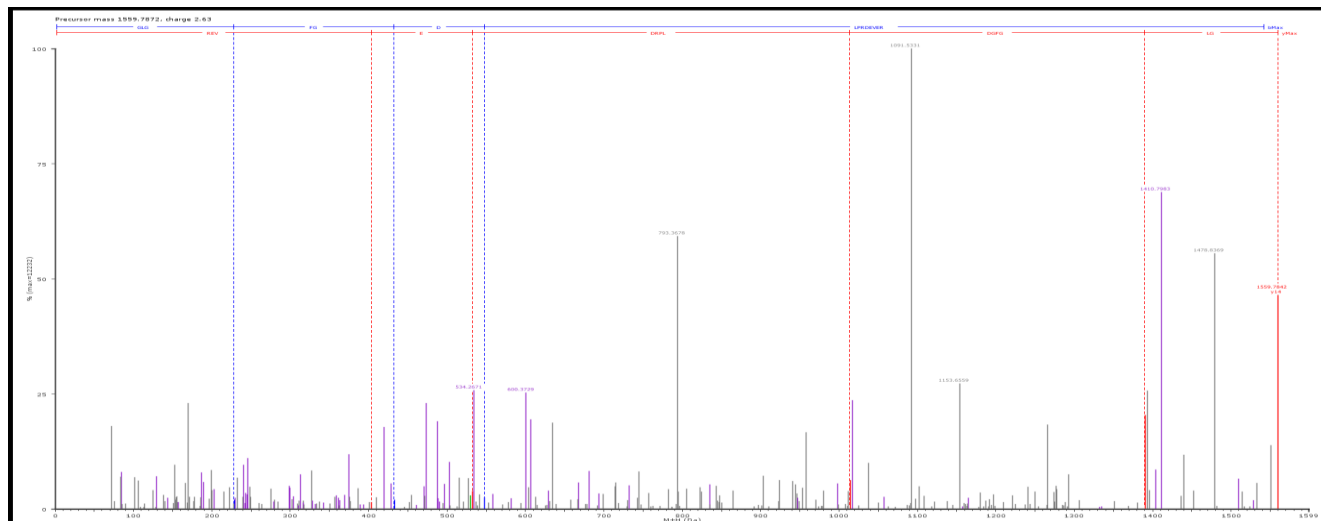
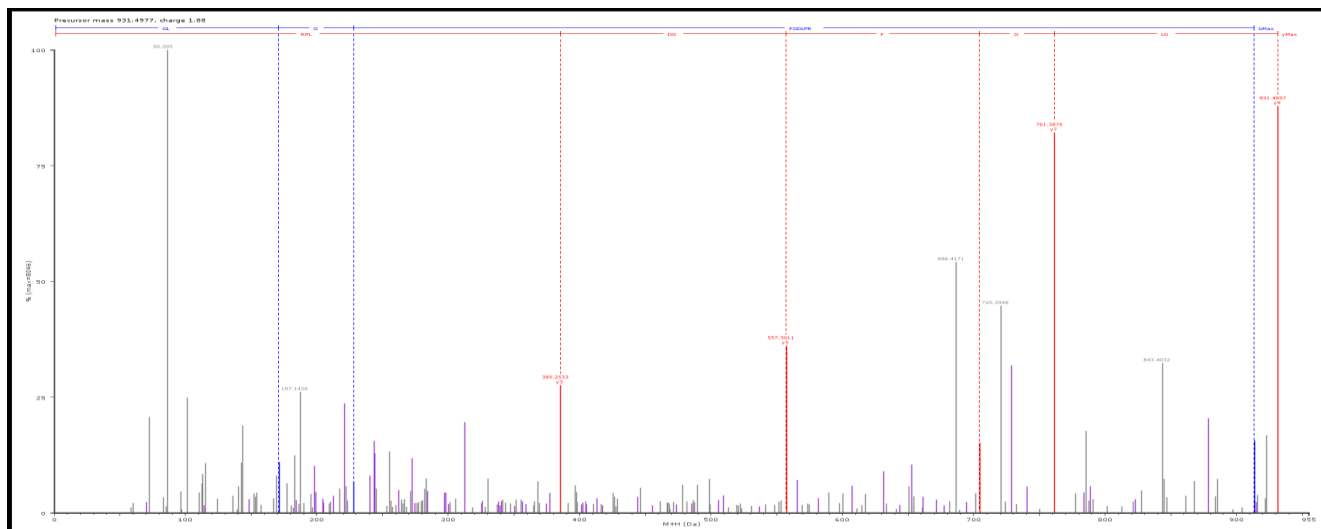
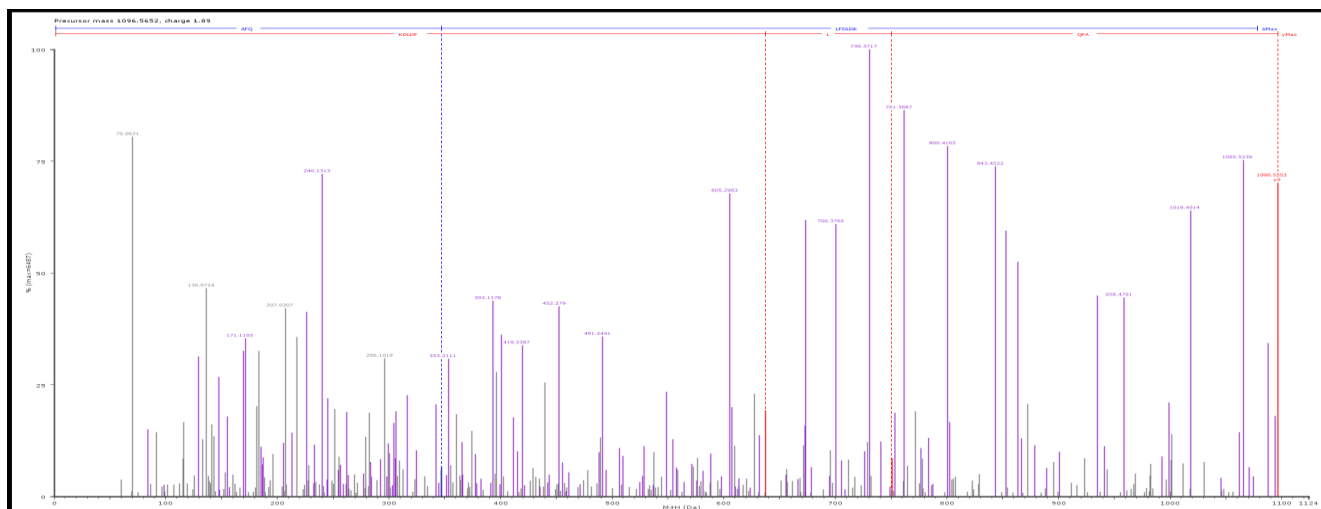
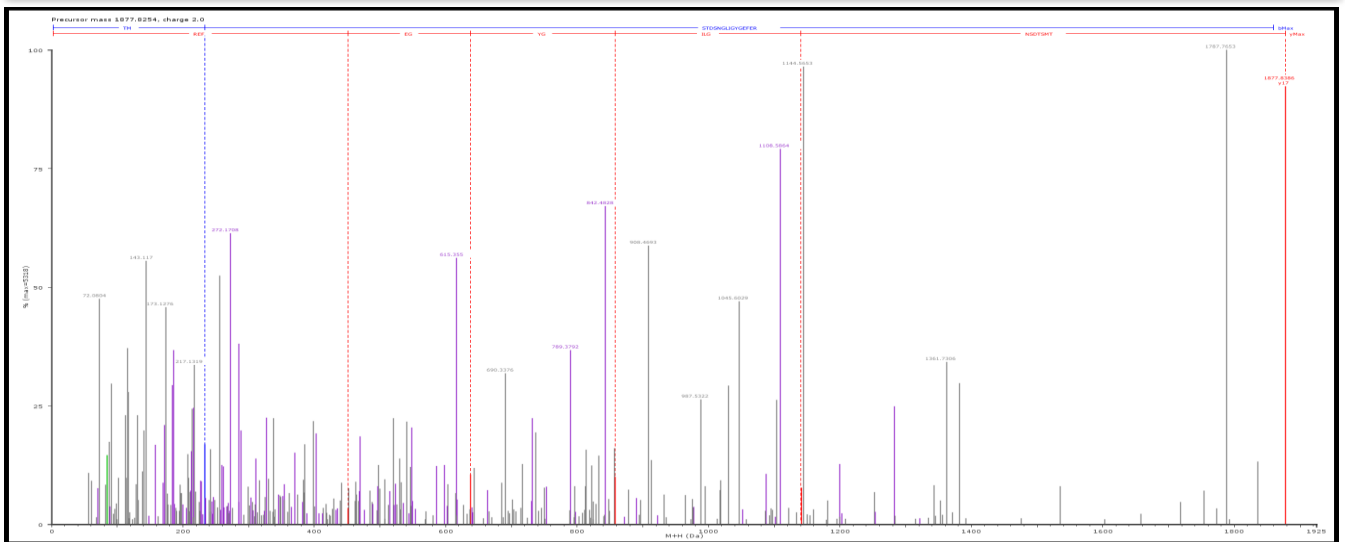
