

SUPPORTING INFORMATION

Fragment-Based Screening of a Natural Product Library against 62 Potential Malaria Drug Targets Employing Native Mass Spectrometry

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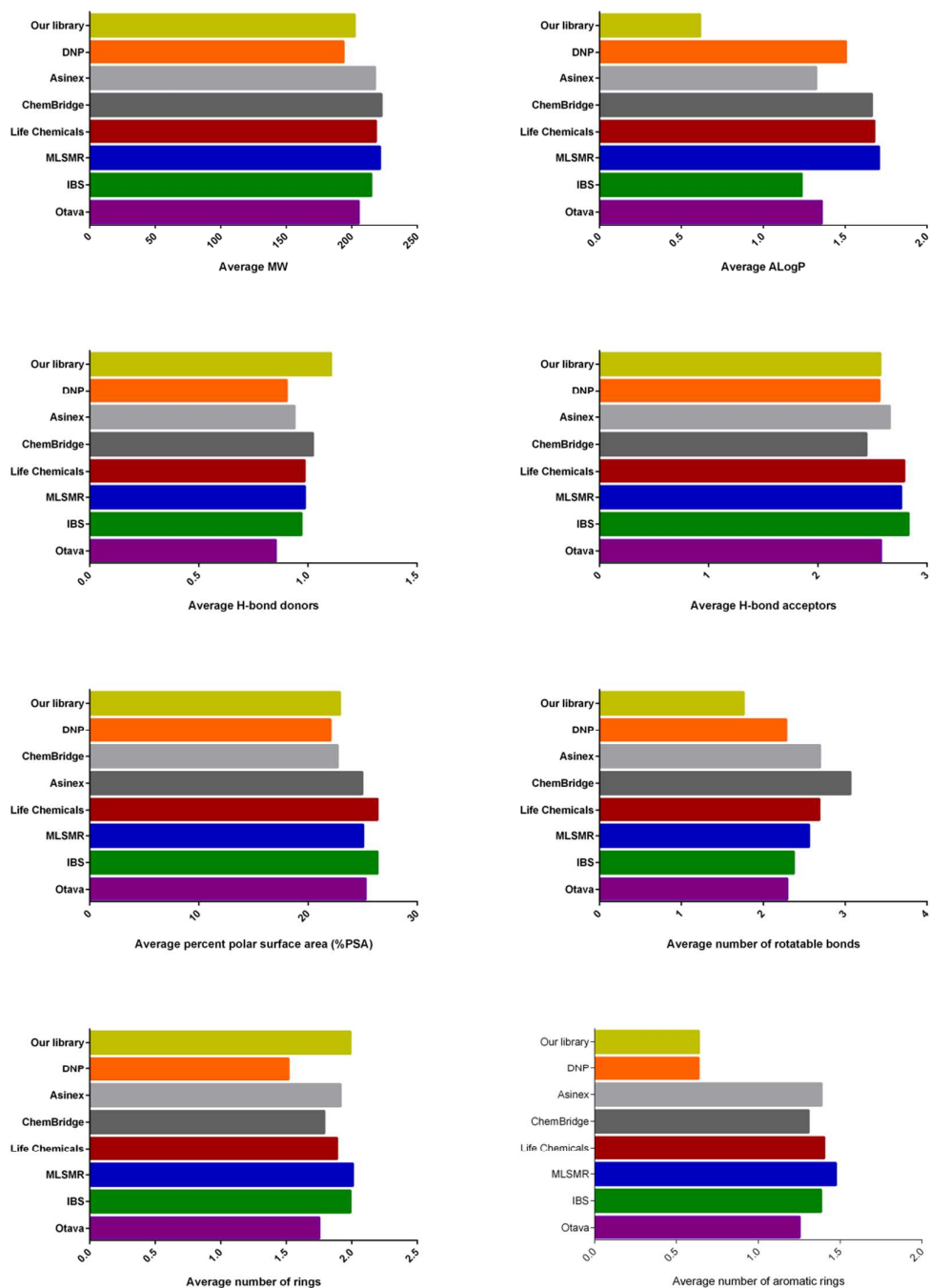


Figure S1. Mean physicochemical properties of different fragment libraries. Dictionary of Natural Products (DNP) and our fragment libraries are natural product-based fragment libraries. Asinex, ChemBridge, InterBioscreen (IBS), Life Chemicals and Otava fragment libraries are from commercially available libraries. Molecular Libraries Small Molecule Repository (MLSMR) fragment library is from the US national screening collection where most of the molecules are commercially sourced.

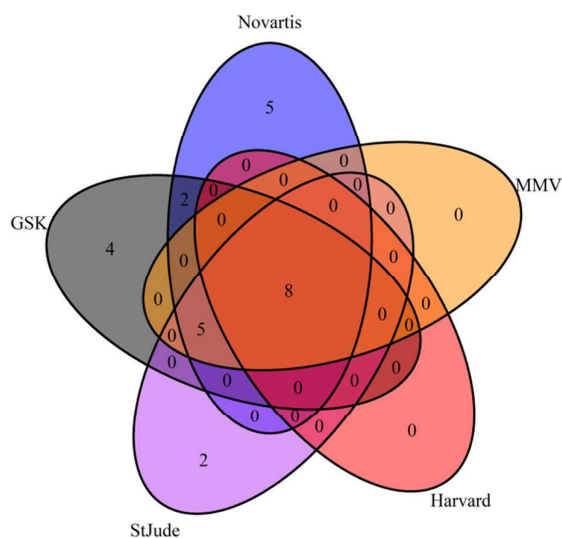


Figure S2. Venn diagram depicting the number of fragment hit scaffolds present within the molecular structures of GSK, Harvard Medical School (Harvard), St. Jude’s Children’s Research Hospital (StJude), MMV and Novartis library members. Diagram produced with R software (<http://www.R-project.org/>)¹, package “VennDiagram” (<http://CRAN.R-project.org/package=VennDiagram>)²

Table S1. List of forty-nine proteins studied for cloning and expression. Thirteen out of the forty-nine proteins were successfully cloned, expressed and purified.

Protein name/function	PlasmoDB ID	Purified
3-oxoacyl-(acyl-carrier-protein) reductase (FabG)	PF3D7_0922900	No
6-pyruvoyltetrahydropterin synthase (6PTPS)	PF3D7_0628000	Yes
Actin-related protein (ARP)	PF3D7_0103800	No
Adenylate cyclase-associated protein, putative (C-terminal actin binding domain)	PF3D7_0105300	No
ADP-ribosylation factor	PF3D7_1316200	No
Alpha-tubulin 2	PF3D7_0422300	No
ARK1	PF3D7_0605300	No
Beta-hydroxyacyl-ACP dehydratase precursor (FabZ)	PF3D7_1323000	No
Beta-ketoacyl-acyl carrier protein synthase III precursor (FabH/FabB/KASIII)	PF3D7_0211400	No
cAMP-dependent protein kinase (PKA)	PF3D7_0934800	No
Casein kinase 2	PF3D7_1108400	No
CDK-related protein kinase 6 (PfPK6)	PF3D7_1337100	Yes
CDP-DAG synthase (CDS)	PF3D7_1409900	No
CelTOS (cell traversal protein for ookinetes and sporozoites)	PF3D7_1216600	Yes
Chitinase	PF3D7_1252200	No
Conserved Plasmodium protein, unknown function	PF3D7_0818600	No
Conserved Plasmodium protein, unknown function (CITH; CAR-1/Trailer Hitch Homolog)	PF3D7_1474900	No
CSP (circumsporozoite surface protein)	PF3D7_0304600	Yes
Dihydrofolate reductase (DHFR)	PF3D7_0417200	Yes
Dihydroorotase	PF3D7_1472900	No
DOZI (development of zygote inhibited)	PF3D7_0320800	No
dUTPase	PF3D7_1127100	Yes
ETRAMP10.3	PF3D7_1016900	No
Fructose-bisphosphate aldolase	PF3D7_1444800	No
Glyceraldehyde-3-phosphate dehydrogenase	PF3D7_1462800	Yes
HMGB2 (high mobility group protein 2) -- transcription factor	PF3D7_0817900	Yes
HU homolog	PF3D7_0904700	Yes
Hypothetical protein	PF3D7_0313200	No
Hypothetical secreted protein	PF3D7_1360000	No
Hypoxanthine phosphoribosyltransferase	PF3D7_1012400	Yes
MDV-1/Peg3	PF3D7_1216500	Yes
Membrane-associated histidine-rich protein-2 (MAHRP2)	PF3D7_1353200	No
Methionine tRNA synthetase	PF3D7_1442300	No

Nek2 (NIMA-related kinase 2)	PF3D7_0525900	No
Nek4 (NIMA-related kinase 4)	PF3D7_0719200	No
Nucleosome assembly protein NapS	PF3D7_0919000	No
Orotate phosphoribosyltransferase	PF3D7_0512700	No
pbs36 homologue (P36)	PF3D7_0404400	No
PK7 (Serine/threonine protein kinase)	PF3D7_0213400	No
O-sialoglycoprotein endopeptidase	PF3D7_0708300	No
Protein kinase	PF3D7_1316000	No
Rhomboid protease ROM1	PF3D7_1114100	No
Secreted ookinete protein, putative (PSOP13, putative secreted ookinete protein 13)	PF3D7_0518800	No
Serine hydroxymethyltransferase	PF3D7_1235600	No
Sexual stage-specific precursor (Pfs16)	PF3D7_0406200	Yes
Signal Peptide Peptidase (SPP)	PF3D7_1457000	No
SPECT1 (sporozoite microneme protein essential for cell traversal)	PF3D7_1342500	No
Spermidine synthase	PF3D7_1129000	Yes
Thymidylate synthase	PF3D7_0417200	Yes*

*Same sequence with dihydrofolate reductase (DHFR).

Table S2. List of seventy-nine proteins studied by native mass spectrometry. Sixty-two out of the seventy-nine proteins were visible at native MS.

No	Protein Name	ID	EC number	MW
1	Thioredoxin peroxidase 2 (Trx-Px2)	PF3D7_1215000	1.11.1.15	18,526
2	Thioredoxin	PF3D7_1330000	1.8.1.8	24,275
3	NADP-specific glutamate dehydrogenase (GDH1)	PF3D7_1416500	1.4.1.4	50,240
4	Glycerol-3-phosphate dehydrogenase	PF3D7_1216200	1.1.1.8	42,133
5	Superoxide dismutase (SOD)	PKH_142350	1.15.1.1	24,976
6	Branched-chain alpha keto-acid dehydrogenase	PF3D7_1312600	1.2.4.4	55,125
7	Thioredoxin 1 (TRX1)	PF3D7_1457200	1.8.1.8	12,607
8	Glyceraldehyde-3-phosphate dehydrogenase	PF3D7_1462800	1.2.1.12	37,530
9	N-myristoyltransferase (NMT)	PF3D7_1412800	2.3.1.97	49,935
10	Ribosomal RNA methyltransferase	PF3D7_1309600	2.1.1.56	25,539
11	Phosphomethylpyrimidine kinase (PMPK)	PF3D7_0520500	2.7.4.7	35,861
12	Guanylate kinase	PVX_099895	2.7.4.8	23,544
13	Spermidine synthase	PF3D7_1129000	2.5.1.16	34,421
14	Uridine phosphorylase (purine nucleoside phosphorylase)	PF3D7_0513300	2.4.2.1	19,554
15	CDK-related protein kinase 6 (PfPK6)	PF3D7_1337100	2.7.11.22	37,995
16	Phosphoglycerate kinase (PGK)	PF3D7_0922500	2.7.2.3	46,325
17	Aspartate carbamoyltransferase	PVX_083135	2.1.3.2	47,102
18	RAB18 GTPase	PF3D7_0807300	3.6.5.2	21,757
19	Adenosine deaminase (ADA)	PVX_111245	3.5.4.4	42,816
20	dUTPase	PF3D7_1127100	3.6.1.23	20,340
21	6-phosphogluconolactonase	PF3D7_1453800	3.1.1.31	41,005
22	RAB1a GTPase	PF3D7_1320600	3.6.5.2	23,241
23	Ubiquitin carboxyl-terminal hydrolase isozyme L3(UCHL3)	PF3D7_1460400	3.4.19.12	27,790
24	Methionine aminopeptidase 1b (METAP1b, MAP1)	PF3D7_1015300	3.4.11.18	37,932
25	S-adenosylhomocysteine hydrolase (SAHH)	PF3D7_0520900	3.3.1.1	54,733
26	ATP-dependent DNA helicase Q1 (RECQ1)	PF3D7_0918600	3.6.4.12	12,576
27	Adenosine deaminase (ADA)	PF3D7_1029600	3.5.4.4	44,666
28	ATP-dependent Clp protease proteolytic subunit (ClpP)	PF3D7_0307400	3.4.21.92	23,552
29	Glyoxalase I	PF3D7_1113700	4.4.1.5	23,775
30	Adenylosuccinate lyase	PVX_003765	4.3.2.2	47,094
31	Deoxyribose-phosphate aldolase	PY02252	4.1.2.4	31,101
32	Orotidine 5' monophosphate decarboxylase (OMPDC)	PF3D7_1023200	4.1.1.23	38,760
33	6-pyruvoyltetrahydropterin synthase	PF3D7_0628000	4.2.3.12	20,953
34	2C-methyl-D erythritol 2,4-cyclodiphosphate synthase (IspF)	PVX_003920	4.6.1.12	22,388
35	Phosphoglycerate mutase (PGM)	PF3D7_1120100	5.4.2.1	29,662
36	Cyclophilin	PF3D7_0528700	5.2.1.8	24,263
37	Ribose 5-phosphate epimerase	PF3D7_0514600	5.3.1.6	27,072
38	D-ribulose- 5-phosphate 3-epimerase	PF3D7_1219900	5.1.3.1	25,780
39	Prolyl tRNA ligase (aPRS)	PF3D7_0925300	6.1.1.15	58,213
40	Ubiquitin conjugating enzyme E2	PF3D7_0305700	6.3.2.19	17,886
41	Adenylosuccinate synthetase (AdSS)	PF3D7_1354500	6.3.4.4	51,461
42	Tryptophan tRNA ligase	PF3D7_1336900	6.1.1.2	47,746
43	Ubiquitin conjugating enzyme E2	PF3D7_1203900	6.3.2.19	17,431
44	Ubiquitin conjugating enzyme E2 N	PF3D7_0527100	6.3.2.19	19,176
45	Ubiquitin conjugating enzyme	PF3D7_0812600	6.3.2.19	19,554
46	Ubiquitin conjugating enzyme	PF3D7_1033900	6.3.2.19	19,288
47	Ubiquitin conjugating enzyme	PF3D7_1356300	6.3.2.19	23,776
48	MTIP	PF3D7_1246400	NA	16,482
49	Bacterial histone-like protein (HU homolog)	PF3D7_0904700	NA	17,403
50	Phosphatidylethanolamine-binding protein	PVX_123630	NA	45,604
51	Dynein light chain 1	PF3D7_1213600	NA	11,544
52	CSP (circumsporozoite surface protein)	PF3D7_0304600	NA	26,787
53	Conserved protein, unknown function	PF3D7_1351100	NA	19,603
54	Vacuolar protein-sorting protein (VPS2)	PF3D7_0816200	NA	25,721
55	Eukaryotic translation initiation factor 5A (eIF5a)	PF3D7_1204300	NA	18,529
56	6-pyruvoyltetrahydropterin synthase	PVX_114505	4.2.3.12	21,975
57	Sexual stage-specific protein precursor (Pfs16)-PF0310w	PF3D7_0406200	NA	9,200
58	Cell traversal protein for ookinetes and sporozoites (CelTOS)	PF3D7_1216600	NA	17,683
59	Hypoxanthine-guanine phosphoribosyltransferase (HGPR1)	PF3D7_1012400	2.4.2.8	26,660
60	High mobility group protein B2 (HMGB2)	PF3D7_0817900	NA	10,107
61	Orotate phosphoribosyltransferase, putative	PKNH_1020400	2.4.2.10	34,802
62	Serine/threonine-protein kinase NEK4, putative	PVX_096360	2.7.11.1	37,077
63	Eukaryotic translation initiation factor 2b (eIF2b), alpha subunit	PF3D7_0828500	NA	NA
64	Thioredoxin reductase	PF3D7_0923800	1.8.1.9	NA
65	Ribonucleotide reductase, small subunit	PVX_086155	1.17.4.1	NA
66	Proline carboxylate reductase	PF3D7_1357900	1.5.1.2	NA
67	Dihydrofolate reductase (DHFR)	PF3D7_0417200	1.5.1.3	NA

68	Superoxide dismutase [Fe] (SOD)	PBANKA_1422700	1.15.1.1	NA
69	S-adenosylmethionine synthetase (SAMS)	PF3D7_0922200	2.5.1.6	NA
70	Ornithine aminotransferase	PY00104	2.6.1.13	NA
71	Nucleoside diphosphate kinase B (NDK)	PF3D7_1366500	2.7.4.6	NA
72	Protein-L-isoaspartate O-methyltransferase	PF3D7_1432700	2.1.1.77	NA
73	Choline Kinase	PF3D7_1401800	2.7.1.32	NA
74	Geranylgeranyl pyrophosphate synthase (GGPPS)	PVX_092040	2.5.1.29	NA
75	Nucleosome assembly protein 1	PKNH_1303500	NA	NA
76	Conserved protein, unknown function	PF3D7_0618200	NA	NA
77	Male development gene 1 (MDV1/MDV-1/Peg3)	PF3D7_1216500	NA	NA
78	Male development gene 1 (MDV1/MDV-1/Peg3)	PVX_123505	NA	NA
79	Protein phosphatase 1, regulatory (inhibitor) subunit	PVX_111140	NA	NA

Table S3. List of 96 detected hits and their anti-malarial activity.

Compound	Name	% Inh at 100 μ M	IC ₅₀ (μ M) <i>P. falciparum</i> asexual (3D7)	IC ₅₀ (μ M) <i>P. falciparum</i>	Sources
1	Indole-3-butyric acid				Widespread in plants ³
2	6-Acetamido-6-deoxy-castanospermine	44			<i>Castanospermum austral</i> , <i>Alexa leiopetala</i> (castanospermine) ⁴
3	Patulin	98	91.6		<i>Aspergillus clavatus</i> , <i>Aspergillus terreus</i> , <i>Aspergillus varians</i> , <i>Penicillium patulum</i> , <i>Penicillium griseofulvum</i> , <i>Byssoschlamys nivea</i> ⁵
4	Santonin	9	-		<i>Artemisia</i> spp. ^{4,6}
5	Piceatannol	92	>100	120	<i>Vouacapoua macropetala</i> , <i>Pericopsis angolensis</i> <i>Vaccinium</i> sp. <i>Picea abies</i> ^{7,8}
6	Resveratrol	60	>100	60	<i>Veratrum grandiflorum</i> , <i>Pinus sibirica</i> , <i>Vitis vinifera</i> , <i>Arachis hypogaea</i> , <i>Polygonum</i> spp., <i>Nothofagus</i> spp., <i>Cudrania javanensis</i> , <i>Eucalyptus</i> spp. ⁹
7	Acivicin	94	5.4		<i>Streptomyces sviveus</i> ¹⁰
8	Citrinin	63	>100		<i>Penicillium citrinum</i> , <i>Guanomyces polythrix</i> , <i>Aspergillus terreus</i> , <i>Monascus</i> spp. ¹¹
9	Cantharidin	61	>100	9	<i>Epicauta</i> spp., <i>Lytta vesicatoria</i> ¹²
10	trans-parthenolide	96	>100		<i>Chrysanthemum parthenium</i> , <i>Michelia champaca</i> , <i>Michelia nilagirica</i> , <i>Michelia compressa</i> ¹³⁻¹⁵
11	(-)-Huperazine				<i>Lycopodium serratum</i> (<i>Huperzia serrata</i>), <i>Lycopodium selago</i> , <i>Lycopodium saururus</i> , <i>Lycopodium erythraeum</i> , <i>Lycopodium gnidioides</i> , <i>Paecilomyces tenuis</i> ^{16,17}
12	2,5-Dihydro-4-hydroxy-5-oxo-3-phenyl-2-furanpropanoic acid	9	-		<i>Acremonium</i> sp., <i>Chaetomella raphigera</i> ¹⁸
13	Phaeofuran B	19	-		<i>Phaeoacremonium</i> sp. ¹⁹
14	Curcumenol	66	>100		<i>Curcuma zedoaria</i> , <i>Curcuma longa</i> ²⁰
15	2 – Hydroxylantolactone	98	44.3		<i>Francoeuria crispa</i> , <i>Inula helenium</i> , <i>Inula royleana</i> ²¹
16	(+)-reynosin	98	32.3		<i>Magnolia grandiflora</i> , <i>Tanacetum praeterium</i> , <i>Saussurea lappa</i> , <i>Ambrosia confertiflora</i> ²²
17	Asperilin	98	23.2		<i>Iva texensis</i> , <i>Telekia speciosa</i> , <i>Iva asperifolia</i> , ²³
18	Graveolide	84	15.6		<i>Inula graveolens</i> ²¹
19	Xanthinosin (or tomentosin)	65	62.8		<i>Dittrichia</i> spp., <i>Inula</i> spp., <i>Pulicaria</i> spp., <i>Xanthium</i> spp. ²⁴
20	Pinnatifidin	92	66.5		<i>Helenium pinnatifidum</i> ²⁵
21	Homononactic acid	30	-		<i>Streptomyces</i> spp. ²⁶

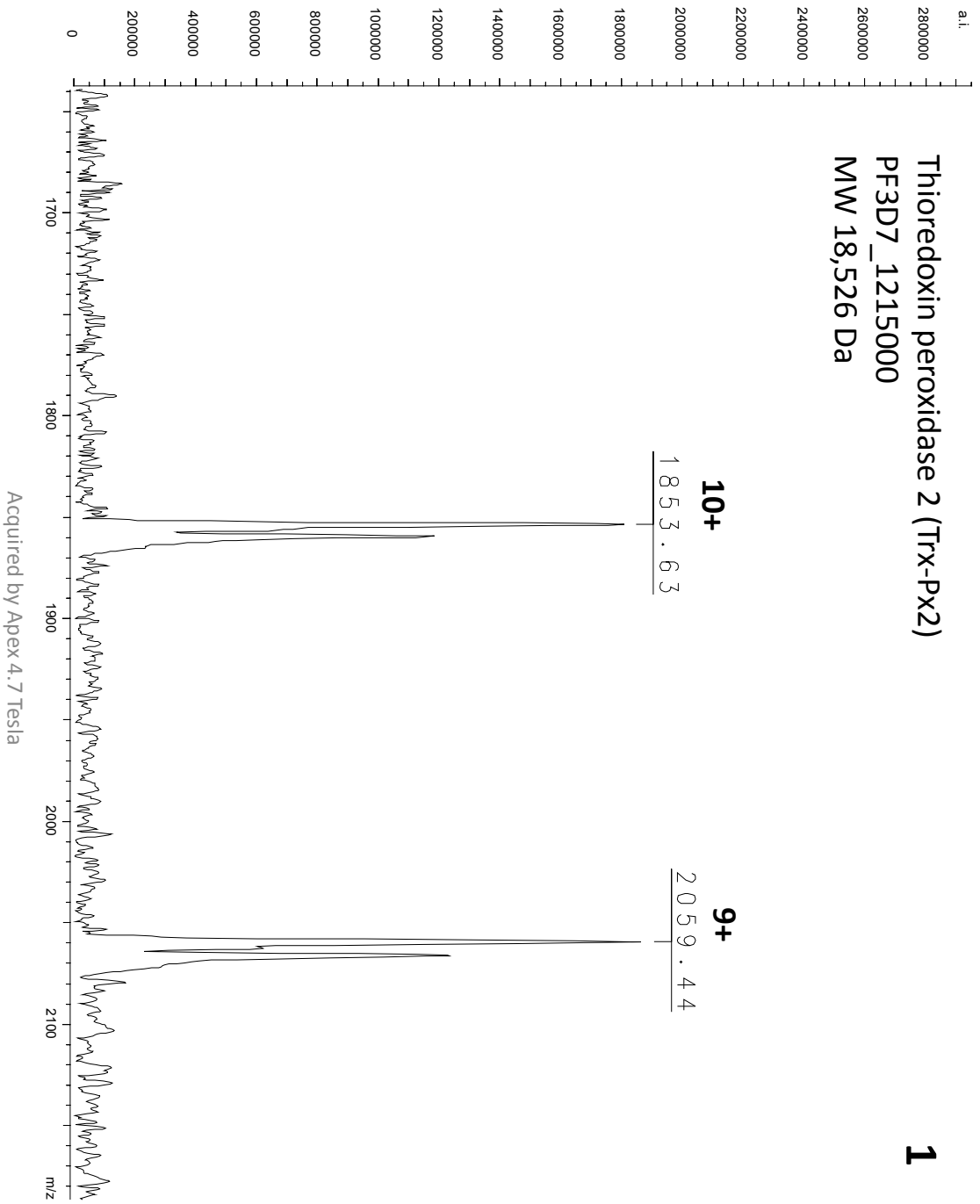
22	Curdionolide B	42	-	<i>Curcuma wenyujin</i> ²⁷
23	Neolitacumone B	27	-	<i>Neolitsea acuminatissima</i> ²⁸
24	β -Santonin	37	-	<i>Artemisia</i> spp. ²⁹
25	Acremine I	43	-	<i>Acremonium byssoides</i> strain A20 ³⁰
26	Asteriscunolide A	68	34.2	<i>Asteriscus aquaticus</i> , <i>Asteriscus graveolens</i> , <i>Asteriscus vogelii</i> ³¹
27	Arteannuin B	98	6.6	<i>Artemisia annua</i> ³²
28	Xerantholide	96	38.3	<i>Anthemis austriaca</i> , <i>Decachaeta thieleana</i> , <i>Xeranthemum cylindraceum</i> ³³
29	1-Epi-inuviscolide	100	79.4	<i>Dittrichia graveolens</i> ³⁴
30	Camphanic acid	23	-	<i>Fagaropsis glabra</i> ³⁵
31	Fraxinellonone	31	-	<i>Fagaropsis glabra</i> ³⁵
32	Dihydrokawain-5-ol	14	-	<i>Piper methysticum</i> ³⁶
33	Costunolide	35	-	<i>Saussurea lappa</i> , <i>Artemisia balchanorum</i> , <i>Talauma ovata</i> , <i>Talauma mexicana</i> , <i>Liriodendron tulipifera</i> , <i>Frullania tamarischii</i> , <i>Frullania nisquallensis</i> , <i>Michelia</i> spp., <i>Magnolia grandiflora</i> , <i>Magnolia sieboldii</i> , <i>Gochnatia foliosa</i> , <i>Chrysanthemum boreale</i> , <i>Cosmos caudatus</i> , <i>Cosmos sulphureus</i> ³⁷⁻³⁹
34	Antibiotic 927A	23	-	<i>Streptomyces griseus</i> ⁴⁰
35	Methyl ester of 3-Chloro-1,5-dihydroxy-4-oxo-2-(1-propenyl)-2-cyclopentene-1-carboxylic acid	98	67.3	<i>Cryptosporiopsis</i> spp. ⁴¹
36	Santamarine	96	61.3	<i>Magnolia grandiflora</i> L. ⁴²
37	Zaluzanin C	99	93.6	<i>Conocephalum conicum</i> , <i>Zaluzania</i> spp., <i>Zaluzania robinsonii</i> ⁴³
38	Derivative of Antibiotic EI 1941-2	20	-	Antibiotic EI 1941-2 was found in <i>Phomopsis</i> sp. (strain No. 7233) ⁴⁴
39	Alantolactone	92	92.7	<i>Inula helenium</i> , <i>Inula racemosa</i> , <i>Inula grandis</i> , <i>Inula magnifica</i> , <i>Saussurea lappa</i> , <i>Codonocephalum grande</i> , <i>Abutilon indicum</i> ⁴⁵
40	Isoalantolactone	50	39.8	<i>Inula helenium</i> , <i>Inula racemosa</i> , <i>Abutilon indicum</i> , <i>Telekia speciosa</i> ⁴⁶
41	5-Fluoro-3,4-dihydro-2,4-dioxo-1(2H)-pyrimidineacetic acid	27	-	<i>Phakellia fusca</i> ⁴⁷
42	Biotin	44	-	yeast, eggs, liver, various microorganisms, sweet corn seedlings, leaves of <i>Raphanus sativus</i> ⁴⁸
43	12-Methoxycarbonylcytisine			<i>Leontice watereri</i> , <i>Petteria ramentacea</i> ⁴⁹
44	Lepiotin C	25	-	<i>Macrolepiota neomastoidea</i> ⁵⁰
45	N-Benzoyl of 4-Piperidinone	33	-	<i>Dichilus strictus</i> , <i>Dichilus reflexus</i> , <i>Dichilus lebeckioides</i> , <i>Dichilus pilosus</i> , <i>Dichilus gracilis</i> ⁵¹

46	cis-parthenolide	76	20.5		<i>Chrysanthemum parthenium</i> , <i>Michelia champaca</i> , <i>Michelia nilagirica</i> , <i>Michelia compressa</i> , <i>Chrysanthemum parthenium</i> , <i>Ambrosia dymosa</i> , <i>Magnolia grandiflora</i> , <i>Artemisia</i> spp., <i>Tanacetum parthenianum</i> , <i>Paramichelia baillonii</i> , <i>Acanthopanax koreanum</i> ¹⁴
47	Isofischeric acid				<i>Neolitsea fischeri</i> ⁵²
48	Thymidine	34	-		<i>Phaseolus vulgaris</i> , <i>Acanthaster planci</i> , <i>Cryptochiton stelleri</i> , <i>Pavonaria finmarchica</i> ⁵³
49	1-Aminocyclopropanecarboxylic acid				<i>Ethylene</i> spp. ⁵⁴
50	7-amino deacetoxy cephalosporanic acid	40	-		<i>Acremonium chrysogenum</i> ⁵⁵
51	Fenchone	48	-		<i>Foeniculum vulgare</i> , <i>Blumea lacera</i> , <i>Prunella vulgaris</i> ^{56, 57}
52	Eseroline	100	20.2		Physostigmine was found in <i>Physostigma venenosum</i> , <i>Streptomyces</i> sp. AH-4, <i>Streptomyces pseudogriseolus</i> ⁵⁸
53	N-tert-butylloxycarbonyl derivative of homoproline	30	-		<i>Arnica montana</i> , <i>Arnica chamissonis</i> ssp. <i>foliosa</i> , <i>Arnica amplexicaulis</i> , <i>Arnica sachalinensis</i> , <i>Tussilago farfara</i> ⁵⁹
54	N-tert-butylloxycarbonyl derivative of 2-piperidinecarboxylic acid	22	-		<i>Amphiroa beauvoisii</i> , <i>Acacia rigidula</i> ⁶⁰
55	Aloperine	43	-		<i>Sophora alopecuroides</i> ⁶¹
56	Lycoperodine 1	49	-		<i>Aleurites fordii</i> , <i>Allium tuberosum</i> , <i>Asterias rollestoni</i> , <i>Lycopersicon esculentum</i> ⁶²
57	N,N-dipropylsuccinamic acid				
58	Derivative of matlystatins, 4-oxo-4-(piperidin-1-yl)butanoic acid	24	-		<i>Actinomadura atramentaria</i> ⁶³
59	Derivative of pyroglutamic acid, (S)-1-(2-amino-2-oxoethyl)-5-oxopyrrolidine-2-carboxamide	24	-		Ginseng (pyroglutamic acid) ⁶⁴
60	1,2,3,4-Tetrahydro-6,7-dihydroxy-3-isoquinolinecarboxylic acid	26	-		<i>Mucuna mutisiana</i> (Leguminosae), <i>Mucuna urens</i> , <i>Mucuna deeringiana</i> , <i>Mucuna andreana</i> , <i>Mucuna holtoni</i> , <i>Mucuna pruriens</i> , <i>Mucuna sloanei</i> , <i>Stizolobium hassjoo</i> ⁶⁵
61	Derivative of almazolone, (Z)-2-methyl-4-(4-methylbenzylidene)oxazol-5(4H)-one	33	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶

62	<i>O</i> -Methyl-L-tyrosine				<i>Cylindrocladium scoparium</i> ⁶⁷
63	<i>N,N</i> -diethyl-4-hydroxy- benzamide				Synthetic? ⁶⁸
64	Diethylamide of 3-hydroxybenzoic acid				Synthetic? ⁶⁹
65	Derivative of dysinosin A, 5-(<i>S</i>)-acetyl-9-(<i>S</i>)-hydroxy- (3 <i>R</i> , 7 <i>R</i>)-octahydro-1 <i>H</i> -Indole	31	-		Dysinosin A was found in <i>Citronia astra</i> ⁷⁰
66	<i>N,N</i> -diethylsuccinamic acid	40	-		<i>Candida tropicalis</i> , <i>Candida</i> sp., <i>Saccharomyces cerevisiae</i> , <i>Xanthomonas campestris</i> , <i>Pseudomonas putida</i> , <i>Xanthomonas campestris oryzae</i> , <i>Gerbera</i> sp. ^{71, 72}
67	Derivative of almazolone, 4-[(2-methyl-5-oxo-4(5 <i>H</i>)-oxazolylidene)methyl] - benzoic acid	41	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
68	Derivative of almazolone, 4-methyl-2-(<i>p</i> -hydroxyphenylmethylene)- 5(2 <i>H</i>)-Oxazolone	35	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
69	Derivative of almazolone, (<i>E</i>)-methyl 4-((2-methyl-5-oxooxazol-4(5 <i>H</i>)-ylidene)methyl)benzoate	22	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
70	Derivative of almazolone, (<i>E</i>)-4-(benzodioxol-5-ylmethylene)-2-methyloxazol-5(4 <i>H</i>)-one	15	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
71	Derivative of almazolone, (<i>E</i>)-4-(4-chlorobenzylidene)-2-methyloxazol-5(4 <i>H</i>)-one	29	-		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
72	Derivative of almazolone, (<i>E</i>)-4-(3,4-dimethoxybenzylidene)-2-methyloxazol-5(4 <i>H</i>)-one	41	>100		Almazolone was found in <i>Haraldiophyllum</i> sp. ⁶⁶
73	5-Methoxytryptophan methyl ester	65	>100		<i>Griffonia simplicifolia</i> , <i>Panaeolus sphinctrinus</i> ⁷³
74	Derivative of cichorine, <i>N</i> -(2-(3-	32	-		Cichorine was found in <i>Aspergillus silvaticus</i> , <i>Alternaria cichorii</i> ⁷⁴

