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Erratum to 'Combining bioinformatics, cheminformatics, functional genomics and whole organism approaches for identifying epigenetic drug targets in Schistosoma mansoni' [IJP Drugs Drug Resist. 8 (2018) 559–570]

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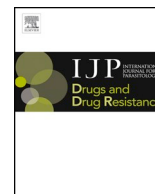
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Erratum to ‘Combining bioinformatics, cheminformatics, functional genomics and whole organism approaches for identifying epigenetic drug targets in *Schistosoma mansoni*’ [IJP Drugs Drug Resist. 8 (2018) 559–570]

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The publisher regrets to inform that there was errors in the column headers of table 1 published in the original article. The corrected table 1 is as below:

Table 1
L8 and L10 antischistosomal activity and cytotoxicity summary.

Compound	EC ₅₀ on schistosomula phenotype	EC ₅₀ on schistosomula motility	EC ₅₀ on juveniles	EC ₅₀ on adult worms	CC ₅₀ on HepG2 cells	SI CC ₅₀ /EC ₅₀ schistosomula (phenotype/motility)	SI CC ₅₀ /EC ₅₀ juveniles	SI CC ₅₀ /EC ₅₀ adult worms
L8	4.19 μM	5.86 μM	3.91 μM	> 50 μM	17.79 μM	4.25/3.04	4.55	< 0.36
L10	2.49 μM	1.46 μM	2.28 μM	14.18 μM	21.05 μM	8.45/14.42	9.23	1.48

The publisher would like to apologise for any inconvenience caused.

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