Erratum to: Predicting the functions of a protein from its ability to associate with other molecules

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Upon publication of the original article [1] it was found that author corrections for Table 1 were not correctly implemented. You can now find the correct Table 1 below and also updated in the original article.

Table 1 The distribution of semantically related and semantically unrelated co-occurrences of molecule \( m_i \) and Protein \( p \) Pair in an Abstract \( A_j \)

|                      | Yes | No  | Total |
|----------------------|-----|-----|-------|
| \( m_i \) and \( p \) co-occur in the same sentence | \( o_{11} \) | \( o_{12} \) | \( R_1 \) |
| \( m_i \) and \( p \) are semantically related      |     |     |       |
| No                   | \( o_{21} \) | \( o_{22} \) | \( R_2 \) |
| Total                | \( c_1 \)   | \( c_2 \)   | \( N \)   |

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References

1. Taha K, Yoo PD. Predicting the functions of a protein from its ability to associate with other molecules. BMC Bioinformatics. 2016;17:34.