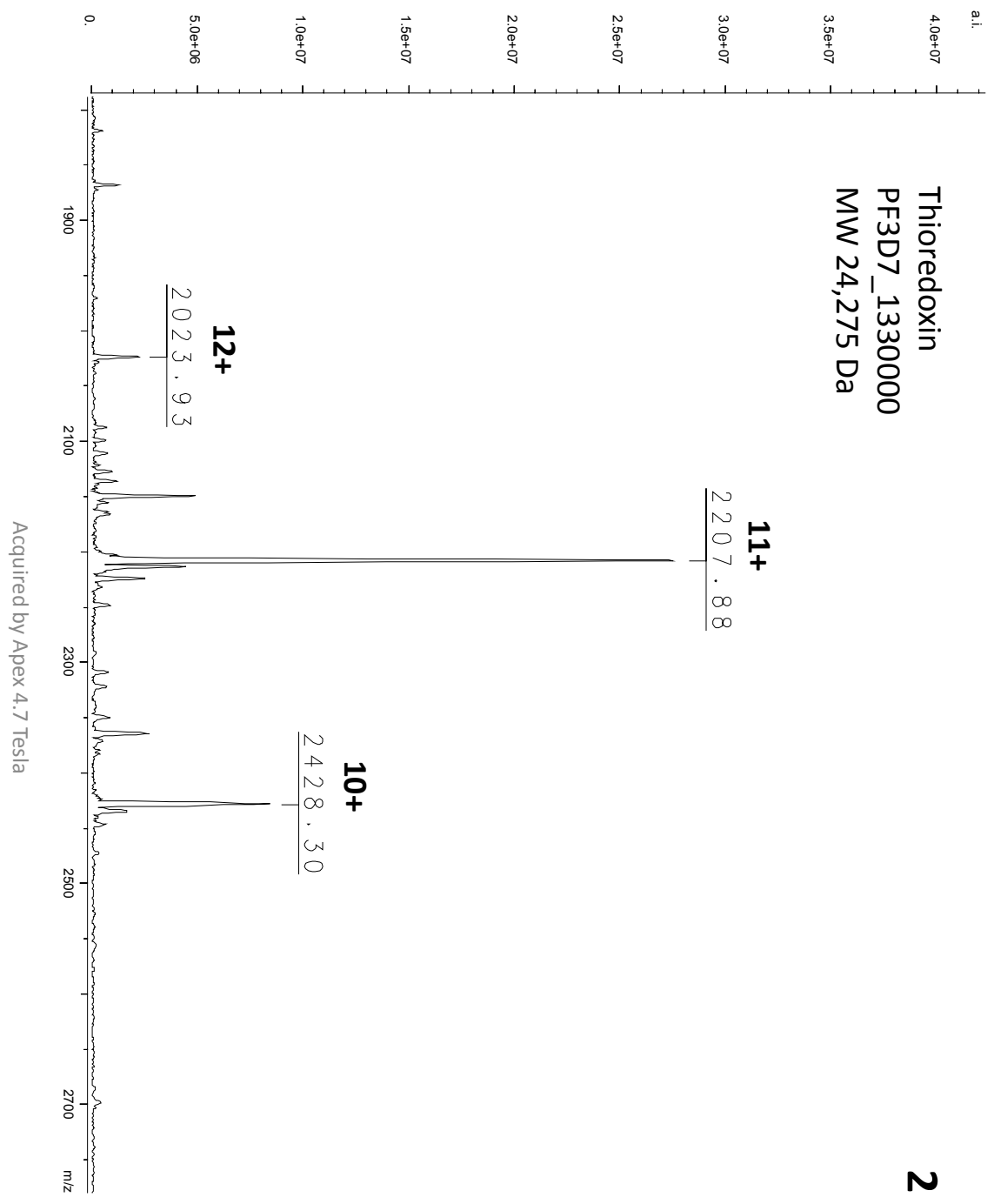
	oxoisindolin-1-yl)ethyl)acetamide				
75	Derivative of astrocasine, 2,3-dihydro-1 <i>H</i> -benzo ³⁶ azepin-1-one	42	-		Astrocasine was found in <i>Astrocasia phyllanthoides</i> ⁷⁵
76	Derivative of astrocasine, 2-acetyl-7-methoxy-2,3,4,5-tetrahydro-1 <i>H</i> -benzo ³⁶ azepin-1-one	45	-		Astrocasine was found in <i>Astrocasia phyllanthoides</i> ⁷⁵
77	(S)-Stachydine	26	-		<i>Capparis tomentosa</i> , <i>Chrysanthemum</i> spp., <i>Citrus</i> spp., <i>Galeopsis grandiflora</i> , <i>Lagochilus hirtus</i> , <i>Medicago sativa</i> , <i>Stachys</i> spp., <i>Aspergillus oryzae</i> ⁷⁶
78	Preclathridine A	100	24.5		<i>Notodoris gardineri</i> ⁷⁷
79	<i>N</i> -Phenylmethylglutamine	57	>100		<i>Dysoxylum</i> sp. ⁷⁸
80	(-)-Norsecurinine	25	40.3		<i>Flueggea virosa</i> *, <i>Securinega virosa</i> ^{79, 80}
81	Allosecurinine				<i>Securinega</i> spp., <i>Securinega suffruticosa</i> ^{81, 82}
82	L-tryosine				Widespread in plants
83	<i>N</i> -(1-carboxyethyl)-5-oxoproline	49	-		<i>Timonius</i> sp. ⁷⁸
84	New compound 1	30	-		<i>Lindera gambleana</i> * ⁷⁸
85	Methyl gallate	18	-	38	<i>Barklya syringifolia</i> *, <i>Koelreuteria paniculata</i> , <i>Euphorbia</i> spp., <i>Geranium</i> spp., <i>Sapium sebiferum</i> , <i>Acacia</i> spp., <i>Spirogyra</i> spp., <i>Rhus glabra</i> , <i>Pelargonium sidoides</i> ⁸³
86	Phyllanthine	90	59.3		<i>Margaritaria dubium-traceyi</i> *, <i>Margaritaria indica</i> , <i>Flueggea leucopyra</i> , <i>Securinega suffruticosa</i> , <i>Phyllanthus discoides</i> ⁸⁴
87	4-Hydroxyallosecurine	51	>100		<i>Margaritaria dubium-traceyi</i> *, <i>Phyllanthus niruri</i> ⁸⁵
88	Securinine	39	20.6		<i>Margaritaria dubium-traceyi</i> *, <i>Margaritaria indica</i> , <i>Securinega suffruticosa</i> , <i>Phyllanthus discoides</i> ⁸⁶
89	Securinine <i>N</i> -oxide				Natural Product Fragment Library ⁸⁷
90	<i>N</i> -Methyltryptophan	41	79.2		<i>Gastrolobium grandiflorum</i> *, <i>Erythrina caffra</i> , <i>Abrus precatorius</i> , <i>Desmodium tiliaefolium</i> ⁸⁸
91	1-Prenyl-6 <i>N</i> -methylisoguanine (new)	35	-		<i>Glochidion zeylanicum</i> ⁷⁸
92	New compound 2				⁷⁸
93	Gentianol				<i>Swertia</i> spp. ⁸⁹
94	Derivative of grandisine E				Grandisine E was found in <i>Elaeocarpus grandis</i> ⁹⁰
95	5-Methoxydictamnine				<i>Ruta chalepensis</i> var. <i>latifolia</i> (Rutaceae) ⁹¹
96	2-Amino-1-phenylethanol				<i>Halostachys caspica</i> (Chenopodiaceae) ⁹²

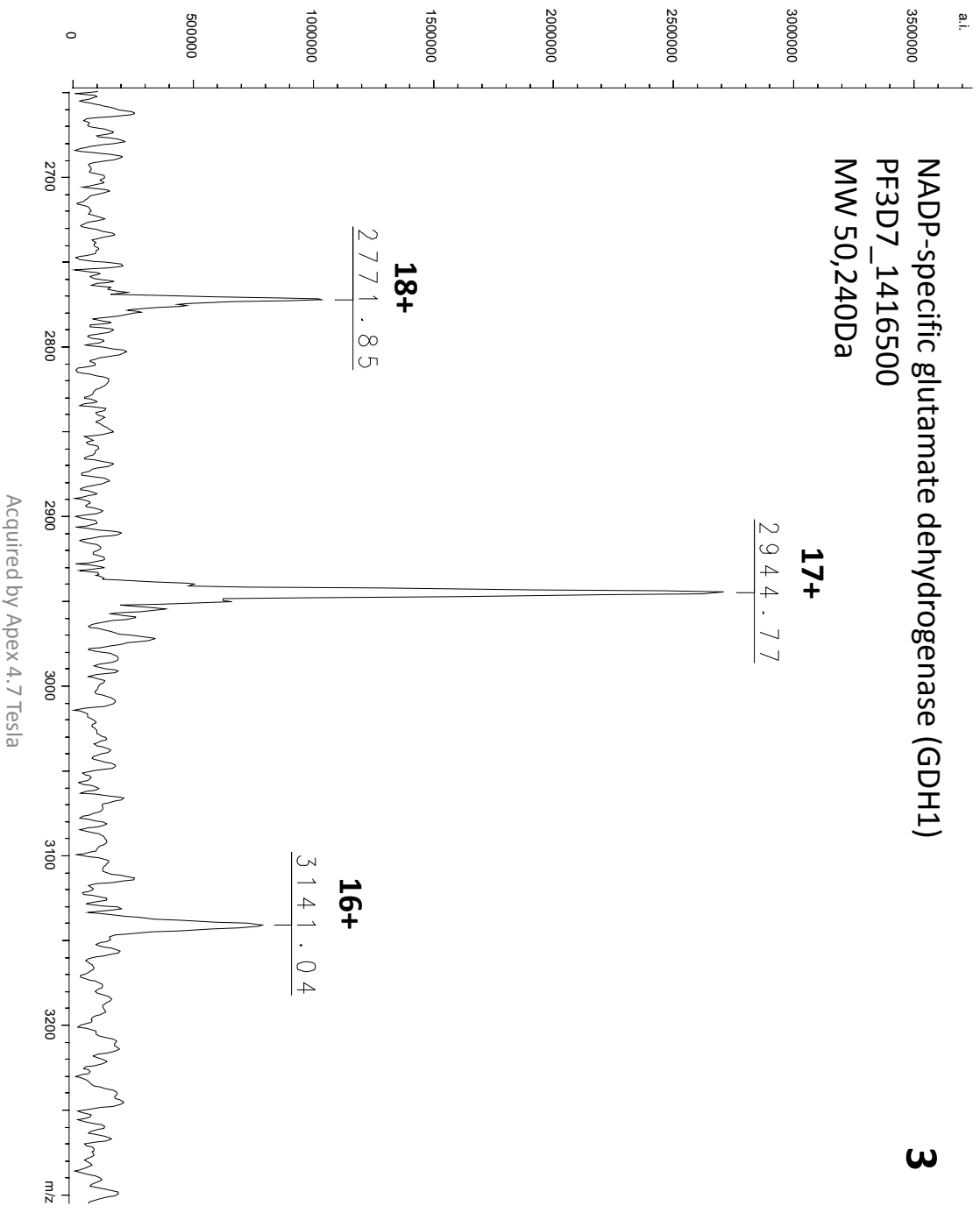
Figure S3. Mass spectra of 62 *Plasmodium* proteins



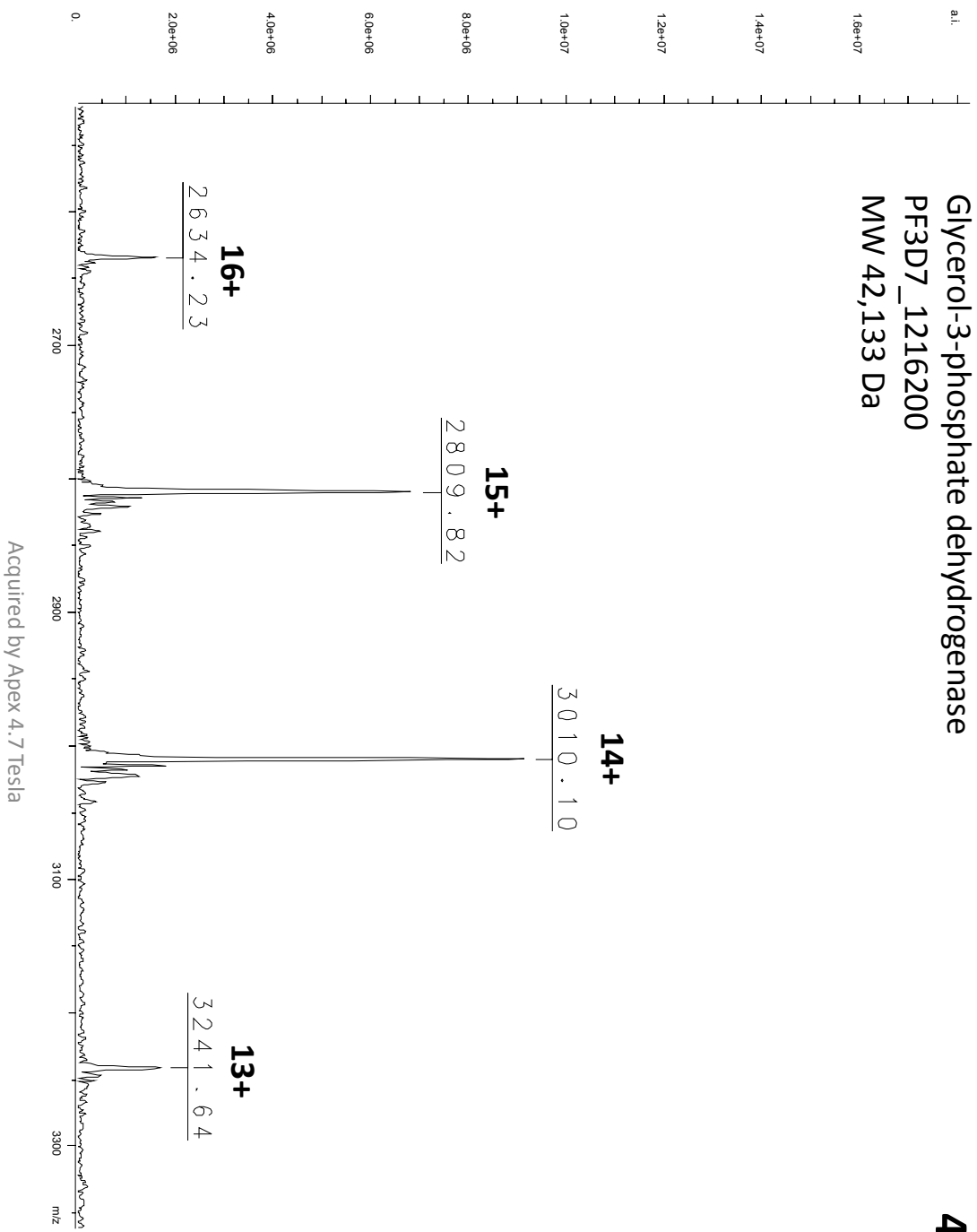
2

Thioredoxin
PF3D7_1330000
MW 24,275 Da



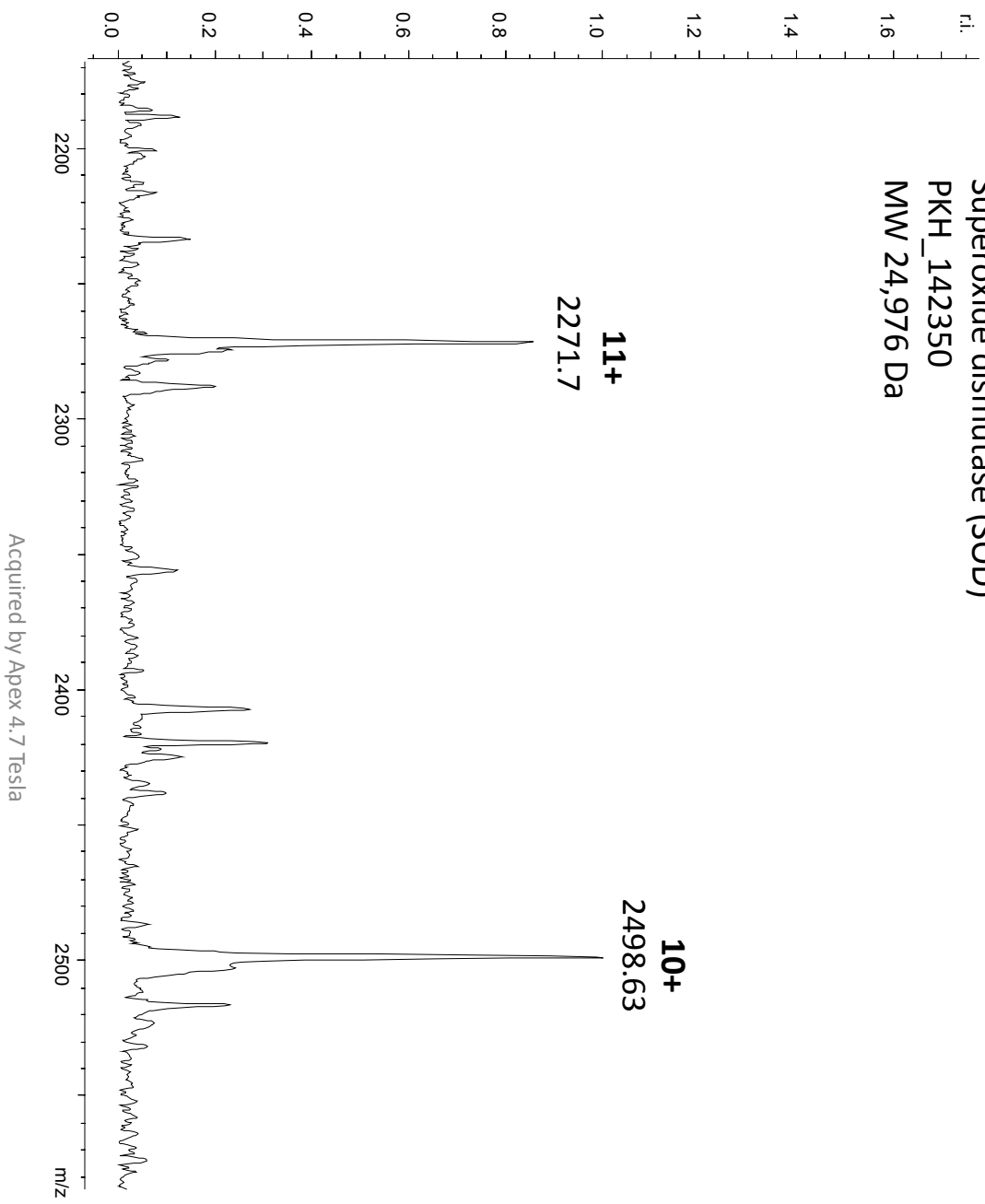


Glycerol-3-phosphate dehydrogenase
PF3D7_1216200
MW 42,133 Da

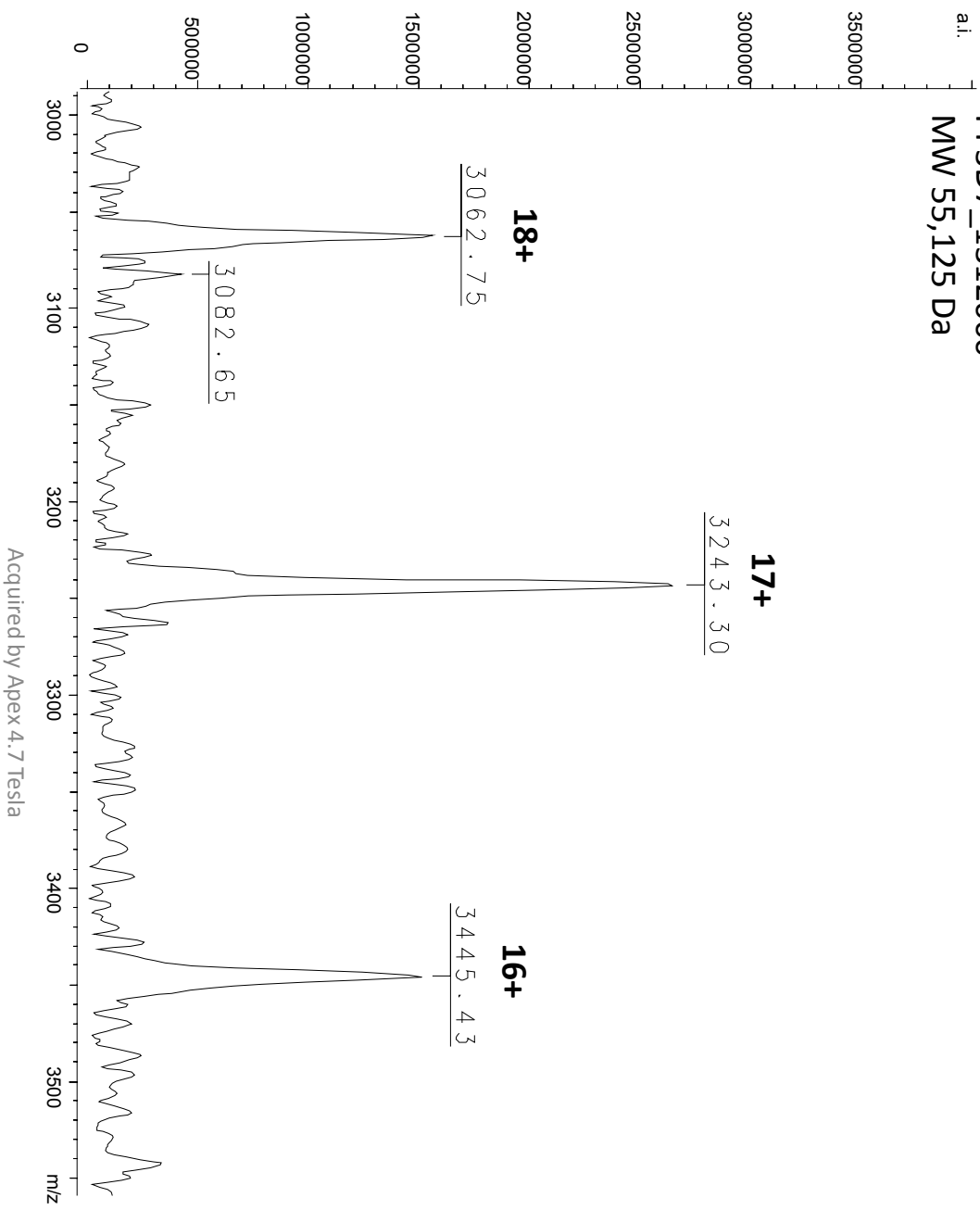


5

Superoxide dismutase (SOD)
PKH_142350
MW 24,976 Da



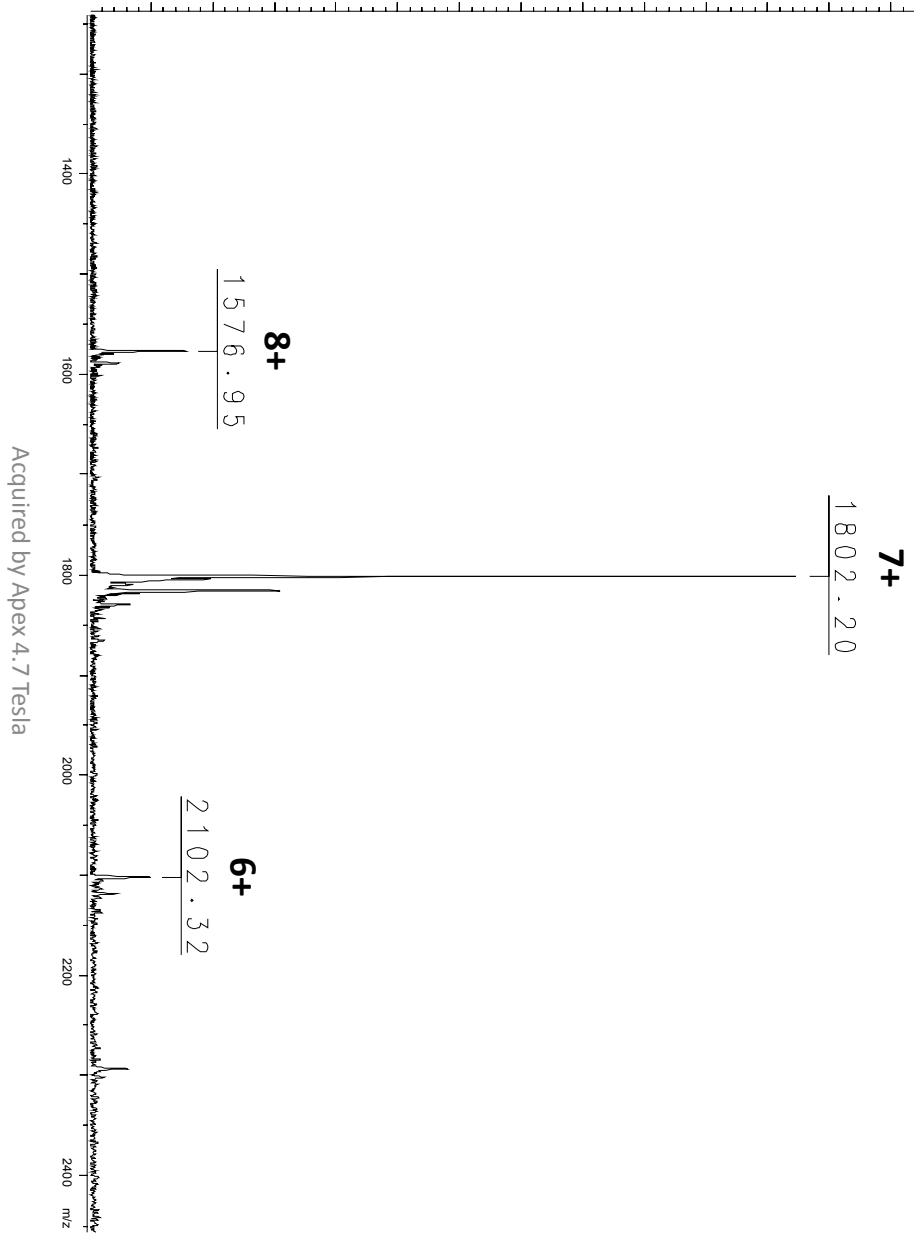
branched-chain alpha keto-acid dehydrogenase (BCKDHA)
PF3D7_1312600
MW 55,125 Da



a.l.
1.5e+07
1.4e+07
1.3e+07
1.2e+07
1.1e+07
1.0e+07
9.0e+06
8.0e+06
7.0e+06
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4.0e+06
3.0e+06
2.0e+06
1.0e+06
0

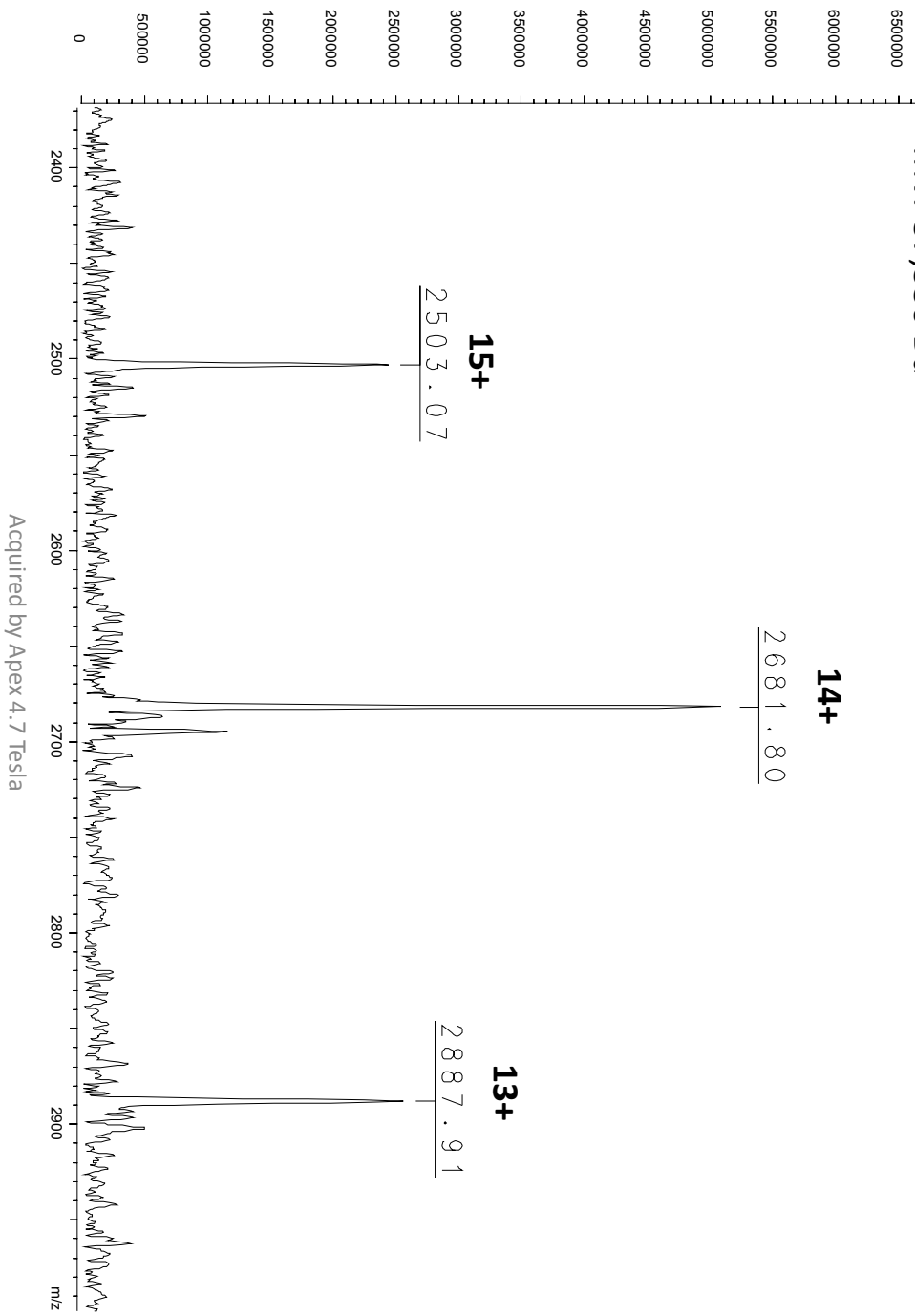
Thioredoxin 1 (TRX1)
PF3D7_1457200
MW 12,607 Da

