P1.05.25 Prognostic implications of oncogenetic pathway alterations in advanced male breast cancer



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- Results in the HR+/HER2-MaBC subgroup were consistent with those reported in the overall cohort. Figure 1.
- The number of HR+/HER2- MaBC with mutations in ESR1 (n=7), AKT1 (n=2), PTEN (n=8), mTOR (n=0), CDK4 (n=0), Rb1 (n=2) were very infrequent.

We identified **226 (1.3%) MaBC** and 17,533 (98.7%) FeBC. in FeBC.

- GATA3 mutations (MT) were associated with better OS post-endocrine therapy (mOS: 49.6 vs. 43.1 months, HR 0.83, 95% CI 0.71-0.9, p<0.01) compared to GATA3 wild type (WT), whereas in MaBC, GATA3-MT was not associated with survival post-endocrine therapy (mOS: 54.8 vs. 60.8 months, HR 1.4, 95% CI 0.63-3.2, p=0.39) compared to GATA3-WT. Figure 1.
- The presence of **TP53** mutations was adversely associated in FeBC but not in MaBC. Poorer outcomes in TP53-MT vs WT were observed in FeBC patients post-endocrine therapy (FeBC: Δ mOS 25.2 months, HR 1.7, 95% CI 1.6-1.7, p<0.01 vs MaBC: Δ mOS 16.3 months, HR 1.08 95% CI 0.43-2.7, p=0.85), with CDK4/6 inhibitor (FeBC: Δ mOS 18.2 months, HR 1.7, 95% CI 1.6-1.9, p<0.01 vs MaBC: Δ mOS 19.5 months, HR 1.35, 95% CI 0.46-2.7, p=0.57) and in the post-chemotherapy setting (FeBC: Δ mOS 18 months HR 1.3, 95% CI 1.3-1.4, p<0.01 vs MaBC: Δ mOS 34 months HR 1.7, 95% CI 0.93-3.4, p=0.07).
- CDH1, BRCA2 and PIK3CA mutations appeared to have no prognostic role in both MaBC and FeBC.

CONCLUSIONS

MaBC may recapitulate a unique tumorigenic trajectory that might differ from FeBC, as suggested by the dissimilar prognostic significance of selected genomic alterations. The data warrant further confirmation, to understand the **sex-defined differences in BC**, thus yield tailored therapeutic strategies.