
crystal Documentation

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Brent S. Pedersen

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crystal is a python module to assign significance to clusters of correlated data. The most likely use-case is for methylation data, but it can be use for any correlated data.

New users should check-out the introductory IPython notebook *here* <<http://nbviewer.ipython.org/github/brentp/crystal/blob/master/notebooks/introductory.ipynb>>

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