Correction: Current status of use of high throughput nucleotide sequencing in rheumatology


The article has been corrected since it was published online. The funding statement has been updated as follows.

We wish to acknowledge the RARENET EU-Interreg for supporting this study. Furthermore, SB wishes to acknowledge and thank the Mainz Research School of Translational Biomedicine (TransMed) for support.

**Open access** This is an open access article distributed in accordance with the Creative Commons Attribution 4.0 Unported (CC BY 4.0) license, which permits others to copy, redistribute, remix, transform and build upon this work for any purpose, provided the original work is properly cited, a link to the licence is given, and indication of whether changes were made.

See: https://creativecommons.org/licenses/by/4.0/.

© Author(s) (or their employer(s)) 2021. Re-use permitted under CC BY. Published by BMJ.