

Project Name: con_sp.hsa.canonical.isoforms_junc.exons_peptides

Normalyzer (ver 1.4.0)

Report created on: 2020-05-07

Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

Documentation for analyzing this report can be found at <http://quantitativeproteomics.org/normalyzer/help.php>





























