Manuscript, Supplementary material.
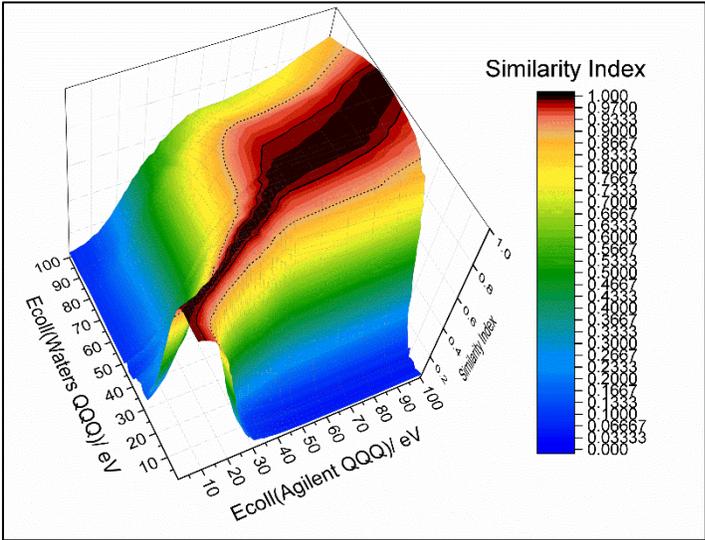
Supplementary material

Fig.S1. Similarity indices show as a 3D contour map for all combinations of collision energies determined on Waters and Agilent QQQ instruments (0-100 eV range).
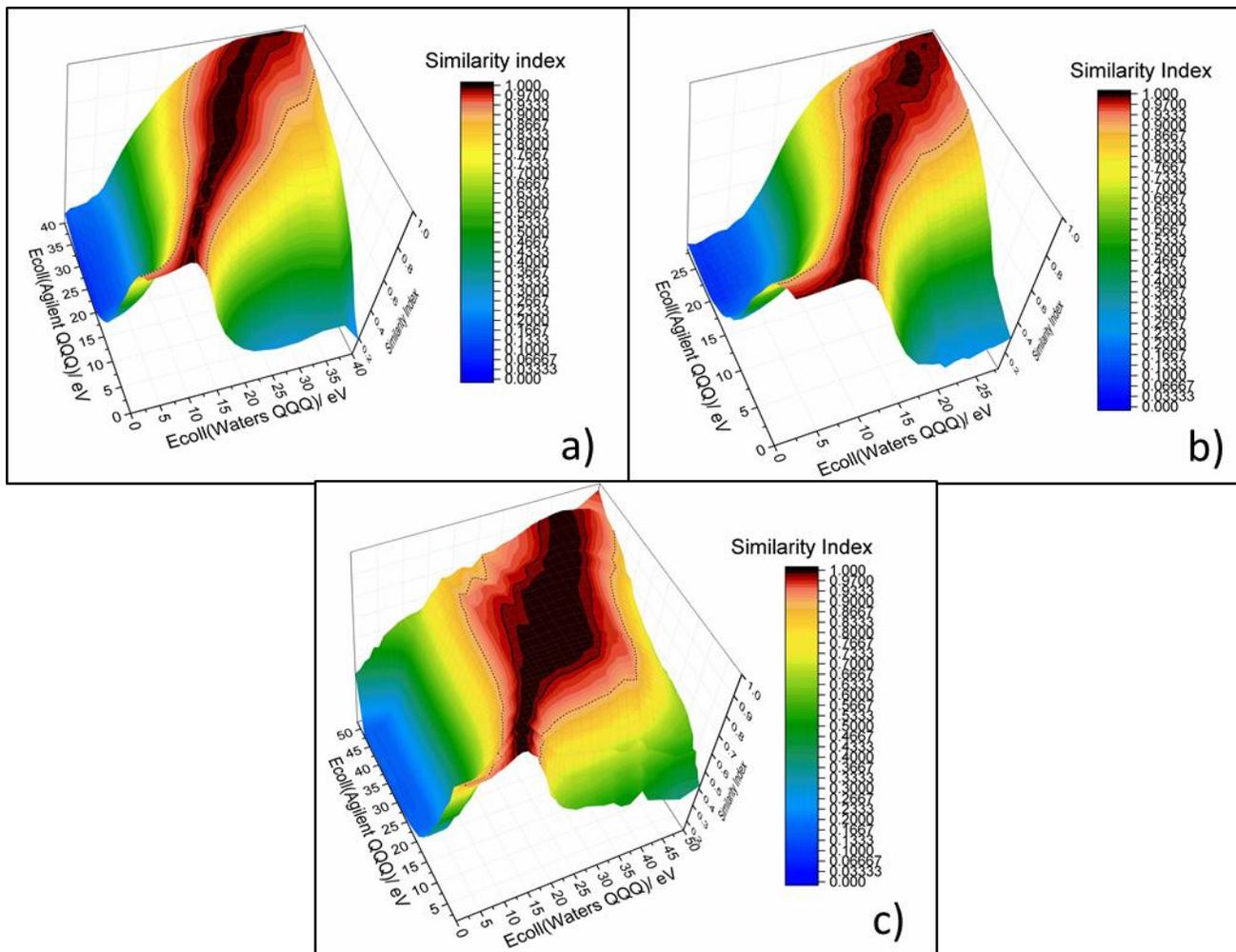
Fig. S2. Similarity indices show for all combinations of collision energies