



Correction: Macrocyclic peptides as inhibitors of WDR5–lncRNA interactions

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Correction for 'Macrocyclic peptides as inhibitors of WDR5–lncRNA interactions' by Jen-Yao Chang *et al.*, *Chem. Commun.*, 2023, **59**, 10656–10659, <https://doi.org/10.1039/D3CC03221C>.

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The authors regret that the K_D values given for entries 14 and 15, in column 3 of Table 1, were incorrectly given in the original article. The authors would like to clarify the correct K_D values for entries 14 and 15 are $0.53 \pm 0.06 \mu\text{M}$ and $5.2 \pm 0.3 \mu\text{M}$ respectively. These errors do not affect the conclusions of the work and the updated version of Table 1 is shown below.

The authors also regret the omission of an acknowledgements section from the article at the time of publication. The acknowledgements section for the article is:

We would like to acknowledge Christiane Heitbrink (TU Dortmund) for measuring HRMS data. We acknowledge the European Synchrotron Radiation Facility for provision of synchrotron radiation facilities and we would like to thank P. Carpentier for assistance in using beamline ID30A-3. We also thank the Ministry of Education (Taiwan) for funding Jen-Yao Chang with a Government Scholarship for Overseas Study. Open Access funding provided by the Max Planck Society.

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.



Table 1 Sequences of WBM site binding peptides and their affinity for WDR5 as determined by fluorescence polarization. Residues used for side-chain macrocyclization are highlighted in bold. See Fig. S1 (ESI) for full structural details of the peptides

No.	Sequence		K_D (μM)
	371	381	
1	FITC-PEG-EDEEVDVTSVD-NH ₂		2.05 ± 0.42
2	FITC-PEG-EDEEVDVTSV-NH ₂		7.11 ± 2.80
3	FITC-PEG-EDEEVDVTS-NH ₂		20.92 ± 8.72
4	FITC-PEG-EDEEVDVT-NH ₂		29.91 ± 15.13
5	Ac-EDEEVDVTSVD-PEG-FITC		2.42 ± 0.14
6	Ac-DEEVDVTSVD-PEG-FITC		6.74 ± 1.04
7	Ac-EEVDVTSVD-PEG-FITC		23.45 ± 6.89
8	Ac-EVDVTSVD-PEG-FITC		>37.5
9	Ac-VDVTSVD-PEG-FITC		>37.5
10	FITC-O2Oc-DEEVDVTSV-NH ₂		10.39 ± 2.69
11	FITC-O2Oc-DEEVDV Dap V-NH ₂		21.86 ± 5.29
12	FITC-O2Oc-DEEVDV Dab V-NH ₂		1.18 ± 0.07
13	FITC-O2Oc-DEEVDV Orn V-NH ₂		2.67 ± 0.10
14	258	268	0.53 ± 0.06
	FITC-O2Oc-DEEEIDVVSVE-NH ₂		
15	407	417	9.87 ± 4.78
	FITC-O2Oc-FSDDL DVVGDG-NH ₂		
16	FITC-O2Oc-DEEEIDV Dab V-NH ₂		0.33 ± 0.02
17	FITC-O2Oc-DEEEIDV Orn V-NH ₂		1.96 ± 0.18
18	FITC-O2Oc-DEEEIDV Dab VE-NH ₂		0.11 ± 0.02
19	FITC-O2Oc-DEEEIDV Orn VE-NH ₂		0.56 ± 0.04
20	FITC-O2Oc-DEEEIDV Dab VE-NH ₂		0.23 ± 0.01
21	FITC-O2Oc-DEEEIDV Dab VE-NH ₂		0.18 ± 0.01
22	FITC-O2Oc-DEEEID I T Dab VE-NH ₂		0.43 ± 0.02
23	FITC-O2Oc-DEEEI E V Dap VE-NH ₂		>37.5
24	FITC-O2Oc-DEEEI Dab V D VE-NH ₂		>37.5
25	FITC-O2Oc-DEEEI Dap V V EVE-NH ₂		>37.5

