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Corrigendum to “Discovery of small molecules through pharmacophore modeling, docking and molecular dynamics simulation against Plasmodium vivax Vivapain-3 (VP-3)” [Heliyon 4(5), May 2018, Article e00612]

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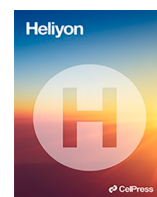
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Corrigendum



Corrigendum to “Discovery of small molecules through pharmacophore modeling, docking and molecular dynamics simulation against *Plasmodium vivax* Vivapain-3 (VP-3)” [Heliyon 4(5), May 2018, Article e00612]

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The authors apologize for any potential confusion caused by the phrase “blunder rate” in the 3.1 3D-model construction and validation section in the original article “Discovery of small molecules through pharmacophore modeling, docking and molecular dynamics simulation against *Plasmodium vivax* Vivapain-3 (VP-3)” that was published in Heliyon Journal, 2018 May 5; 4(5):e00612.

The program affords an “overall quality factor” assessment that is outlined because the share of the macromolecule that the considered “**error rate**” collapse less than the 95 % statistical refusal border (lines 18–21 of page no 10).

To improve the article, the authors correct the phrase from “blunder rate” to “error rate” in line 19 of page no. 10, section 3.1.

The authors would like to make apologies for any inconvenience caused.

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