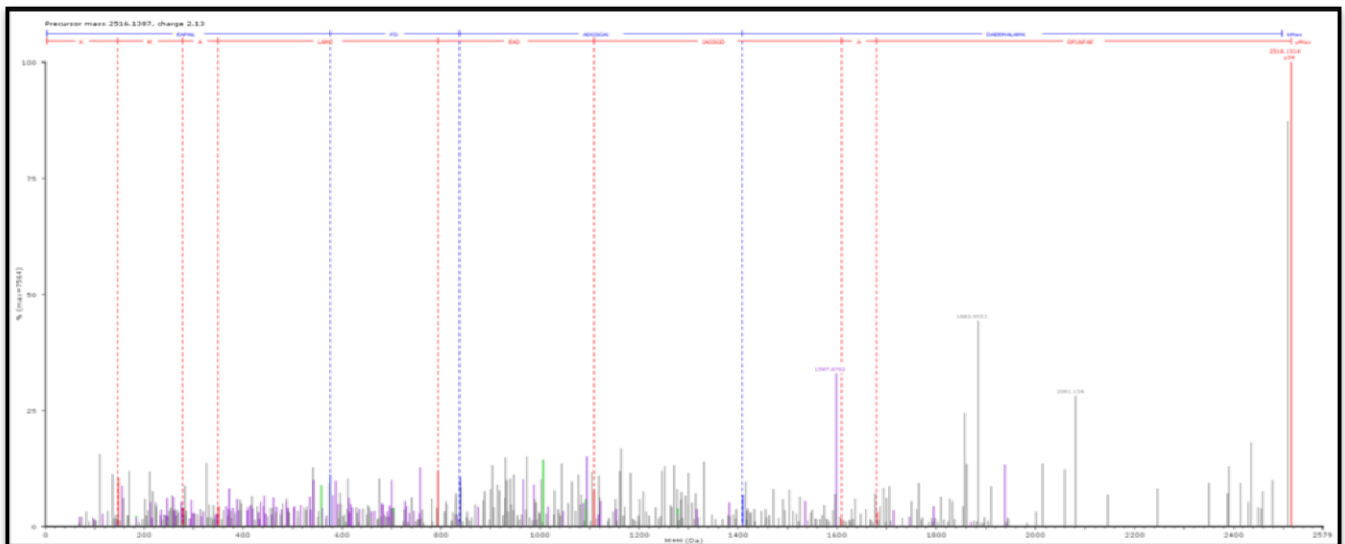
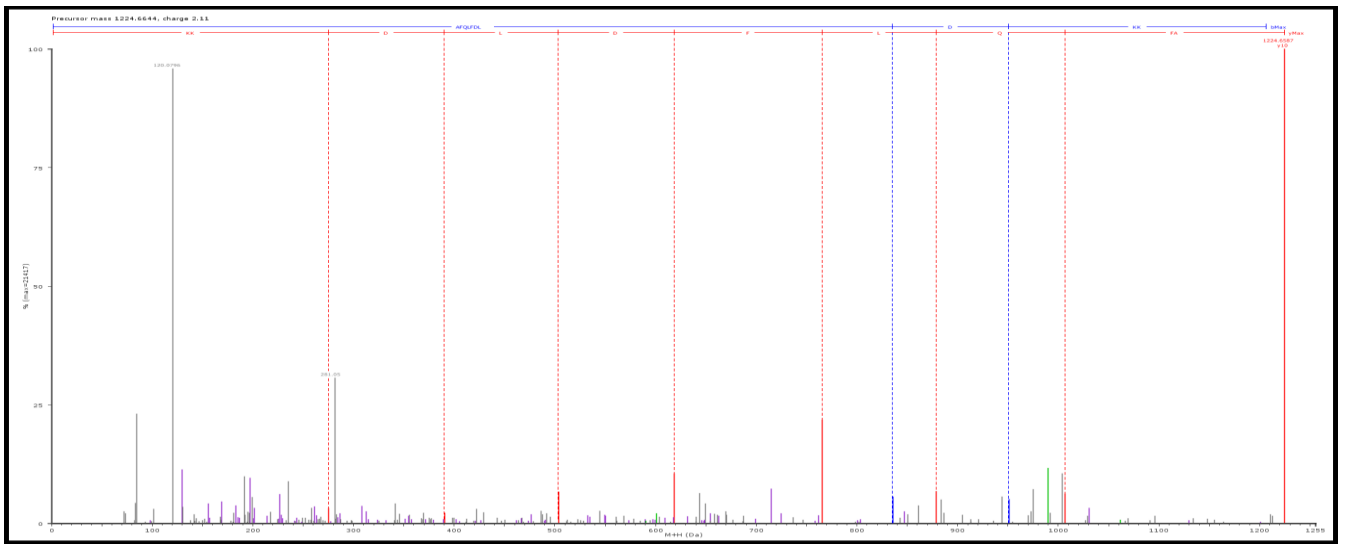
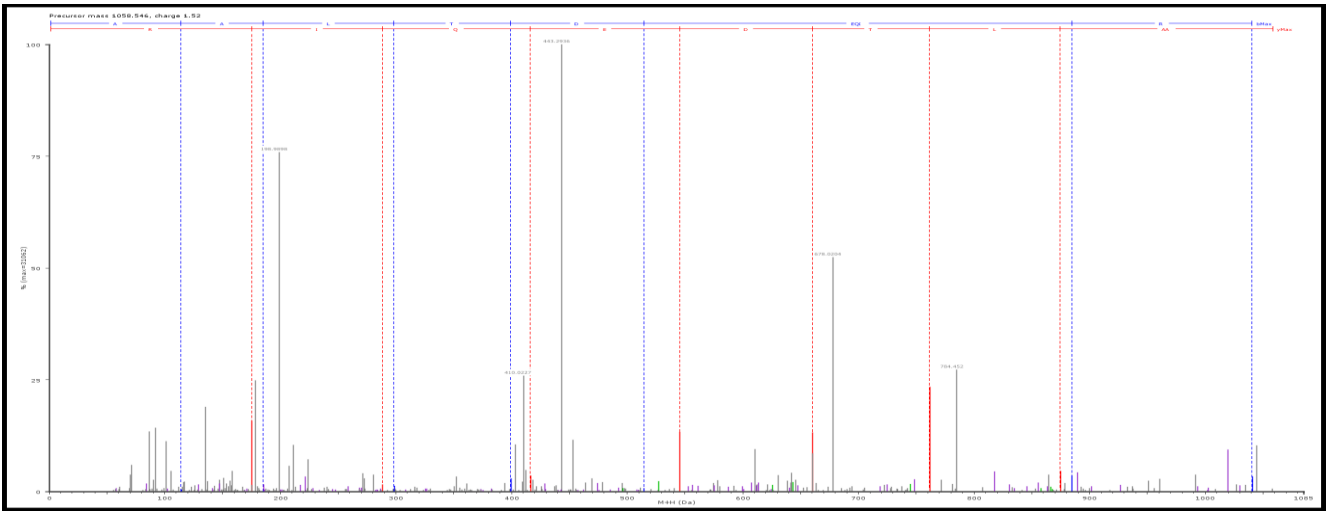


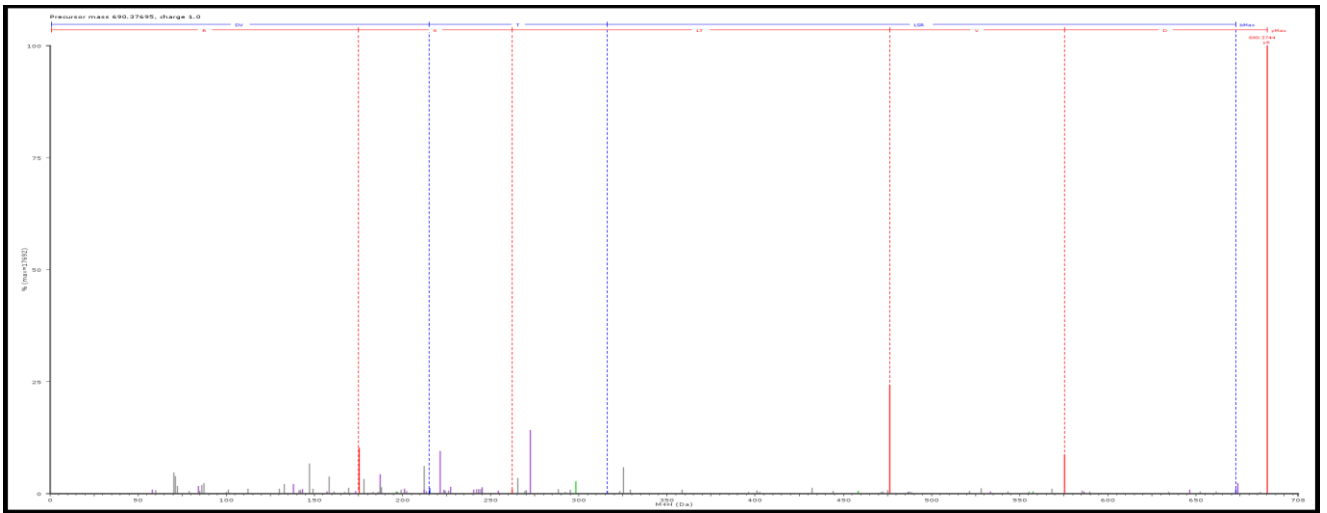
