

addenda and errata

Production, crystallization and X-ray diffraction analysis of two nanobodies against the Duffy binding-like (DBL) domain DBL6ε-FCR3 of the *Plasmodium falciparum* VAR2CSA protein. Corrigendum

Anneleen Vuchelen,^{a,b} Els Pardon,^{a,b} Jan Steyaert,^{a,b}
Benoît Gamain,^{c,d,e,f} Remy Loris,^{a,b*} Nico A. J. van
Nuland^{a,b} and Stéphanie Ramboarina^{a,b*}

^aStructural Biology Brussels, Department of Biotechnology (DBIT), Vrije Universiteit Brussel, Pleinlaan 2, B-1050 Brussels, Belgium, ^bMolecular Recognition Unit (MoRe), Department of Structural Biology, VIB, Pleinlaan 2, B-1050 Brussels, Belgium, ^cInserm, UMRS665, F-75015 Paris, France, ^dUniversité Paris Diderot, F-75013 Paris, France, ^eInstitut National de la Transfusion Sanguine, F-75015 Paris, France, and ^fLaboratoire d'Excellence Gr-Ex, France

Correspondence e-mail: reloris@vub.ac.be, stephanie.ramboarina@cea.fr

A correction is made to Fig. 1 and the text of the article by Vuchelen *et al.* (2013, *Acta Cryst. F* **69**, 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.

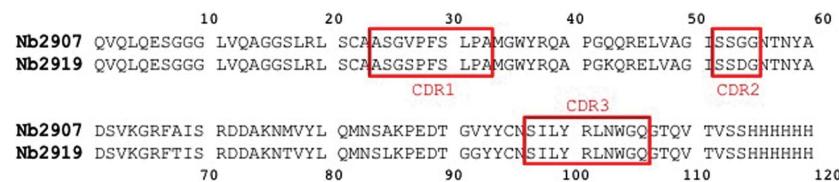


Figure 1

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. F* **69**, 270–274.