

OPTIMA: Optimal Personalised Treatment of early breast cancer using Multi-parameter Analysis, an international randomized trial of tumor gene expression test-directed chemotherapy treatment in a largely node-positive population.

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Background

- Multi-parameter tumor gene expression assays (MPAs) are widely used to mandating • OPTIMA, ovarian by estimate individual patient residual risk in ER-positive HER2-negative nodenegative early breast cancer, allowing patients at low risk to safely avoid chemotherapy. • TAILORx provides evidence for MPA use in node-negative breast cancer. induced menopause.
- Data to support test use for 1-3 node-positive postmenopausal patients are suggestive but definitive level 1 evidence remains lacking.
- No data exist to support test use for patients with ≥ 4 involved nodes.
- MPA trials conducted to date have demonstrated chemotherapy benefit for premenopausal women in lower risk MPA categories. This is potentially explained by imbalance arising from chemotherapy-induced menopause.
- OPTIMA is a UK-initiated RCT of test-directed chemotherapy use with a noninferiority design that recruits patients at high clinical risk of recurrence.
- OPTIMA is currently recruiting in the UK, Norway & Sweden; additional international expansion is anticipated during 2021.

Patient characteristics (at 1 May 2021)

Characteristic		Invasive tumor size	
Number of participants	2123	<30mm	55%
Age: median (range)	56 (40-83)	≥30mm	45%
Menopausal status		Nodal status	
Premenopausal	36%	pN0 & pN1 mi	7%
Postmenopausal	63%	1N+	40%
Male	1%	2N+	23%
Grade		3N+	11%
Grade 1	5%	4N+	8%
Grade 2	62%	5-9 N+	11%
Grade 3	33%	Low Prosigna Score (≤60)	68%

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- OPTIMA is the only MPA trial that recruits
- OPTIMA will add to data generated by other trials tumors.
- OPTIMA welcomes additional collaborators from all nations



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Additional secondary outcome measures *Requires control-arm testing • DRFI, BCSS, OS, Health Resource use & Quality of Life

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