

Production, crystallization and X-ray diffraction analysis of two nanobodies against the Duffy binding-like (DBL) domain DBL6 ϵ -FCR3 of the *Plasmodium falciparum* VAR2CSA protein.

Corrigendum

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A correction is made to Fig. 1 and the text of the article by Vuchelen *et al.* (2013, *Acta Cryst.* **F69**, 270–274).

In the article by Vuchelen *et al.* (2013) there is a mistake in Fig. 1. The second line of Nb19 is a copy of the second line of Nb07, instead of the correct Nb19 line. This results in a difference of four amino acids. In §3, it is mentioned that there is one amino-acid substitution between the two nanobodies but this should be five (to include the four from the second line). Also the sequence identity should be 94% instead of 98%. A correct version of Fig. 1 is published here.

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      10      20      30      40      50      60
Nb2907 QVQLQESGGG LVQAGGSLRL SCAASGVVFFS LPAMGWYRQA PGQORELVAG ISSGGNTNYA
Nb2919 QVQLQESGGG LVQAGGSLRL SCAASGSPPFS LPAMGWYRQA PGKQRELVAG ISSDGNNTNYA
                                CDR1          CDR3          CDR2

Nb2907 DSVKGRFAIS RDDAKNMVYL QMNSAKPEDT GYYCNSILY RLNWGQSTQV TVSSHHHHHH
Nb2919 DSVKGRFTIS RDDAKNTVYL QMNSLKPEDT GYYCNSILY RLNWGQSTQV TVSSHHHHHH
      70      80      90      100     110     120
  
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Figure 1

Corrected sequence alignment of Nb2907 and Nb2919. The complementary determining regions (CDR) are shown in red boxes.

References

Vuchelen, A., Pardon, E., Steyaert, J., Gamain, B., Loris, R., van Nuland, N. A. J. & Ramboarina, S. (2013). *Acta Cryst.* **F69**, 270–274.