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CORRECTION

Correction to: Biallelic Variants in the Nuclear Pore Complex Protein NUP93 Are Associated with Non-progressive Congenital Ataxia

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The original version of this article unfortunately contained mistake in Fig. 3 image.

The online version of the original article can be found at https://doi.org/10.1007/s12311-019-1010-5

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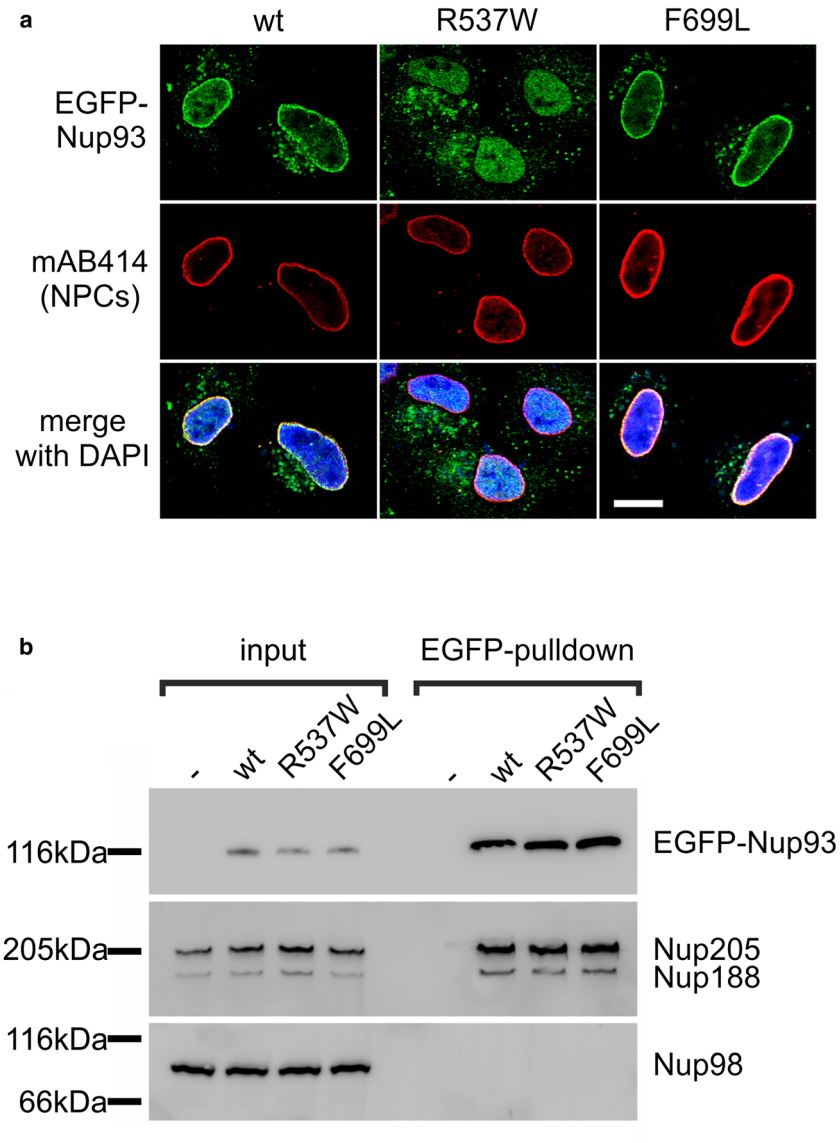

Fig. 3 The NUP93 p.R537W variant interferes with integration into nuclear pore complexes. **a** HeLa cells were transfected with different EGFP-NUP93 fusion constructs, fixed 24 h post transfection and stained with the NPC marker mAB414 (red) and DAPI. The EGFP-NUP93 signal is shown in green. Scale bar 10 μm. **b** HEK297 cells were mock transfected or with indicated EGFP-NUP93 fusion constructs. Immunoprecipitates and 5% of the inputs from cell lysates generated 24 h post-transfection were analyzed by western blotting for the NUP93 interactors NUP188 and NUP205. NUP98 served as a negative control for a noninteracting nucleoporin

The western blot presented in Figure b shows white lines in the published version not seen in the original picture. With this, the correct Fig. 3 is hereby published.

Also the authors would like to replace the ESM3 of the article.

The original article has been corrected.