

CORRECTION

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Correction to: Prevalence of *RECQL* germline variants in Pakistani early-onset and familial breast cancer patients

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Correction to: *Hered Cancer Clin Pract* 18, 25 (2020)
<https://doi.org/10.1186/s13053-020-00159-6>

Following publication of the original article [1], a type-setting error was identified. Figure 1 was not published in full. The complete Fig. 1 is given in this correction article and the original article [1] has been corrected.

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Published online: 07 January 2021

Reference

1. Rashid MU, Muhammad N, Khan FA, et al. Prevalence of *RECQL* germline variants in Pakistani early-onset and familial breast cancer patients. *Hered Cancer Clin Pract.* 2020;18:25. <https://doi.org/10.1186/s13053-020-00159-6>.

The original article can be found online at <https://doi.org/10.1186/s13053-020-00159-6>.

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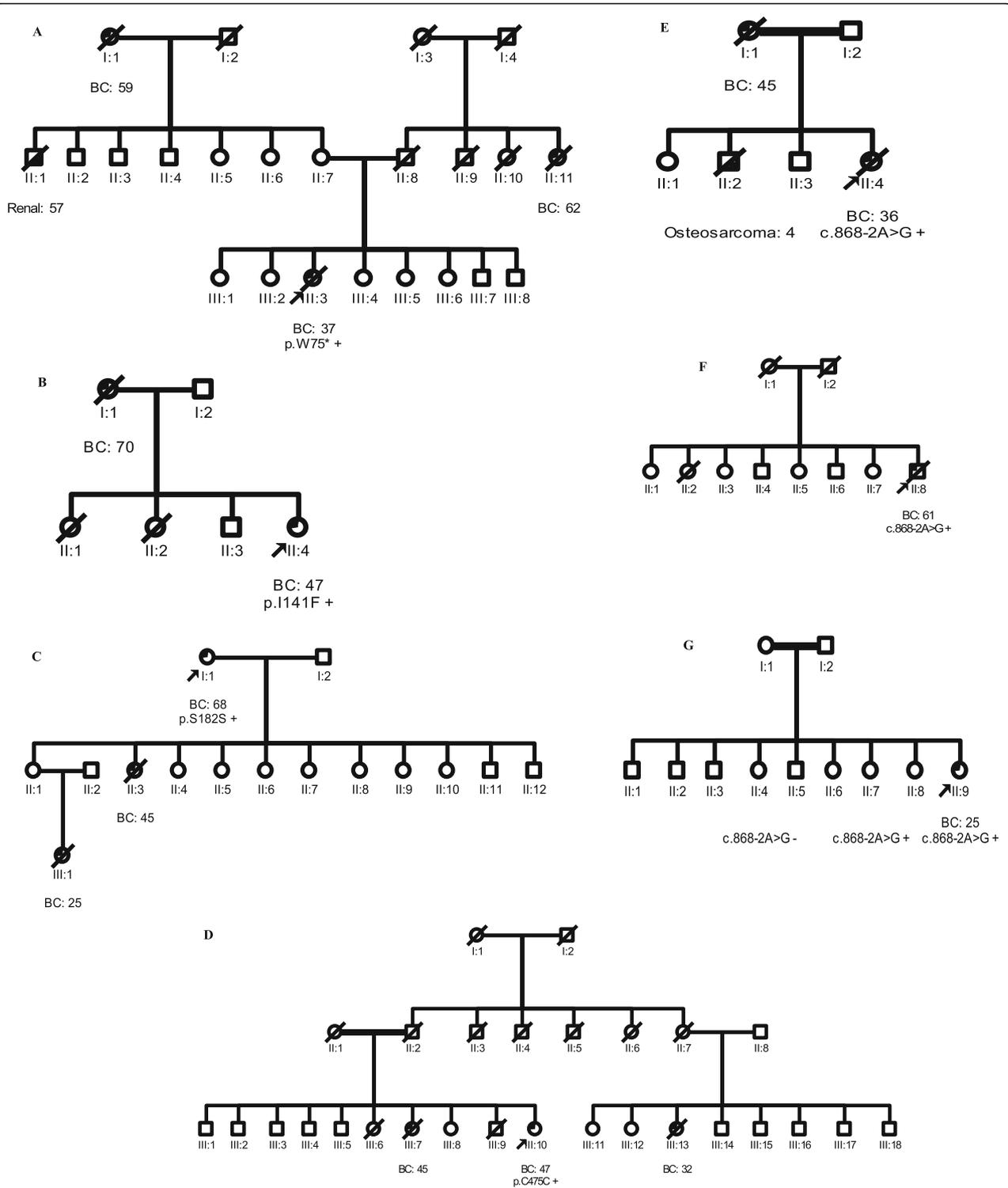


Fig. 1 Pedigrees of breast cancer patients with *RECQL* variants. **a** Family 282 carrying the pathogenic variant p.W75*. **b-d** Families 565, 649, and 625 carrying the VUS p.I141F, p.S182S, and p.C475C, respectively. **e-g** Families 471, 577 and 595 carrying the benign variant c.868-2A > G. Circles are females, squares are males, and a diagonal slash indicates a deceased individual. Symbols with filled left upper quadrant: unilateral breast cancer. Symbols with filled right lower quadrant: cancer other than breast, the name of that cancer is indicated. Double line between spouses: consanguineous marriage. Identification numbers of individuals are below the symbols. The index patient is indicated by an arrow. BC: breast cancer. The numbers following these abbreviations indicate age at cancer diagnosis. +: carrier, -: non-carrier