

## Erratum

### High-throughput inference of protein–protein interfaces from unassigned NMR data

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**The following sentence was published in the Summary of the above paper:**

We introduce a geometric optimization problem where we must cluster the cells in an arrangement on the boundary of a 3-manifold. The arrangement is induced by a spherical quadratic form, which in turn is parameterized by a  $SO(3)\times R^2$ .

**It should read as follows:**

We introduce a geometric optimization problem where we must cluster the cells in an arrangement on the boundary of a 3-manifold, where the arrangement is induced by a spherical quadratic form.

**The publisher wishes to apologise for this error.**