7

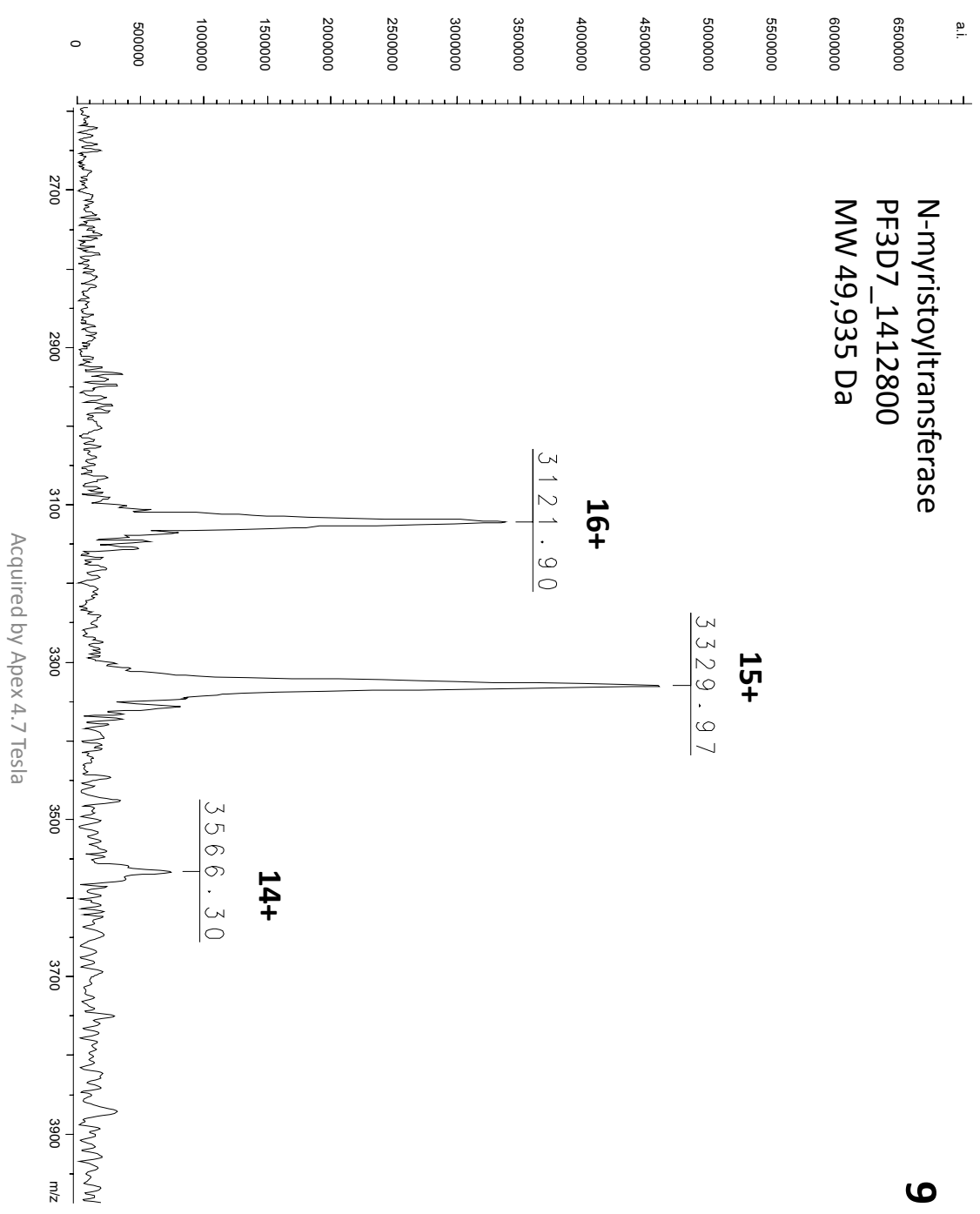


a.i. **Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)** **8**

PF3D7_1462800
MW 37,530 Da

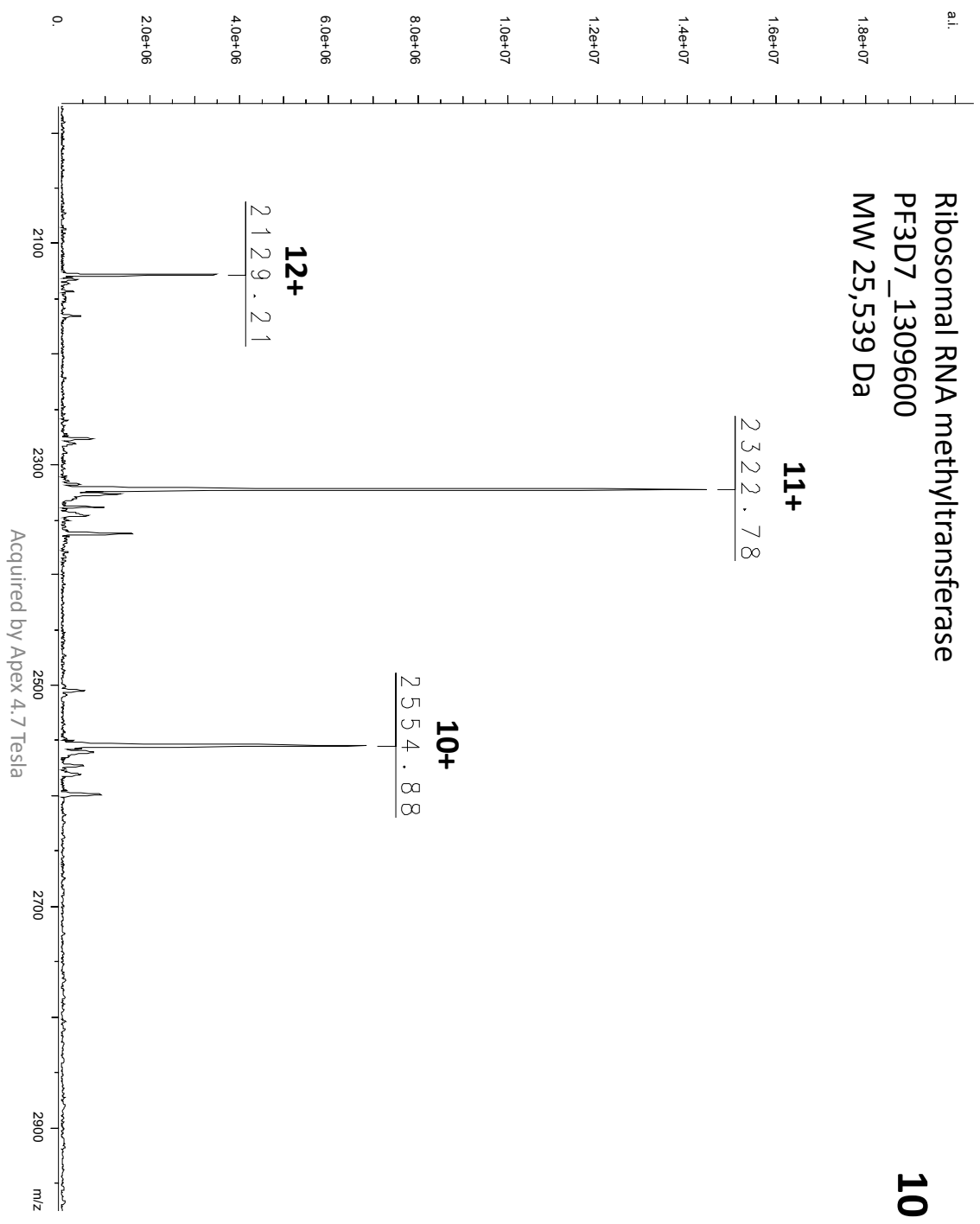


N-myristoyltransferase
PF3D7_1412800
MW 49,935 Da



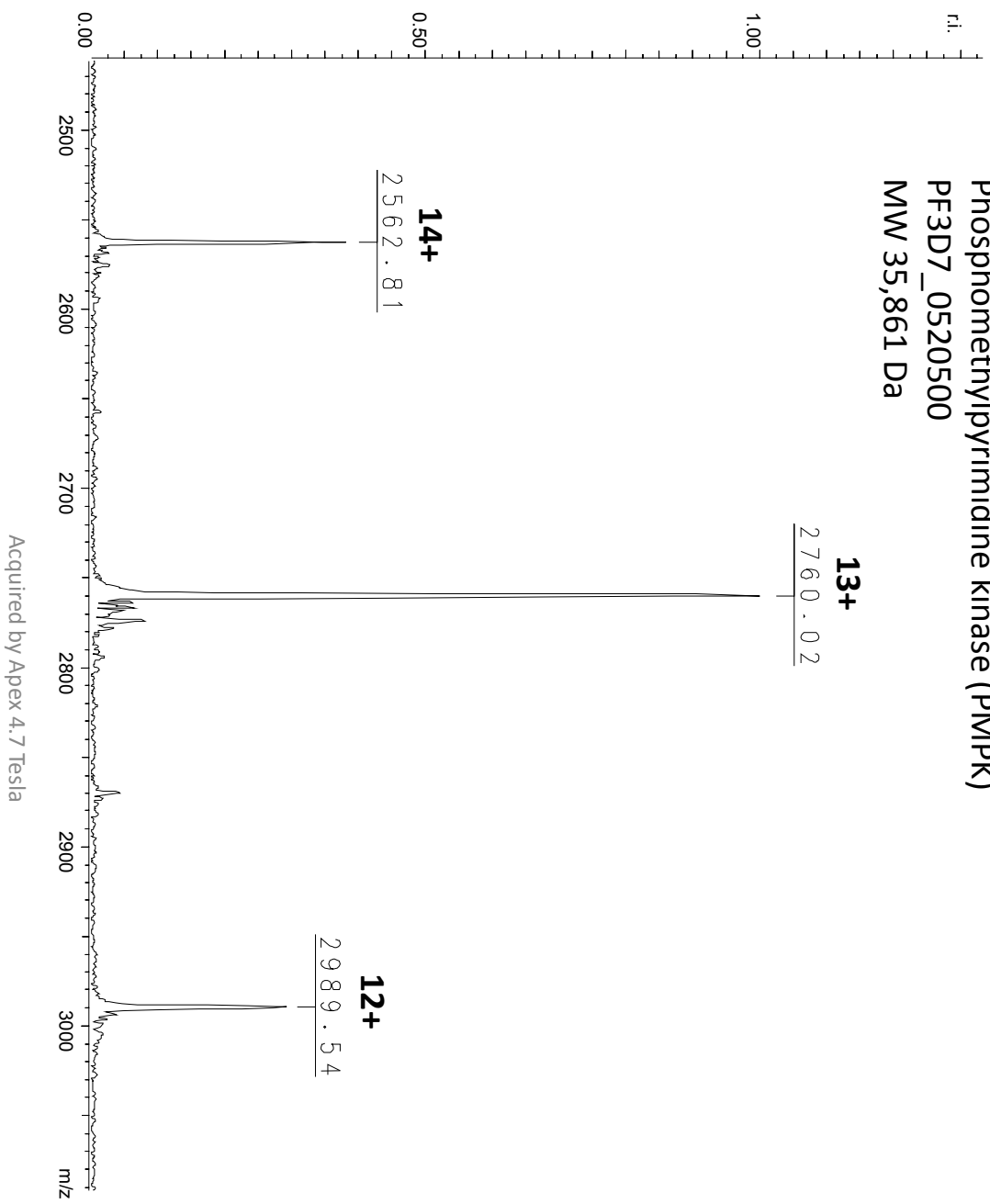
10

Ribosomal RNA methyltransferase
PF3D7_1309600
MW 25,539 Da



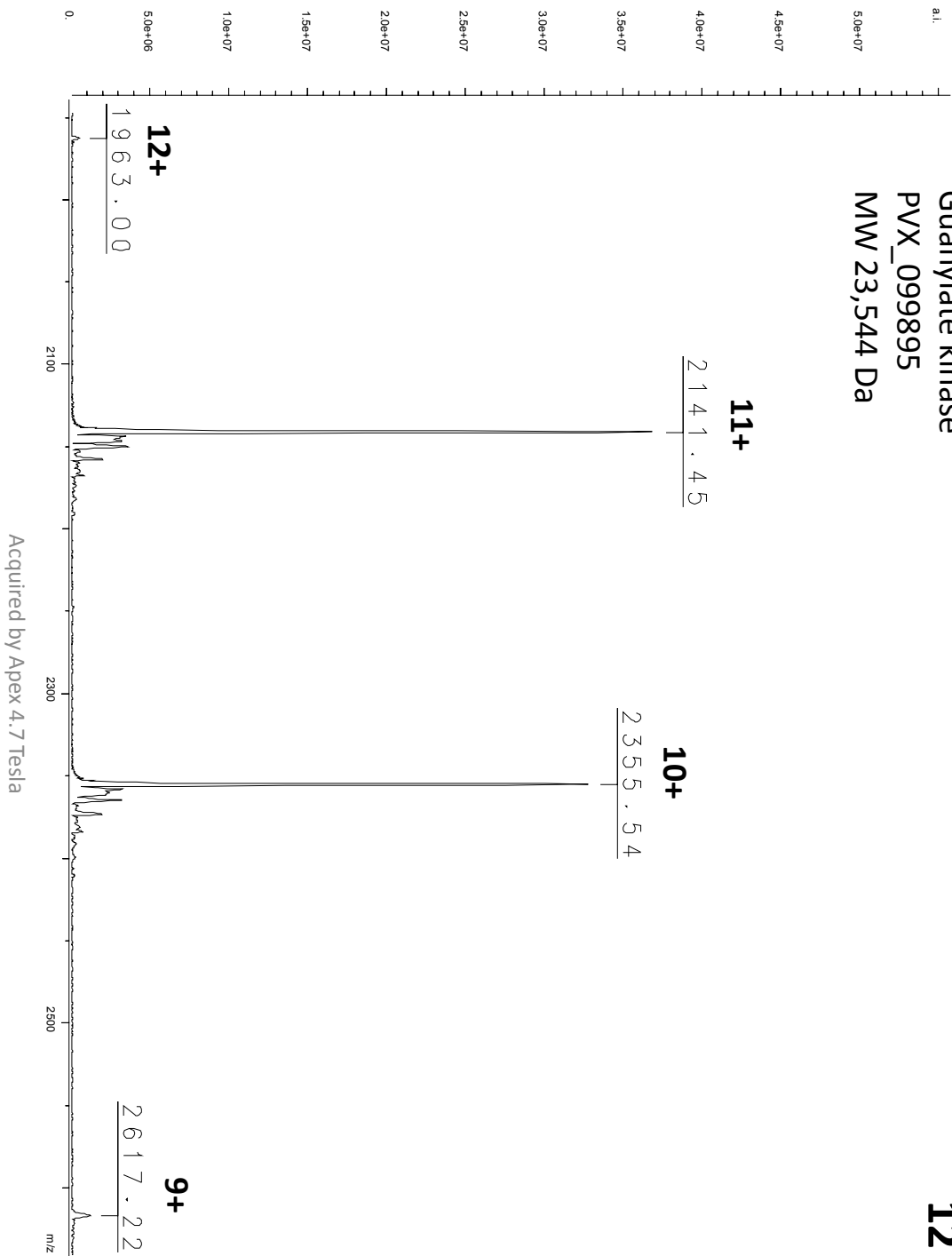
Phosphomethylpyrimidine kinase (PMPK)
PF3D7_0520500
MW 35,861 Da

11

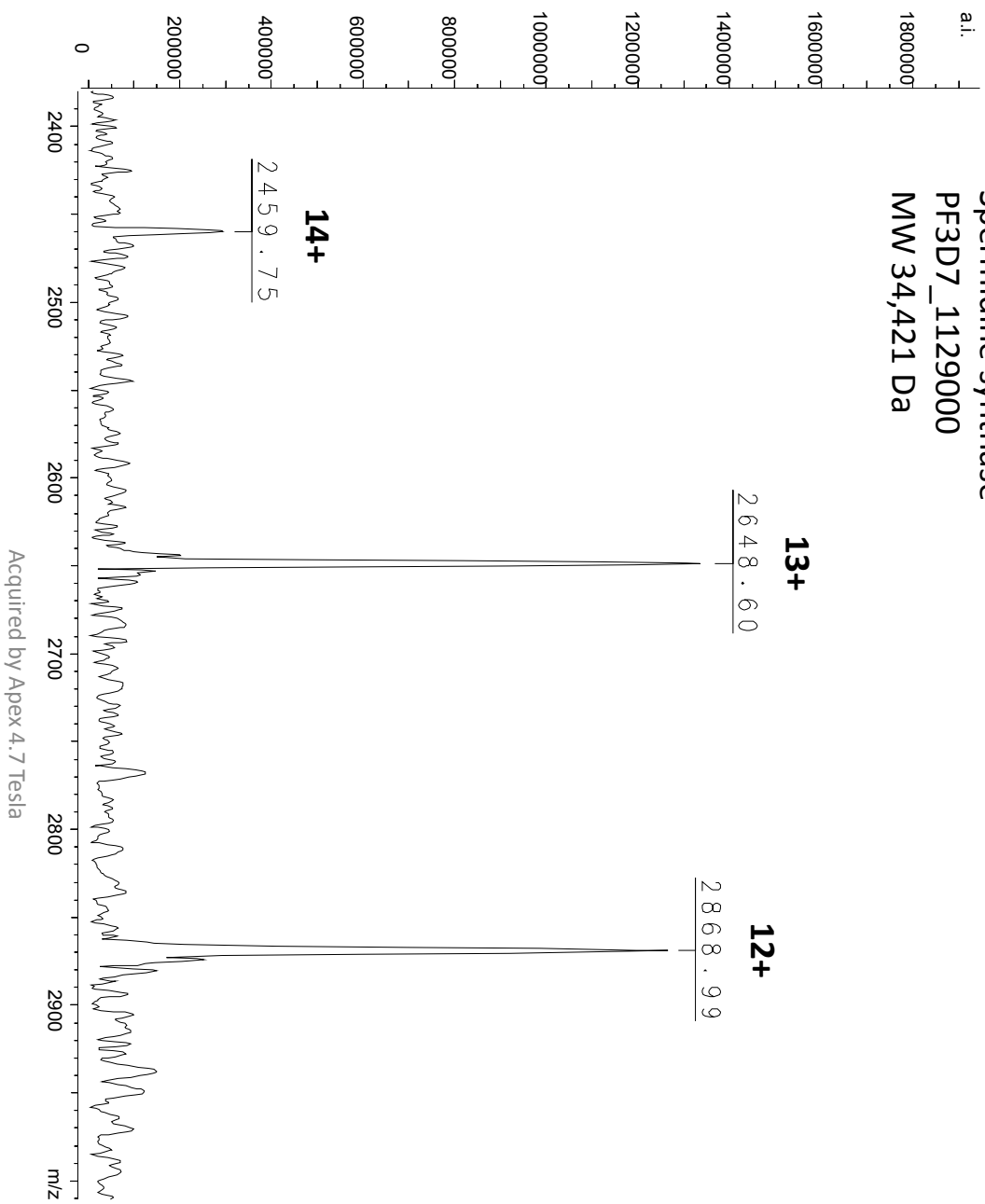


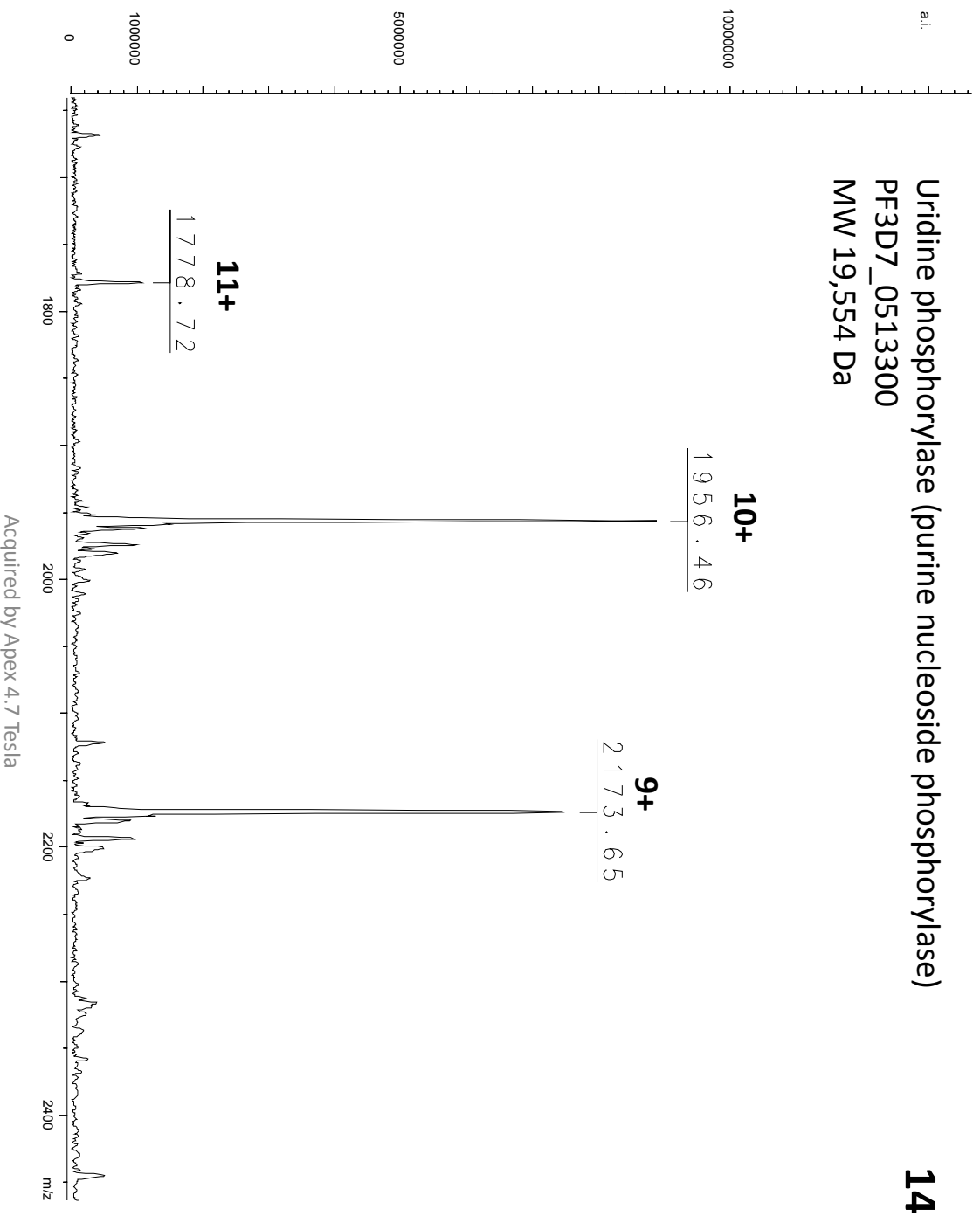
Guanylate kinase
PWX_099895
MW 23,544 Da

12



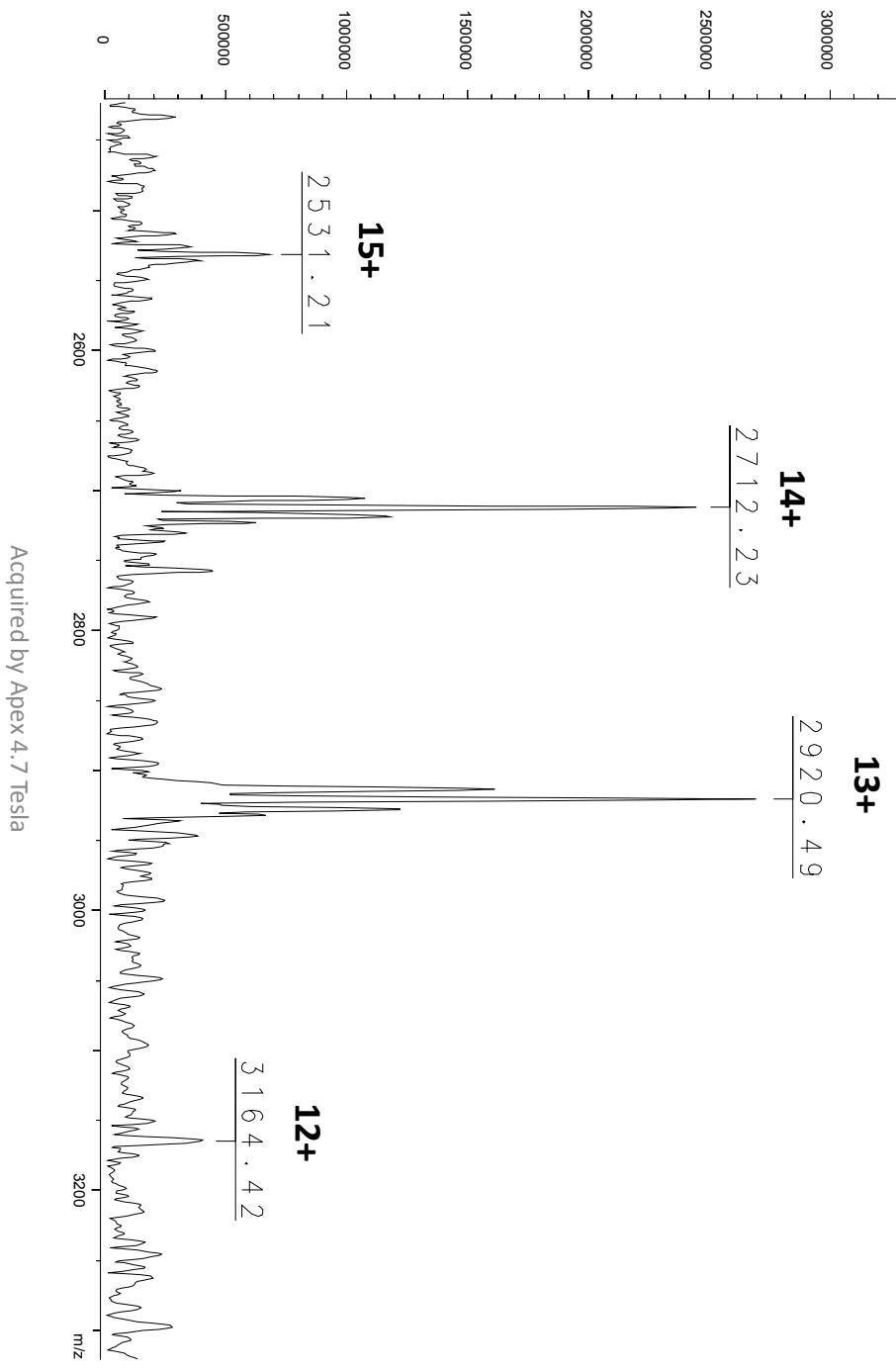
Spermidine synthase
PF3D7_1129000
MW 34,421 Da





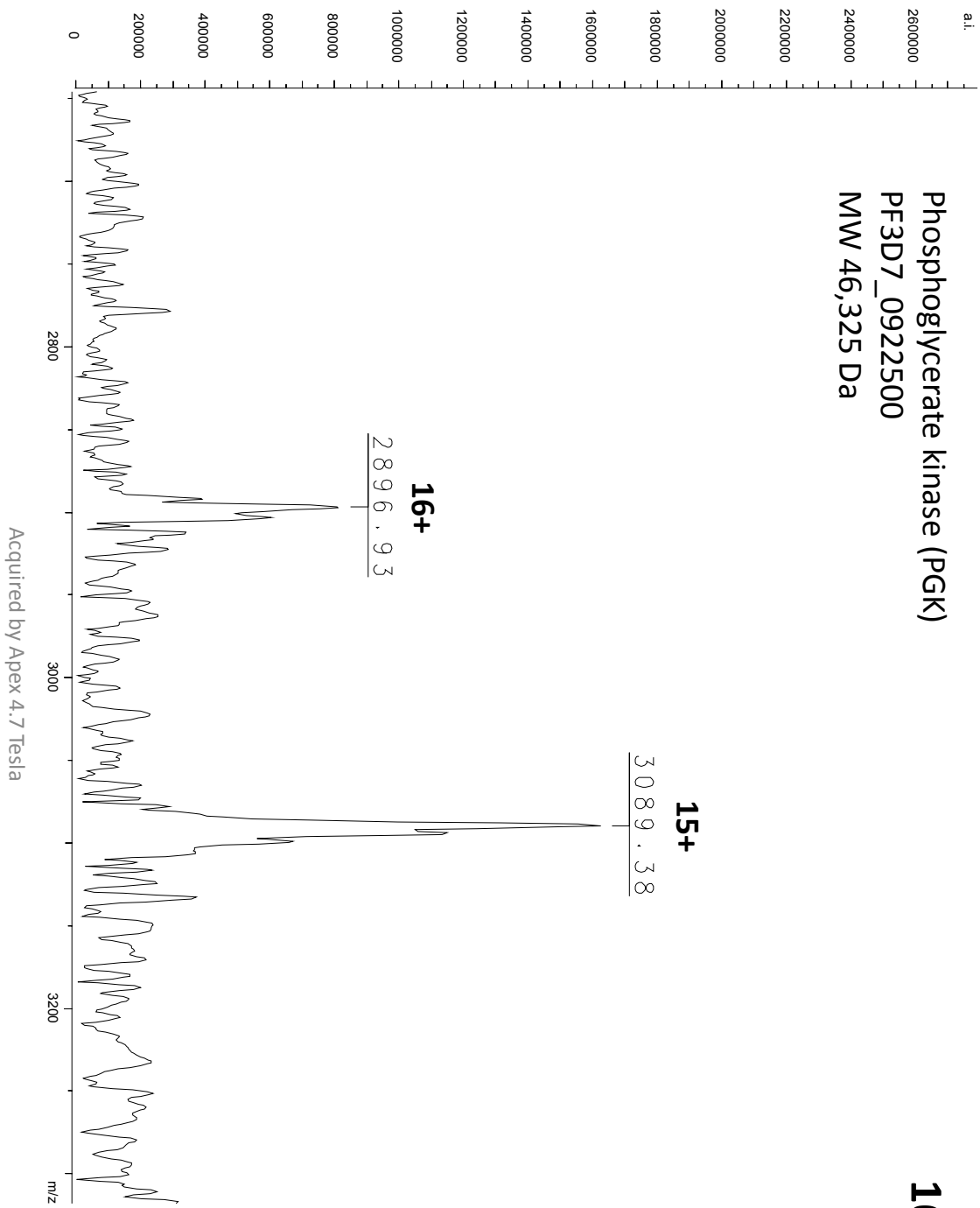
a.i.
CDK-related protein kinase 6 (PK6)
PF3D7_1337100
MW 37,995 Da

15



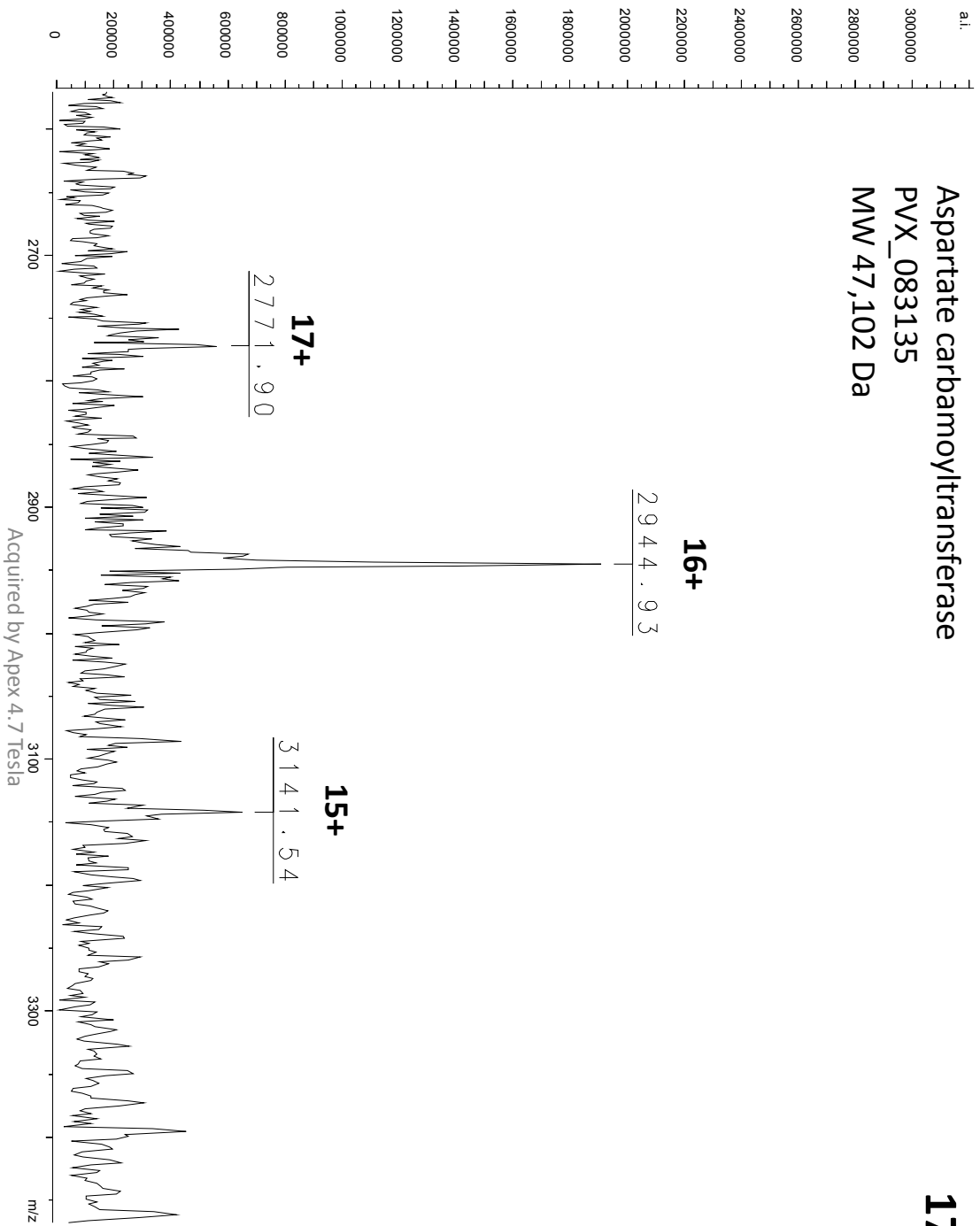
Phosphoglycerate kinase (PGK)
PF3D7_0922500
MW 46,325 Da