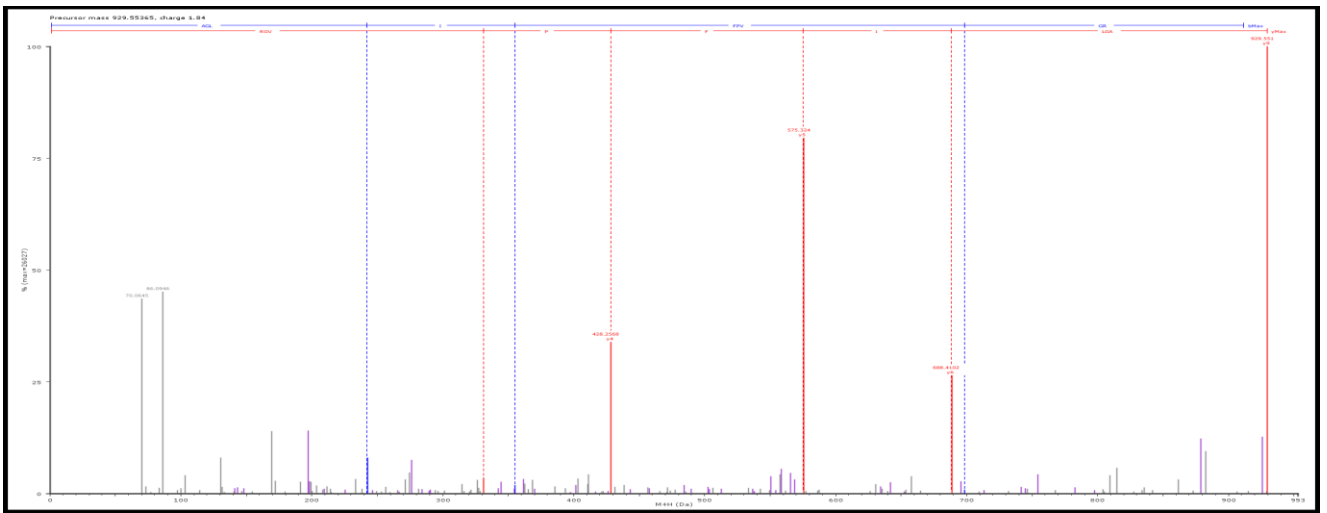
# S1 File. Examples of peptide mass spectra (MS/MS): Centrin and Histones

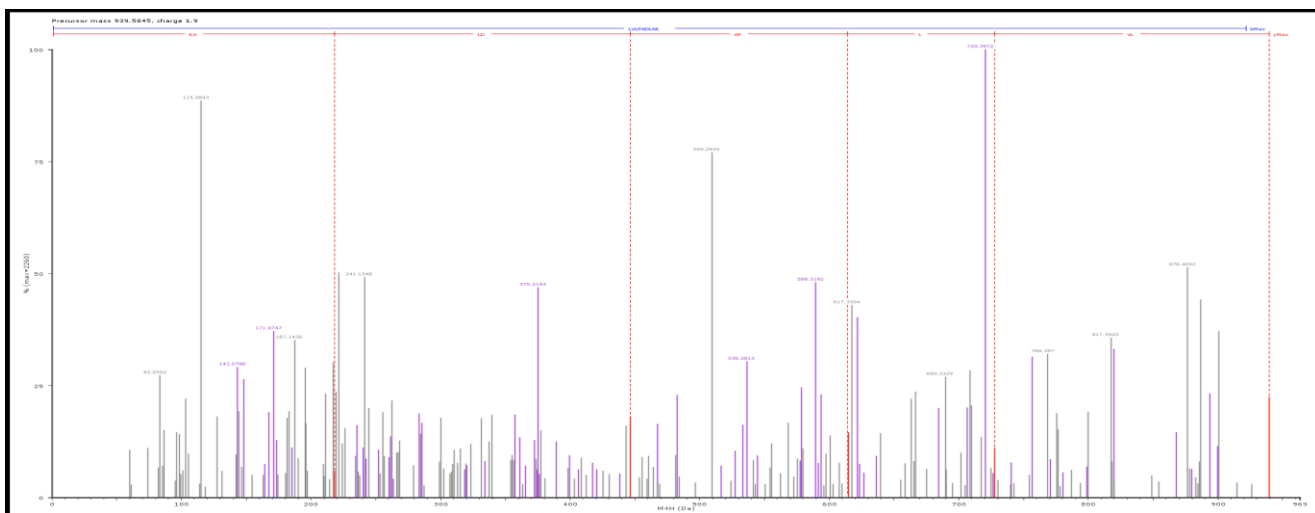
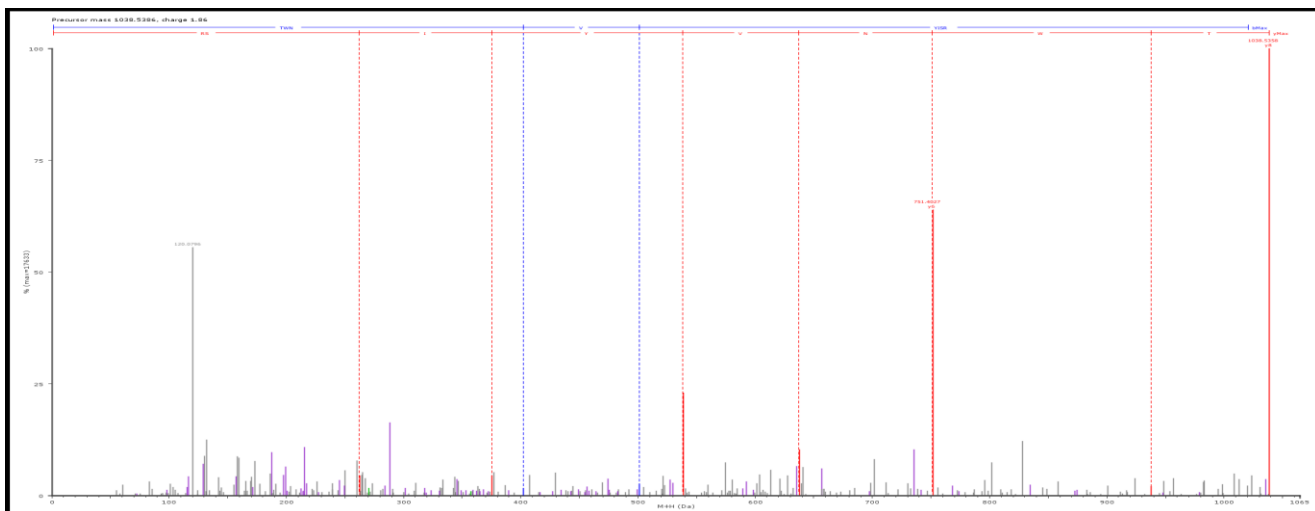
