

Sisl

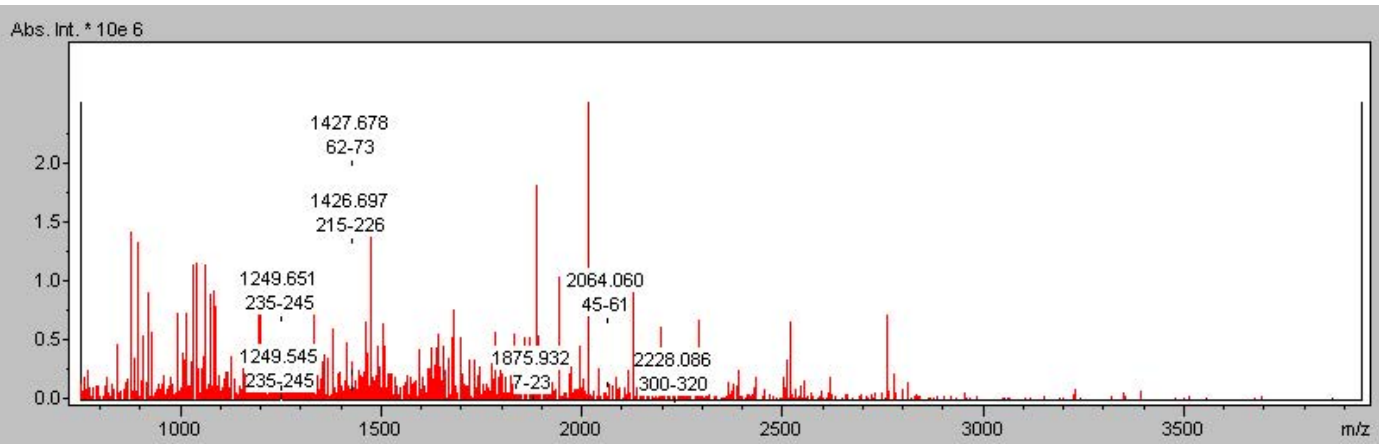
Sequence Name:Protein SIS1 OS=Saccharomyces cerevisiae (strain ATCC 204508 / S288c) GN=SIS1 PE=1 SV=1 SIS1_YEAST

MH+ (avg):1.008

MH+ (mono):1.008

Number of Peaks:1266

Tolerance (Da):0.900



Sequence data:

Intensity Coverage:0.3 % (472052 cnts)

Sequence Coverage MS/MS:25.6%

Sequence Coverage MS:25.6%

pl (isoelectric point):9.4

10	20	30	40	50	60	70	80	90	100
MVKETKLYDL	LGVSPSANEQ	ELKKGYRKAA	LKYHPDKPTG	DTEKFKEISE	AFEILNDPQK	REIYDQYGLE	AARSGGPSFG	PGGPGGAGGA	GGFPGGAGGF
110	120	130	140	150	160	170	180	190	200
SGGHAFSNE	AFNIFSQFFG	GSSPFGGADD	SGFSFSSYPS	GGGAGMGGMP	GGMGGMHGGH	GGMPGGFRSA	SSSPTYPEEE	TVQVNLVSL	EDLFVGKKKS
210	220	230	240	250	260	270	280	290	300
FKIGRKGPHG	ASEKTQIDIQ	LKPGWKAGTK	ITYKNQGDYN	PQTGRRRTLQ	FVIQEKSHPN	FKRDGDDLIY	TLPLSFKESL	LGFSKTIQTI	DGRTLPLSRV
310	320	330	340	350	360				
QPVQPSQTST	YPGQGMPTPK	NPSQRGNLIV	KYKVDYPISL	NDAQKRAIDE	NF				