Author’s response to reviews

Title: Targeted capture-based NGS is superior to multiplex PCR-based NGS for hereditary BRCA1 and BRCA2 gene analysis in FFPE tumor samples

Authors:

Falk Zakrzewski (falk.zakrzewski@uniklinikum-dresden.de)
Laura Gieldon (Laura.Gieldon@uniklinikum-dresden.de)
Andreas Rump (Andreas.Rump@uniklinikum-dresden.de)
Michael Seifert (Michael.Seifert@tu-dresden.de)
Konrad Grützmann (Konrad.Gruetzmann@uniklinikum-dresden.de)
Alexander Krüger (Alexander.Krueger@uniklinikum-dresden.de)
Sina Loos (Sina.Loos@uniklinikum-dresden.de)
Silke Zeugner (Silke.Zeugner@uniklinikum-dresden.de)
Karl Hackmann (Karl.Hackmann@uniklinikum-dresden.de)
Jospeh Porrmann (Joseph.Porrmann@uniklinikum-dresden.de)
Johannes Wagner (Johannes.Wagner@uniklinikum-dresden.de)
Karin Kast (Karin.Kast@uniklinikum-dresden.de)
Pauline Wimberger (Pauline.Wimberger@uniklinikum-dresden.de)
Gustavo Baretton (Gustavo.Baretton@uniklinikum-dresden.de)
Evelin Schroeck (Evelin.Schroeck@uniklinikum-dresden.de)
Daniela Aust (Daniela.Aust@uniklinikum-dresden.de)
Barbara Klink (Barbara.Klink@uniklinikum-dresden.de)

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REQUEST
1- Please check whether your manuscript falls under the category of requiring consent to publish information relating to human participants.

Manuscripts can fall into this category if they report any details relating to individual participants including family trees, description of symptoms and individual reports of their clinical details. For all manuscripts that include details, images, or videos relating to individual participants, written informed consent for the publication of these must be obtained from the participants (or their parent or legal guardian in the case of children under 16) and a statement to this effect should appear in the manuscript. If the participant has died, then consent for publication must be sought from the next of kin of the participant. Further details on this policy can be found here:

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Your manuscript includes two family trees in Figure 4. Please either remove the trees or obtain consent to publish.

ANSWER

We obtained written consent to publish family trees displayed in Figure 4 of this work from respective participants. We also included this statement in the declarations section of the manuscript. Because of significant contribution of Johannes Wagner (Institute for Clinical Genetics, Carl Gustav Carus Faculty of Medicine, Technische Universität Dresden, Dresden, Germany) in obtaining this data we included him into the author list as co-author of this study.