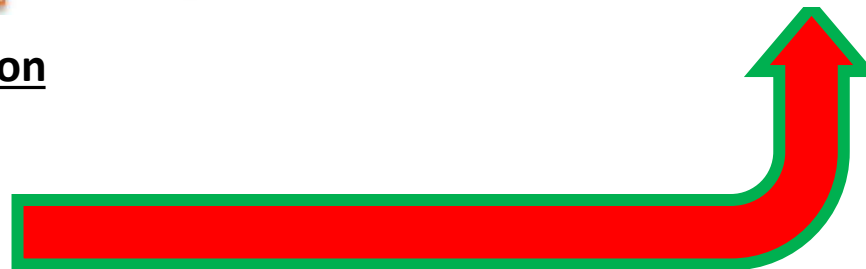
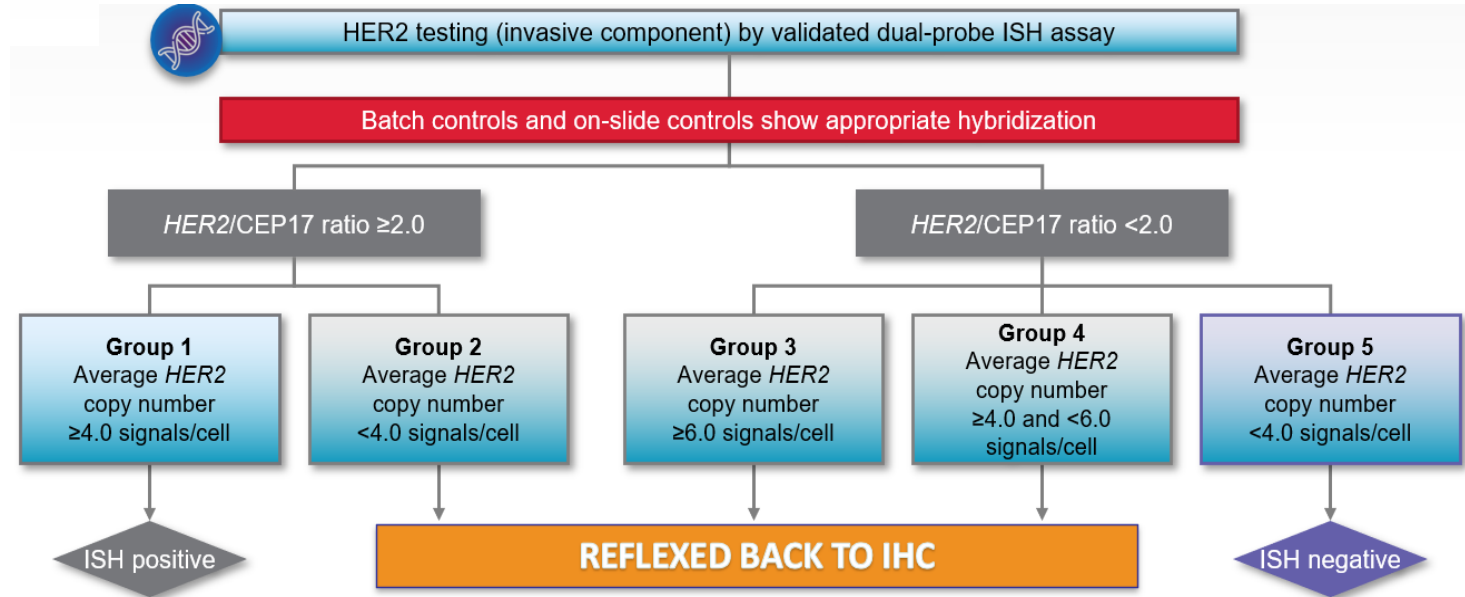
Immunohistochemistry



2+ cases are “reflexed” to ISH (in situ hybridization)

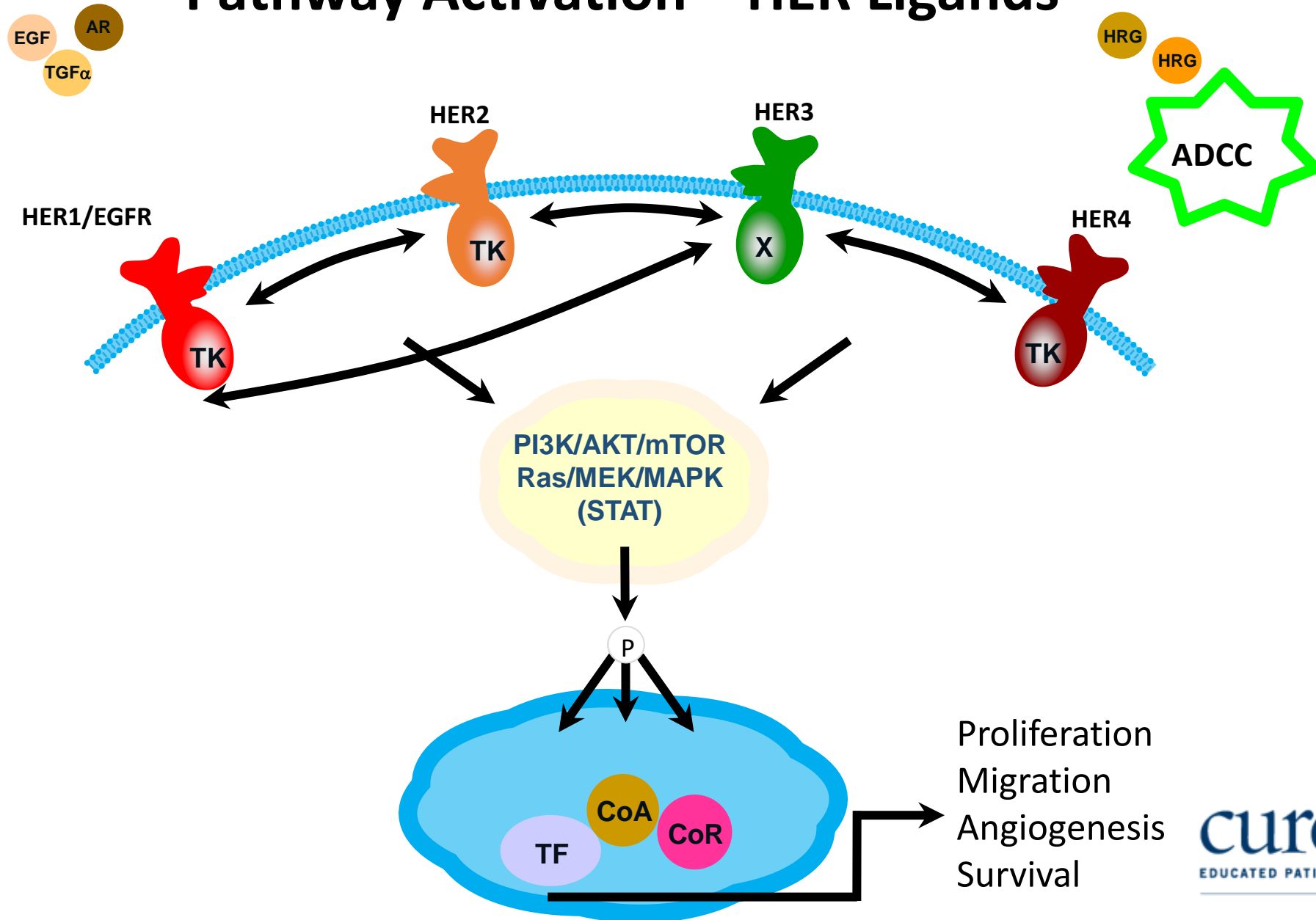


Interpretation of Dual Probe ISH based



*Note – Next-generation tumor DNA sequencing can also call HER2 amplification, but both antibody and ISH analysis should be done to confirm