determined on Waters and Agilent QQQ instruments using a) α-aminoadipic acid , b) aminocaproic acid, c) adenosine.
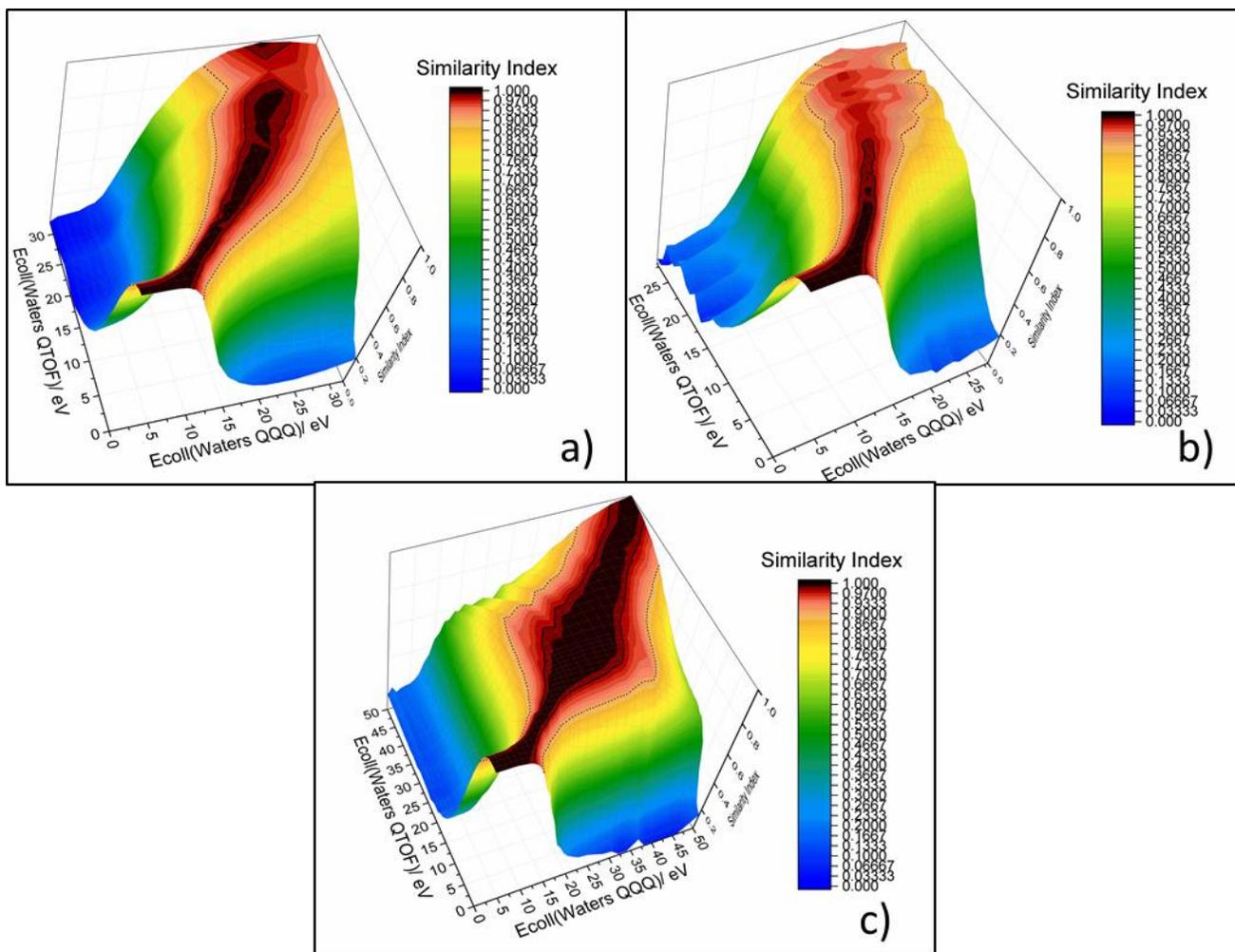
Fig. S3. Similarity indices shown as a 3D contour map and for all combinations of collision energies determined on Waters QTOF and Agilent QQQ instruments using a) α-aminoadipic acid , b) aminocaproic acid, c) adenosine.
