### **ESMO BREAST CANCER** VIRTUAL MEETING

### THE RIGHT BIOMARKER FOR THE RIGHT PATIENT

From immunohistochemistry to predictive molecular pathology of breast cancer.

#### Nicola Fusco

European Institute of Oncology (IEO) University of Milan, Italy



ESVO GOOD SCIENCE BETTER MEDIC BEST PRACTICE www.esmo.org www.ieo.it

### DISCLOSURES

I have received honoraria for consulting/advisory role from Merck Sharp & Dohme (MSD) and Boehringer Ingelheim





### Hormone receptors: ER & PgR

75%-80% of invasive breast cancers are ER+/PgR+

- Rationale for clinical testing: to identify patients who may benefit from <u>hormonal</u> <u>therapy</u>
  - >> substantial survival benefits in ER<sup>+</sup> >> weak prognostic factor

METHOD:

- IHC on FFPE tissue sections
- Only nuclear staining

ER<sup>+</sup>/PgR<sup>+</sup> moderately differentiated (G2) invasive ductal carcinoma



- Single-gene expression assays are not recommended
- False-negative results: still ~15% of the cases

>> patients may not receive effective therapy

>> internal and external controls





### **ER<sup>+</sup>/PgR<sup>-</sup> invasive breast cancers**

#### 5% of all invasive breast cancers

- Subset of Luminal B tumors
- Preferentially post-menopause
- Clinically heterogeneous
- Larger tumor size than PgR<sup>+</sup>
- Worse prognosis than PgR<sup>+</sup>
- Higher response but also worse longterm outcome after neoadjuvant chemotherapy
- Genomic instability

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>> Enriched for mutations in cancer genes (e.g. TP53, PIK3CA, CDH1, HER2, BRAF)



### ER+/PgR- moderately differentiated (G2) invasive ductal carcinoma

### **ER-low invasive breast cancers**

Invasive carcinomas with low level (1-10%) of ER expression

2-3% of ER<sup>+</sup> invasive breast cancers

Clinically challenging

>> Heterogeneous behavior and biology
>> Gene expression profiles more similar to ERcancers

>> Eligible for HT but limited data on the benefit

#### **Diagnostically challenging**

>> Usually weak/very weak nuclear staining
>> Pre-analytical issues
>> Inter-observer reproducibility
>> An additional comment should be provided in the pathology report







#### ~15-20% of invasive breast cancers overexpress HER2

Rationale for clinical testing: to determine patient eligibility for anti-HER2 therapy

METHOD:

- IHC on FFPE tissue sections



>>Only membrane staining

- In situ hybridization (ISH) in IHC 2+
- In both IHC and ISH the pre-analytic phase is crucial
- "HER2-enriched" by transcriptomic analysis

#### >> super-responders

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Metzeger Filho et al. ASCO 2019; Prat et al. J. Natl. Cancer Inst. 2019; Perez et al. BMC Cancer 2019; Song et al; Rye et al. Mol. Oncol. 2018; Modi et al. JCO 2020; Modi et al. NEJM 2019; Modi et al. SABCS 2019; Banerji et al. Lancet Oncol 2019; Fehrenbacher et al. Cancer Res 2018; von Minckwitz et al. NEJM 2019





Score 0 (40x)

Score 1 (40x)





Score 2 (40x)

Score 3 (40x)





HER2/CEP17 = 3.1 (AMPLIFIED) (HER2copy number 6.6)

### **HER2** intra-tumor heterogeneity

2% of HER2<sup>+</sup> breast cancers show intra-tumor heterogeneity of HER2 expression

Patterns of HER2 heterogeneity:

>> "clustered", topographically distinct HER2<sup>+</sup> and HER2<sup>-</sup> tumor clones

>> "scattered", isolated HER2+ cells in a HER2- tumor >> "mosaic", diffuse intermingling of cells with different HER2 statuses (ISH)

- Lower pCR after neoadjuvant treatment with TTZ+chemo
- No pCR in stage II/III after neoadjuvant T-DM1 and pertuzumab





### **HER2-low invasive breast cancers**

Spectrum of carcinomas with different degrees of HER2 expression (1+ to 2+/ISH<sup>NEG</sup>)

- 45%-55% of all invasive breast cancers
- Poorer prognosis compared to HER2negative breast carcinomas
- TTZ duocarmazine (SYD-985) and TTZ deruxtecan (DS-8201) have shown encouraging response rates in HER2low breast cancer



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### **Programmed death-ligand 1 (PD-L1)**

- PD-L1 is expressed in 40-65% of TNBC
- Expression is restricted, in most cases, to immune cells
- PD-L1 expression is predictive of response to Atezolizumab (anti-PD-L1)
- Chemotherapy may enhance tumorantigen release and antitumor responses to immune checkpoint inhibition
- IMpassion130 Study: Atezolizumab + nab-paclitaxel prolonged PFS in PD-L1 TNBC patients



### **PD-L1 testing method**

PD-L1 20x





TNBC tissue showing dark brown punctate and linear IC staining.





IHC staining with VENTANA PD-L1 SP142 Assay (CDx) demonstrates staining in TILS and occasionally in tumor cells

The PD-L1 tumor-infiltrating immune cell (IC) status is defined by the percentage of tumor area occupied by PD-L1-positive ICs

'NBC tissue showing moderate to strong circumferential TC membrane staining.