Pathway Activation – HER Ligands



Trastuzumab

Humanized Anti-HER2 Monoclonal Antibody

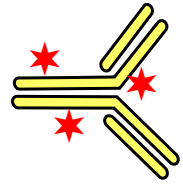


- Targets HER2 protein
- Selectively binds with high affinity ($K_d \leq 0.5$ nM)
- 95% human, 5% murine
- IgG1 isotype able to activate antibody-dependent cytotoxicity

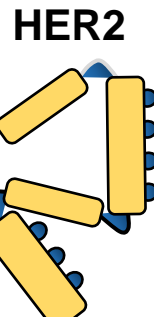
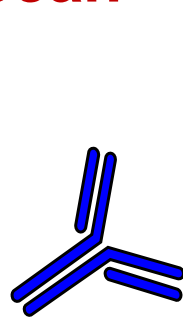
Trastuzumab, Pertuzumab, Lapatinib and T-DM1: Complementary Mechanisms

T-DM1

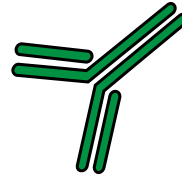
Trastuzumab Deruxtecan



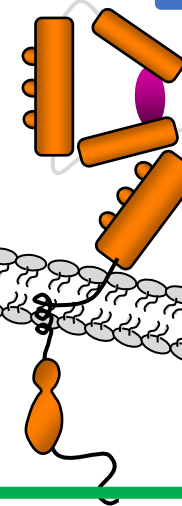
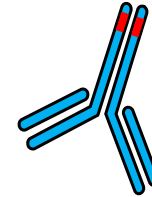
Trastuzumab



Pertuzumab



Margetuximab

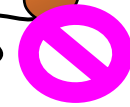


HER1/3/4

Subdomain IV

Dimerization domain

Lapatinib
Neratinib
Tucatinib



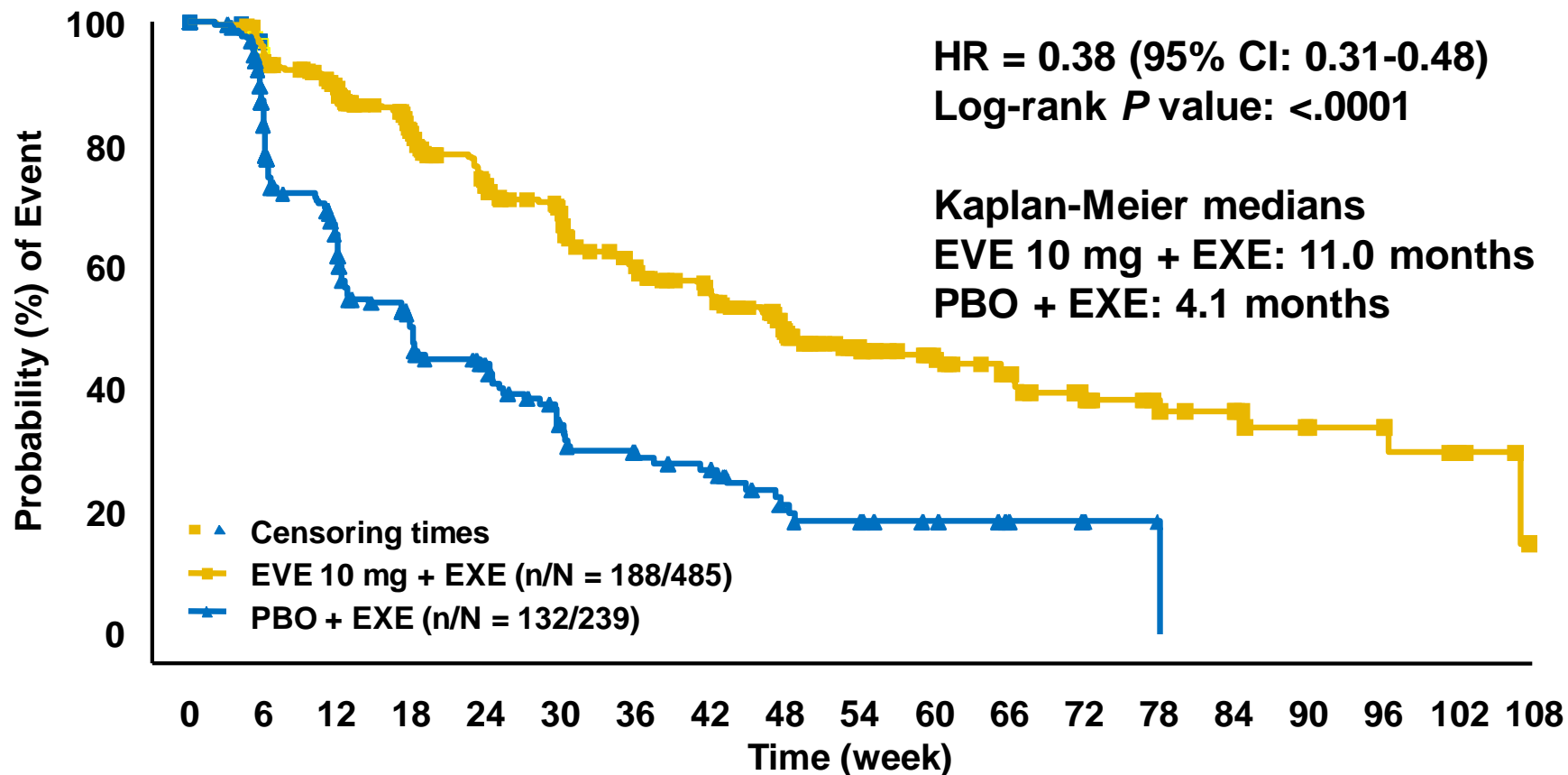
- Trastuzumab:**
 - Inhibits ligand-independent HER2 signaling
 - Activates ADCC
 - Prevents HER2 ECD shedding

