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Author Correction: Exploring the Sequence-based Prediction of Folding Initiation Sites in Proteins

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Correction to: *Scientific Reports* https://doi.org/10.1038/s41598-017-08366-3, published online 18 August 2017

This Article contains errors in the Methods section under subheading 'Code availability'.

"The EFoldMine code is available from https://www.dropbox.com/s/eslk4bkpflsgiia/code.zip (note: this is a temporary link, will be replaced when accepted), for academic use only as a stand-alone package with the following dependencies: Python2.7 including numpy ($>=1.6.1$), ScipPy ($>=0.9$) and the scikit-learn package (http://scikit-learn.org/stable/install.html)."

should read:

"The EFoldMine code is available from https://figshare.com/articles/EFoldMine_code/5649373, for academic use only as a stand-alone package with the following dependencies: Python2.7 including numpy ($>=1.6.1$), ScipPy ($>=0.9$) and the scikit-learn package (http://scikit-learn.org/stable/install.html)."

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