

Correction to Quantitative Analysis of the Human AKR Family Members in Cancer Cell Lines Using the mTRAQ/MRM Approach

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Supporting Information

The authors have added the following statement to the Supporting Information associated with the original article (DOI: 10.1021/pr301153z): MRM data sets associated with this paper have been deposited to both the GigaScience database¹ and the PeptidesAtlas SRM Experiment Library (PASSEL) (<http://www.peptideatlas.org/passel/>, dataset identifier: PASS00087)². The complete Supporting Information associated with the original article has been replaced and also is available with this Correction.

■ ASSOCIATED CONTENT

Supporting Information

Corrected Supporting Information associated with the original article (pr301153z). This material is available free of charge via the Internet at <http://pubs.acs.org>.

■ REFERENCES

(1) Zhang, S.; Wen, B.; Zhou, B.; Yang, L.; Cha, C.; Xu, S.; Qiu, X.; Wang, Q.; Sun, H.; Lou, X.; Zi, J.; Zhang, Y.; Lin, L.; Liu, S. Quantitative proteomics data using mTRAQ/MRM looking at human AKR family members in cancer cell lines GigaScience Database. <http://dx.doi.org/10.5524/100047>, 2013.

(2) Farrah, T.; Deutsch, E. W.; Kreisberg, R.; Sun, Z.; Campbell, D. S.; Mendoza, L.; Kusebauch, U.; Brusniak, M. Y.; Huttenhain, R.; Schiess, R.; Selevsek, N.; Aebersold, R.; Moritz, R. L. PASSEL: the PeptideAtlas SRM experiment library. *Proteomics* **2012**, *12* (8), 1170–5.