- Pertuzumab:**
 - Inhibits ligand-dependent HER2 dimerization and signaling
 - Activates ADCC

- Lapatinib, Neratinib, Tucatinib:**
 - Inhibit intracellular kinase domain of HER2, HER1 (EGFR)

- Margetuximab:**
 - Like trastuzumab, but higher affinity for Fc receptor may contribute more immune effect

Exemestane +/- mTOR Inhibitor Everolimus: BOLERO-2 18 Month Followup



Number of patients still at risk

EVE 10 mg + EXE	485	427	359	292	239	211	166	140	108	77	62	48	32	21	18	11	10	5	0
PBO + EXE	239	179	114	76	56	39	31	27	16	13	9	6	4	1	0	0	0	0	0

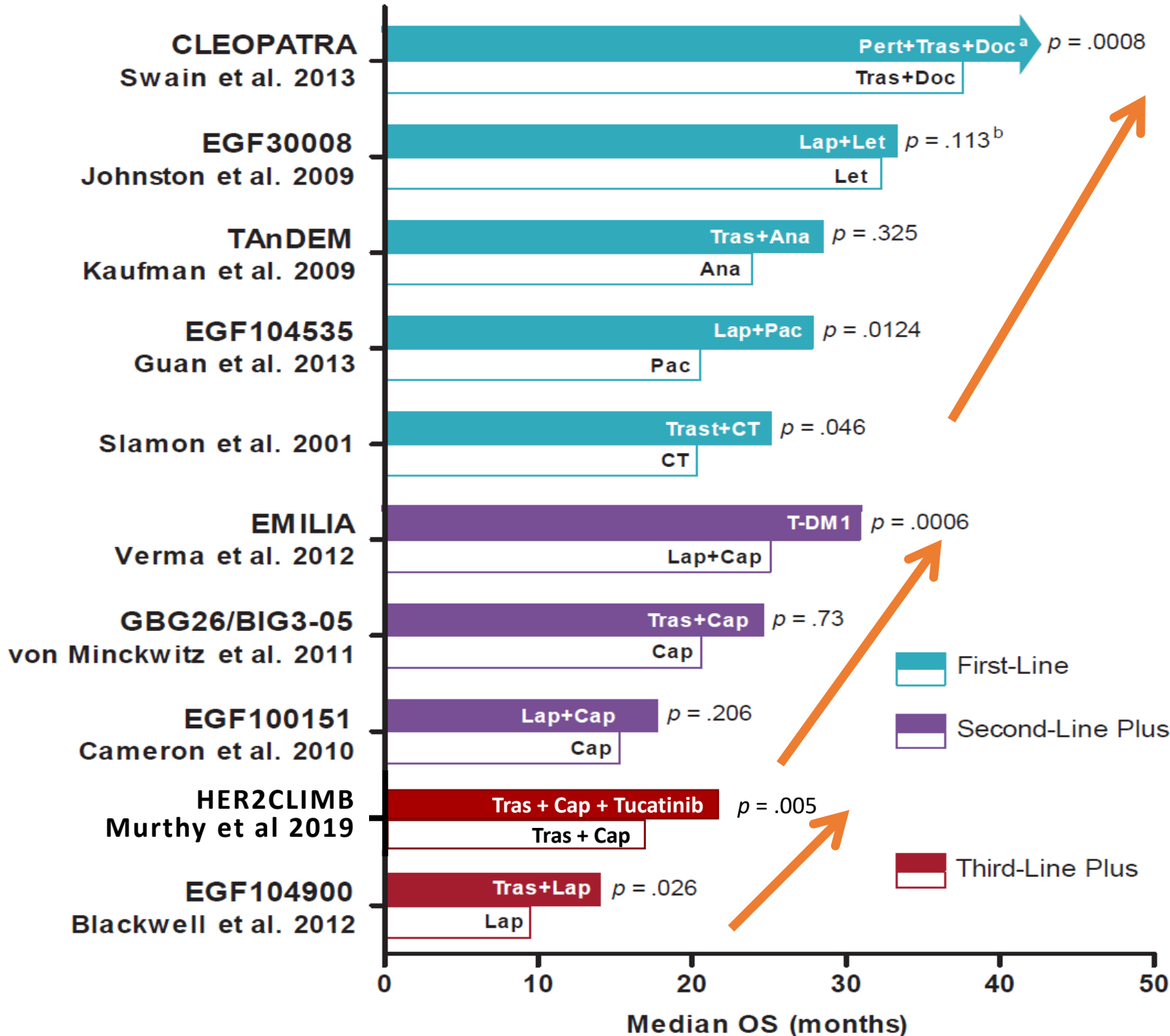
Results for Pivotal CDK 4/6 Inhibitor Trials

Trial	CDK Inhibitor	Line of Therapy (Endocrine Rx)	Menopausal Status	PFS HR	Statistical Significance	OS HR	Statistical Significance
PALOMA-2	Palbociclib	1 st Line/AI	Post	0.56	Yes	0.96	No
MONALEESA-2	Ribociclib	1 st Line/AI	Post	0.57	Yes	0.76	Yes
MONALEESA-7*	Ribociclib	1 st Line/AI or Tam	Pre/Peri	0.55	Yes	0.70	Yes
MONARCH-3	Abemaciclib	1 st Line/AI	Post	0.54	Yes	0.75	No (close)
PALOMA-3	Palbociclib	2 nd Line/Fulv	Pre/Post	0.46	Yes	0.81	No
MONARCH-2	Abemaciclib	2 nd Line/Fulv	Pre/Post	0.55	Yes	0.78	Yes
MONALEESA-3	Ribociclib	1 st /2 nd Line/Fulv	Pre/Post	0.59	Yes	0.72	Yes

*PFS/OS data reported for approved AI subset. Abbreviations: CDK=Cyclin-dependent kinase; Rx=therapy; PFS=progression-free survival; HR=hazard ratio; OS=overall survival; AI=aromatase inhibitor; Fulv=fulvestrant; NR=not reported

References:

- i. PALOMA-2: Finn R, et al. New Engl J Med 2016; Rugo H, et al. Breast Cancer Res Treat, 2019; Finn R, et al. ASCO 2022.
- ii. MONALEESA-2: Hortobagyi G, et al. New Engl J Med 2016; Hortobagyi G, et al. Ann Oncol 2018; Hortobagyi G, et al. New Engl J Med 2022.
- iii. MONALEESA-7: Tripathy D, et al. Ann Oncol 2018; Im S-A, et al New Engl J Med 2019. [Note PFS/OS data reported for approved AI subset].
- iv. MONARCH-3: Goetz M, et al. J Clin Oncol 2017; Johnson S, et al. npj Breast Cancer 2019; Goetz M, et al. ESMO 2022.
- v. PALOMA-3: Turner N, et al. New Engl J Med 2015; Cristofanilli M, et al. Lancet Oncol 2016; Turner N, et al New Engl J Med 2018.
- vi. MONARCH-2: Sledge G, et al. J Clin Oncol. Sledge G, et al. JAMA Oncol 2019.
- vii. MONALEESA-3: Slamon D, et al. J Clin Oncol 2018; Slamon D, et al New Engl J Med 2020.



