Protocol Update for large-scale genome and gene functi classification system (v.14.0)

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Citation Report

#	Article	IF	Citations
2	Mining cancer biology through bioinformatic analysis of proteomic data. Expert Review of Proteomics, 2019, 16, 733-747.	1.3	21
3	Population-specific long-range linkage disequilibrium in the human genome and its influence on identifying common disease variants. Scientific Reports, 2019, 9, 11380.	1.6	22
4	Comparative transcriptomic analysis and structure prediction of novel Newt proteins. PLoS ONE, 2019, 14, e0220416.	1.1	13
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9	Knockdown of estrogen receptor Î² increases proliferation and affects the transcriptome of endometrial adenocarcinoma cells. BMC Cancer, 2019, 19, 745.	1.1	13
10	Impact of merging commercial breeding lines on the genetic diversity of Landrace pigs. Genetics Selection Evolution, 2019, 51, 60.	1.2	18
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