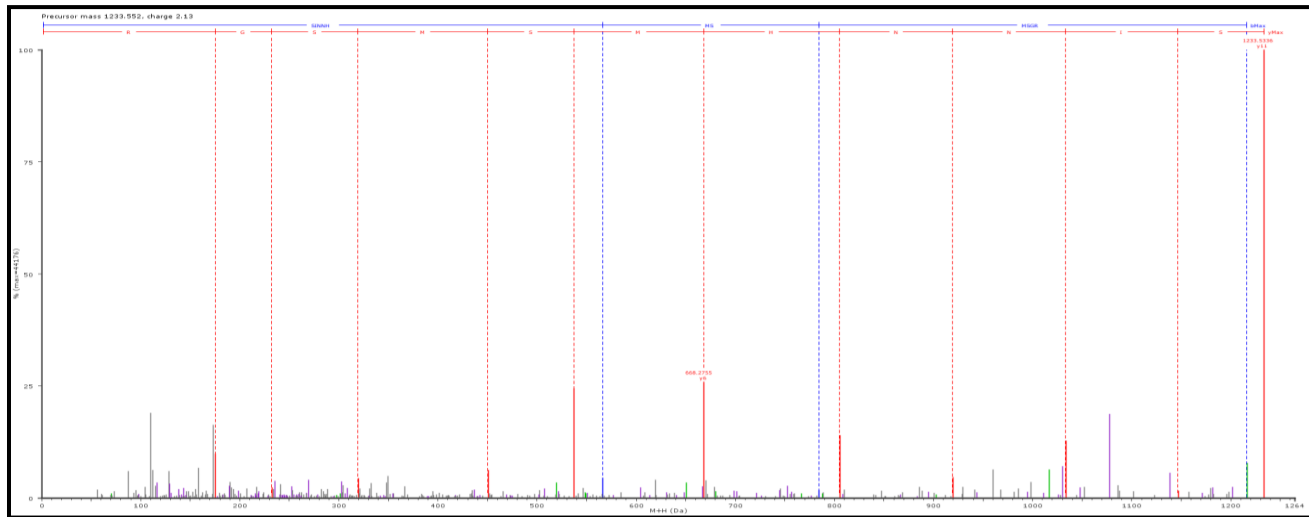
Protein entry	Protein description	# Peptides	Peptide sequence	Peptide score
Q4DQ49	Centrin_ putative Trypanosoma cruzi (strain CL Brener) Tc00.1047053508323.60 4 1	8	AFQLFDLDKK	7,6496
			GLGFGDLPR	7,3367
			GLGFGDLRDEVER	6,7946
			AFQLFDLDK	7,0327
			EAFNLFADGSGAIDAEEM ALAMK	6,7973
			TMSTDSNGLIGYGEFER	6,9267
			AALTDEQIR	7,6967
			TMSTDSNGLIGYGEFER	6,5735













Protein entry	Protein description	Peptide sequence	Peptide score
Q4DEL1	Histone H4 Trypanosoma cruzi (strain CL Brener) Tc00.1047053507941.150 3 1	TVTAVDVVNALR	9,1406
		TVTAVDVVNALRK	7,6205
		SFVEGVVR	8,4695
		ISGVIYDEVK	8,7713
		KTVTAVDVVNALR	7,1371

