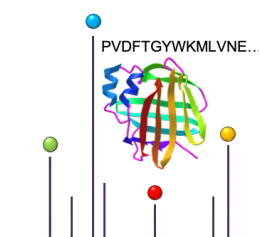


UCI Mass Spectrometry Facility

Peptide/Protein MS Form



Date:

User Information

Name:
UCInetID:
Group:
Location:
Email:

AA Sequence:

Sample Information

Name:
Sequence ID/UniProt/GenBank:
MW:
Buffer/solvent:
Conc (mg/mL):
Estimated purity:

Modifications

Disulfides (oxidized):
Glycans present: PNGaseF treat:
Exp System: Bacterial: Yeast: Insect: Mammalian:
HisTag: Other tag:
Other PTMs:

Digestion

Trypsin (pH7): Pepsin (pH2):

Sample Prep

PAGE gel slice (Coomassie stained):
Co-IP: # Washes: Elution buffer:

Analysis Method

MALDI: SA: CHCA:
LC-MS: Phenyl-hexyl (5 min): C₄ (30 min): SEC:

Experimental Information: goals, data required, etc.