

ESR1-Mutation Testing at First Progression in ER-Positive, HER2-Negative Advanced Breast Cancer

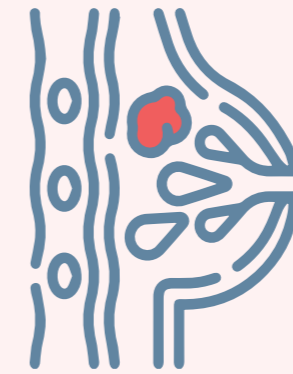
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Background

- Resistance to endocrine therapy in ER+/HER2- advanced breast cancer can be classified as intrinsic or acquired^{2,3}
- Intrinsic mutations include alterations to the PI3K/AKT/MTOR, RAS-MAPK, or FGFR1 pathways; RB1 loss; and TP53 activation⁴
- Mutations of the ER ligand-binding domain (*ESR1* mutations) are acquired predominantly during endocrine therapy in the metastatic setting, and result in constitutively active ER signalling, which promotes tumour growth and resistance to aromatase inhibitors or fulvestrant (Figure 1)^{1,5,7}



- ESR1* mutations are detected in up to 40% of patients with ER+/HER2- advanced breast cancer following initial endocrine therapy⁸
- ESR1* mutations emerge during endocrine therapy in the metastatic setting, becoming drivers of resistance, and are associated with poor outcomes in advanced breast cancer⁹
- Longer exposure to endocrine therapy in the metastatic setting increases the chance of developing an *ESR1* mutation during treatment (Figure 1)⁸⁻¹⁷



- ESR1* mutations are a predictive biomarker of response to treatment with elacestrant⁸
- Testing for *ESR1* mutations is recommended at each progression during the metastatic treatment course to identify patients who are most likely to respond to targeted treatments^{10,12,14,18}
- ASCO, NCCN, and ESMO guidelines recommend testing for *ESR1* mutations at each progression, if not detected previously¹⁹⁻²¹
- Data suggest that greater progression-free survival benefit is achieved in biomarker-selected patient subgroups with targeted agents than with standard-of-care treatment^{8,22,23}

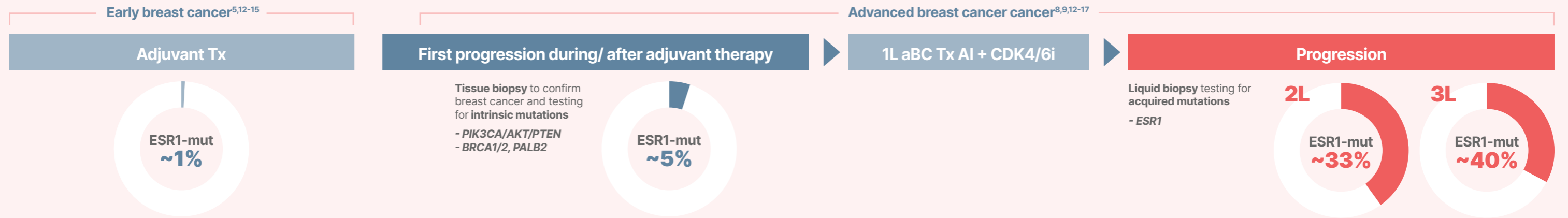


Figure 1: *ESR1*-mutation prevalence rate by treatment line.

As *ESR1* mutations occur almost exclusively after ET in the aBC setting,¹⁴ testing for *ESR1*-mut should occur at each progression if not detected previously^{2,19}



- Guidelines suggest testing for *ESR1* mutations at each progression, preferably in ctDNA (liquid biopsy). Testing in primary archived tissue is not recommended given the acquired nature of *ESR1* mutations during aBC treatment.^{20,21}
- Since acquired *ESR1* mutations are subclonal, they are not always detected with tissue biopsy^{24,25}

- Liquid biopsy confers more sensitivity to identify *ESR1* mutations, especially if ctDNA tumor fraction is >1% (Figure 2)¹⁷
- Archival tissue from the primary tumour should not be used to identify *ESR1* mutations, as they develop mainly during treatment of advanced breast cancer¹⁸

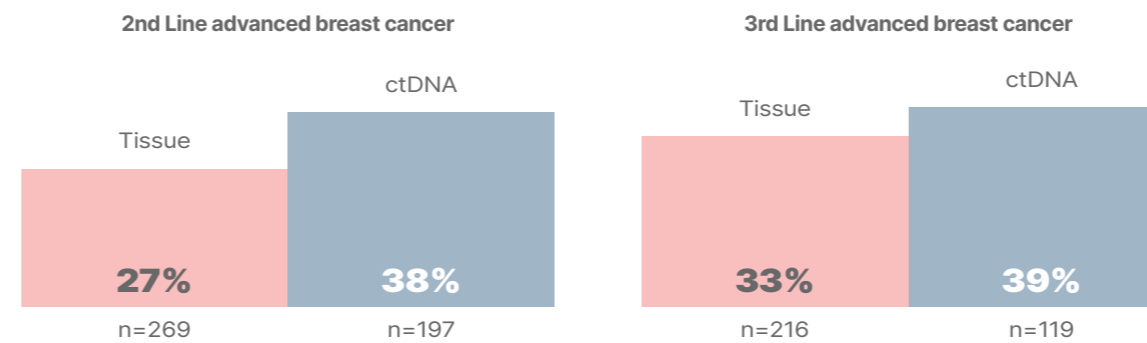


Figure 3: *ESR1*-mutation prevalence rate by treatment line in tissue and liquid biopsy.¹⁷ Percentages have been rounded to the nearest whole number. ctDNA (TF≥1%) showed a markedly higher prevalence of any of the genomic alterations assessed.

***ESR1*-mutation testing, preferably by blood-based ctDNA, is recommended at each progression during the metastatic treatment course to identify patients who are most likely to respond to targeted treatments¹⁹⁻²¹**

Abbreviations

2L: second line; 3L: third line; aBC: Advanced Breast Cancer; AKT: protein kinase B; ASCO: American Society of Clinical Oncology; ctDNA: circulating tumour DNA; ER: oestrogen receptor; ER+: oestrogen receptor positive; *ESR1*: oestrogen receptor 1; ERα: oestrogen receptor-alpha; ESMO: European Society for Medical Oncology; FGFR1: fibroblast growth factor receptor 1; HER2: human epidermal growth factor receptor 2; HER2- human epidermal growth factor receptor negative; MTOR: mechanistic target of rapamycin; mut: mutation; NCCN: National Comprehensive Cancer Network; PI3K: phosphoinositide 3-kinase; RAS-MAPK: rat sarcoma-mitogen-activated protein kinase; RB1: retinoblastoma protein; TF: tumour fraction; TP53: tumour protein p53.

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