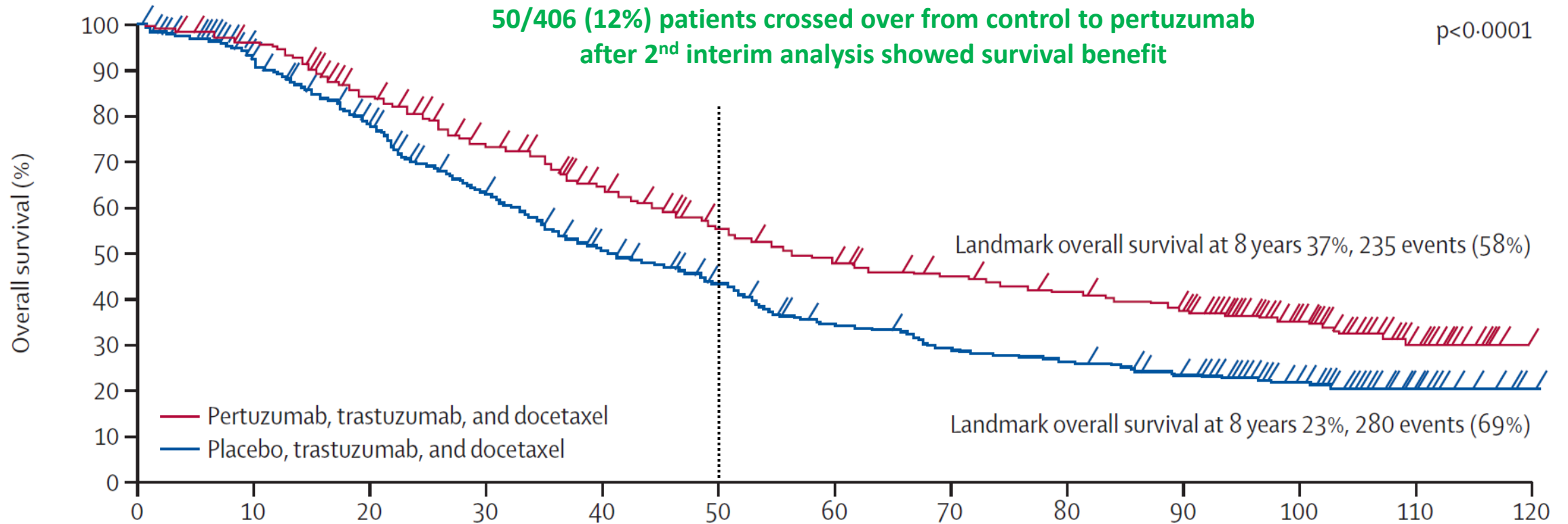
HER2+ Metastatic Breast Cancer:

Serial Improvements in Survival with Newer Agents and Combinations

BUT.... Rare “Cures”

Modified from:
Verma S, et al. Oncologist 2013

Updated Data from CLEOPATRA Trial: Docetaxel + Trastuzumab +/- Pertuzumab in First Line



Number at risk
(number censored)

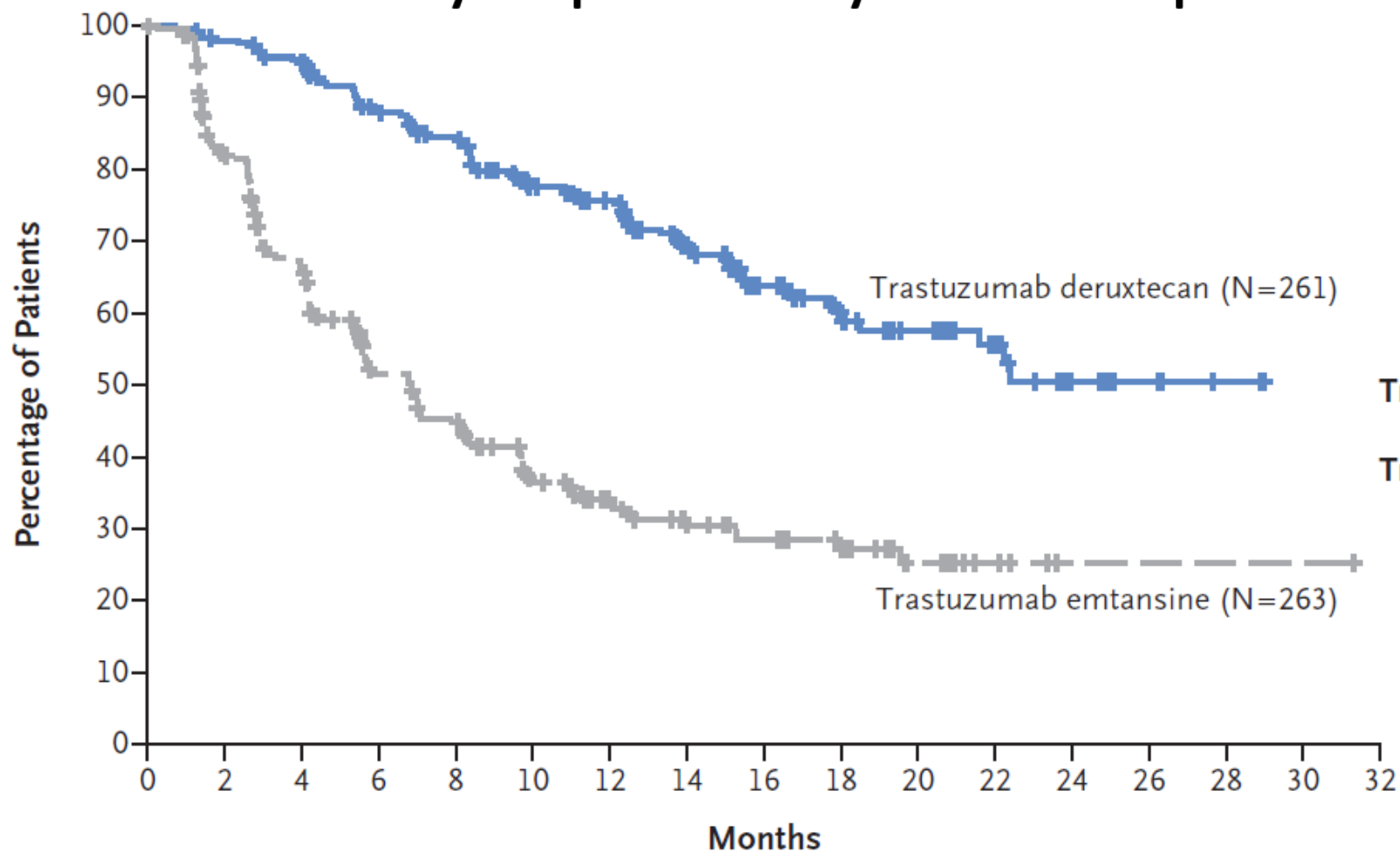
Pertuzumab	402 (0)	371 (14)	318 (23)	269 (32)	228 (41)	188 (48)	165 (50)	150 (54)	137 (56)	120 (59)	71 (102)	20 (147)	0 (167)
Placebo	406 (0)	350 (19)	289 (30)	230 (36)	181 (41)	149 (48)	115 (52)	96 (53)	88 (53)	75 (57)	44 (84)	11 (115)	1 (125)

