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Erratum to: 'NETTAB 2014: From highthroughput structural bioinformatics to integrative systems biology'

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Unfortunately, the original version of this article [1] contained a few errors. The editorial department of BMC Bioinformatics would like to apologize and inform its readers of the following revisions.

The first sentence of the last paragraph "The manuscript "Weighted integration of multi-omic layers of conditions in genome-scale models" [12]" should be "The manuscript "Multiplex methods provide effective integration of multi-omic data in genome-scale models" [12]".

In the reference list, the title of reference [12] "Weighted integration of multi-omic layers of conditions in genome-scale models" should be "Multiplex methods provide effective integration of multi-omic data in genome-scale models".

Reference 7 has 2016;17 Suppl 3:S2 as its year, volume and supplement number and this should be 2016;17(Suppl 4):54 instead.

Reference 8 has 2016;17 Suppl 3:S3 as its year, volume and supplement number and this should be 2016;17(Suppl 4):57 instead.

Reference 9 has 2016;17 Suppl 3:S4 as its year, volume and supplement number and this should be 2016;17(Suppl 4):69 instead.

Reference 11 has 2016;17 Suppl 3:S5 as its year, volume and supplement number and this should be 2016;17(Suppl 4):61 instead.

Reference 12 has 2016;17 Suppl 3:S6 as its year, volume and supplement number and this should be 2016;17(Suppl 4):83 instead.

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Reference

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