16



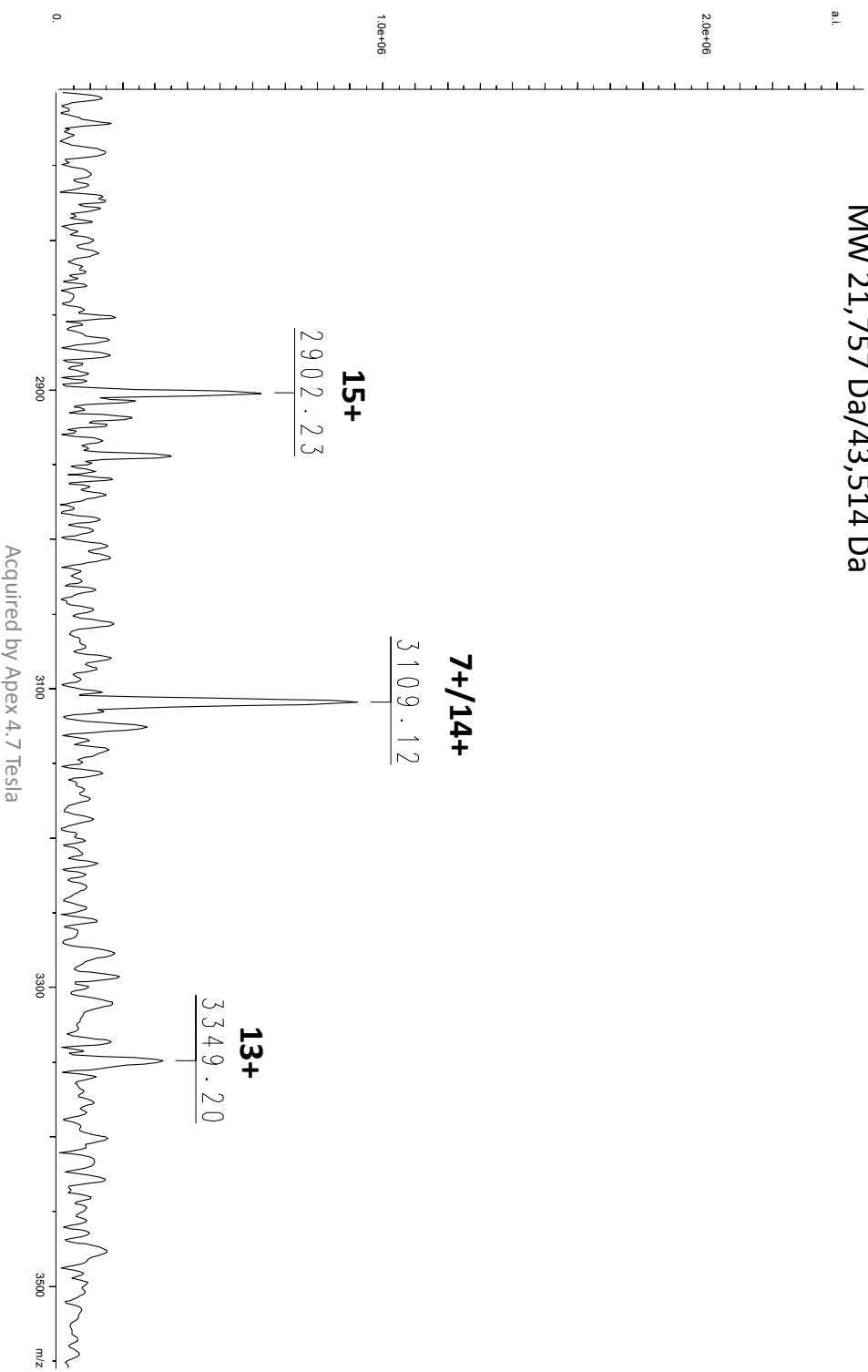
a.i.
Aspartate carbamoyltransferase
PVX_083135
MW 47,102 Da

17

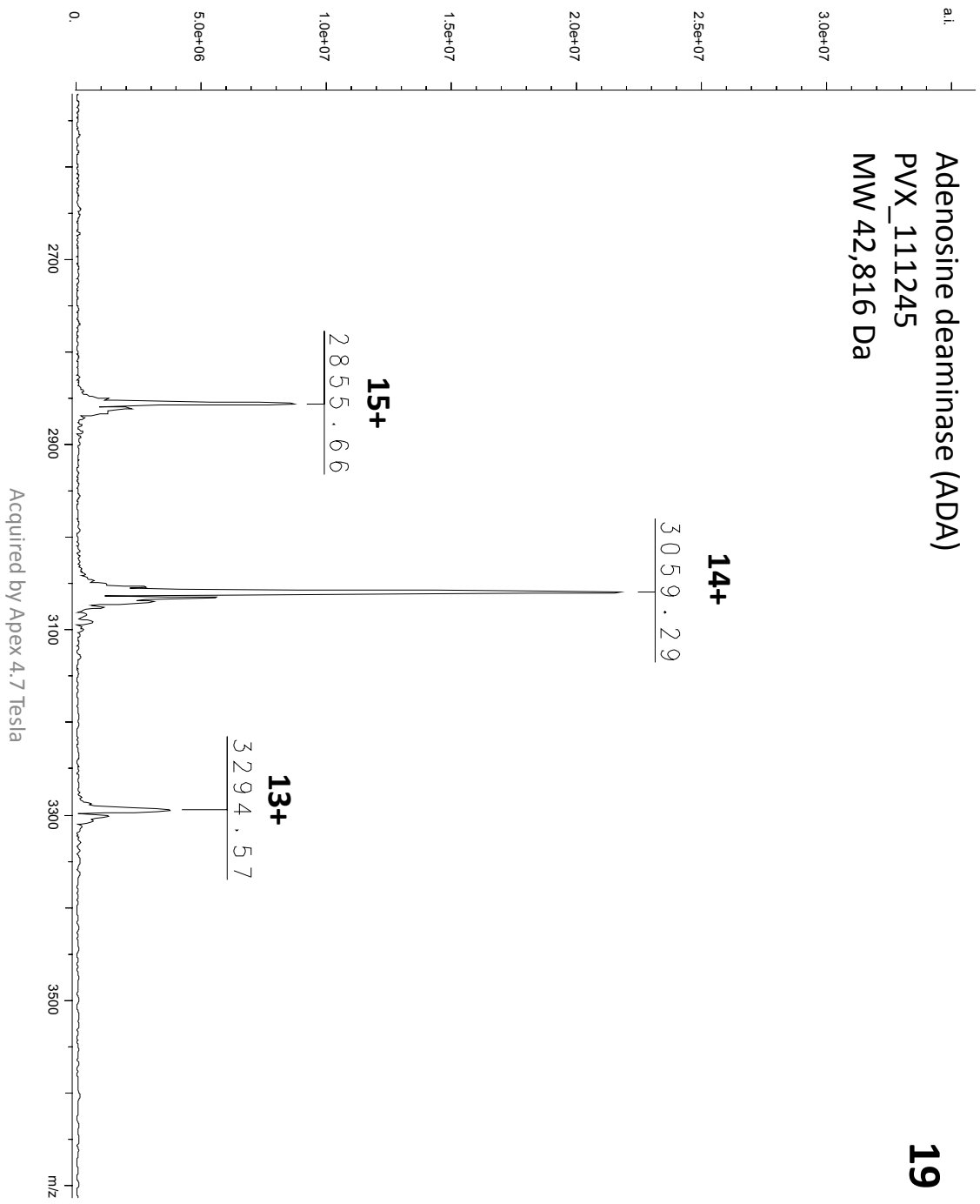


Rab18GTPase
PF3D7_0807300
MW 21,757 Da/43,514 Da

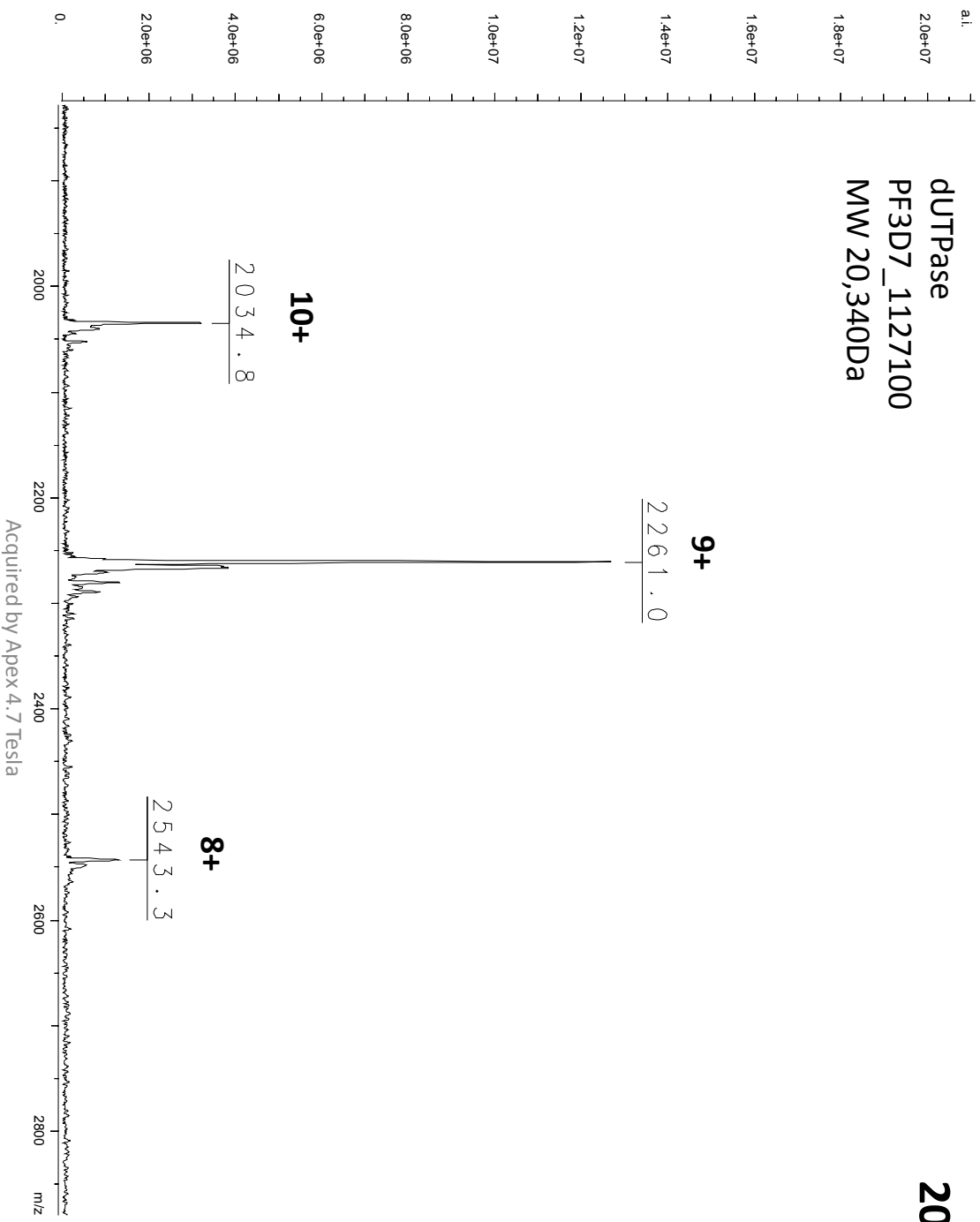
18



Adenosine deaminase (ADA)
PVX_111245
MW 42,816 Da

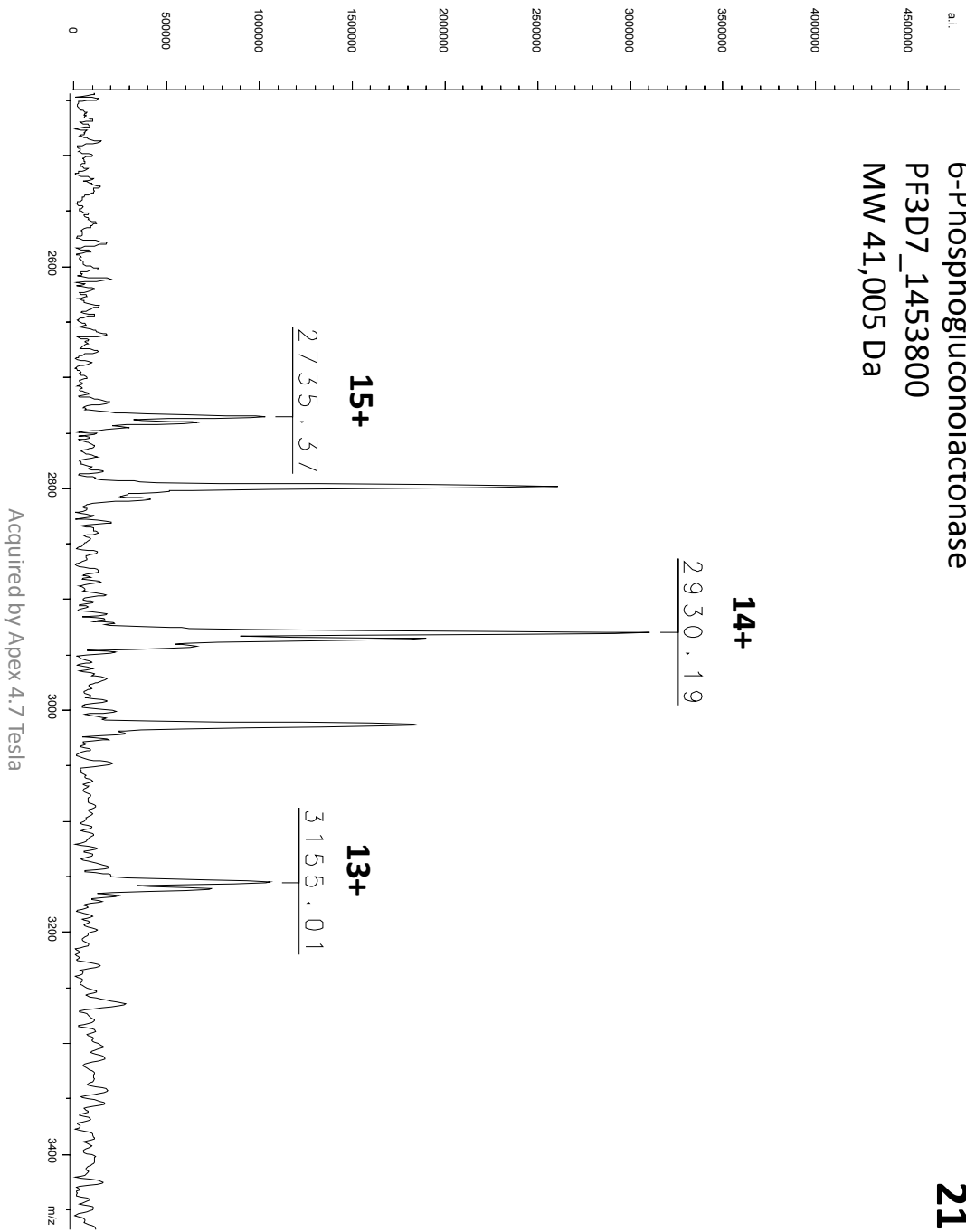


dUTPase
PF3D7_1127100
MW 20,340Da

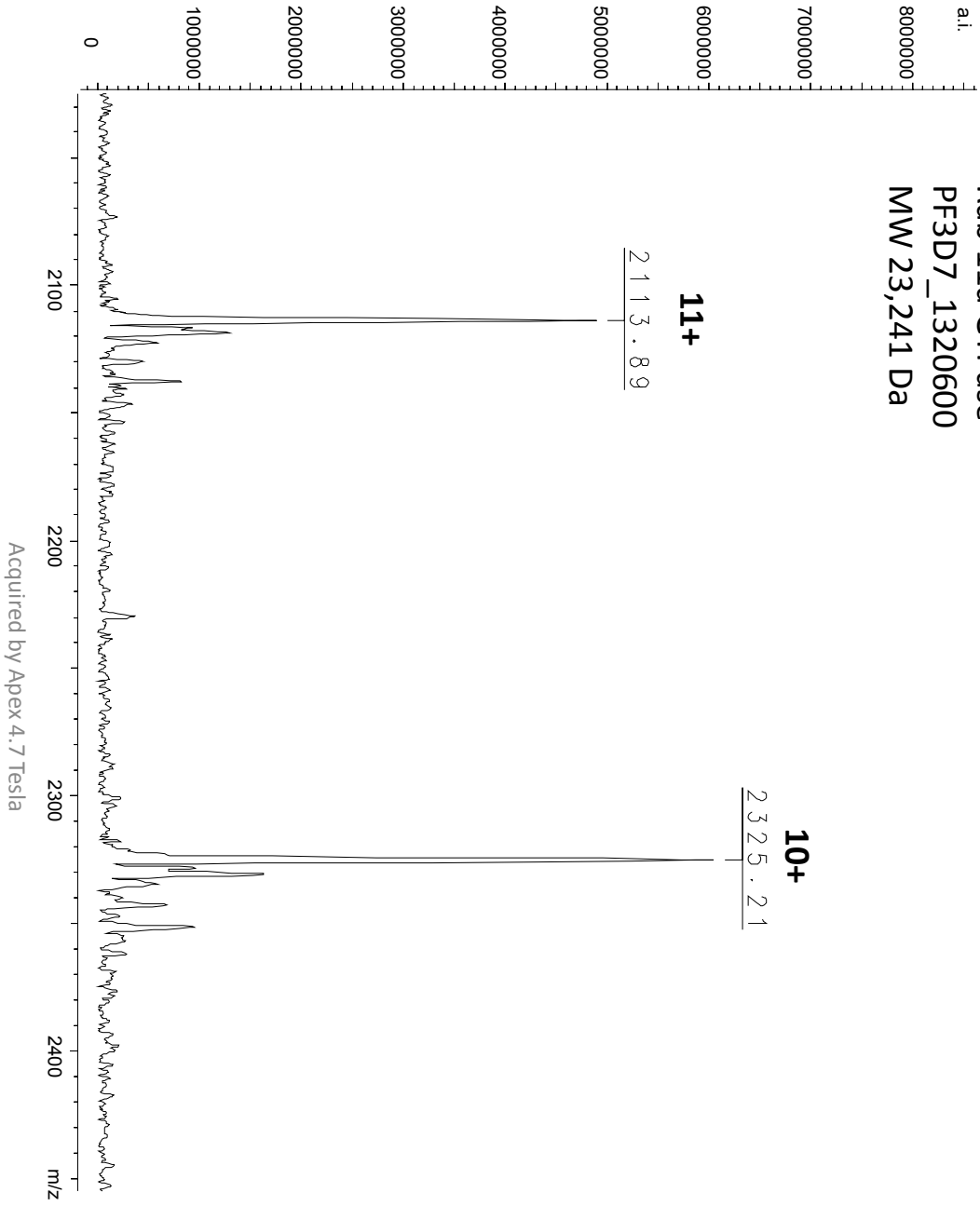


6-Phosphogluconolactonase
PF3D7_1453800
MW 41,005 Da

21

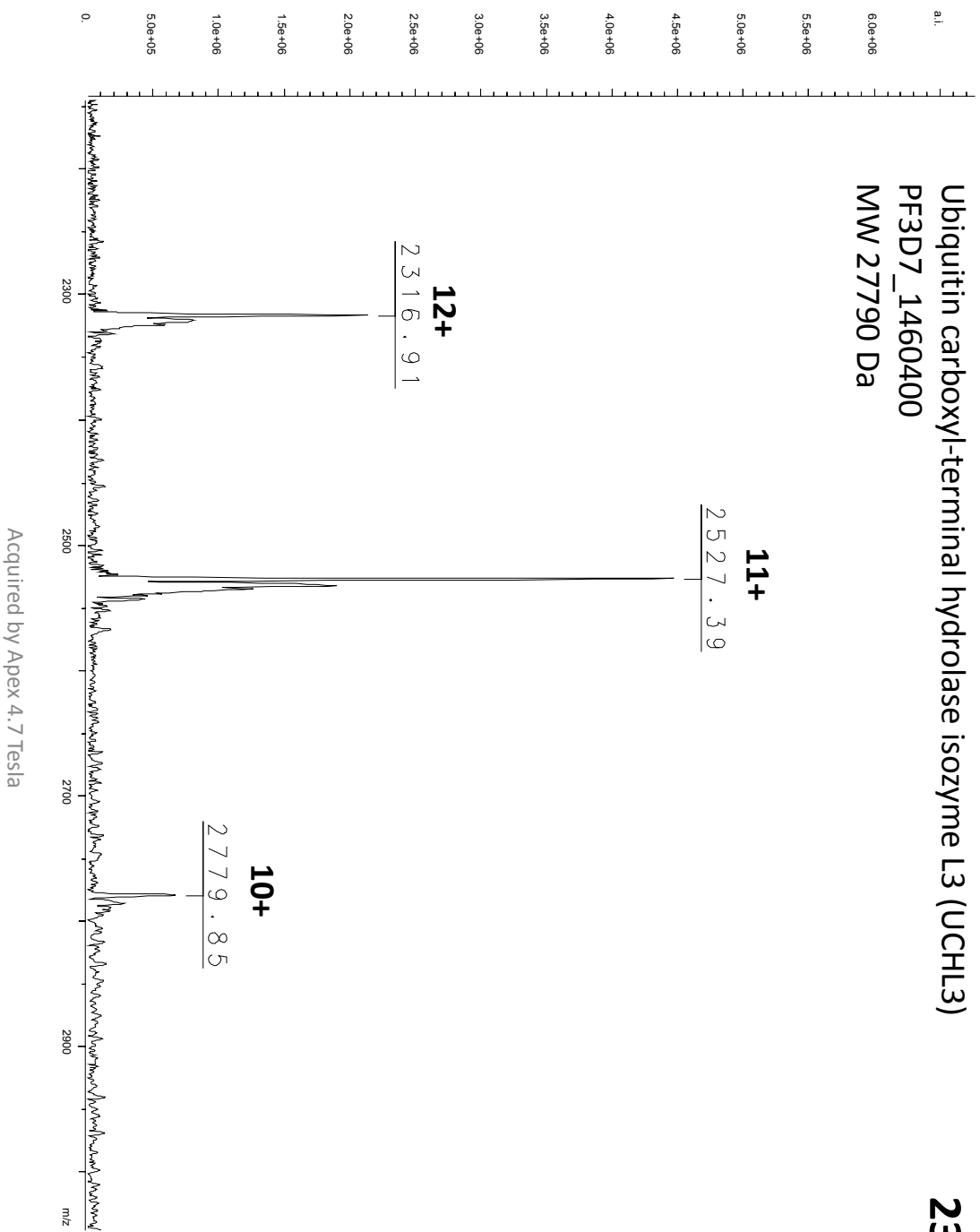


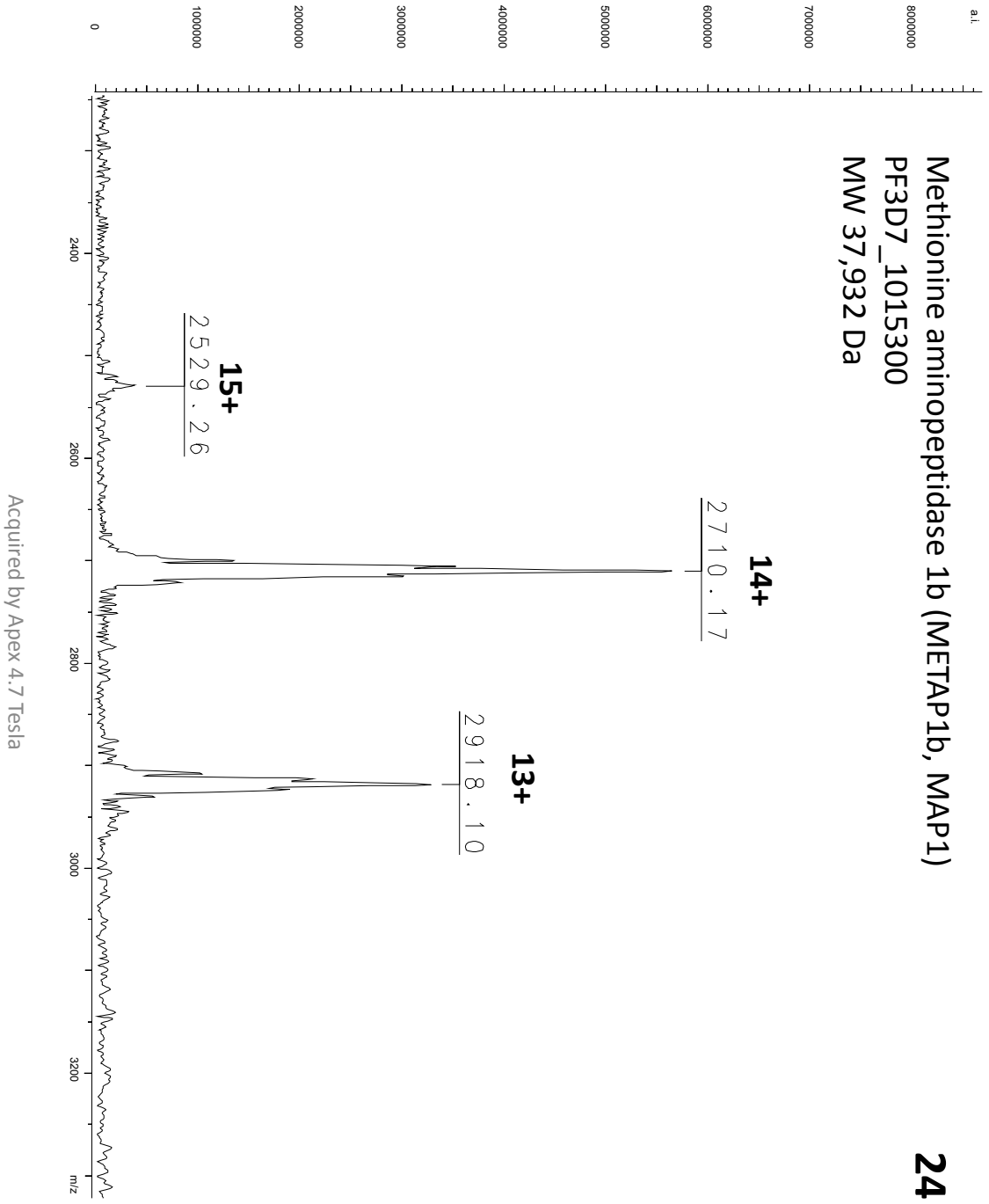
Rab 11a GTPase
PF3D7_1320600
MW 23,241 Da



Ubiquitin carboxyl-terminal hydrolase isozyme L3 (UCHL3)
PF3D7_1460400
MW 27790 Da

23

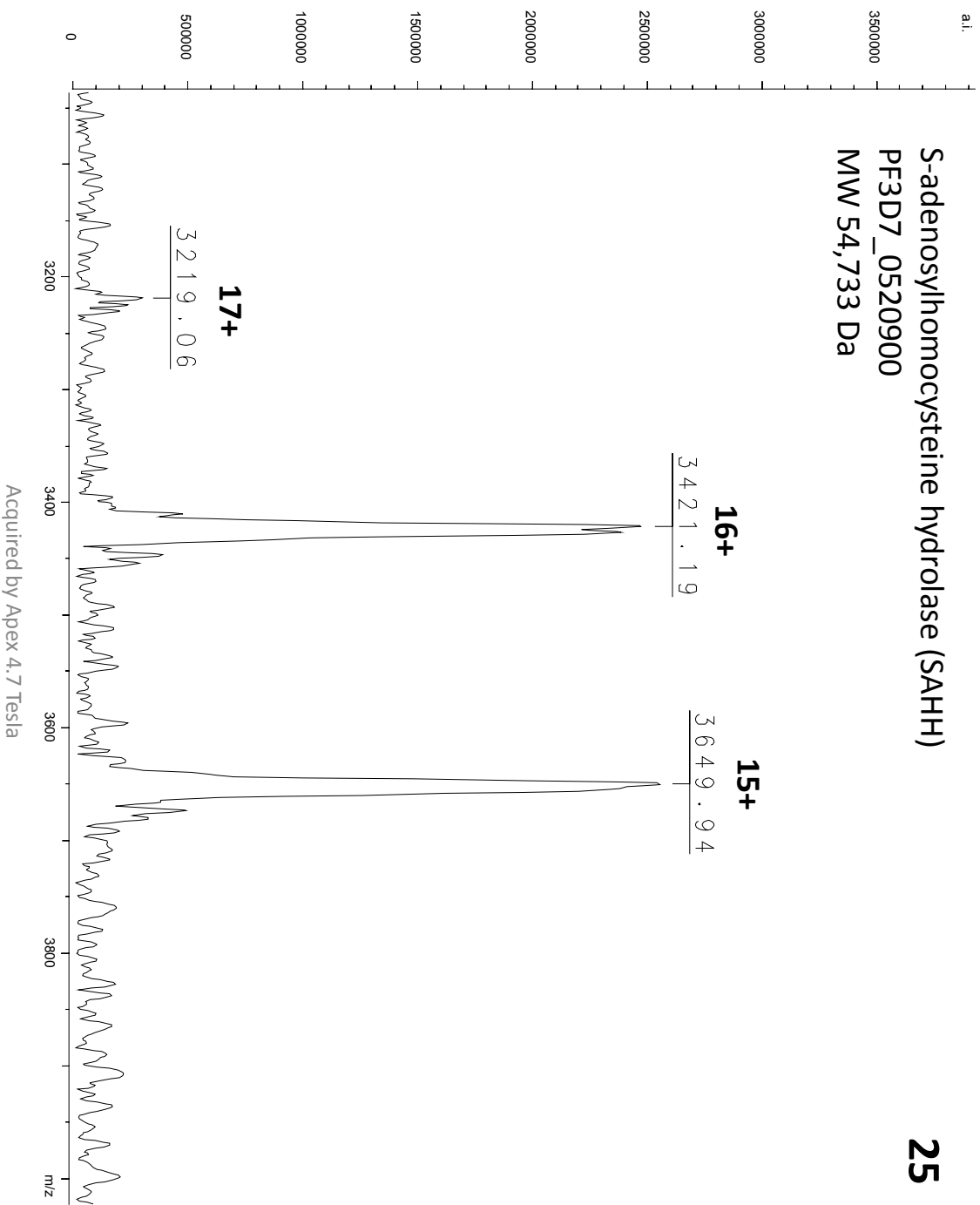




a1.

