

ERRATUM

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Erratum to: ppiPre: predicting protein-protein interactions by combining heterogeneous features

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Erratum

The authors wish to acknowledge that the software package associated with our Research Article [1], under the name 'ppiPre', re-used software code for some of its functions from an existing software package, GOSemSim [2], without proper attribution and in breach of the software's licencing terms. Additionally we neglected to cite the article by Yu et al. [3] describing the GoSemSim software.

The software code from GoSemSim [2] is used in the implementation of two GO semantic similarity measures, TCSS and IntelliGO. ppiPre additionally implements a KEGG-based similarity measure and three topological similarity measures, and integrates features with a support vector machine.

We have now updated our software package such that it is licensed under a compatible GPL version 2 licence, and revised the package to give the appropriate attribution.

We apologize for any inconvenience this oversight may have caused.

Availability and Requirements

- **Project name:** ppiPre.
- **Project home page:** <http://cran.r-project.org/web/packages/ppiPre/index.html>.
- **Operating system(s):** Platform independent.
- **Programming language:** R.
- **Other requirements:** None.
- **License:** GPL-2.
- **Any restrictions to use by non-academics:** None.

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References

1. Yue D, Lin G, Bingbo W. ppiPre: predicting protein-protein interactions by combining heterogeneous features. *BMC Syst Biol.* 2013;7 Suppl 2:S8.
2. Guangchuan Yu. GOSemSim. <http://www.bioconductor.org/packages/release/bioc/html/GOSemSim.html>. (Accessed 7 July 2015)
3. Yu G, Li F, Qin Y, Bo X, Wu Y, Wang S. GOSemSim: an R package for measuring semantic similarity among GO terms and gene products. *Bioinformatics.* 2010;26:976–8.

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