

Proteomic Analysis of the Organ of Corti Using Nanoscale Liquid Chromatography Coupled with Tandem Mass Spectrometry

Supplementary Material

Figure S1. (A) Representative base peak chromatography for peptides obtained with the nanoLC system coupled with LTQ-Orbitrap XL for MS/MS analysis. (B) MS/MS spectrum for a phosphopeptide HAFSPVASVESASGETLHSPK, which was reconciled as the triply charged peptide ion arising from alpha-2-HS-glycoprotein. The phosphorylation site is Ser-314.