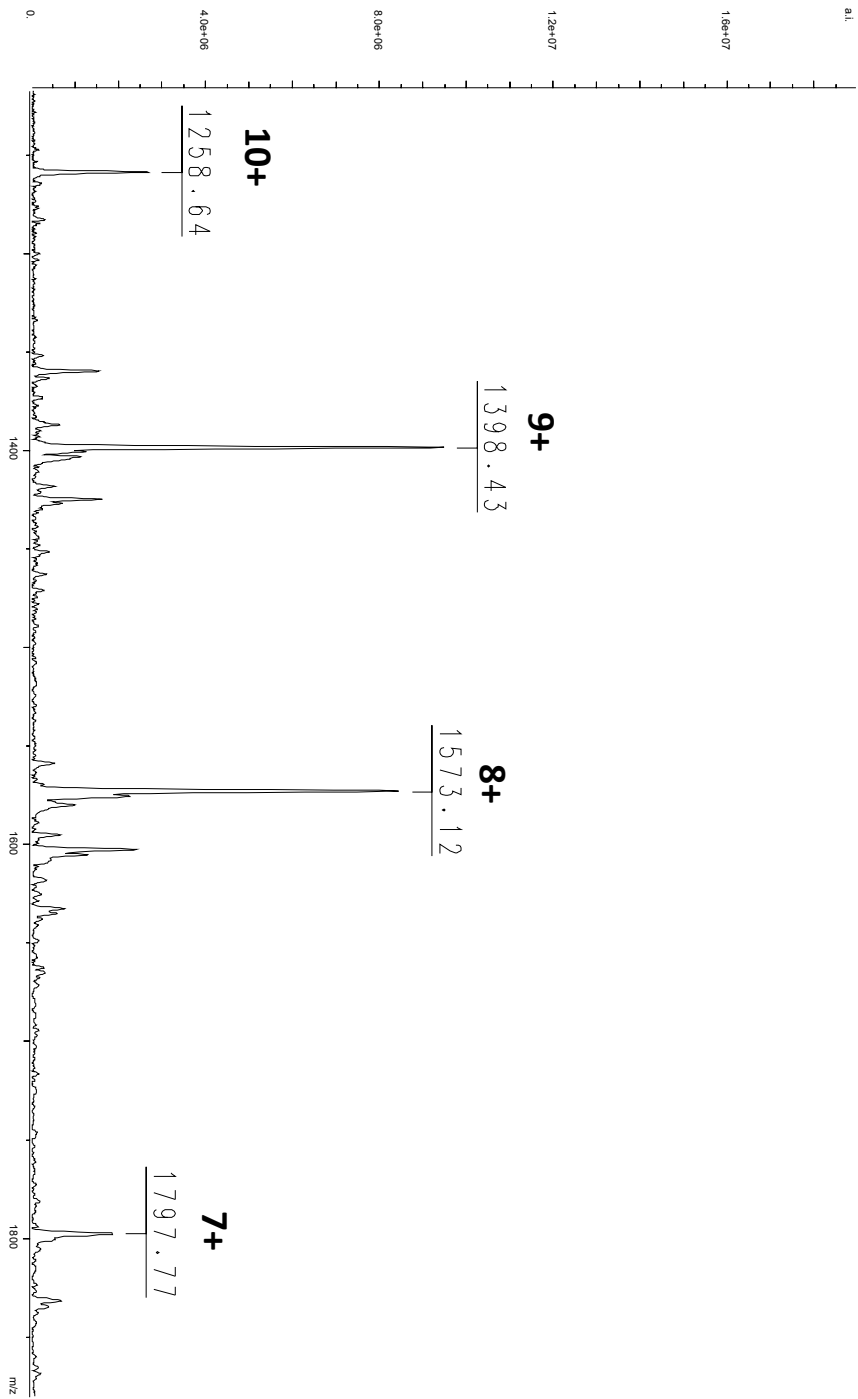
S-adenosylhomocysteine hydrolase (SAHH)
PF3D7_0520900
MW 54,733 Da

25



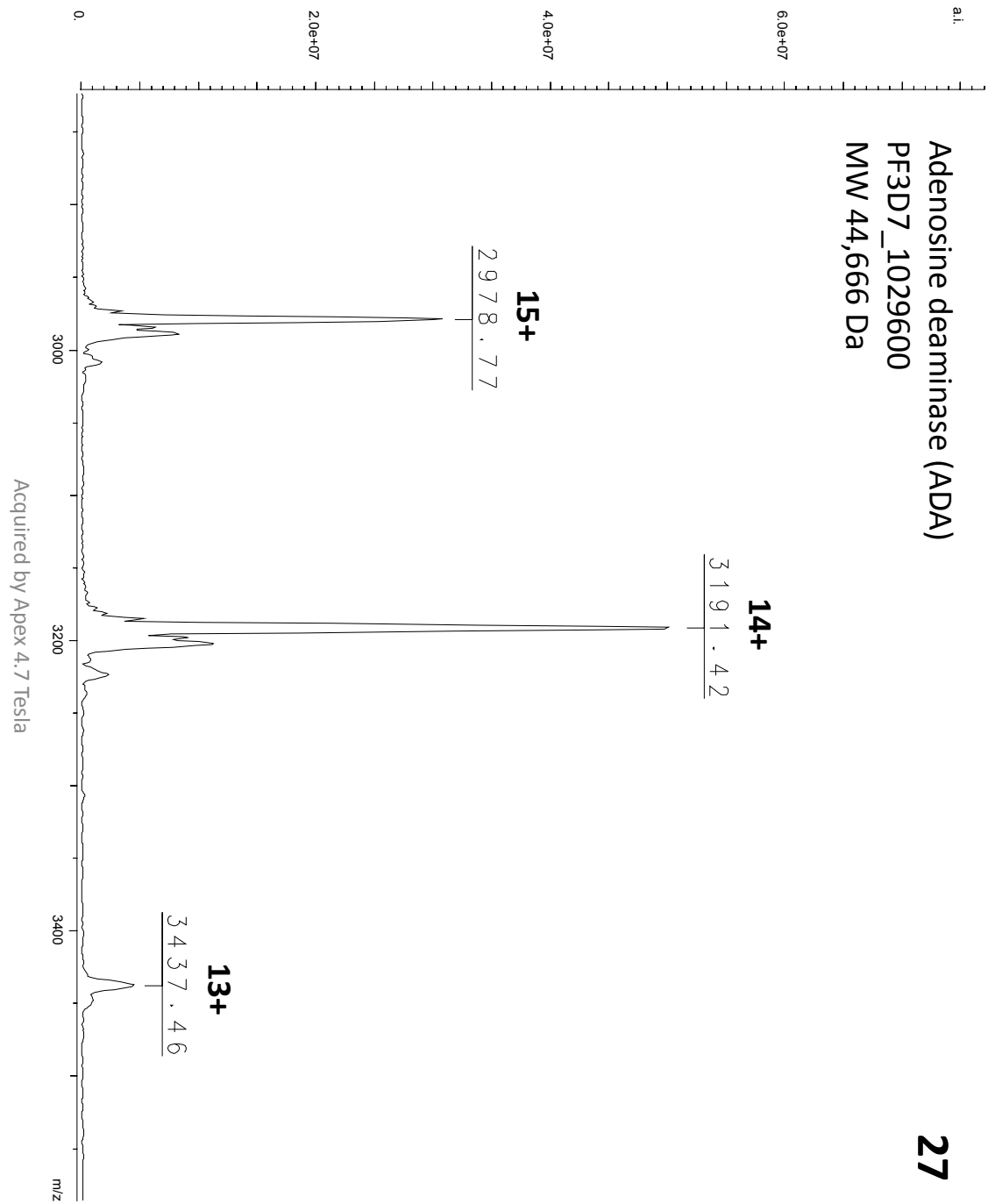
ATP-dependent DNA helicase Q1 (RECQ1)
PF3D7_0918600
MW 12,576 Da

26

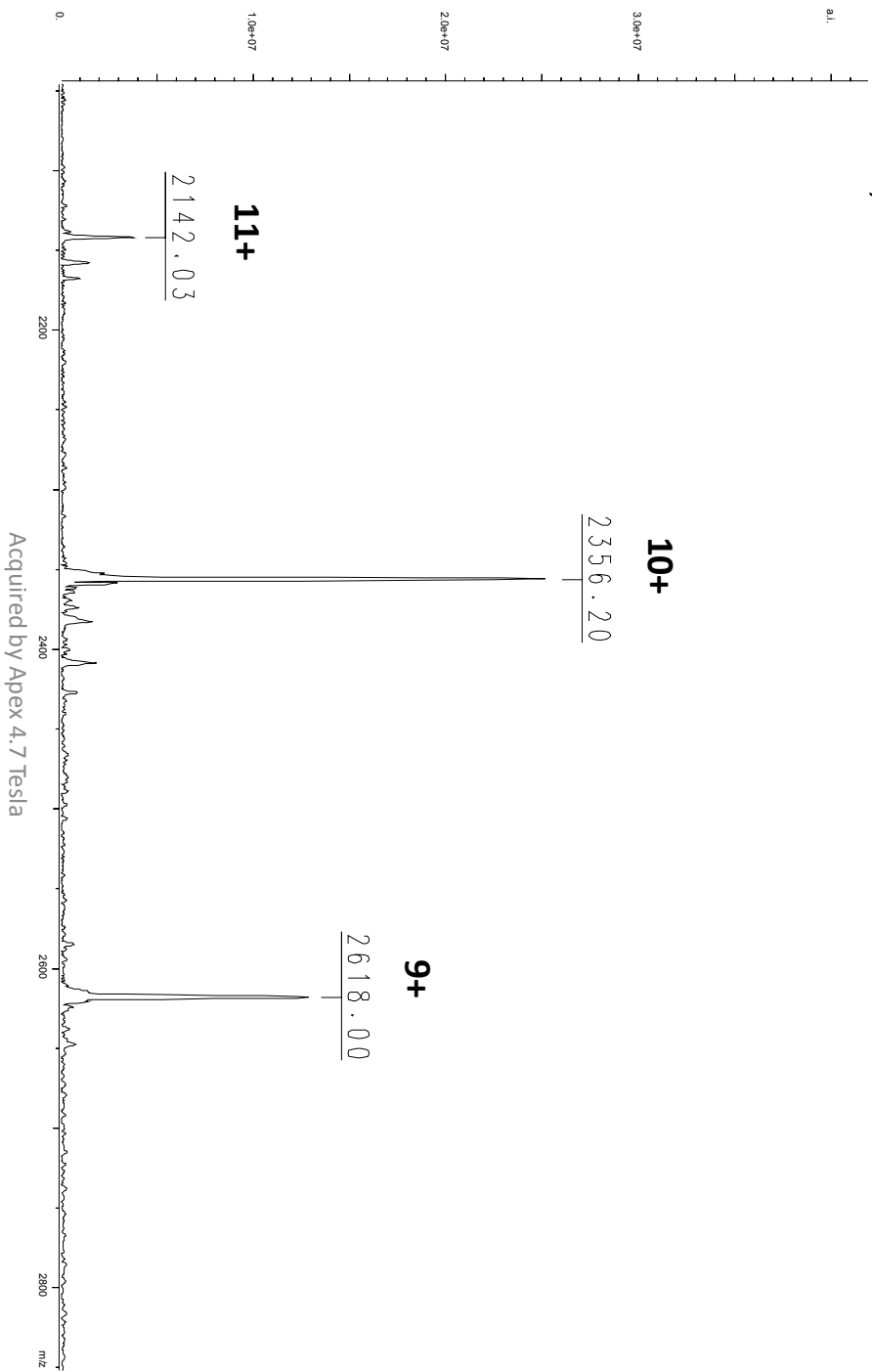


27

Adenosine deaminase (ADA)
PF3D7_1029600
MW 44,666 Da

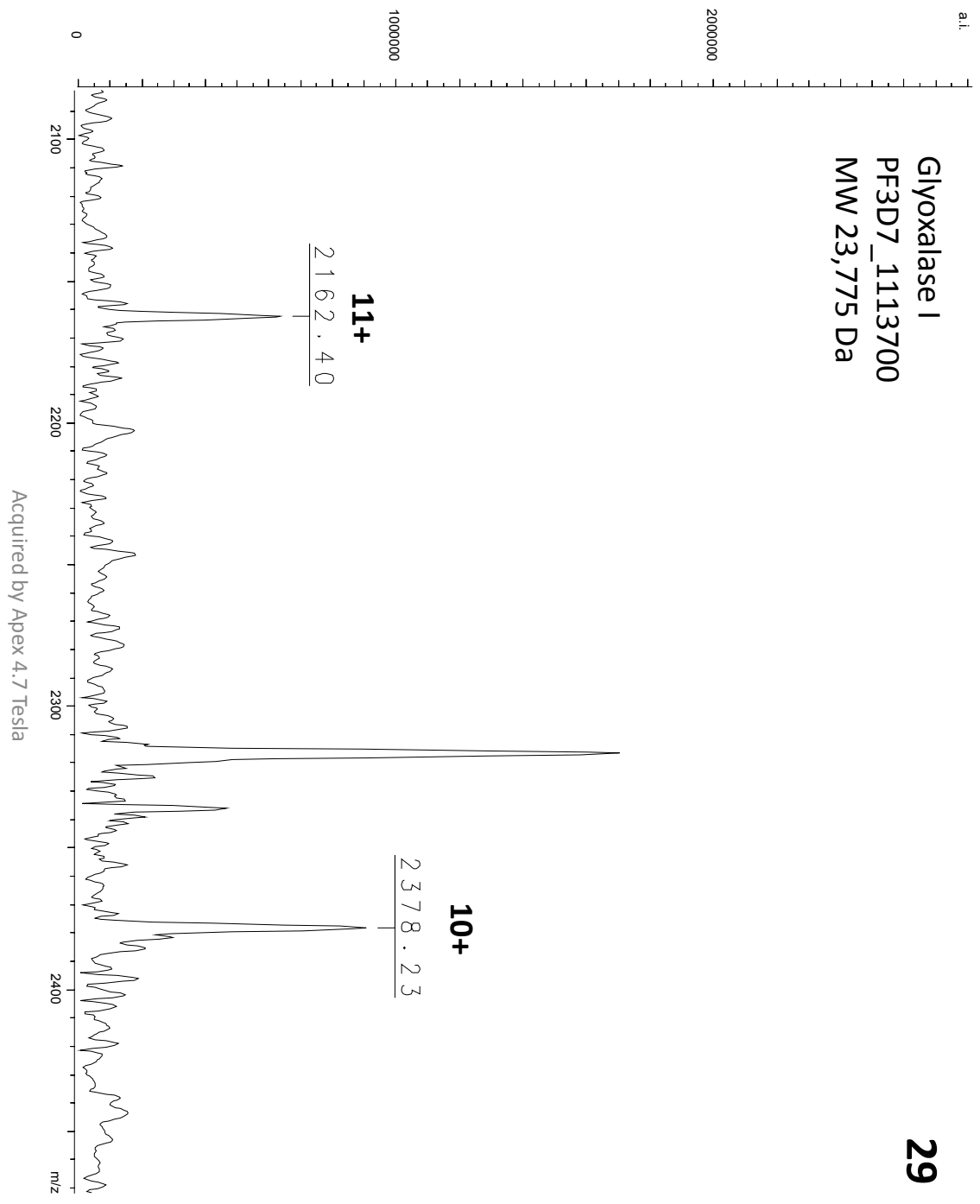


ATP-dependent Clp protease proteolytic subunit (ClpP)
PF3D7_0307400
23,552 Da



29

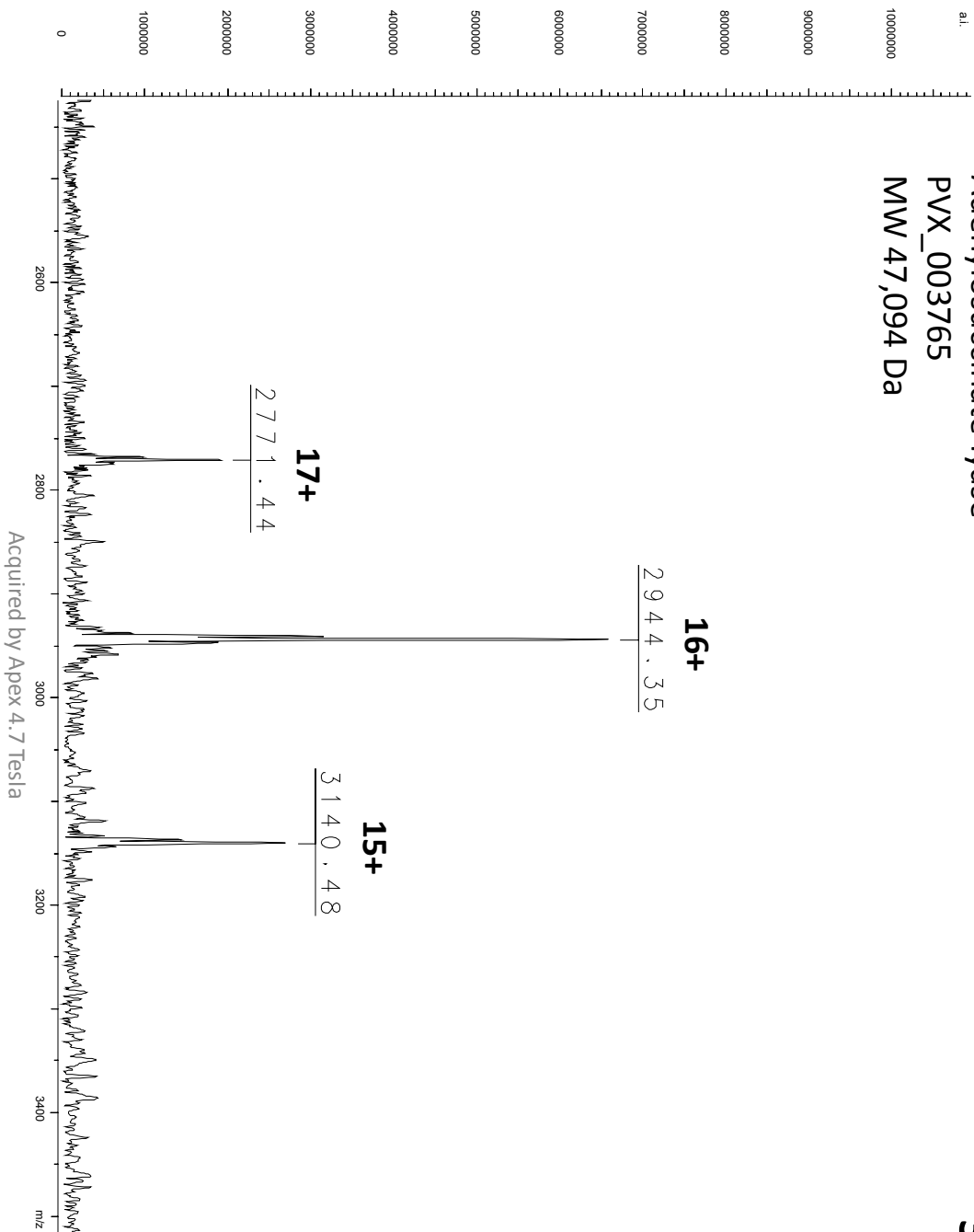
Glyoxalase I
PF3D7_1113700
MW 23,775 Da



a1.

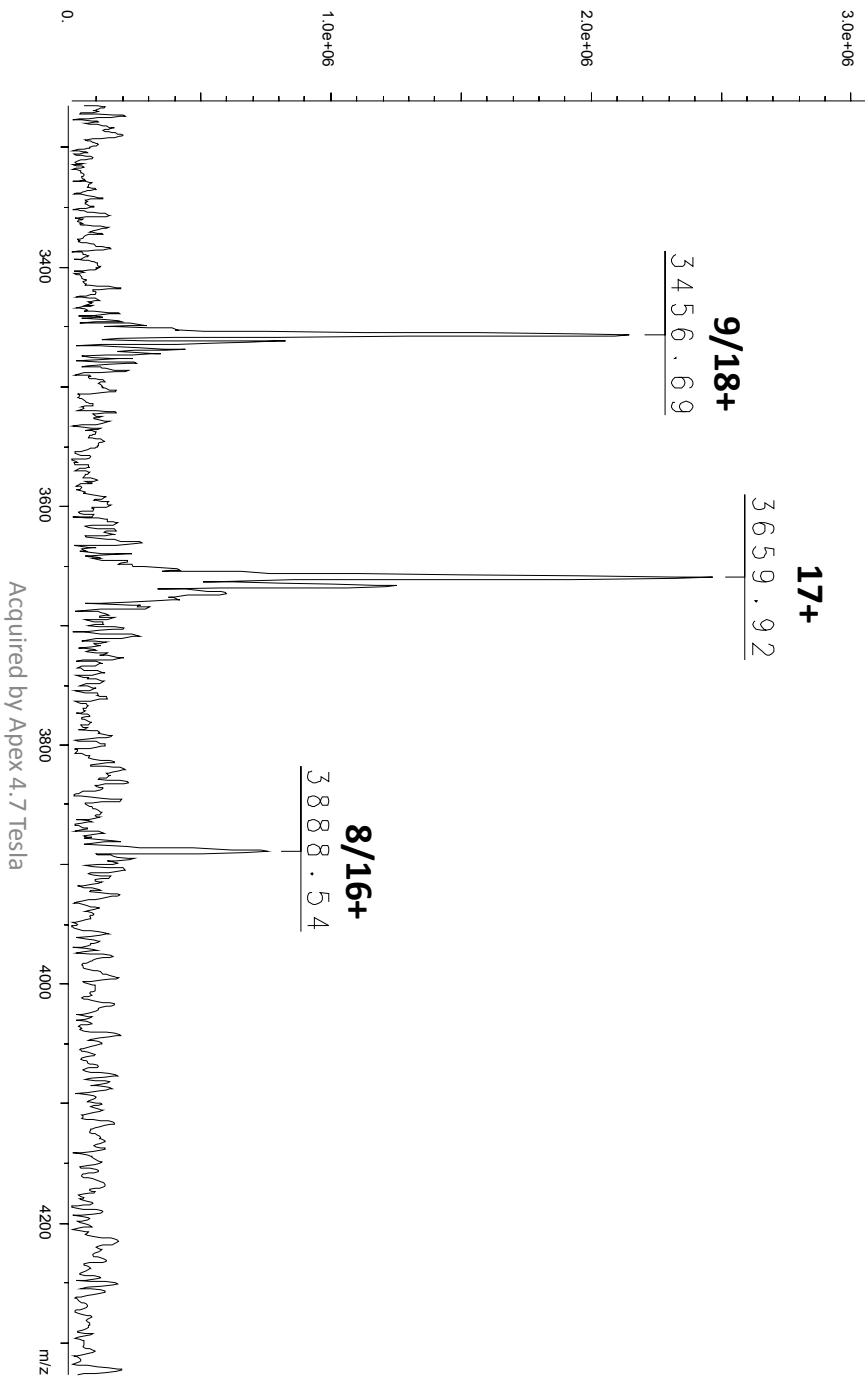
Adenylosuccinate lyase
PVX_003765
MW 47,094 Da

30

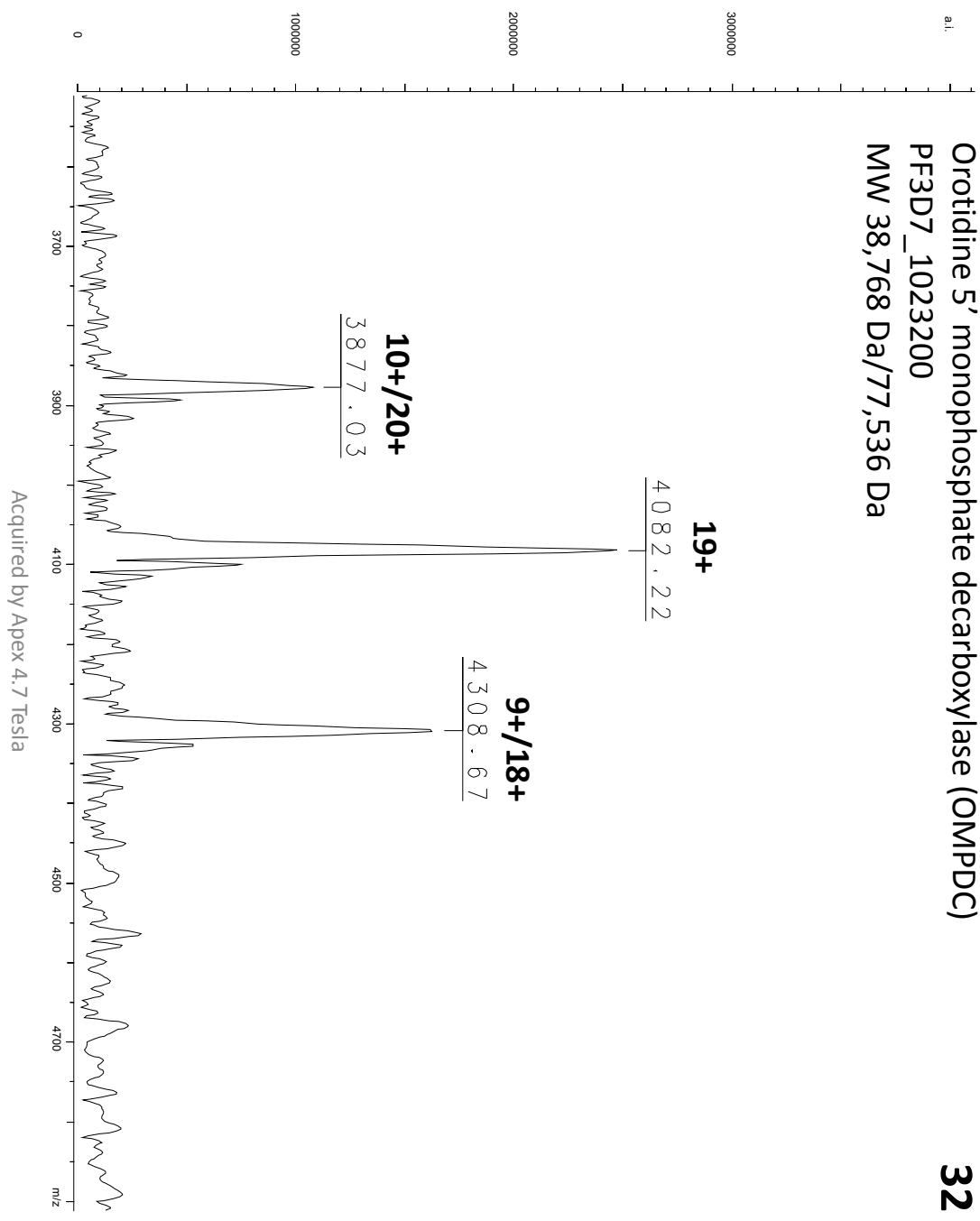


a.i.
Deoxyribose-phosphate aldolase
PY02252
MW 31,101 Da/62,201 Da

31



Orotidine 5' monophosphate decarboxylase (OMPDC) **32**
PF3D7_1023200
MW 38,768 Da/77,536 Da

