

## CORRECTION

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[View Journal](#) | [View Issue](#)Cite this: *Chem. Sci.*, 2021, 12, 10395**Correction: Development of a structure-based computational simulation to optimize the blocking efficacy of pro-antibodies**Bo-Cheng Huang,<sup>a</sup> Yun-Chi Lu,<sup>bc</sup> Jun-Min Liao,<sup>bc</sup> Hui-Ju Liu,<sup>d</sup> Shih-Ting Hong,<sup>d</sup> Yuan-Chin Hsieh,<sup>e</sup> Chih-Hung Chuang,<sup>cf</sup> Huei-Jen Chen,<sup>d</sup> Tzu-Yi Liao,<sup>d</sup> Kai-Wen Ho,<sup>d</sup> Yeng-Tseng Wang<sup>\*h</sup> and Tian-Lu Cheng<sup>\*abcf</sup>

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