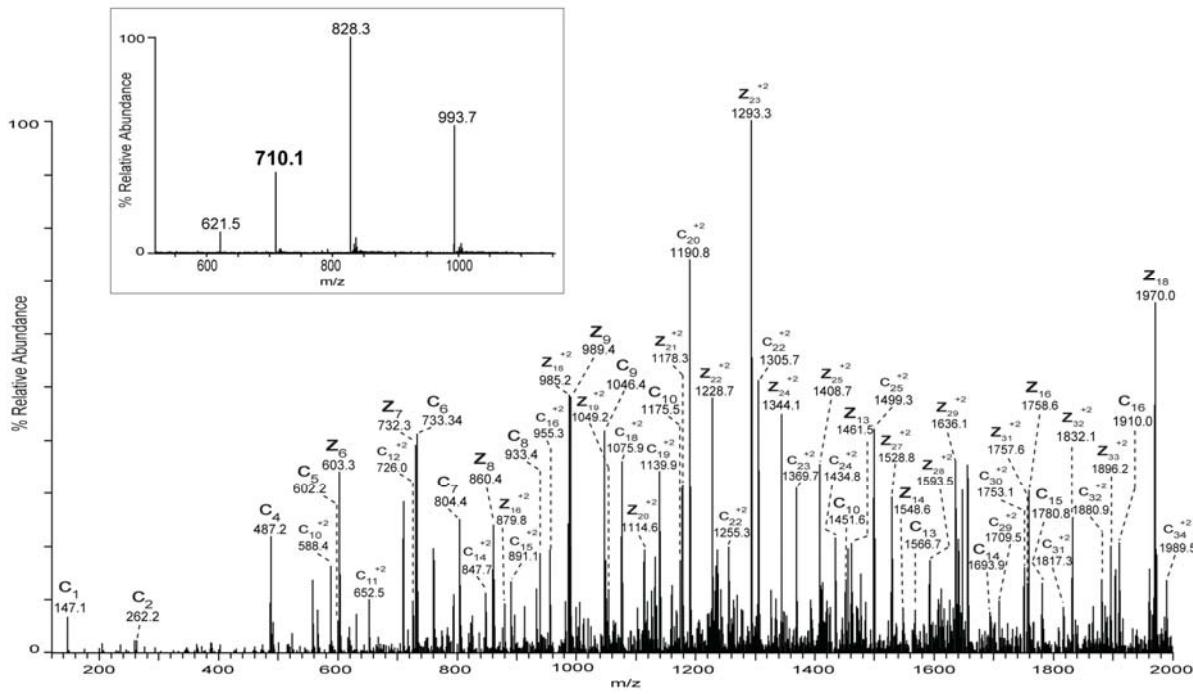


SUPPLEMENTAL INFORMATION**Spatially-Directed Protein Identification from Tissue Sections by Top-Down LC-MS/MS with
Electron Transfer Dissociation****Kevin L. Schey, David M. Anderson, and Kristie L. Rose**

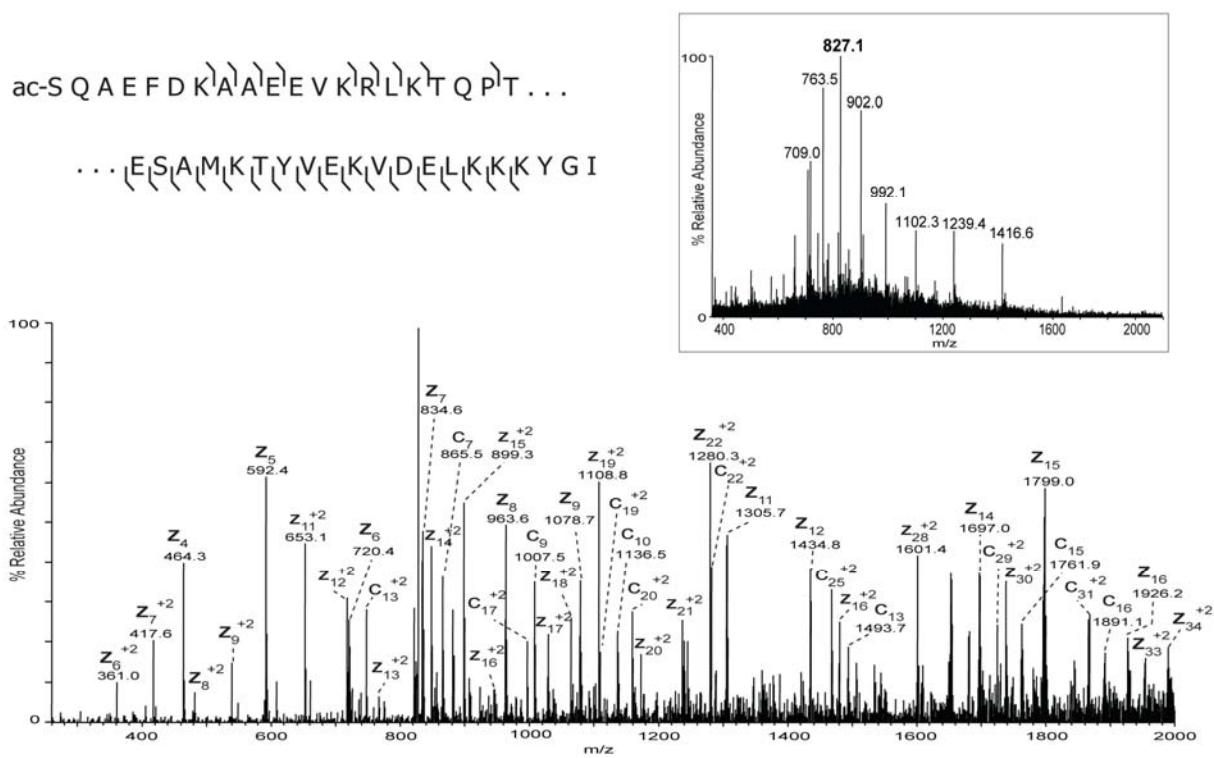
Department of Biochemistry and Mass Spectrometry Research Center,
Vanderbilt University School of Medicine, Nashville, TN

Abstract: The supplemental information provided in this document provides further support of the spatially-directed tissue extraction method. Specifically, this supplemental information contains two ETD tandem mass spectra showing abundant sequence ions used to identify proteins from tissue extracts.

ac-S¹D¹K P¹D¹M¹A¹E¹I¹E¹K¹F¹D¹K¹S¹K¹L¹K¹T¹E¹T¹Q¹E¹K¹N P¹L P¹S¹K¹E¹T¹I¹E¹Q¹E¹K¹Q A G E S



Supplemental Figure 1. ETD MS/MS spectrum of the $[M+7H]^{+7}$ precursor ion (m/z 710.1) of thymosin β 4. The observed distribution of this protein is displayed in blue (m/z 4963) in the MALDI image of mouse brain (Figure 2).



Supplemental Figure 2. ETD MS/MS spectrum of the $[M+12H]^{12+}$ precursor ion (m/z 827.1) of Acyl CoA-binding protein. The observed distribution of this protein is displayed in blue (m/z 9914) in the MALDI image of mouse kidney (Figure 3).