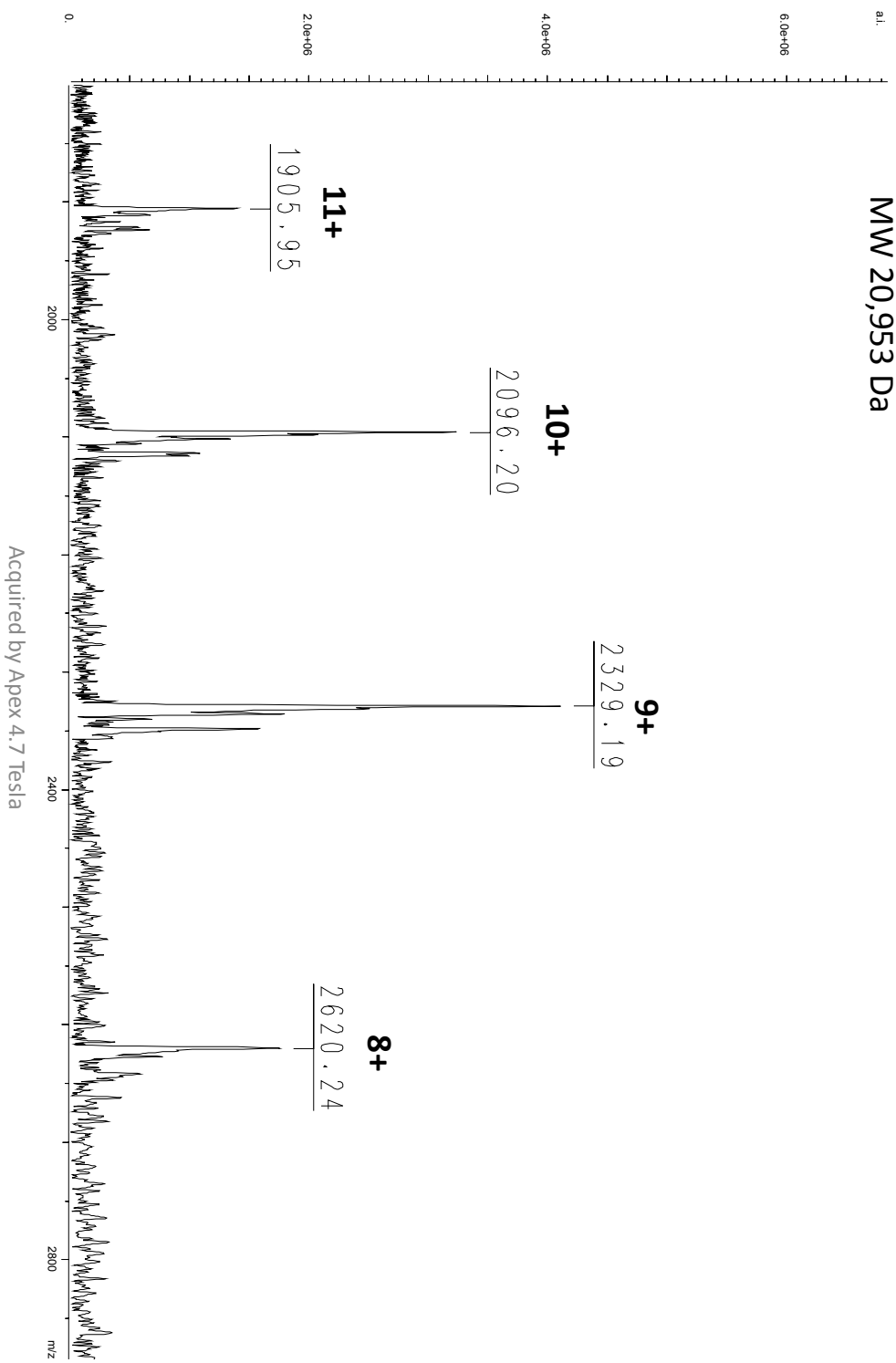


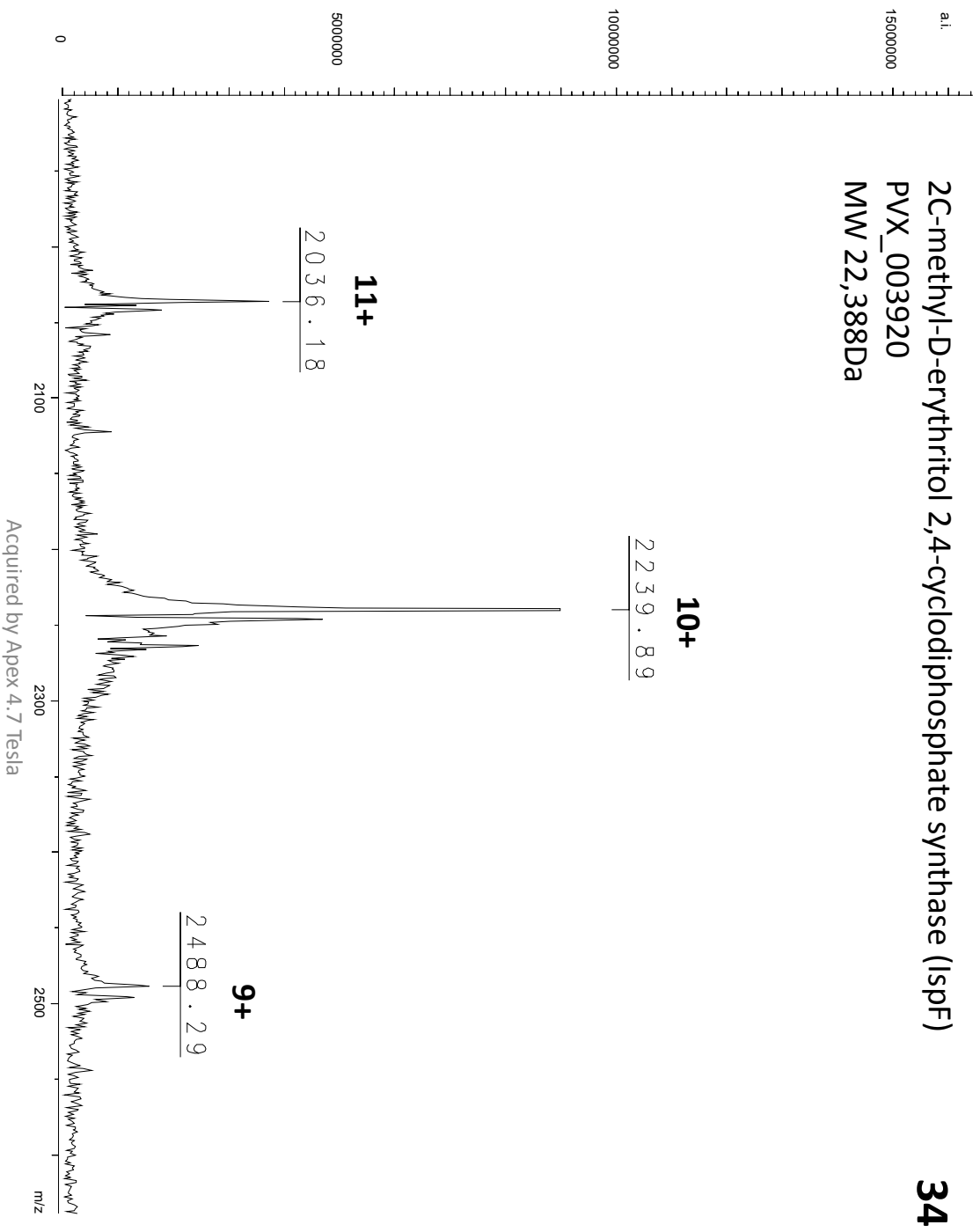
6-pyruvoyltetrahydropterin synthase

PF3D7_0628000

MW 20,953 Da

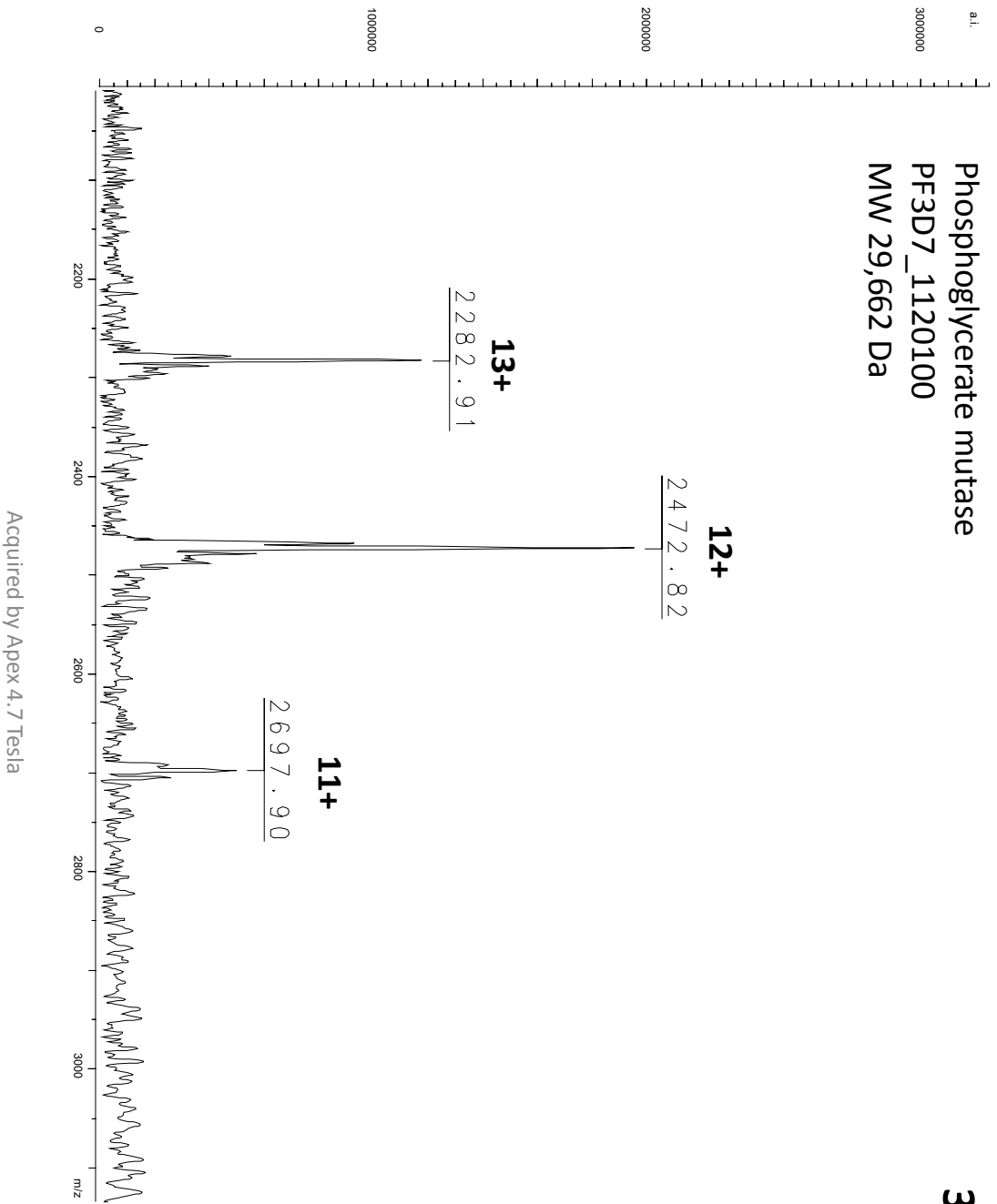
33



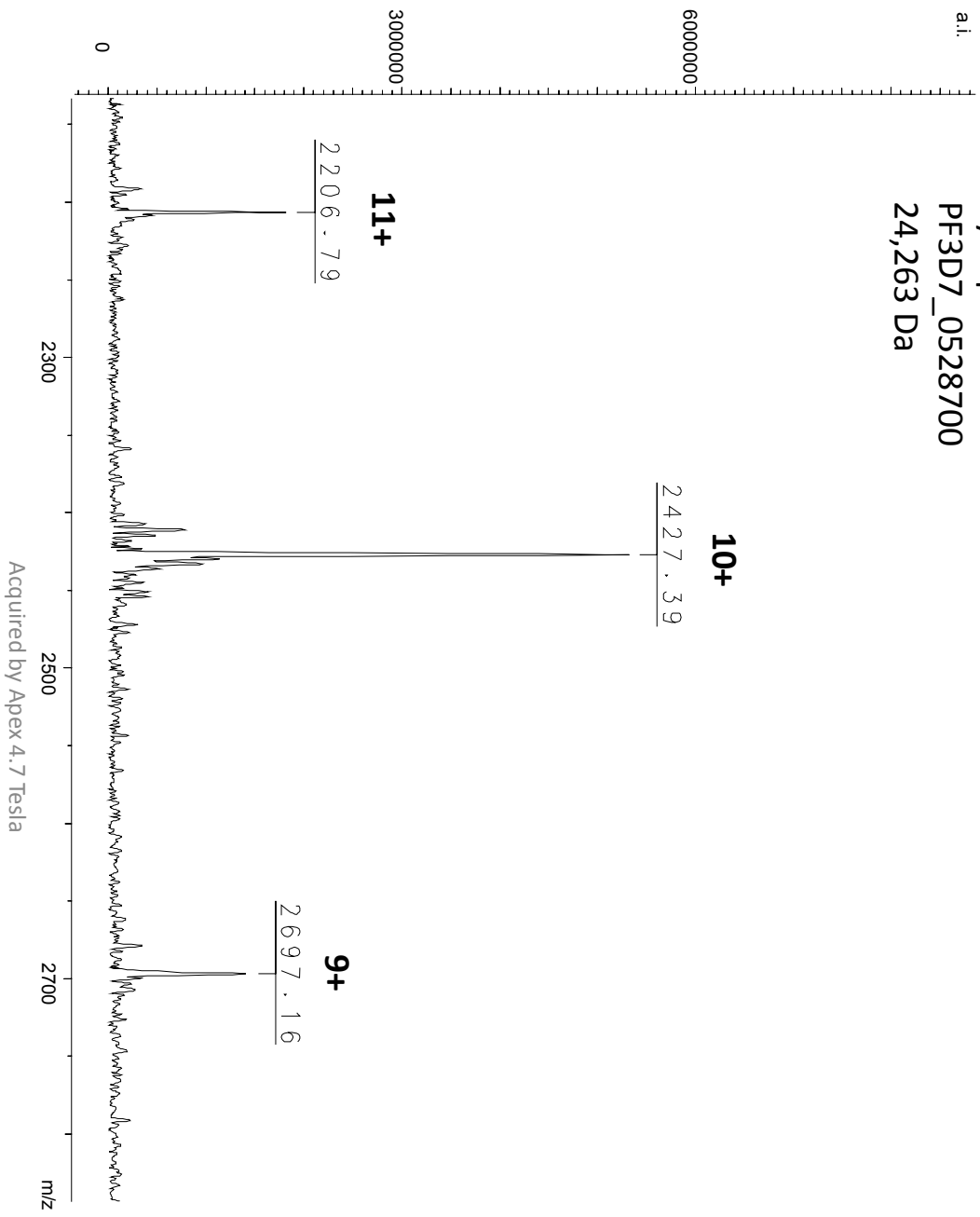


a.1.
Phosphoglycerate mutase
PF3D7_1120100
MW 29,662 Da

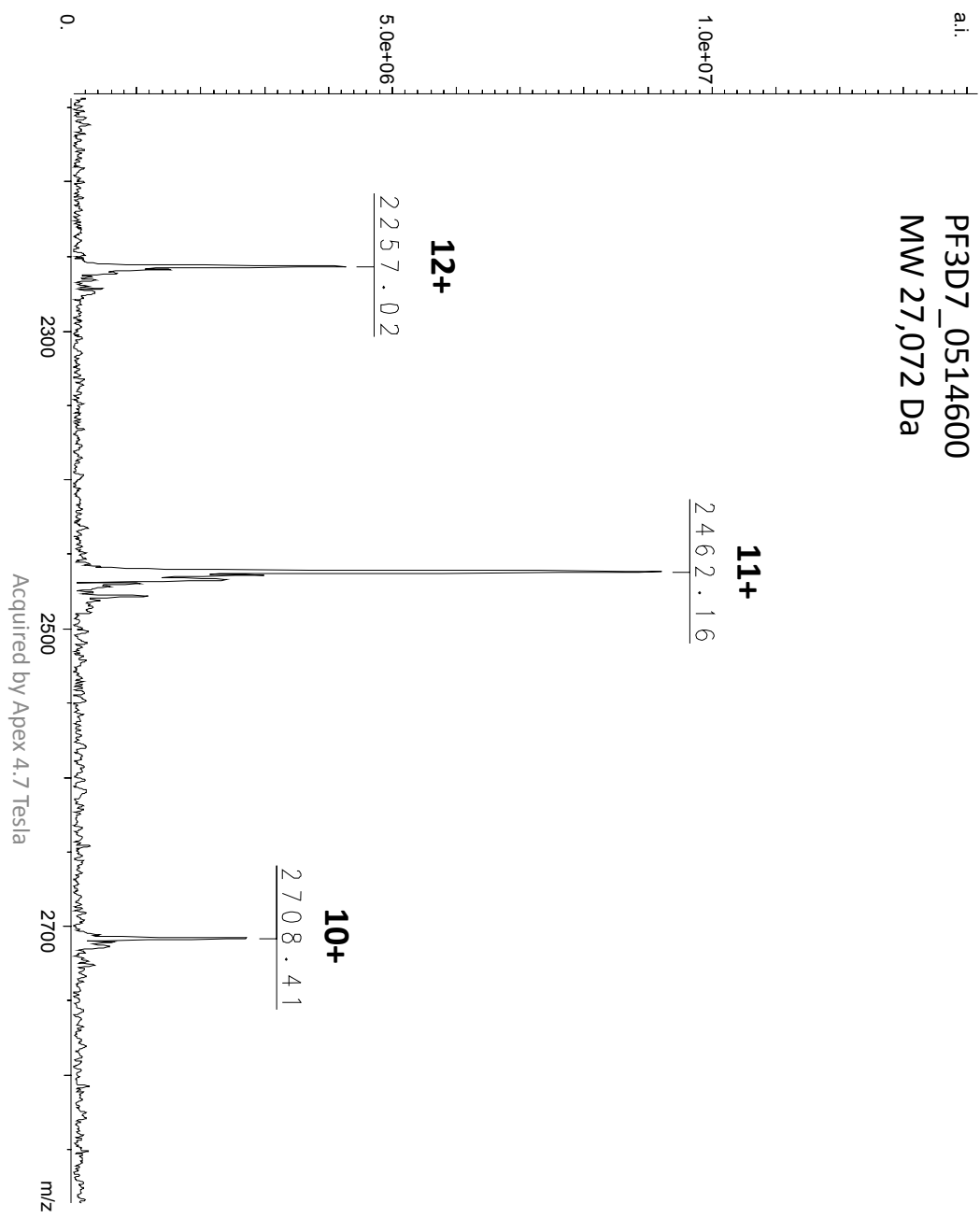
35



Cyclophilin
PF3D7_0528700
24,263 Da

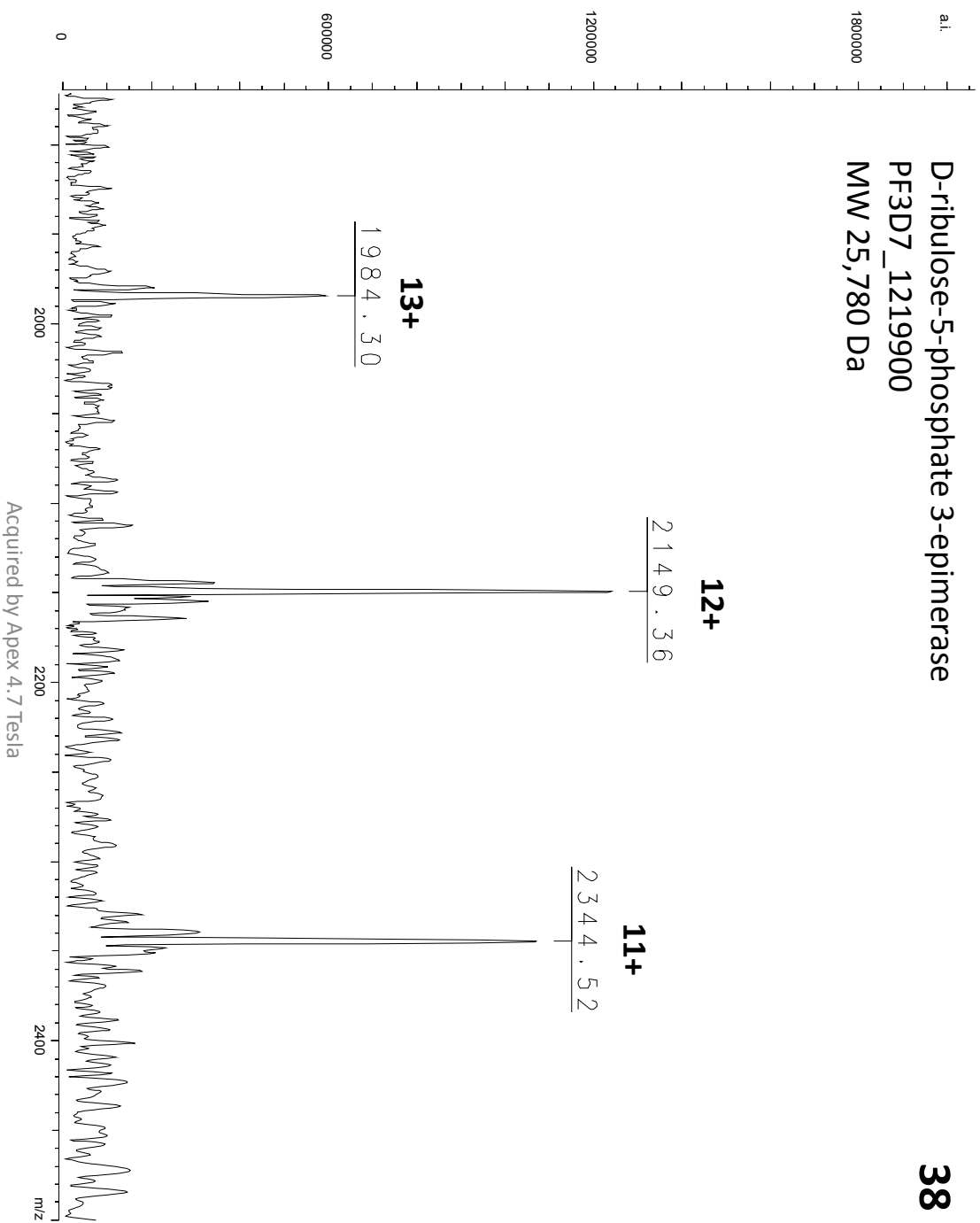


Ribose 5-phosphate isomerase
PF3D7_0514600
MW 27,072 Da



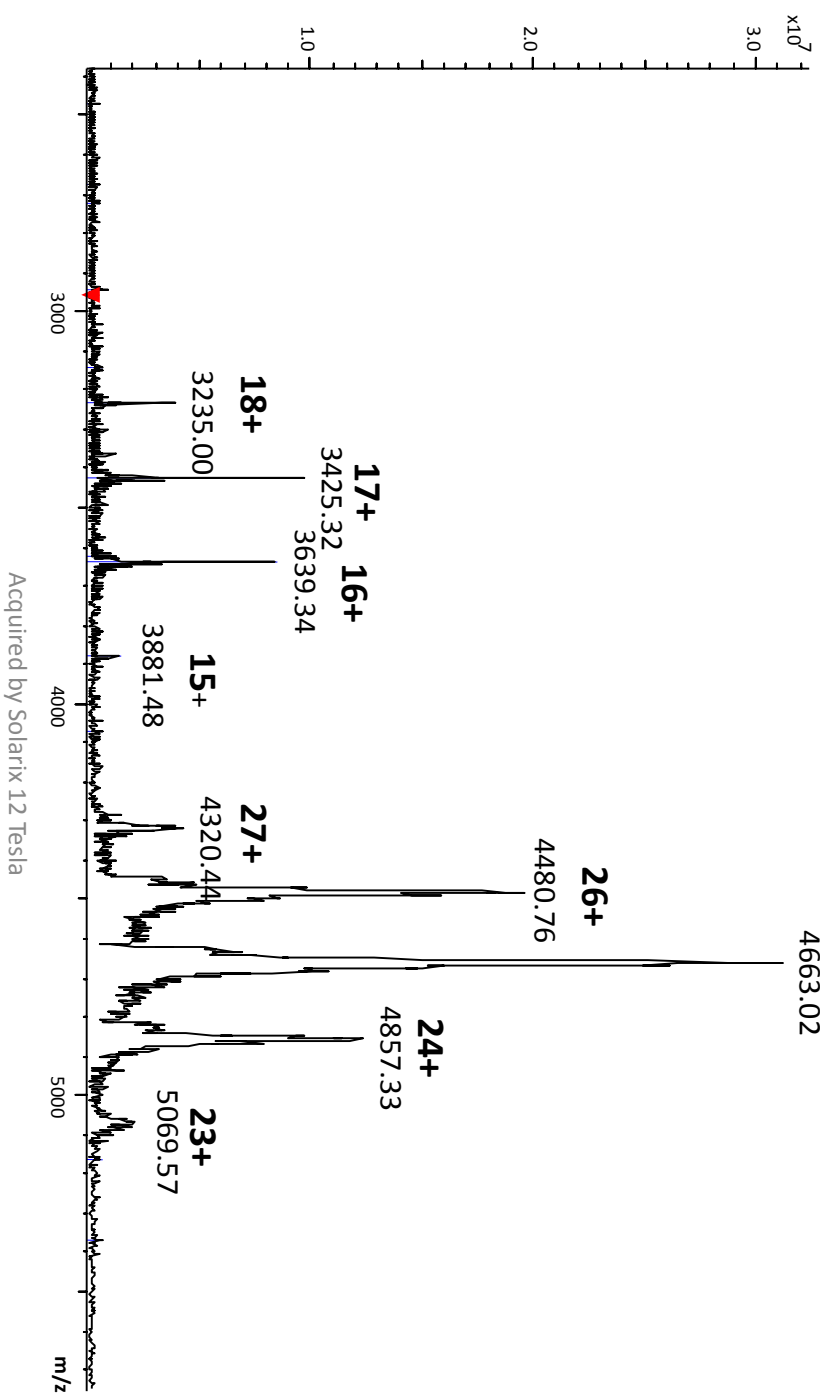
38

a.i.
D-ribulose-5-phosphate 3-epimerase
PF3D7_1219900
MW 25,780 Da



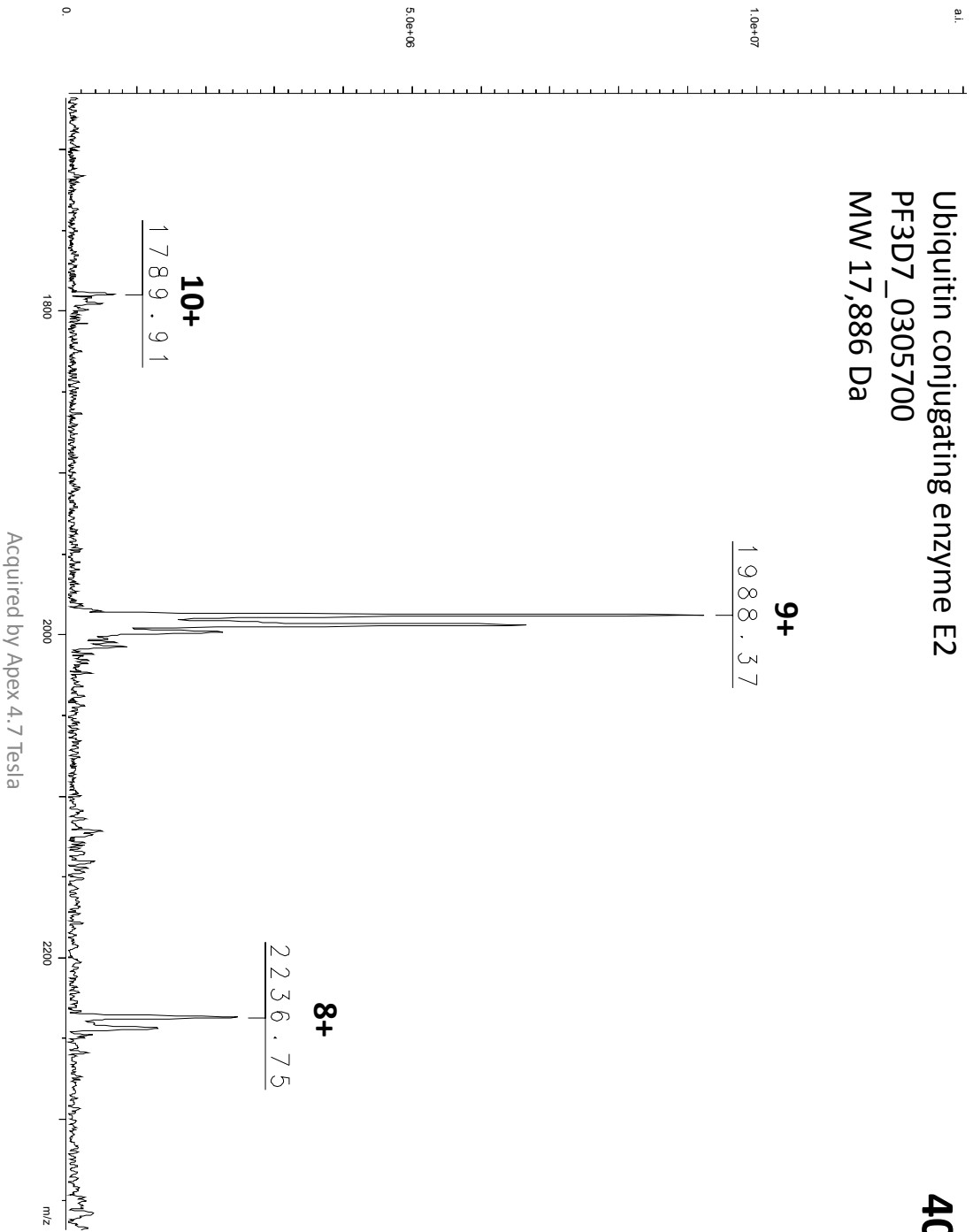
Prolyl- tRNase
PF3D7_0925300
MW 58,213 Da/ 116,426 Da (dimer)

39



Ubiquitin conjugating enzyme E2
PF3D7_0305700
MW 17,886 Da

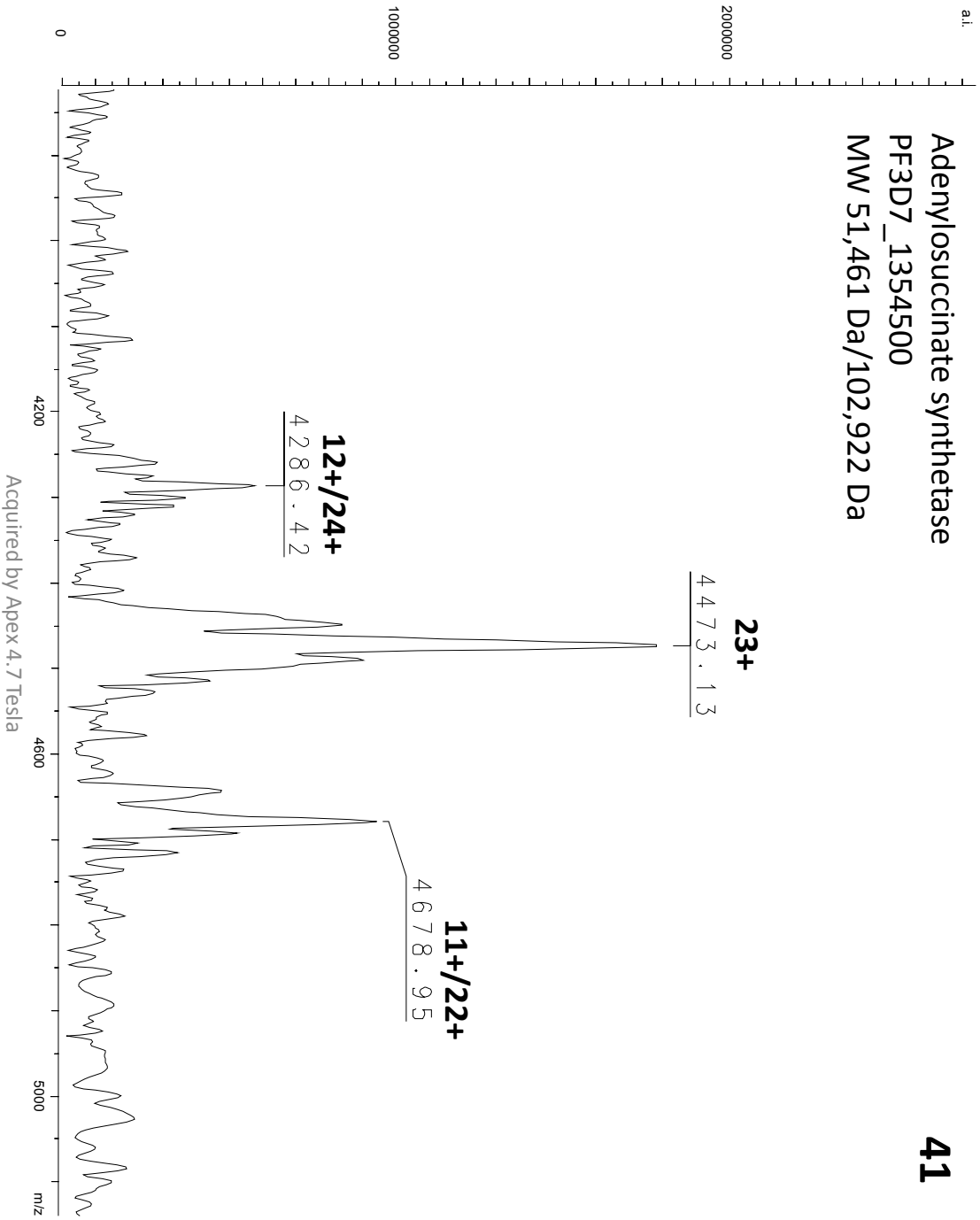
40



a.i.

Adenylosuccinate synthetase
PF3D7_1354500
MW 51,461 Da/102,922 Da

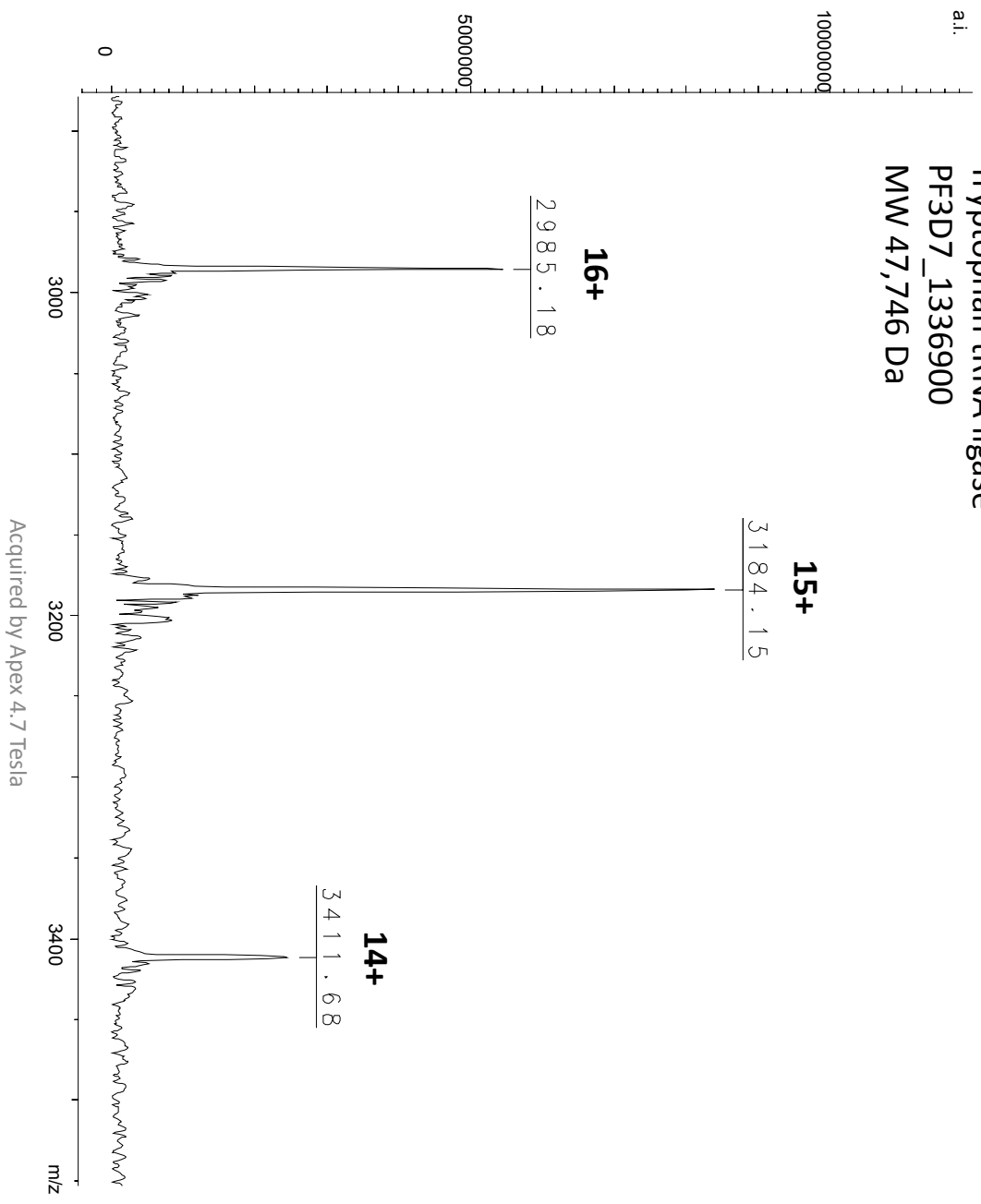
41



a.i.

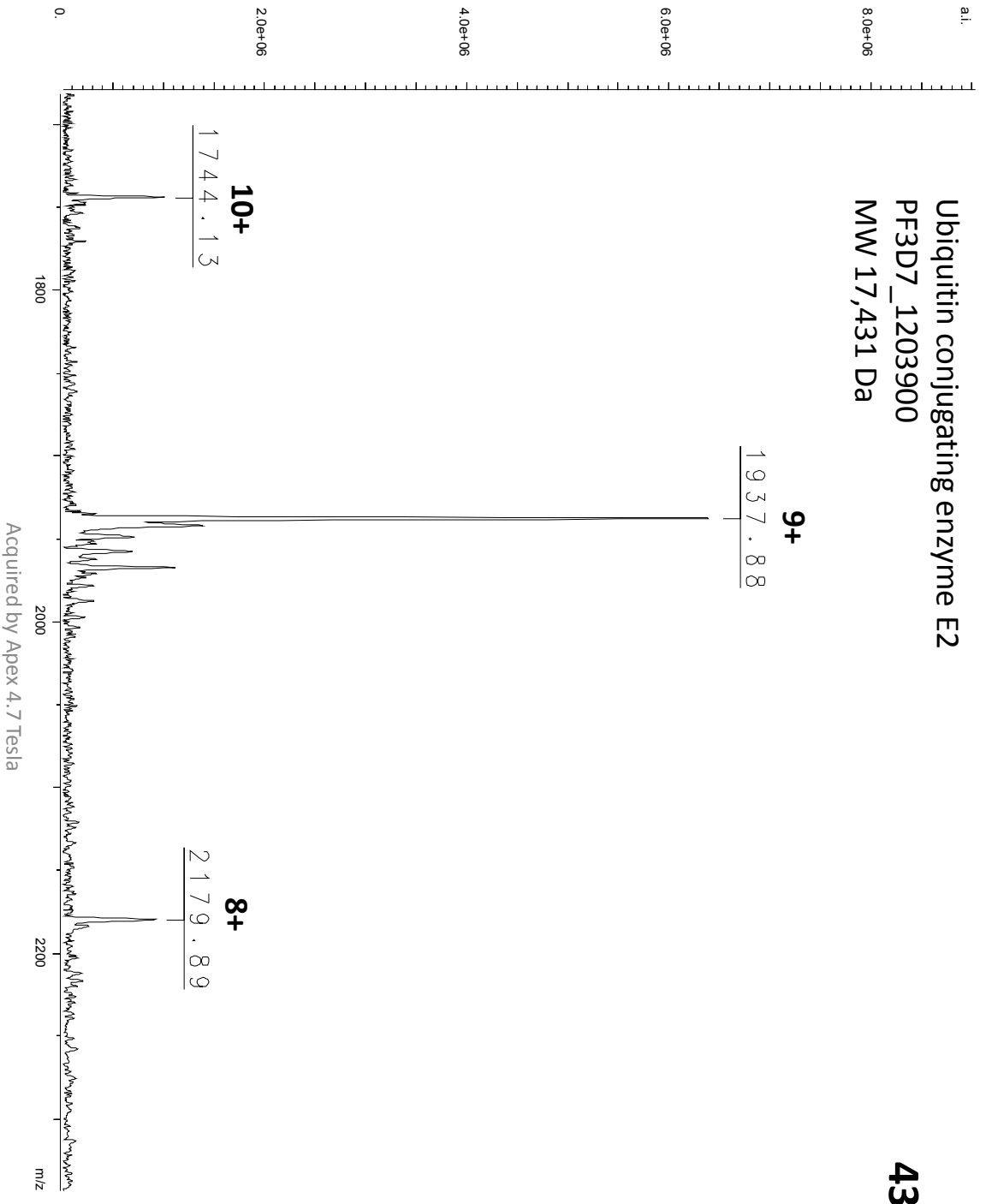
Tryptophan tRNA ligase
PF3D7_1336900
MW 47,746 Da

