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Comparison of three parallelism assessment methods of biomarker quantification by LC-MS/MS: a case study of the bioanalysis of creatinine in human urine samples.

DOI: 10.4155/bio-2021-0255 Bioanalysis, 2022, , .

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4	Development of a UPLC-MRM-based targeted proteomic method to profile subcellular organelle marker proteins from human liver tissues. <i>Scientific Reports</i> , 2022 , 12,	4.9	, O
3	Criteria of slope and endogenous level for selection of matrix in the quantitation of endogenous compounds by LC-MS/MS. 2022 , 14, 807-816		1
2	Multi-laboratory evaluation of immunoaffinity LCMS-based glucagon-like peptide-1 assay. 2023 , 15, 207-218		O
1	Signature peptide selection workflow for biomarker quantification using LCMSD ased targeted proteomics.		0