Hereditary Cancer in Clinical Practice

CORRECTION Open Access

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

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Following publication of the original article [1], a typesetting 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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**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.

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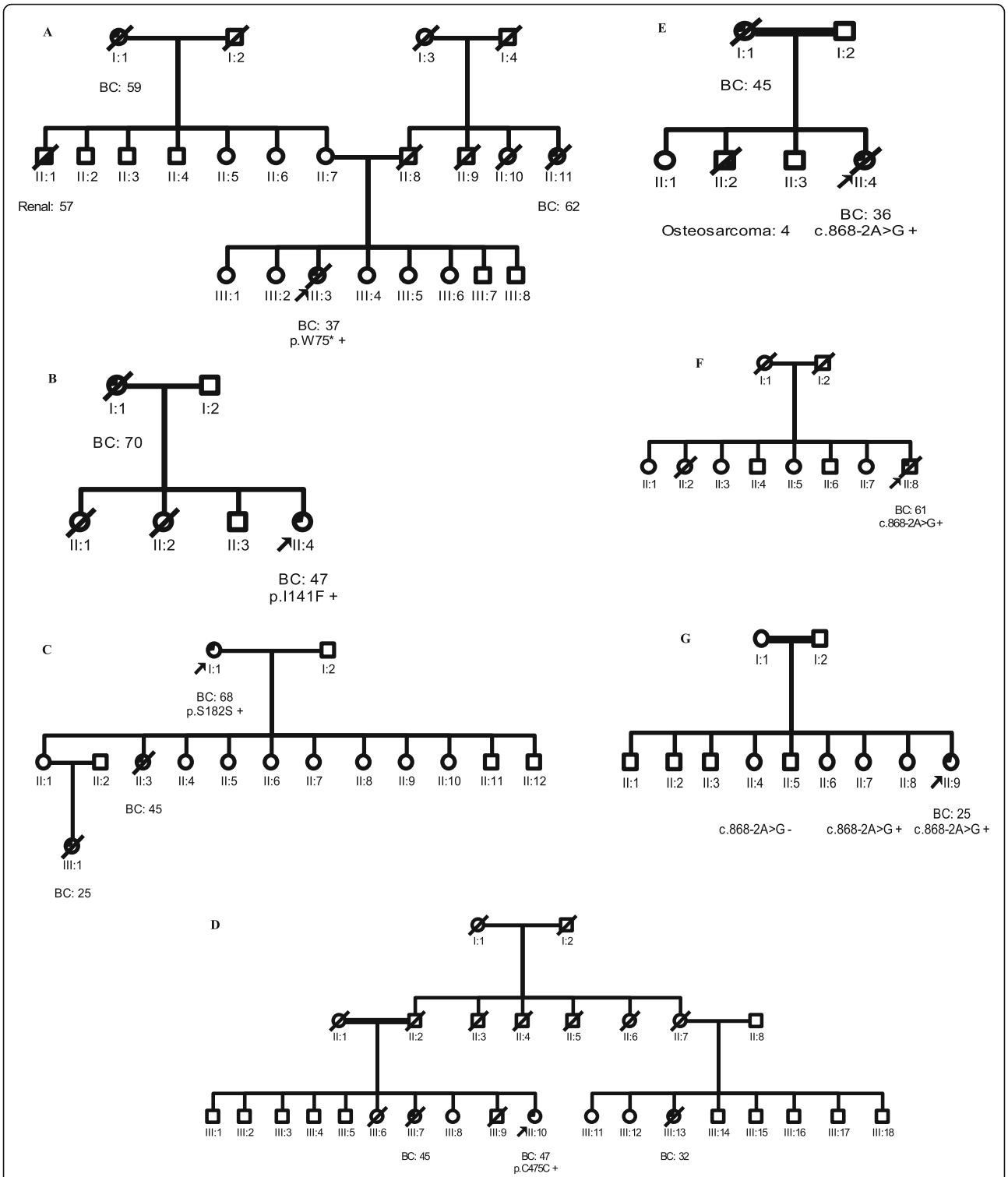


**Fig. 1** Pedigrees of breast cancer patients with *RECQL* variants. **a** Family 282 carrying the pathogenic variant p.W75*. **b-d** Families 565, 649, and 625 carrying the VUS p.I141F, p.S182S, and p.C475C, respectively. **e-g** Families 471, 577 and 595 carrying the benign variant c.868-2A > G. *Circles* 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