42



43

Ubiquitin conjugating enzyme E2
PF3D7_1203900
MW 17,431 Da



a.1
8.0e+06

Ubiquitin conjugating enzyme E2N
PF3D7_0527100
MW 19,176 Da

44

6.0e+06

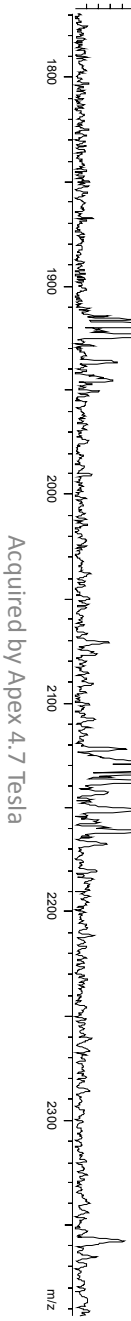
9+
2131.76

4.0e+06

10+
1918.68

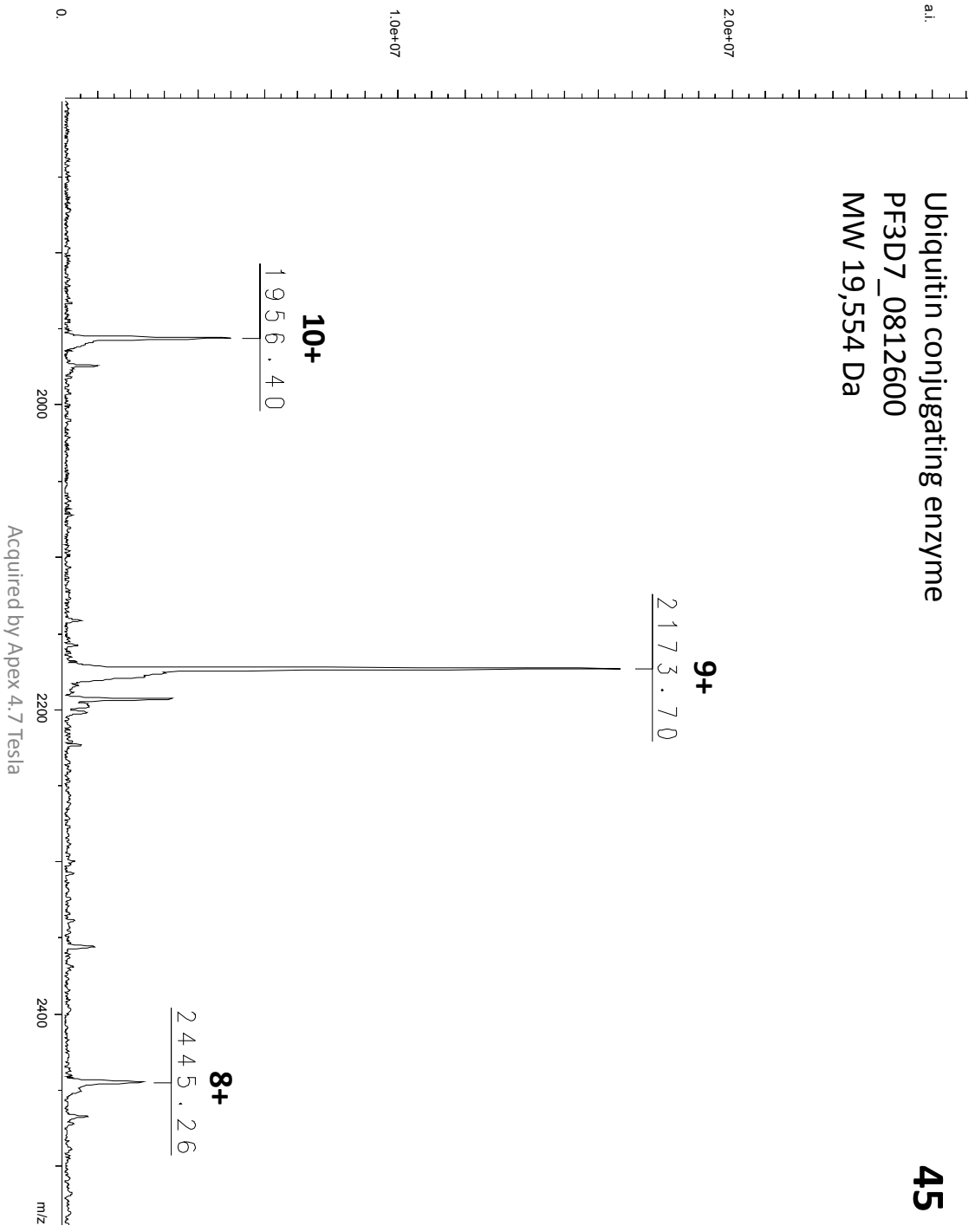
2.0e+06

0



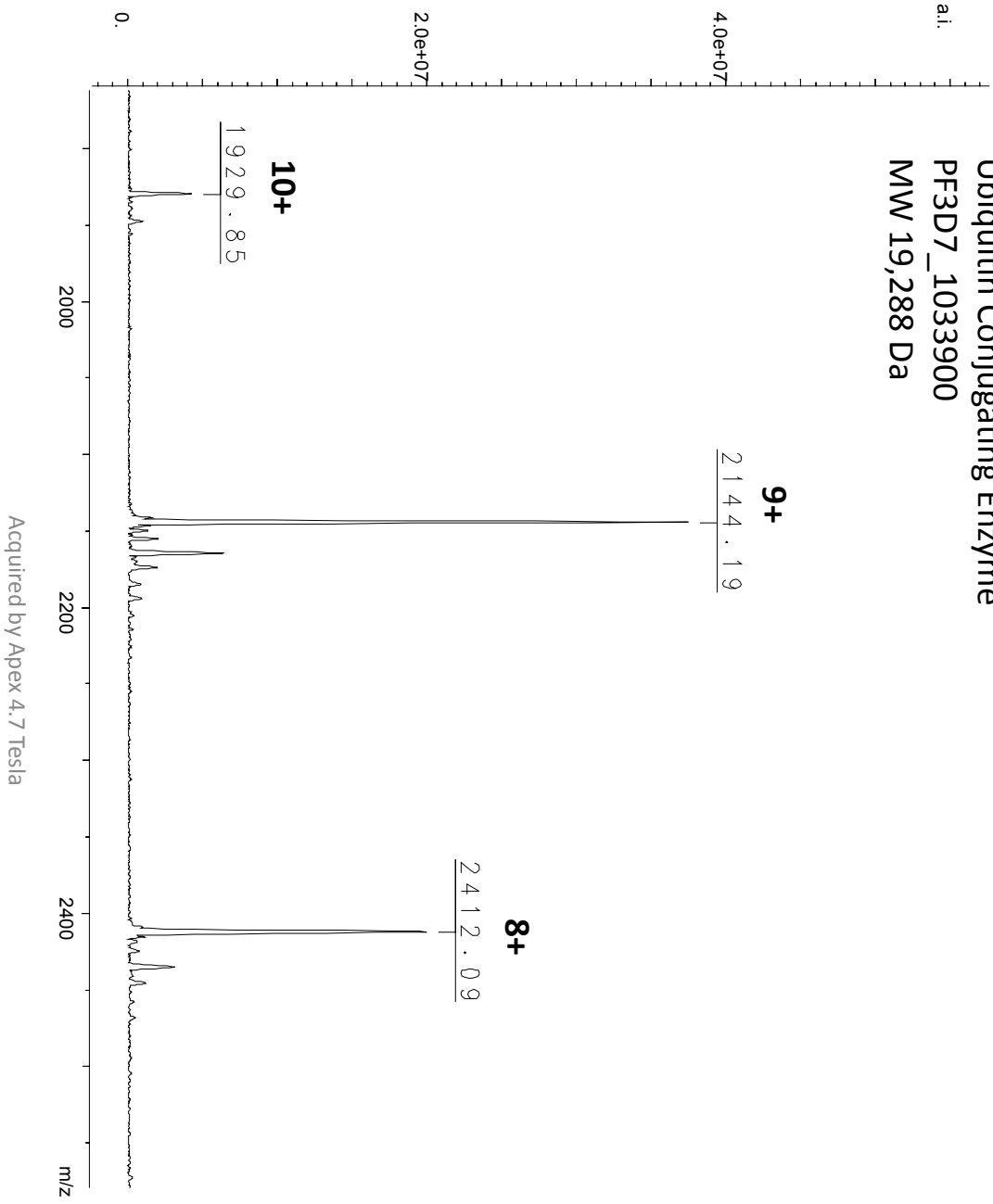
45

Ubiquitin conjugating enzyme
PF3D7_0812600
MW 19,554 Da



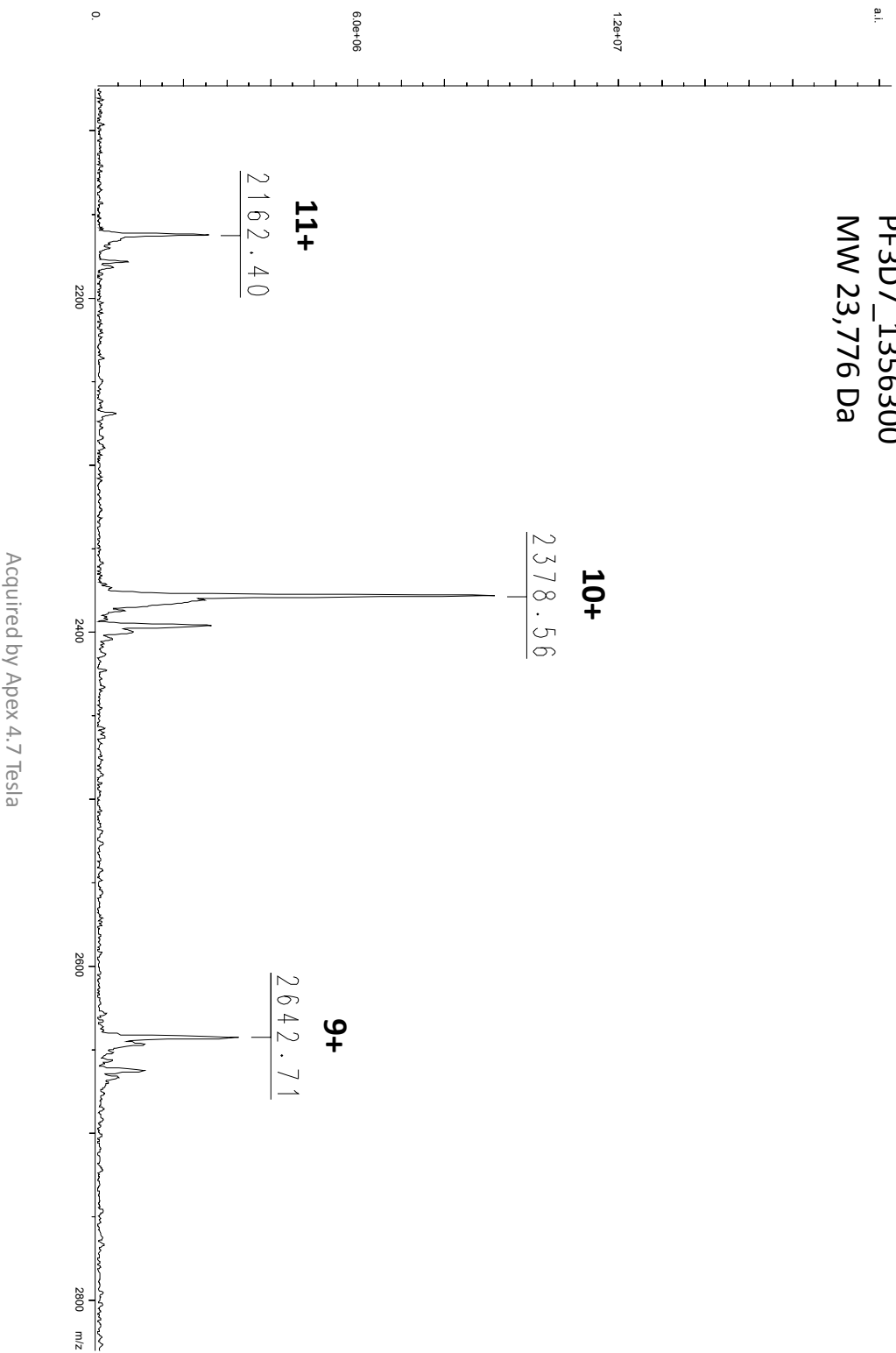
Ubiquitin Conjugating Enzyme
PF3D7_1033900
MW 19,288 Da

46



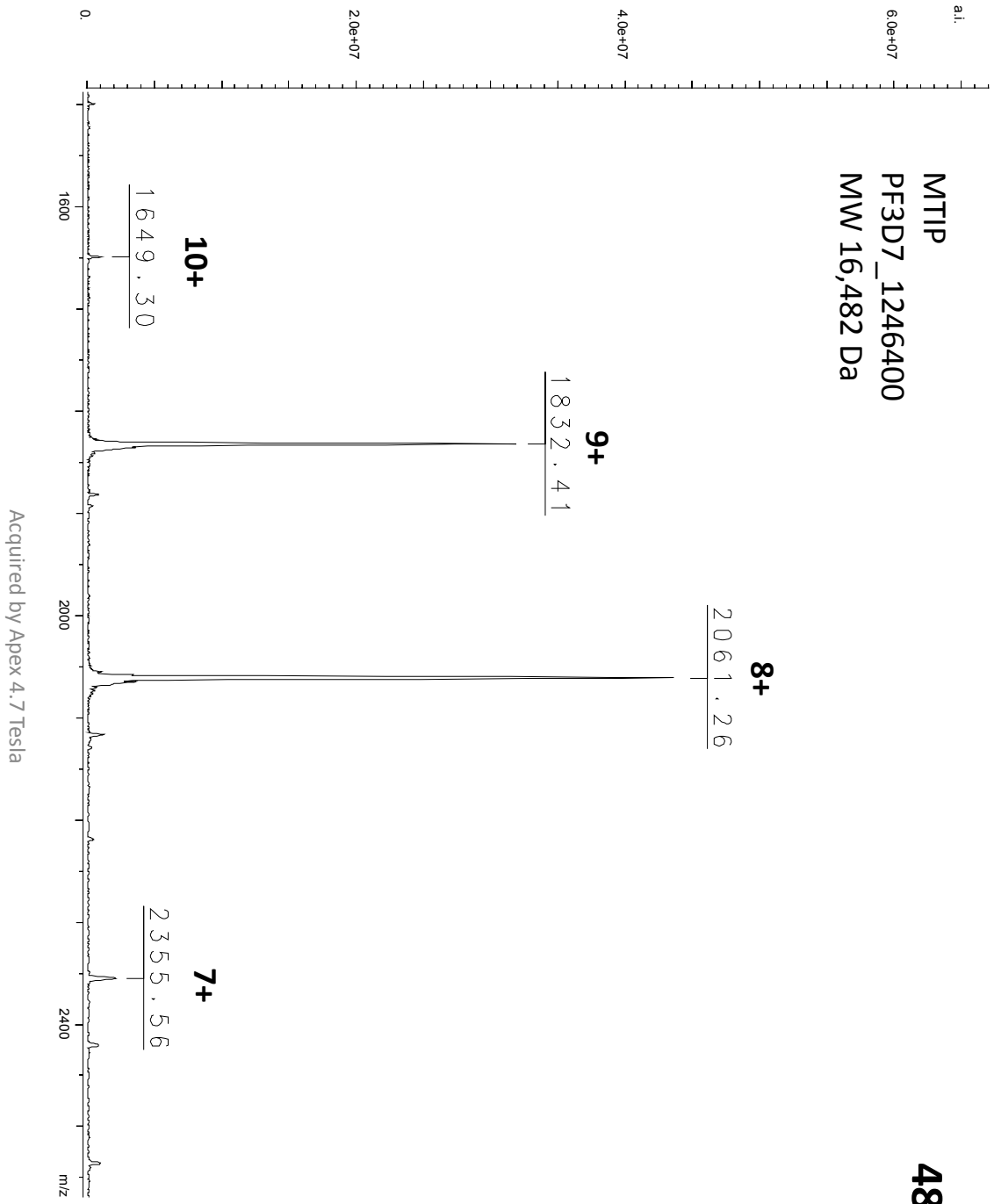
47

Ubiquitin conjugating enzyme
PF3D7_1356300
MW 23,776 Da



48

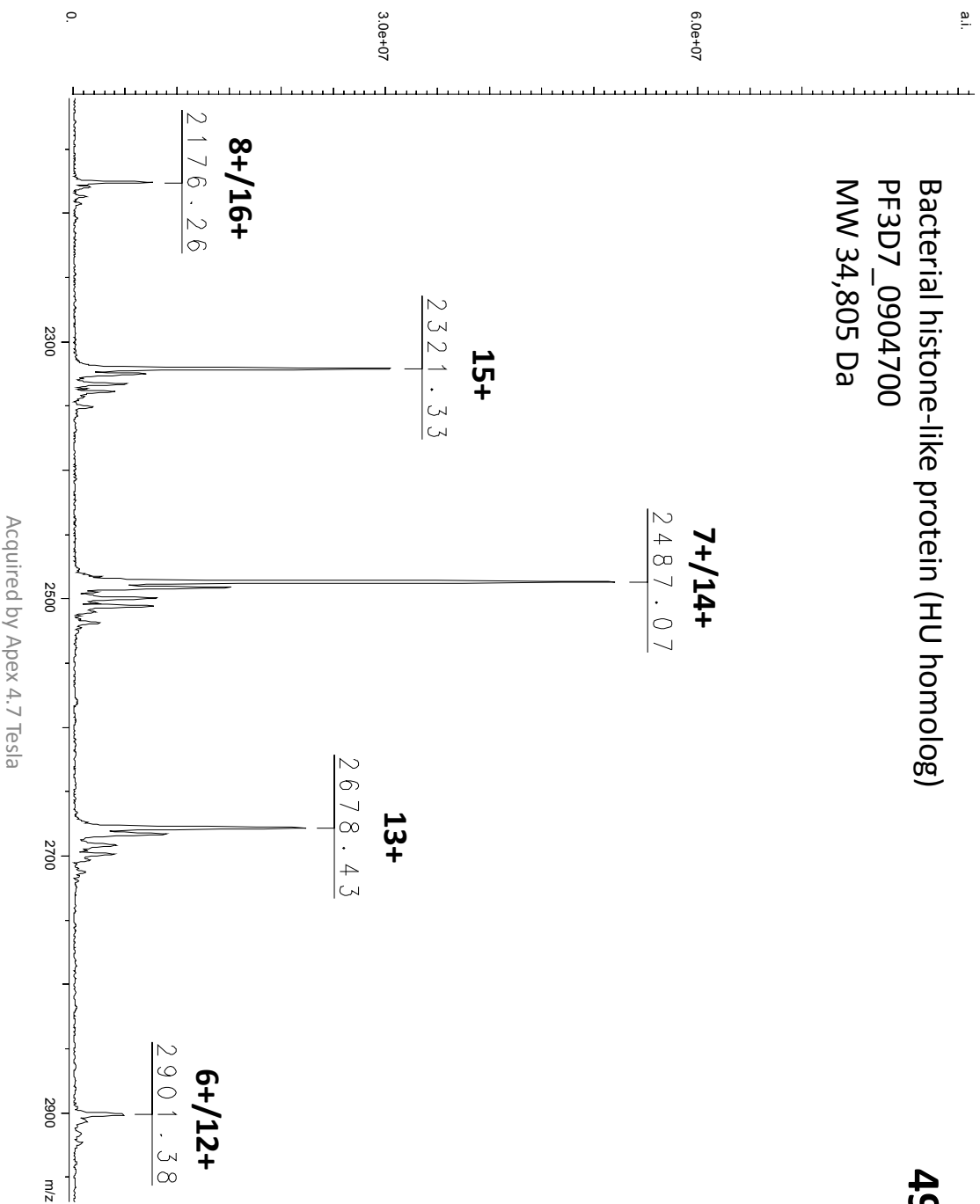
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PF3D7_1246400
MW 16,482 Da



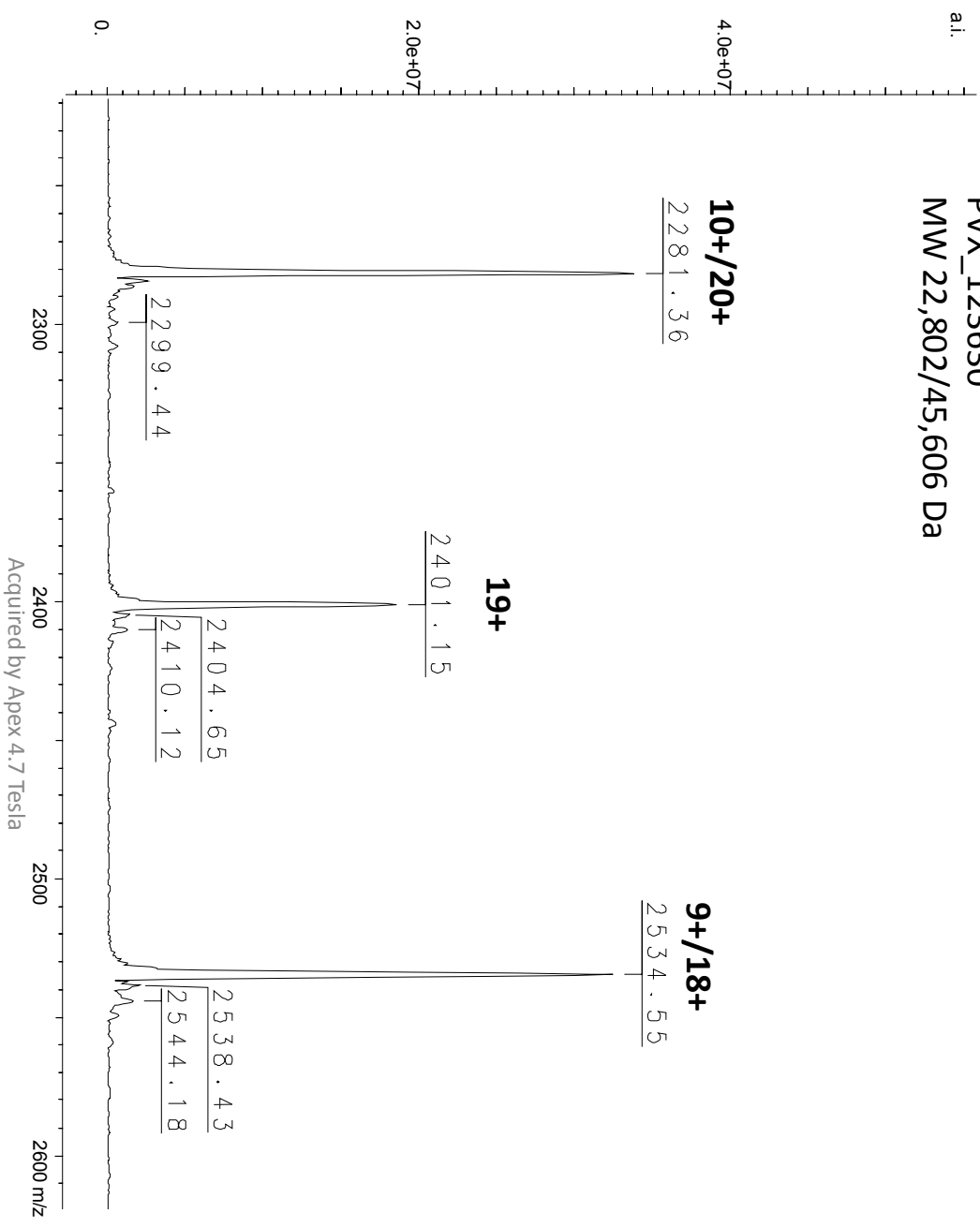
a.i.

Bacterial histone-like protein (HU homolog)
PF3D7_0904700
MW 34,805 Da

49



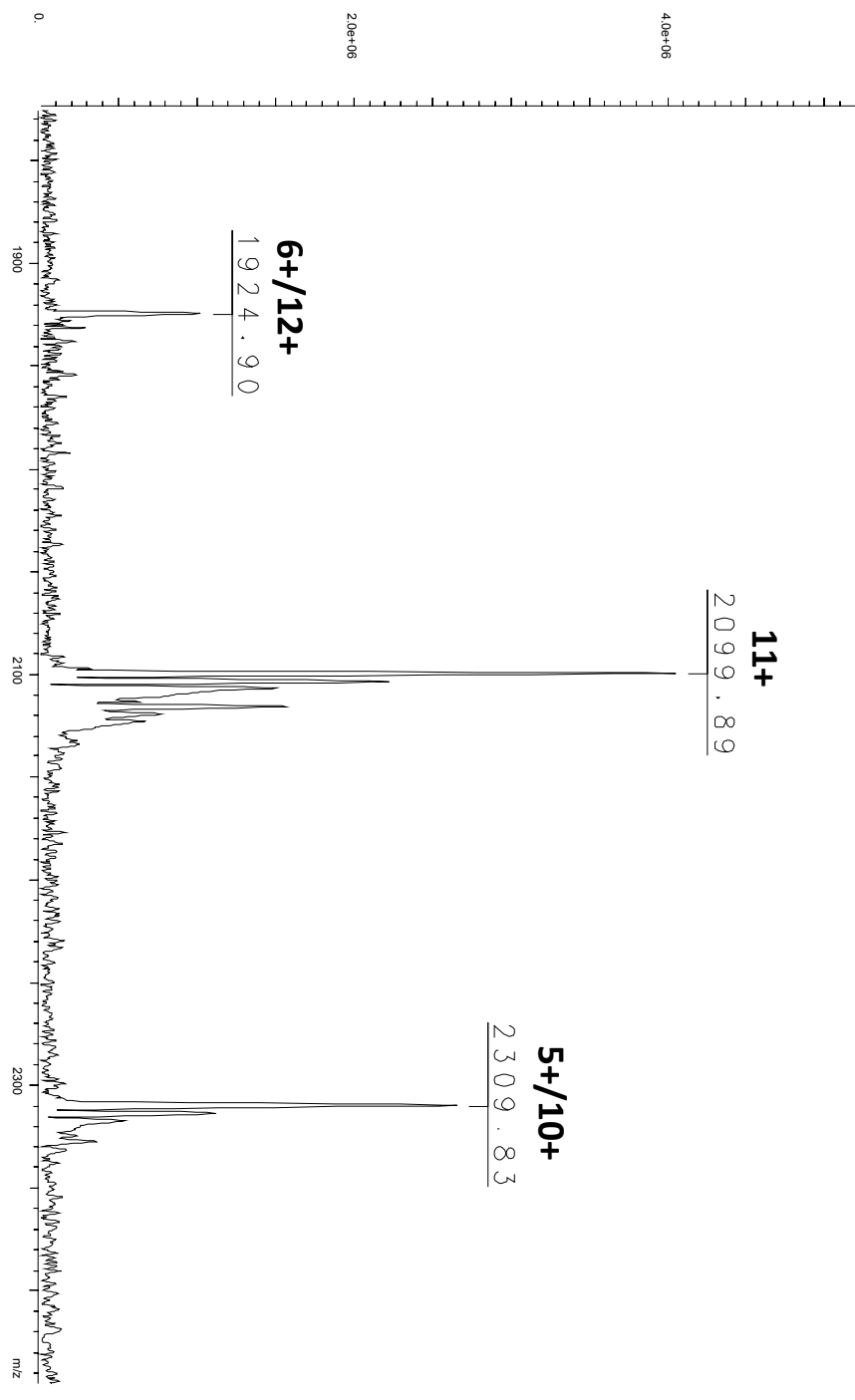
Phosphatidylethanolamine-binding protein
PVX_123630
MW 22,802/45,606 Da



a.i.
6.0e+06

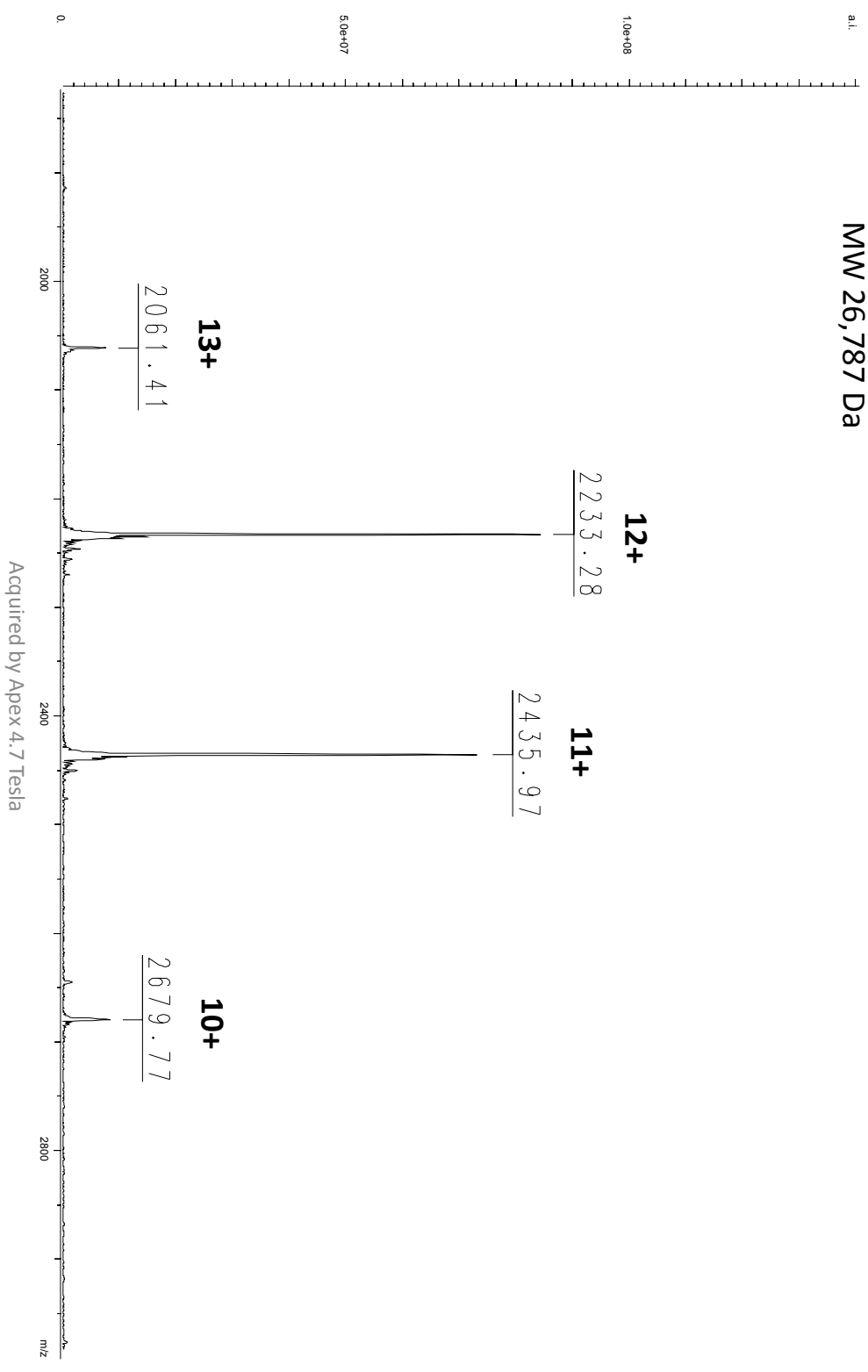
Dynein light chain 1
PF3D7_1213600
MW 11,544 Da/23,088 Da

51



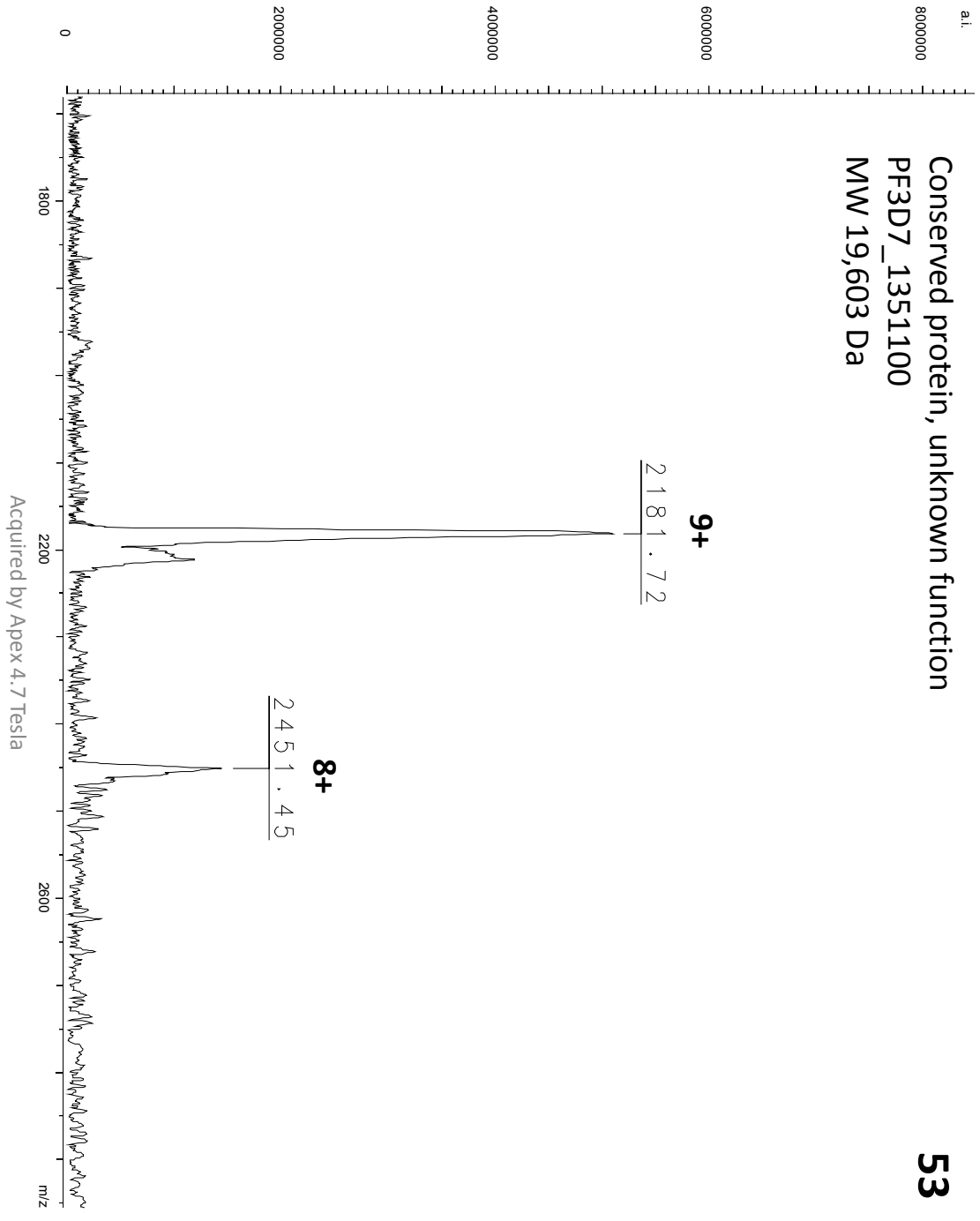
Circumsporozoite surface protein (CSP)
PF3D7_0304600
MW 26,787 Da

52



a.i.
Conserved protein, unknown function
PF3D7_1351100
MW 19,603 Da

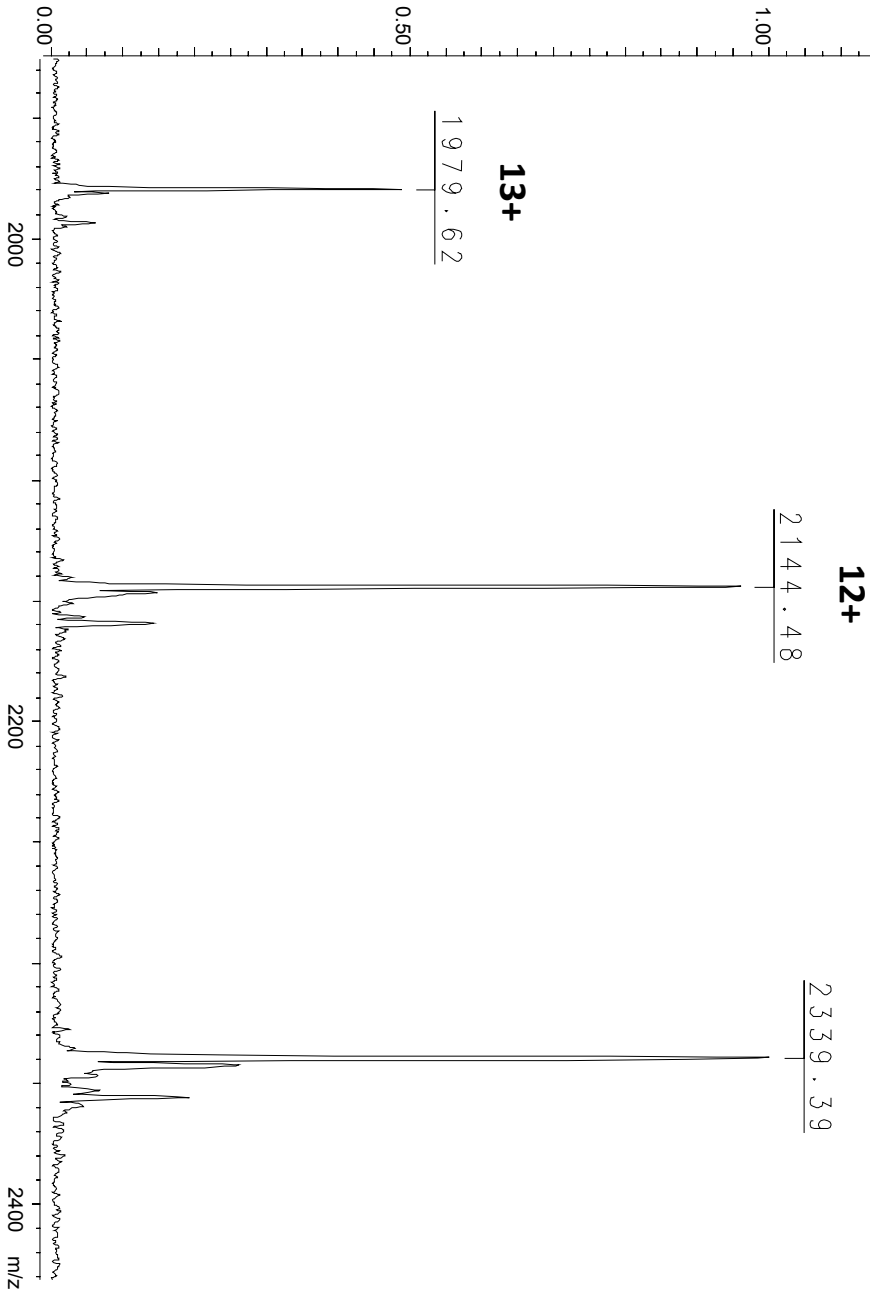
53



ri.