Median Progression-Free Survival: THP=18.8 mo; TH=12.4 mo (HR=0.69; 95% CI 0.59-0.81)

Median Overall Survival: THP=57.1 mo; TH=40.8 mo (HR=0.69; 95% CI 0.58-0.82)

Destiny-Breast03 Randomized trial T-DXd vs. T-DM1

Primary Endpoint: PFS by blinded independent central review



	Median Progression-free Survival (95% CI) <i>mo</i>	12-Mo Progression-free Survival (95% CI) %
Trastuzumab Deruxtecan	NR (18.5–NE)	75.8 (69.8–80.7)
Trastuzumab Emtansine	6.8 (5.6–8.2)	34.1 (27.7–40.5)

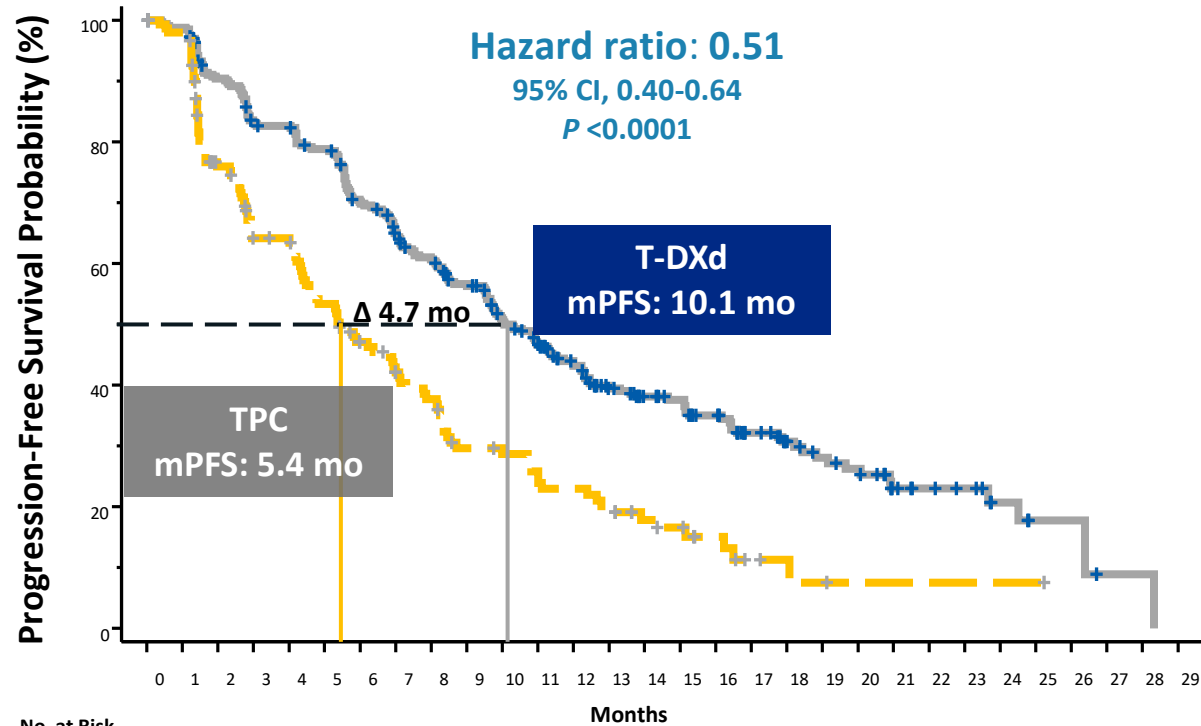
Hazard ratio for disease progression or death, 0.28 (95% CI, 0.22–0.37)
P<0.001

No. at Risk

Trastuzumab deruxtecan	261	250	240	214	200	168	150	112	79	53	36	25	10	5	2		
Trastuzumab emtansine	263	200	155	108	93	65	51	37	29	21	12	6	1	1	1	1	0

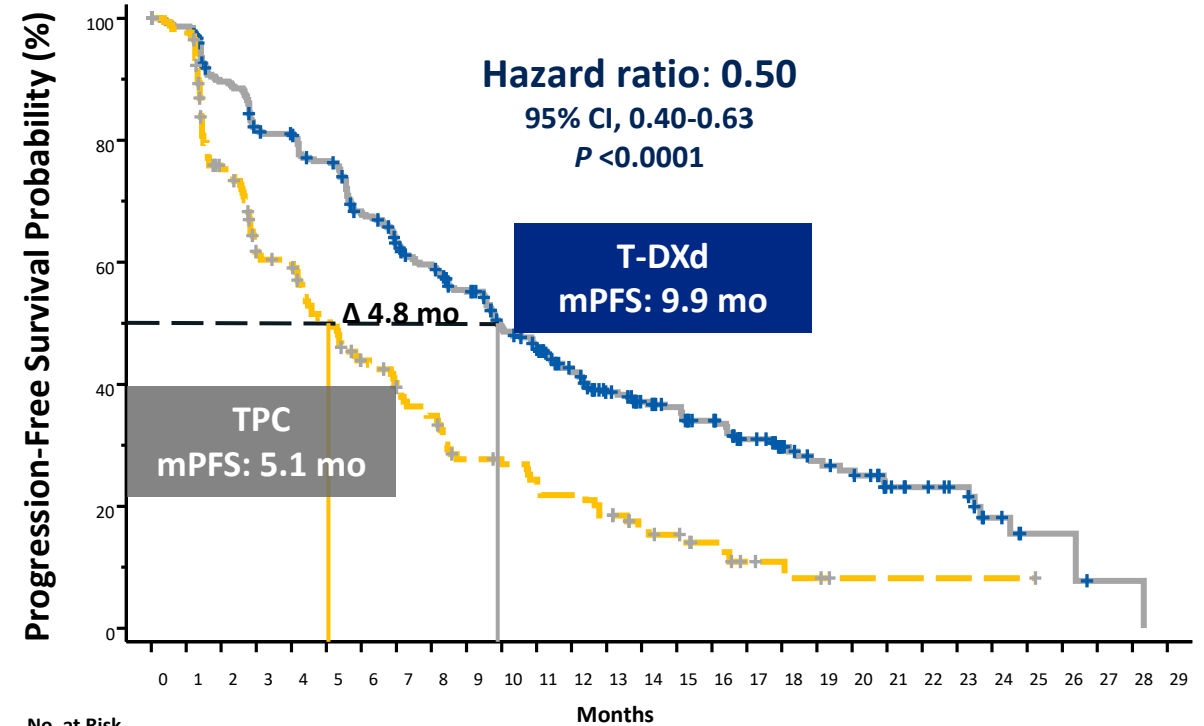
Progression-Free Survival in HR+ and All Patients

