

CORRECTION

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Correction to: Molecular analyses of triple-negative breast cancer in the young and elderly

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After publication of the original article [1], the authors identified an error in Fig. 4A. The correct figure is given below.

The original article has been corrected.

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1. Aine M, et al. Molecular analyses of triple-negative breast cancer in the young and elderly. *Breast Cancer Res.* 2021;23:20. <https://doi.org/10.1186/s13058-021-01392-0>.

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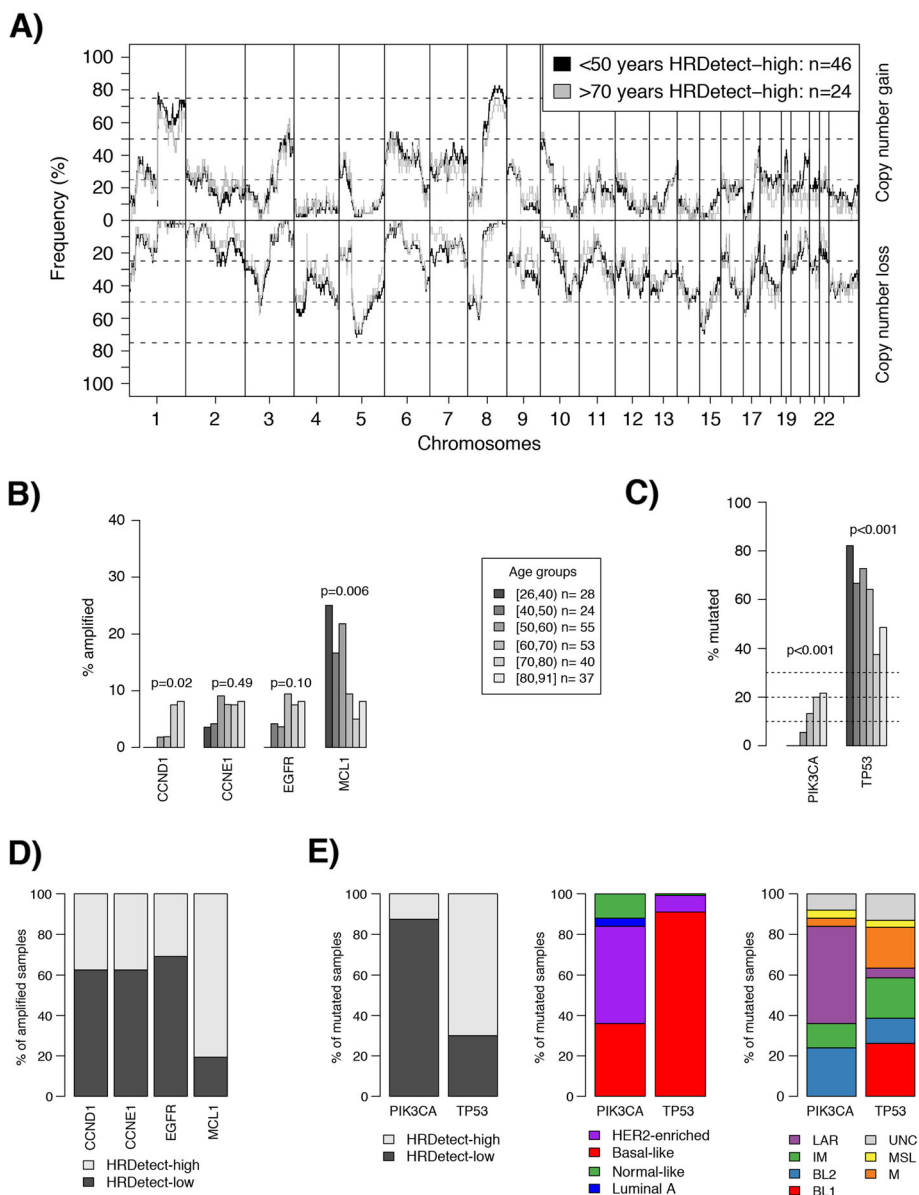


Fig. 4 Copy number alterations versus age at diagnosis in TNBC. **a** Copy number landscape of HRDetect-high patients < 50 years at diagnosis versus > 70 years at diagnosis. **b** Difference in amplification frequency of CCND1, CCNE1, EGFR, and MCL1 with age groups when analyzed in the total SCAN-B cohort. Two-sided *p* values calculated using chi-square test for trends in proportions. **c** Difference in mutation frequency of PIK3CA and TP53 with age groups when analyzed in the total SCAN-B cohort. Two-sided *p* values calculated using chi-square test for trends in proportions. **d** Proportions of amplified cases for CCND1, CCNE1, EGFR, and MCL1 according to HRDetect classification. **e** Proportions of mutated cases for PIK3CA and TP53 according to HRDetect (left), PAM50 (center), and TNBCtype (right) classifications. For age group definitions, “[” equals ≥, “)” equals <, and “]” equals ≤ for the value specified next to it