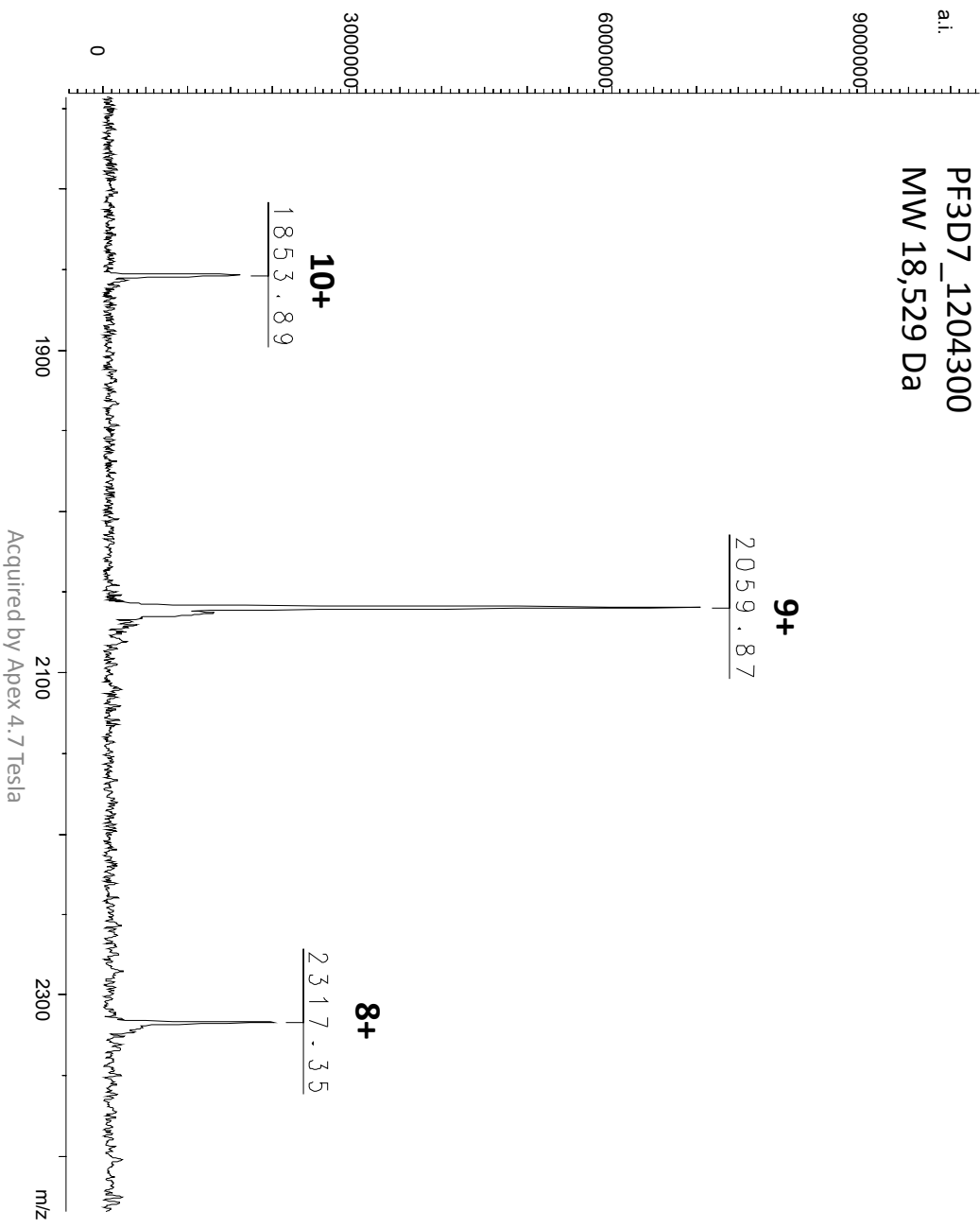
Vacuolar protein-sorting protein (VPS2)
PF3D7_0816200
MW 25,722 Da

54



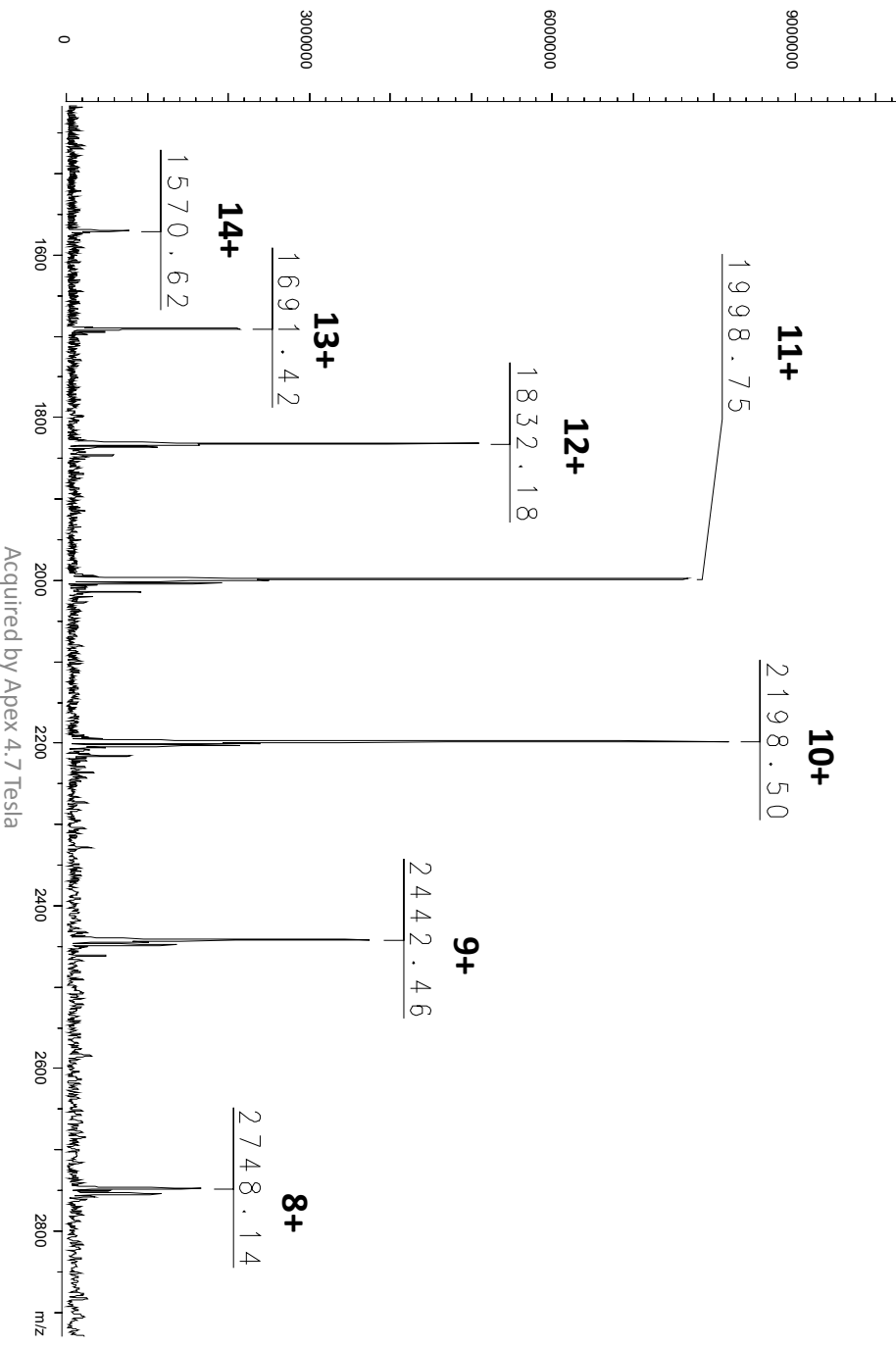
Acquired by Apex 4.7 Tesla

Eukaryotic translation initiation factor 5A (eIF5a)
PF3D7_1204300
MW 18,529 Da



a.i.
6-pyruvoyltetrahydropterin synthase
PvX_114505
MW 21,975 Da

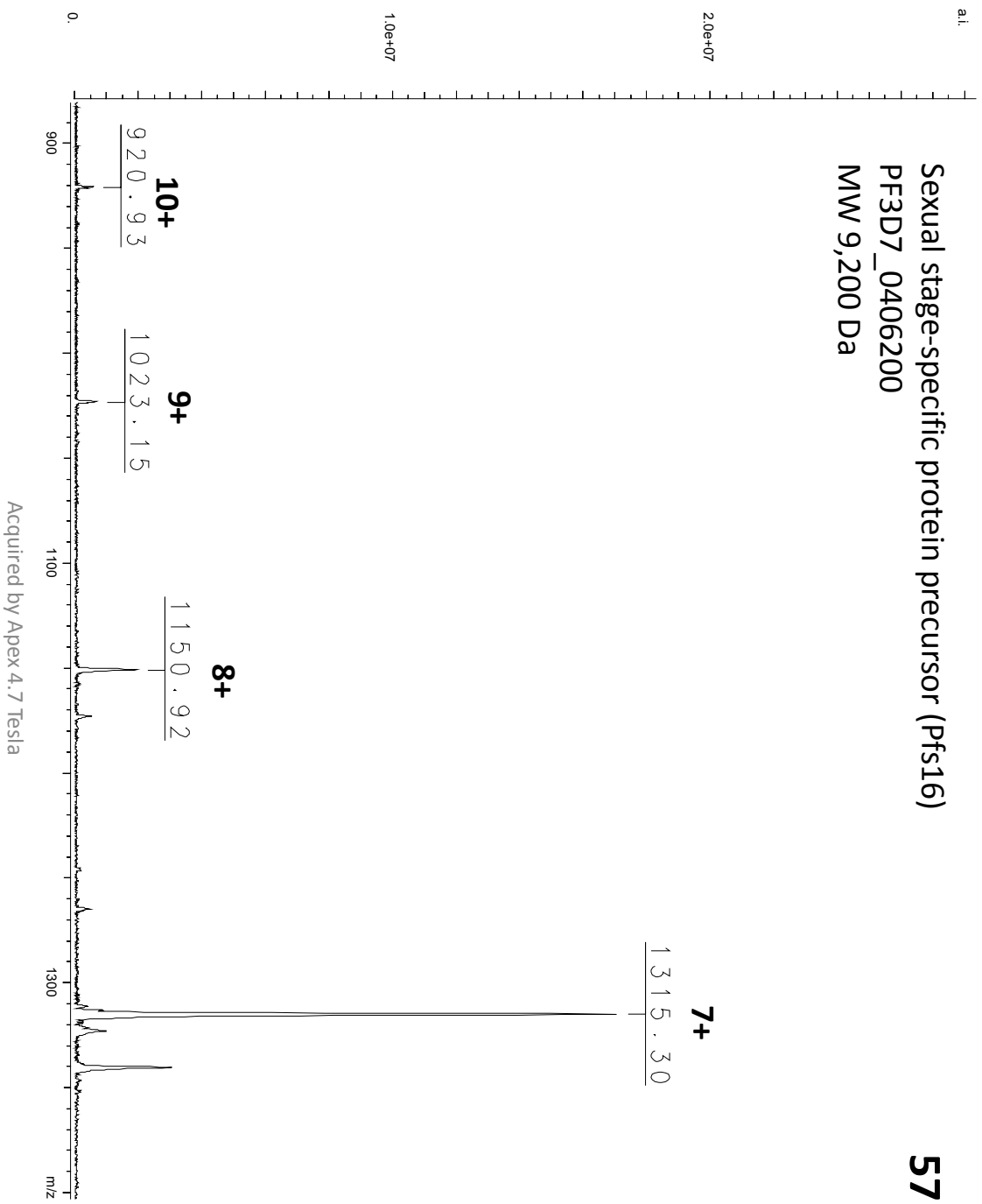
56



a.i.

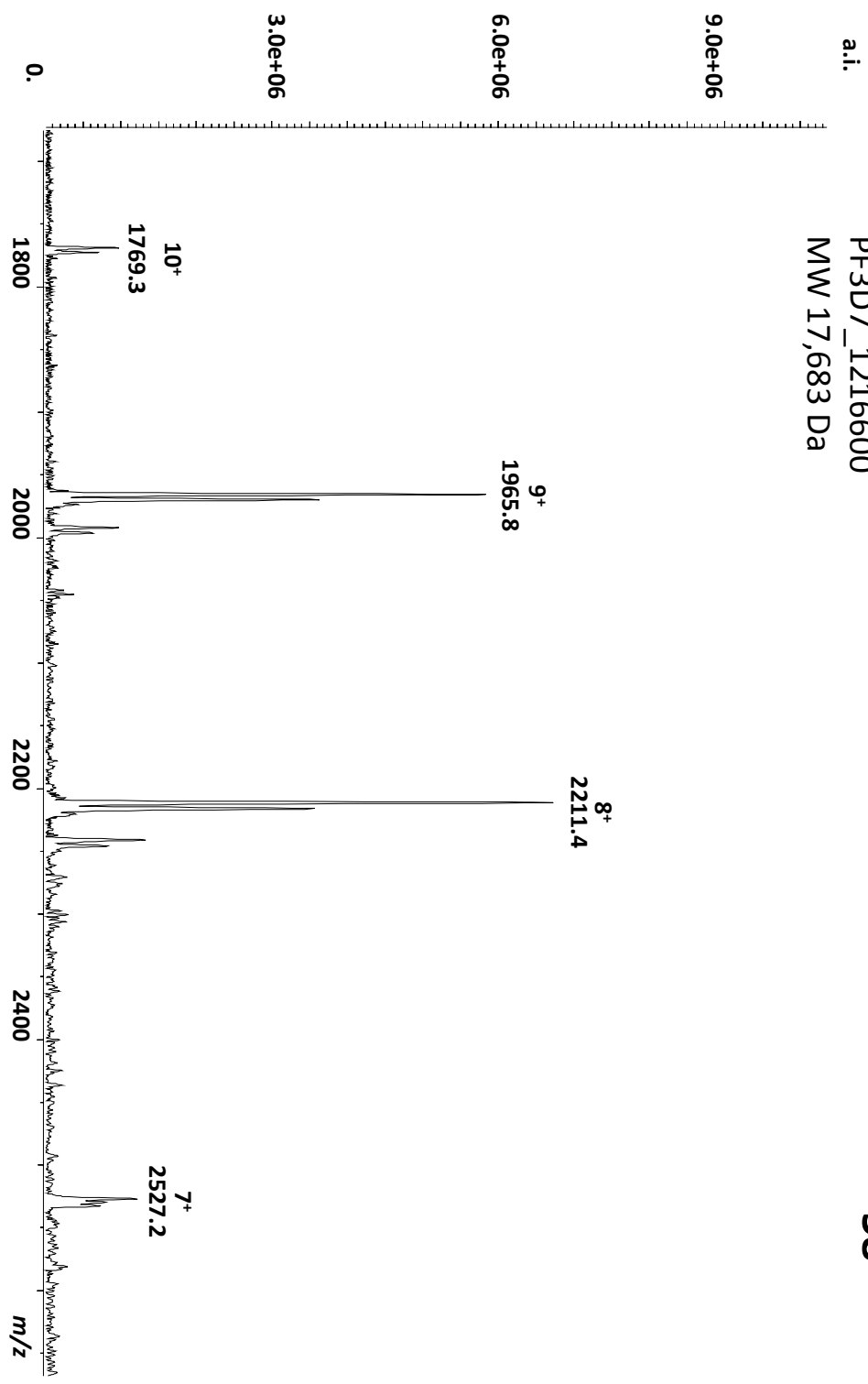
Sexual stage-specific protein precursor (Pfs16)
PF3D7_0406200
MW 9,200 Da

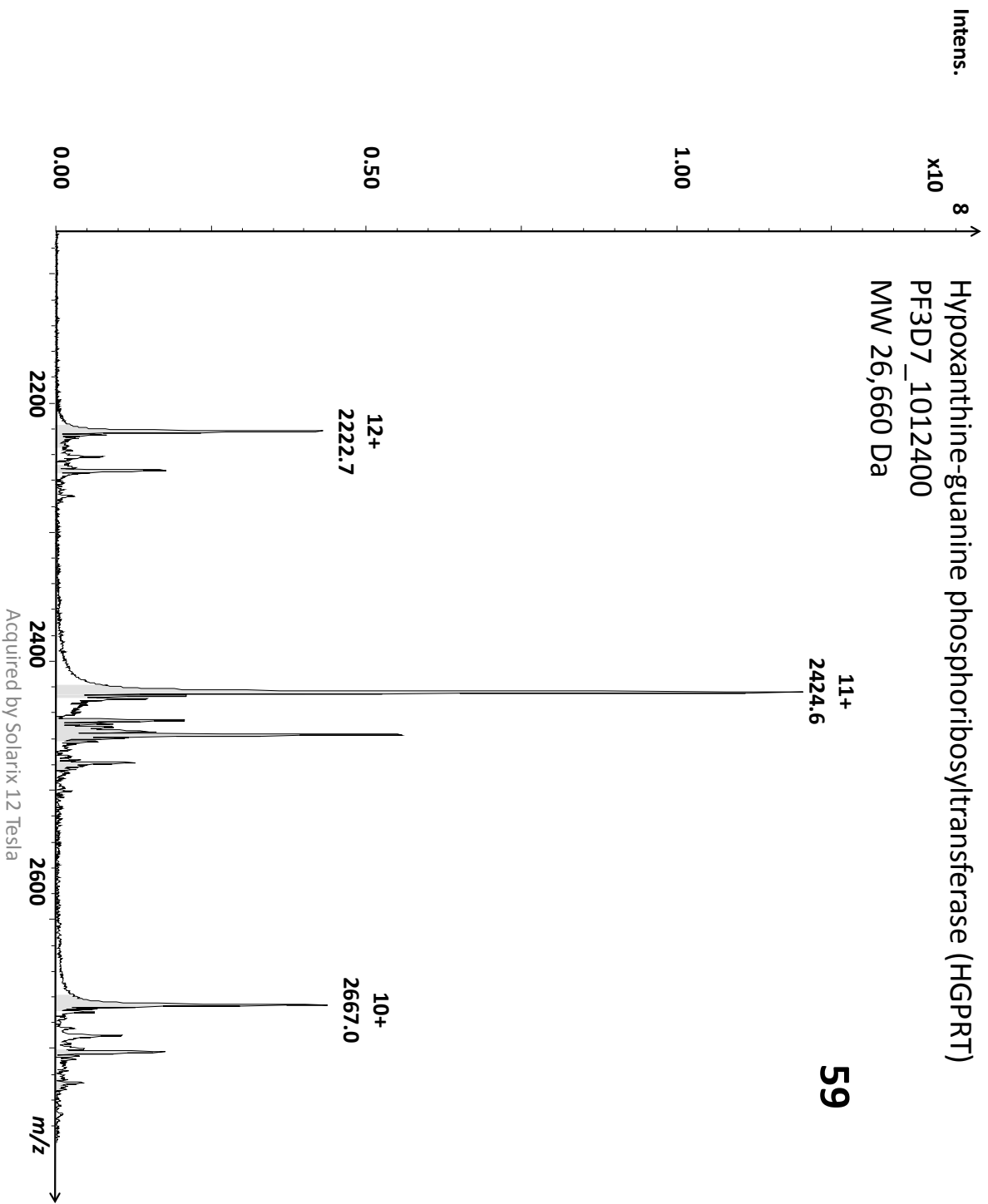
57

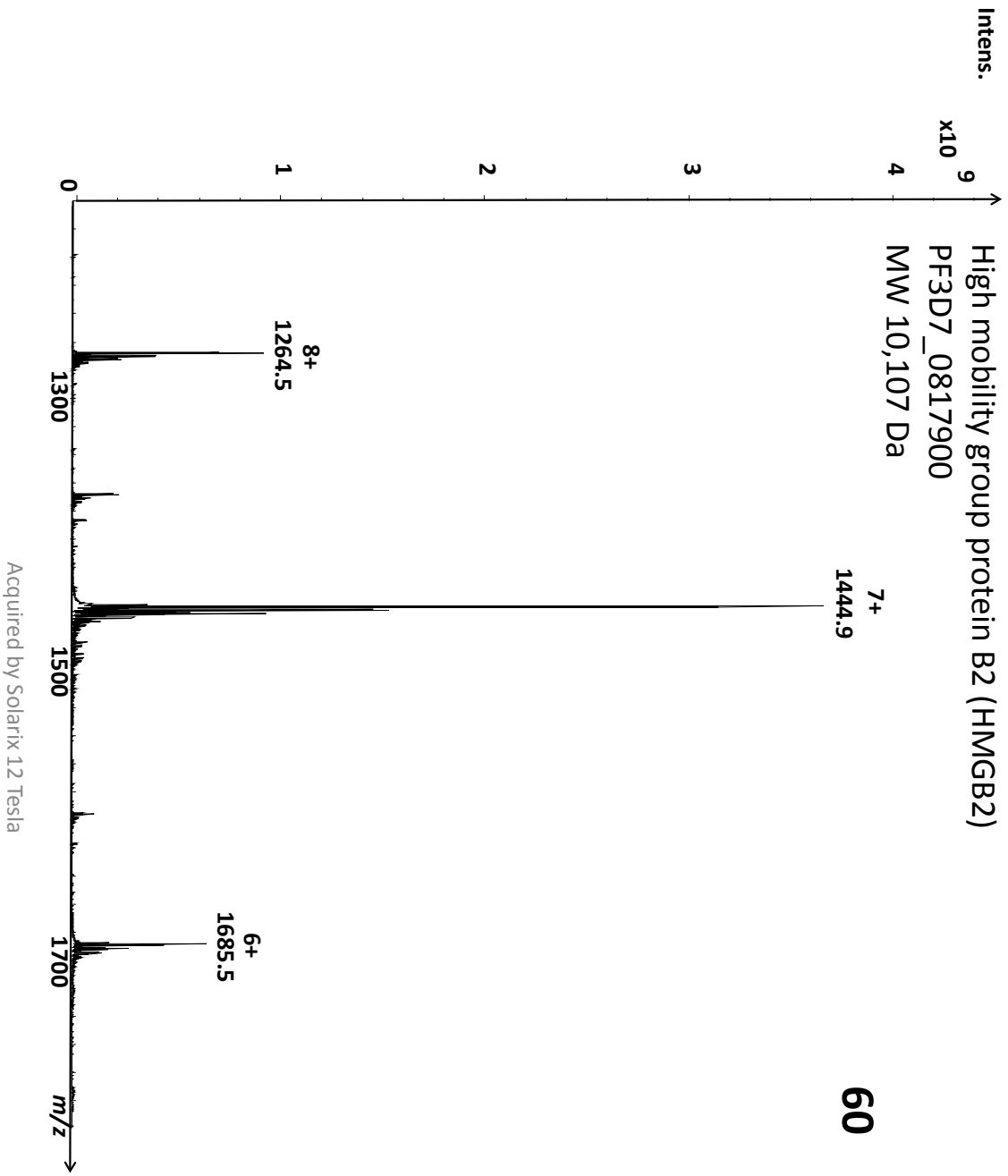


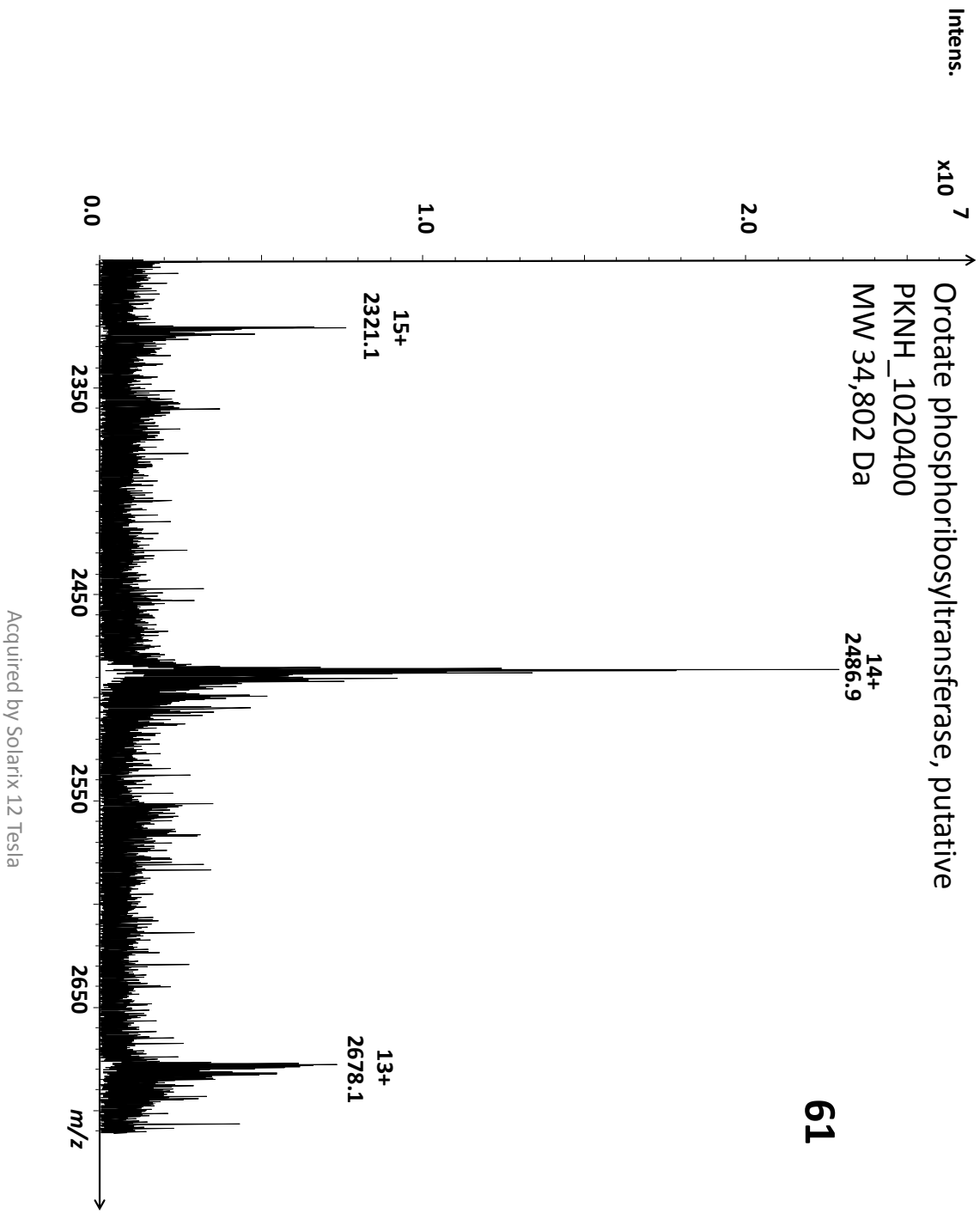
Cell traversal protein for ookinetes and sporozoites (CeTOS)
PF3D7_1216600
MW 17,683 Da

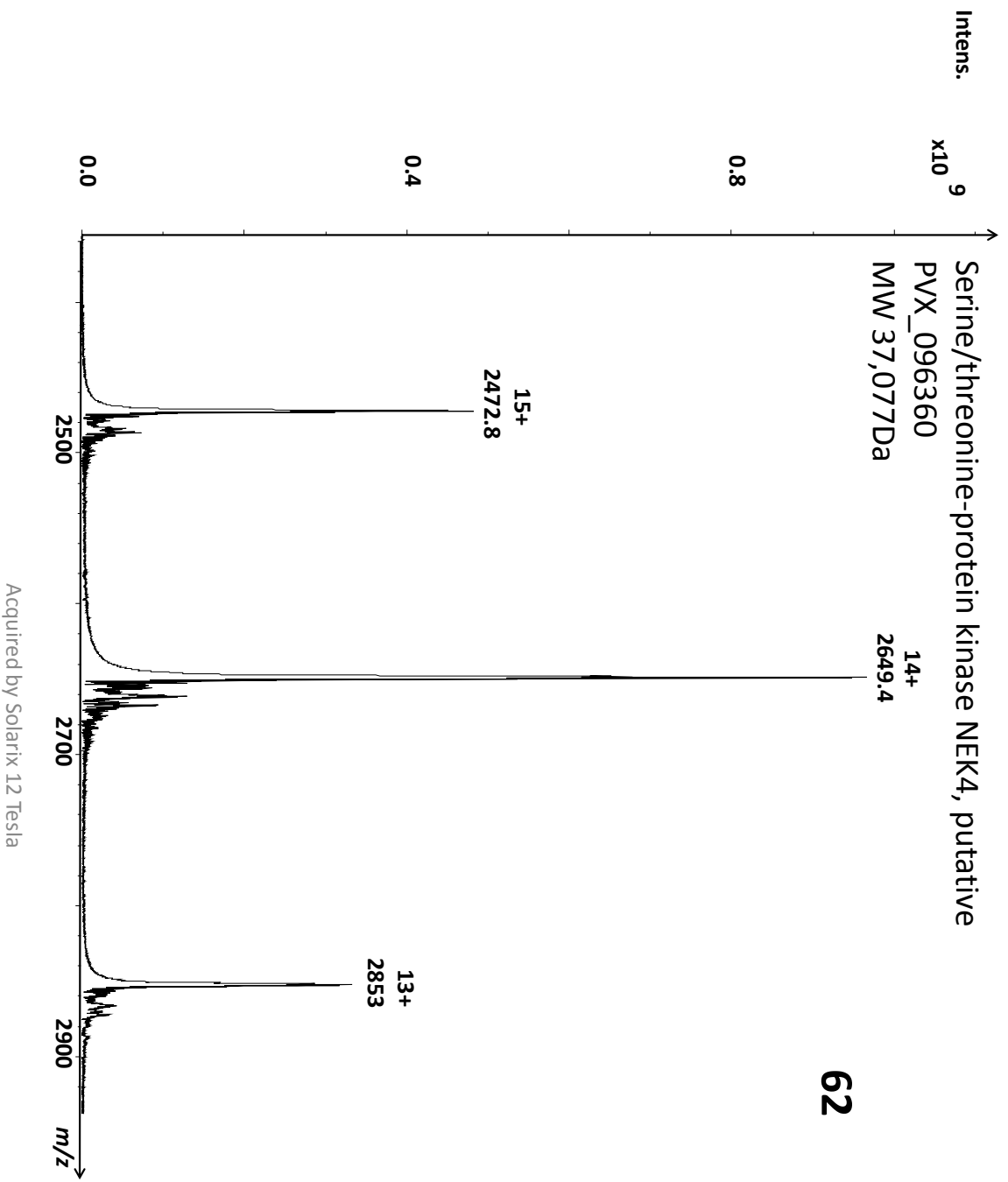
58











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