# IMMMUNE-RELATED MARKERS Tumor-infiltrating lymphocytes (TILs)

- TILs should be routinely characterized in TNBC because of their prognostic value (St Gallen 2019, WHO Breast Tumours 2019)
- Data are inadequate to recommend TILs to guide neo/adjuvant treatment choices in TNBC (St Gallen 2019)
- Stromal TILs are prognostic in TNBC and HER2<sup>+</sup> breast cancer
- Not prognostic in ER<sup>+</sup> tumors



Working category to describe tumors with "more lymphocytes than tymor cells".

Definitions vary across studies with stromal TILs of 50–60% used as a threshold. LPBC can be used for predefined subgroup analyses and for description of tumors with a particularly high immune infiltrate, however, keep in mind that TILs are a continuous parameter and the threshold for LPBC is still arbitrary.

Stromal TILs

Lymphocyte-predominant breast cancer (LPBC)





TILs with direct cell-cell contact with carcinoma cells, might be an indicator of direct cell-based antitumor effects.

Several studies have shown that intratumoral TILs and more difficult to evaluate and do not provide additional predictive/prognostic information compared to stromal TILs.

> Sagado et al. Ann Oncol 2015 Dieci et al. Semin Cancer Biol. 2018



### The mismatch repair (MMR) system

- Major contributor to DNA integrity
- Four main proteins
   >> MLH1, MSH2, MSH6, and PMS2
- Genomes of MMR deficient (dMMR) cancers contain extraordinarily high numbers of somatic mutations

Tumor mutational burden (**TMB**) Microsatellite instability (**MSI**)

 FDA approves pembrolizumab for dMMR and/or MSI-H cancers regardless of the tumor site >> histology agnostic approval
 >> no CDx



### **MMR testing methods**

What is the optimal MMR testing method for breast cancer?

- **IHC** >> MLH1, MSH2, MSH6, PMS2
- MSI >> PCR (BAT25, BAT26, D2S123, D5S346, D17S250, ...) vs. NGS
- Sequencing/methylation assays
- **TMB** >> targeted panels, WES





#### **PROGNOSTIC OR PREDICTIVE?**

### **Multigene Tests**

- Useful complementary information in ER<sup>+</sup> breast cancers.
- Since ER<sup>-</sup> cancers tend to have higher proliferation rates, the prognostic value of current multigene tests in these cancers is limited.
- May help informing chemotherapy decision in ER<sup>+</sup>/HER2<sup>-</sup> N0/N1a breast cancers



NCCN Guidelines Version 3.2020
 e Invasive Breast Cancer
 NCCN Evidence Blocks™

Assay	Predictive	Prognostic
21-gene (Oncotype Dx) (for pN0 or node negative)	Yes	Yes
21-gene (Oncotype Dx) (for pN+ or node positive)	N/A* *awaiting results of RxPONDER study	Yes
70-gene (MammaPrint) (for node negative and 1–3 positive nodes)	Not determined	Yes
50-gene (PAM 50) (for node negative and 1–3 positive nodes)	Not determined	Yes
12-gene (EndoPredict) (node negative and 1–3 nodes)	Not determined	Yes
Breast Cancer Index (BCI)	Not determined	Yes



### PIK3CA

 Activating mutations of *PIK3CA* occur in 40% of ER<sup>+</sup>/HER2<sup>-</sup> breast cancer

>> Hyperactivation of the alpha isoform of phosphatidylinositol 3-kinase (PI3Kα)
>> Real Time PCR (exons 7, 9, and 20)

- Alpelisib is a selective inhibitor of PI3Kα
- SOLAR-1 Trial >> longer PFS and greater response with alpelisib–fulvestrant than with placebo–fulvestrant in patients with *PIK3CA*mutated, ER<sup>+</sup>/HER2<sup>-</sup>advanced breast cancer
- Resistance to Alpelisib can be related to alterations in *PTEN* and *ESR1* genes



### BRCA1&2

- PARP1 inhibition in BRCA-mutated breast cancers >> synthetic lethality
- Olaparib is a PARP-inhibitor with antitumor activity in BRCA-mutated metastatic breast cancers (OlympiAD trial)





Adapted from Liu et al. Nucleic Acids Res. 2014

## ETV6-NTRK3 fusion gene

#### Secretory breast carcinoma







- NTRK fusions occur in many very different tumors
- There are a few tumors like secretory breast cancer and congenital fibrosarcoma for which NTRK fusions are pathognomonic
- TRK inhibitors offer now the possibility to use NTRK fusion as targets in a tumor agnostic fashion

Märkl et al. Pathol Res Prac 2019

Adapted from: Church et al. Mod Pathol 2017



# Coming soon?

#### Phosphoinositide 3-kinases (PI3Ks)

- ER transcriptional activity and signaling through HER2/PI3K/AKT/mTOR increase cyclin D1 levels, activating CDK4/6 and promoting cellular progression to the S phase.
- Inhibition of CDK4/6 in the PI3K pathway can suppress mTORC1

#### Janus kinase 2 (JAK2)

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- JAK2/STAT3 regulates lipid metabolism through fatty acid β-oxidation (FAO), promoting breast cancer stemness and chemoresistance.
- Blocking FAO re-sensitize cancer cells to chemotherapy while reducing cancer stemness in vivo.



#### **PROBLEMS TO BE ADDRESSED**

### **Intra-tumor heterogeneity**



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Martelotto et al. Breast Cancer Res 2014; Lin et al. Cancers 201

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European Society for Medical Oncology (ESMO) Via Ginevra 4, CH-6900 Lugano T. +41 (0)91 973 19 00 esmo@esmo.org