Hormone receptor-positive



No. at Risk	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29
T-DXd (n=331):	331	324	290	265	262	248	218	198	182	165	142	128	107	89	78	73	64	48	37	31	28	17	14	12	7	4	4	1	1	0
TPC (n=163):	163	146	105	85	84	69	57	48	43	32	30	27	24	20	14	12	8	4	3	2	1	1	1	1	1	1	1	0	0	

All patients



No. at Risk	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29
T-DXd (n=373):	373	365	325	295	290	272	238	217	201	183	156	142	118	100	88	81	71	53	42	35	32	21	18	15	8	4	4	1	1	0
TPC (n=184):	184	166	119	93	90	73	60	51	45	34	32	29	26	22	15	13	9	5	4	3	1	1	1	1	1	1	1	0	0	

PFS by blinded independent central review.

HR, hormone receptor; PFS, progression-free survival; T-DXd, trastuzumab deruxtecan; TPC, treatment of physician's choice.

Mechanisms of DNA Repair

Environmental factors
(UV, radiation, chemicals)

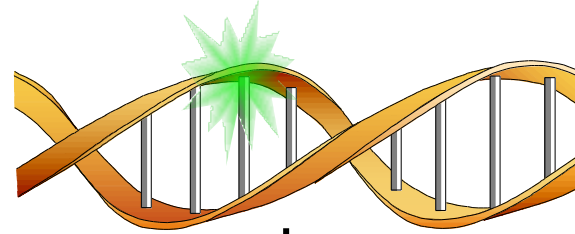
Normal physiology
(DNA replication, ROS)

Chemotherapy
(alkylating agents, antimetabolites)

Radiotherapy



DNA DAMAGE



Cell Death



MAJOR DNA REPAIR PATHWAYS

Single Strand Breaks

- Nucleotide excision repair
 - Base excision repair
- PARP1

Replication Lesions

- Base excision repair
- PARP1
-

Double Strand Breaks

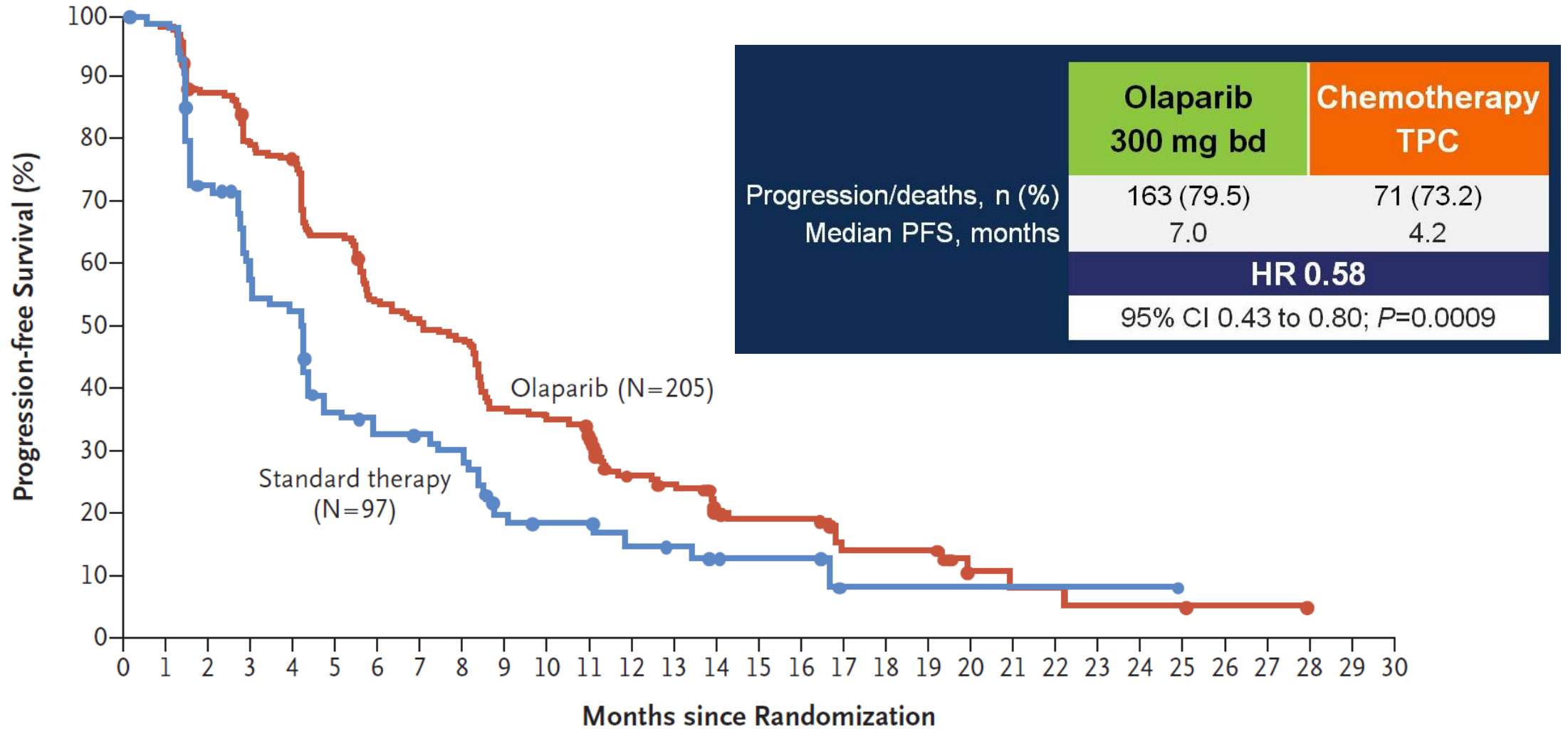
- Non-homologous end-joining
- Homologous recombination
BRCA1/BRCA2
- Fanconi anemia pathway
- Endonuclease-mediated repair

DNA Adducts/Base Damage

- Alkyltransferases
 - Nucleotide excision repair
 - Base excision repair
- PARP1

