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# Erratum: Molecular features of interaction between VEGFA and anti-angiogenic drugs used in retinal diseases: a computational approach

Frontiers Production Office \*

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## An erratum on

**Molecular features of interaction between VEGFA and anti-angiogenic drugs used in retinal diseases: a computational approach**

by Platania, C. B. M., Di Paola, L., Leggio, G. M, Romano, G. L., Drago, F., Salomone, S., et al. (2015)  
*Front. Pharmacol.* 6:248. doi: 10.3389/fphar.2015.00248

### Reason for Erratum:

Due to a typesetting error, the article was published with incorrect values in **Table 4**. The publisher apologizes for this error and the correct version of **Table 4** appears below. This error does not change the scientific conclusions of the article in any way.

## REFERENCES

Papadopoulos, N., Martin, J., Ruan, Q., Rafique, A., Rosconi, M. P., Shi, E., et al. (2012). Binding and neutralization of vascular endothelial growth factor (VEGF) and related ligands by VEGF Trap, ranibizumab and bevacizumab. *Angiogenesis* 15, 171–185. doi: 10.1007/s10456-011-9249-6

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**TABLE 4 | MM-PBSA results compared to experimental binding parameters.**

Complex	Binding parameters			MM-PBSA energy terms (KJ/mol)				
	$K_{on}/10^5$ ( $M^{-1} s^{-1}$ )	$K_{off}/10^{-5}$ ( $s^{-1}$ )	$K_D$ ( $\mu M$ )	$\Delta E_{binding}$	$\Delta E_{VdW}$	$\Delta E_{electrostatic}$	$\Delta G_{Polar}$	$\Delta G_{Apolar}$
Ranibizumab/VEGFA	1.60	0.73	46	$-760 \pm 40$	$-418 \pm 5$	$-160 \pm 20$	$410 \pm 30$	$-592 \pm 7$
Fab-bevacizumab/VEGFA	5.30	3.10	58	$-807 \pm 30$	$-362 \pm 10$	$-252 \pm 20$	$343 \pm 30$	$-536 \pm 7$
VEGFR1d2_R2d3/VEGFA	410	2.01	0.49	$-1440 \pm 90$	$-307 \pm 50$	$-1433 \pm 100$	$1050 \pm 100$	$-750 \pm 40$

*Kinetic and binding parameters are from Papadopoulos et al. (2012).*