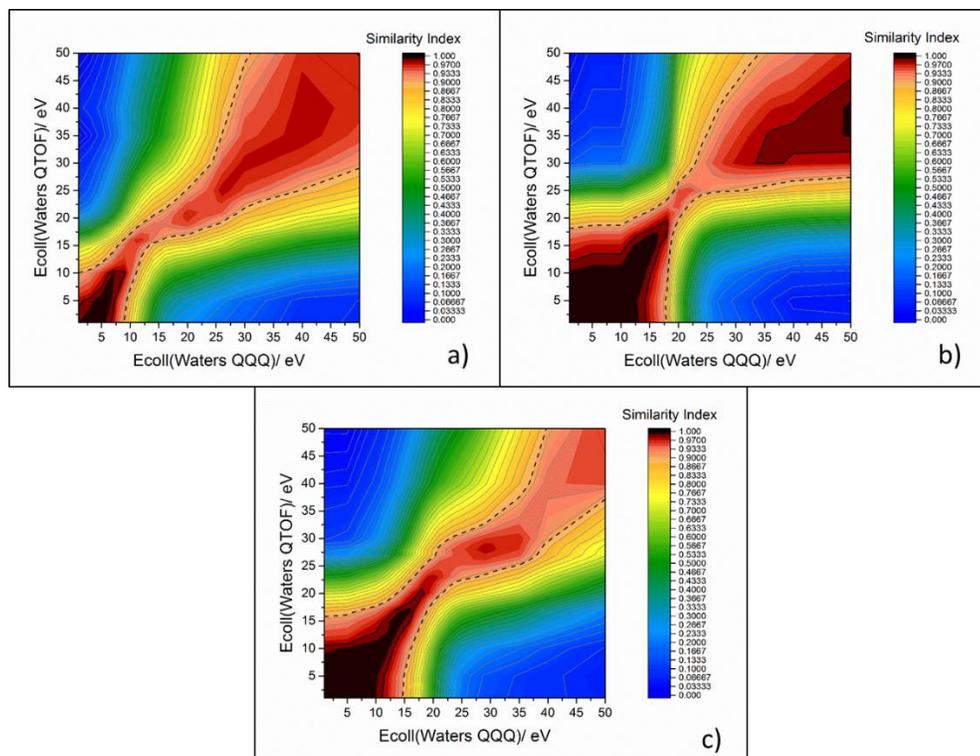
Fig. S4. Similarity indices shown as a 3D contour map and for all combinations of collision energies determined on Waters QTOF and QQQ instruments using studied doubly protonated tryptic peptides derived from bovine serum albumin a) m/z = 582.3189, sequence: LVNELTEFAK , b) m/z= 653.3617, sequence: HLVDEPQNLIK, c) m/z = 740.4013, sequence: LGEYGFQNALIVR.