OlympiAD: Olaparib vs. Chemotherapy of Physician's Choice*

Progression-Free Survival



No. at Risk

Olaparib	205	201	177	159	154	129	107	100	94	73	69	61	40	36	23	21	21	11	11	11	4	3	3	2	2	1	1	1	0
Standard therapy	97	88	63	46	44	29	25	24	21	13	11	11	8	7	4	4	4	1	1	1	1	1	1	1	1	0	0	0	0

*Capecitabine (45%); eribulin (37%); vinorelbine (18%)

Robson M, et al. NEJM 2017

KEYNOTE-355 Study Design (NCT02819518)

Key Eligibility Criteria

- Age ≥ 18 years
- Central determination of TNBC and PD-L1 expression
- Previously untreated locally recurrent inoperable or metastatic TNBC
- Completion of treatment with curative intent ≥ 6 months prior to first disease recurrence
- ECOG performance status 0 or 1
- Life expectancy ≥ 12 weeks from randomization
- Adequate organ function
- No systemic steroids
- No active CNS metastases
- No active autoimmune disease



Stratification Factors:

- Chemotherapy on study (taxane vs gemcitabine/carboplatin)
- PD-L1 tumor expression (CPS ≥ 1 vs CPS < 1)
- Prior treatment with same class chemotherapy in the neoadjuvant or adjuvant setting (yes vs no)

^aPembrolizumab 200 mg intravenous (IV) every 3 weeks (Q3W)

^bChemotherapy dosing regimens are as follows:

Nab-paclitaxel 100 mg/m² IV on days 1, 8, and 15 every 28 days

Paclitaxel 90 mg/m² IV on days 1, 8, and 15 every 28 days

Gemcitabine 1000 mg/m²/carboplatin AUC 2 on days 1 and 8 every 21 days

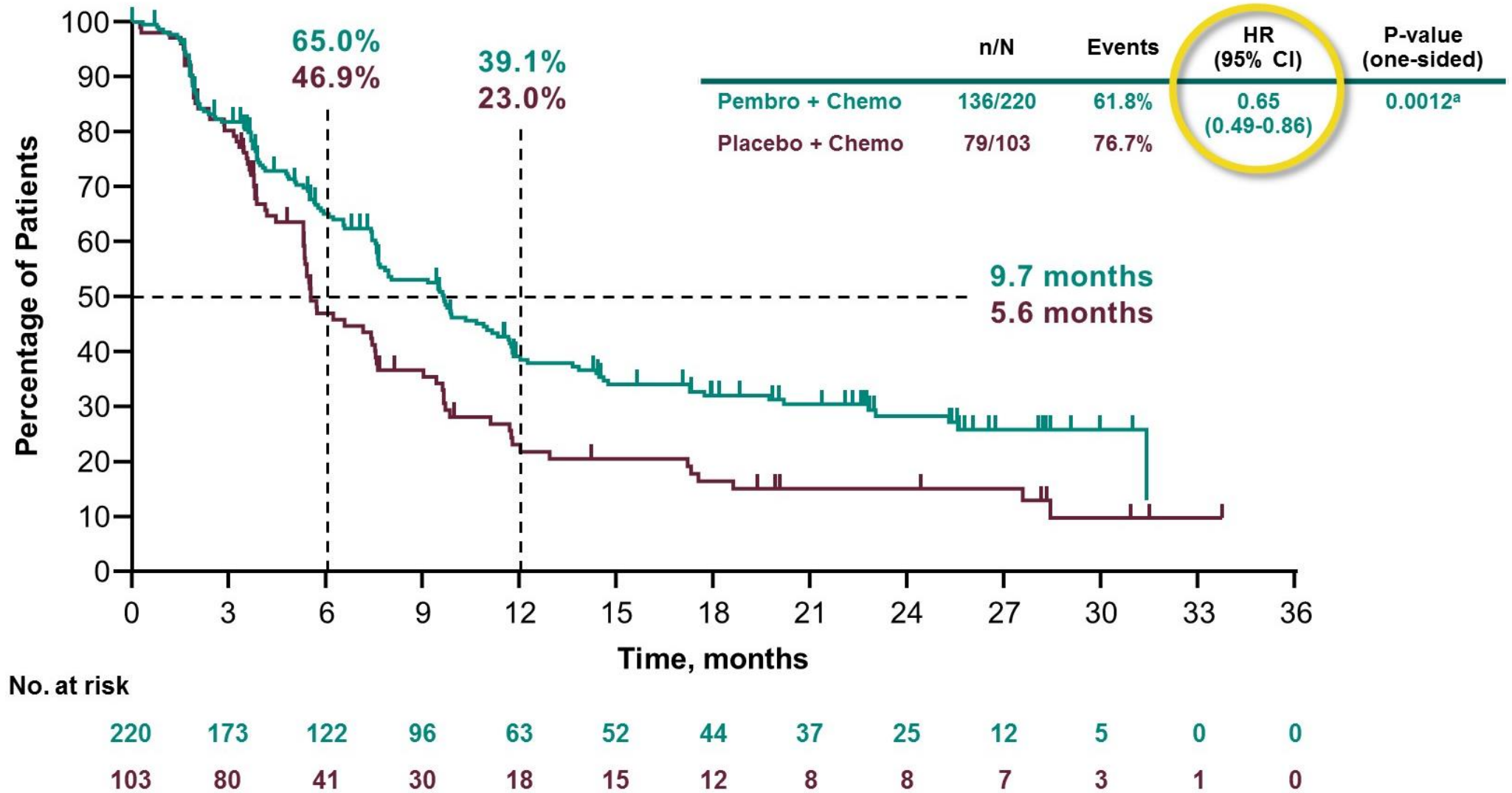
^cNormal saline

^dTreatment may be continued until confirmation of progressive disease

CNS=central nervous system; ECOG=Eastern Cooperative Oncology Group;

PD-L1=programmed death ligand 1; R=randomized; TNBC=triple-negative breast cancer

Progression-Free Survival: PD-L1 CPS ≥ 10



^aPrespecified *P* value boundary of 0.00411 met.

Hazard ratio (CI) analyzed based on a Cox regression model with treatment as a covariate stratified by the randomization stratification factors. Data cutoff December 11, 2019.

Genomic Aberrations in Breast Cancer that Guide Precision Medicine in Breast Cancer

Gene	Aberration	Frequency	Targeted Drug(s)
EVIDENCE BASED (from randomized or Phase II trials)			
HER2	Amplification	20%	Trastuzumab, pertuzumab, T-DM1, lapatinib, neratinib, more
BRCA 1/2	Inactivation mutations (germline)	5%	Olaparib, talozoparib
PIK3CA	Activating mutations	30-40%	Alpelisib
BRAF	Activating mutation (V600E)	2-3% (TNBC)	Dabrafenib and trametinib
Several	Microsatellite instability	1-2%	Pembrolizumab
NTRK	Gene fusion	<1%	Larotrectinib
RET	Gene fusion/rearrangements/SNV	<1.5%	Selpercatinib
HER2	Activating mutations (non-amplified HER2)	2-10%	Neratinib (not FDA-approved)
EMERGING			
FGFR1-4	Amplification	10%	FGFR inhibitors
ESR1	Mutation (after AI exposure)	30-40%	Fulvestrant, other SERDs (elacestrant in randomized trial)
AKT	Activating mutations	2%	AKT, mTOR inhibitors (capivasertib, everolimus)
PTEN	Inactivating mutations/silencing	20%	PI3K (non alpha-selective), Akt, mTOR inhibitors
Myc	Amplification	16%	BET inhibitors
c-MET	Amplification/mutation	15%	Met inhibitors (cabozantinib)
CDH1	Inactivating mutations/silencing	7%	Wnt inhibitors, ALK/ROS inhibitors

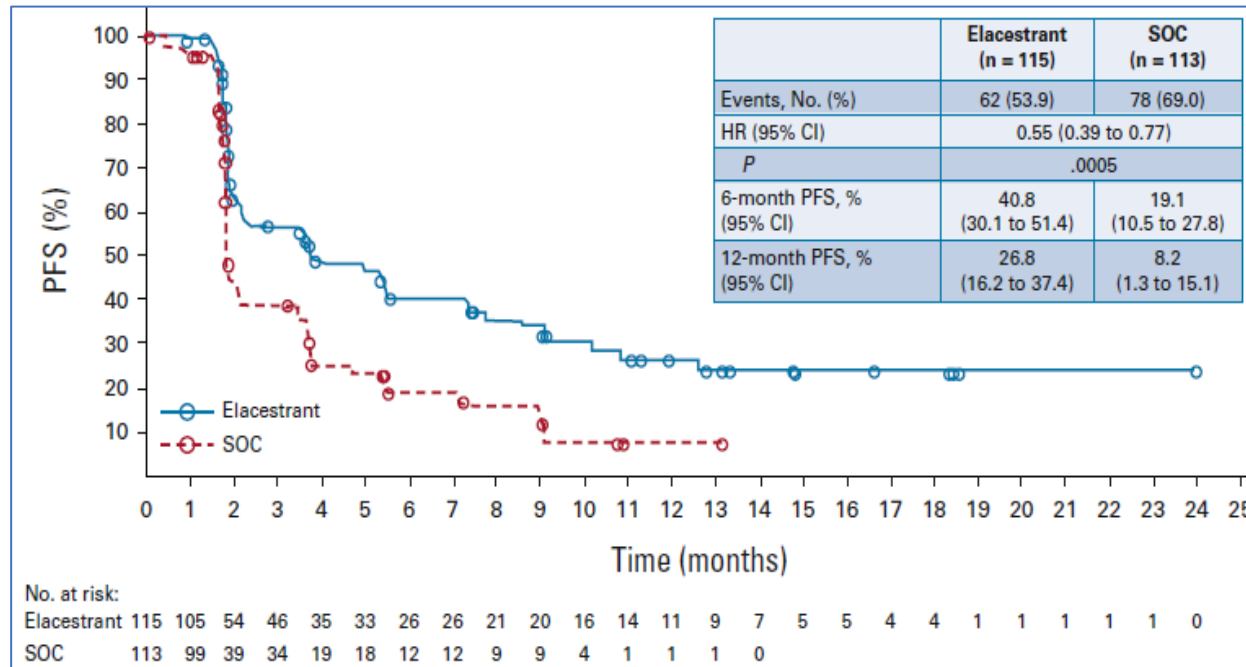
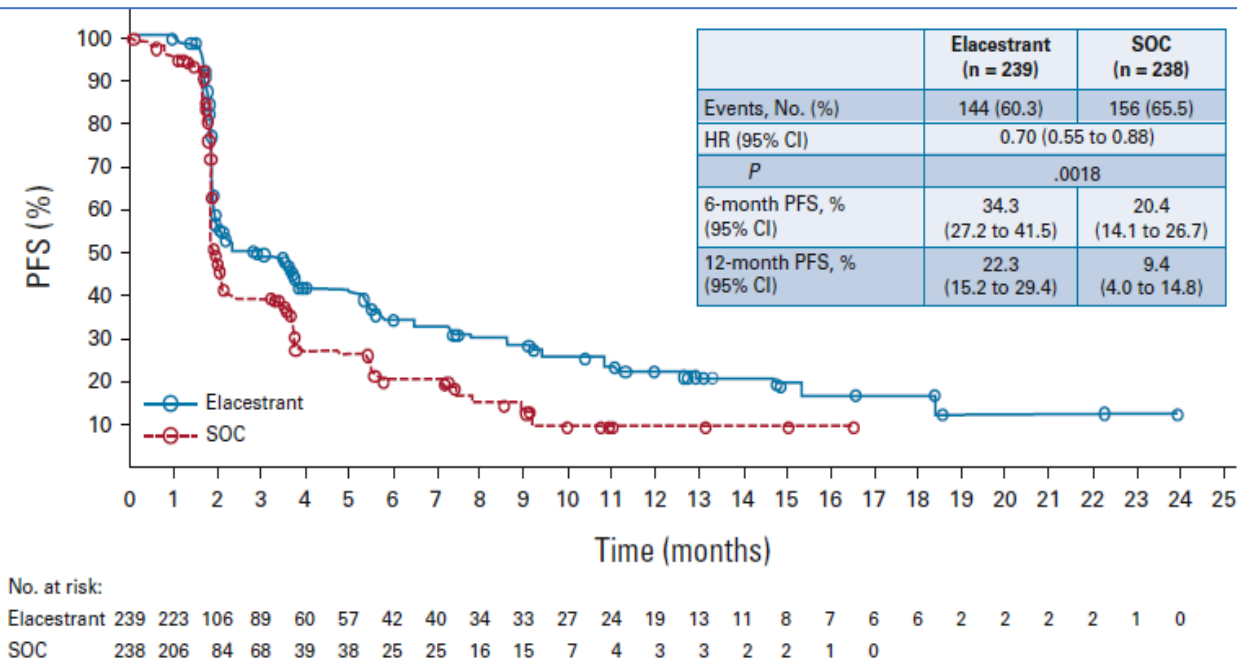
Abbreviations IHC=immunohistochemistry; AI = aromatase inhibitor; SERD=selective estrogen receptor downregulator

EMERALD Trial

PFS: Elacestrant vs Fulvestrant (All Patients and *mESR1* Group)

All Patients

Patients With Tumors Harboring *mESR1*



Elacestrant demonstrated a significant improvement versus Fulvestrant as SOC in patients with ER+/HER2-advanced/metastatic breast cancer and *mESR1* following CDK4/6i therapy

Approved on 1/27/2023 for HR+/HER2, 2nd/3rd line, ESR1-mutated cancer



THANK YOU