

Supplementary Tables

Characterization by LC-MS/MS analysis of KLH vaccine conjugated with a tick antigen peptides.

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Table S1. Diagnostic ions originated by the fragmentation of the peptide bond located at the *N*-terminal side of Pro¹¹ [1] in Cys¹pP0 peptides that were used to validate the assignment of the type 2 peptides performed by Kojak [2] and pLink2 [3] software. These fragment ions provide information on the number of amino acids of the proteolytic fragment of Cys¹pP0 in the structures of all assigned type 2 peptides

#	Cys ¹ pP0	y'' _n (α or β)	m (y'' _n)
1	(1-14)	y'' ₃	374.1563
2	(1-16)	y'' ₅	589.2833
3	(1-17)	y'' ₆	717.3782

Table S2. Diagnostic ions named as C+L correspond to the fragmentation of the pseudopeptide bond between the linker and the epsilon amino group of Lys residues present in all type 2 peptides with transcyclized linker that contain in their structures different proteolytic fragments of Cys¹pP0 peptide (see **Fig. S2** and **Fig. S3**). C means the different proteolytic fragments of the Cys¹pP0 peptide linked by a pseudopeptide bond to the transcyclized linker named L (+C₇H₆NO₃) in this manuscript. C+L fragment ion is considered a b_n ion type, and the 2+ is due to the addition of one proton and the carbonyl group with a positive charge (-C=O[⊕]).

#	Cys ¹ pP0	C+L	
		Global formula	m/z
1	(1-11)	C ₄₀ H ₆₃ N ₁₃ O ₁₅ S ₁	499.7222 (2+) / 998.4366 (1+)
2	(1-14)	C ₅₅ H ₈₄ N ₁₆ O ₂₂ S ₁	677.2911 (2+) / 1353.5744 (1+)
3	(1-16)	C ₆₄ H ₁₀₂ N ₁₉ O ₂₅ S ₁	784.8547 (2+) / 1568.7016 (1+)
4	(1-17)	C ₇₀ H ₁₁₃ N ₂₁ O ₂₆ S ₁	848.9022 (2+) / 1696.7966 (1+)

Table S3. The diagnostic ion named as C+80 corresponds to the fragmentation of the new pseudopeptide bond originated after the transcyclization process that transforms the thiosuccinimide linker into transcyclized (**Fig. S2** and **Fig. S3**). C corresponds to different proteolytic fragments of the Cys¹pP0 peptide. The C+80 fragment ion is exclusive of transcyclized linker, and also provides information on the number of amino acids present in the proteolytic fragment of Cys¹pP0 for all type 2 peptides assigned with transcyclized linker.

#	Cys ¹ pP0	C+80	
		Global formula	m/z
1	(1-11)	C ₃₇ H ₅₈ N ₁₂ O ₁₄ S ₁	464.2036 (2+) / 927.3994 (1+)
2	(1-14)	C ₅₂ H ₇₉ N ₁₅ O ₂₁ S ₁	641.7726 (2+) / 1282.5374 (1+)
3	(1-16)	C ₆₁ H ₉₆ N ₁₈ O ₂₄ S ₁	749.3361 (2+) / 1497.6644 (1+)
4	(1-17)	C ₆₇ H ₁₀₈ N ₂₀ O ₂₅ S ₁	813.3836 (2+) / 1625.7594 (1+)

Table S4. Diagnostic ions named as C+L_n corresponding to the fragmentation of the pseudopeptide bond between the linker and the epsilon amino group of Lys residues present in all type 2 peptides that contain in their structures different proteolytic fragments of Cys¹pP0 peptide and two structural variants of the linker: hydrolyzed thiosuccinimide linkers (see **Fig. S2** and **Fig. S4**). C means the different proteolytic fragments of the Cys¹pP0 peptide linked by a pseudopeptide bond to the hydrolyzed thiosuccinimide linkers named as L (+C₇H₇NO₄) in this manuscript. C+L_n fragment ion is considered a b_n ion type, and the 2+ is due to the addition of one proton and the carbonyl group with a positive charge (-C=O[⊕]).

#	Cys ¹ pP0	C+L _n	
		Global formula	m/z
1	(1-11)	C ₄₀ H ₆₅ N ₁₃ O ₁₆ S ₁	508.7275 (2+) / 1016.4471 (1+)
2	(1-14)	C ₅₅ H ₈₆ N ₁₆ O ₂₃ S ₁	686.2964 (2+) / 1371.5851 (1+)
3	(1-16)	C ₆₄ H ₁₀₄ N ₁₉ O ₂₆ S ₁	794.3638 (2+) / 1587.7198 (1+)
4	(1-17)	C ₇₀ H ₁₁₅ N ₂₁ O ₂₇ S ₁	857.9074 (2+) / 1714.8070 (1+)

Table S5. The diagnostic ion named as C+98 corresponds to the fragmentation of the new pseudopeptide bond originated after the hydrolysis process that transforms the thiosuccinimide linker into hydrolyzed thiosuccinimide linkers (**Fig. S2** and **Fig. S4**). C corresponds to different proteolytic fragments of the Cys¹pP0 peptide. The C+98 fragment ion is exclusive of hydrolyzed thiosuccinimide linkers, and also provides information on the number of amino acids present in the proteolytic fragment of Cys¹pP0 for all type 2 peptides assigned with hydrolyzed thiosuccinimide linkers.

#	Cys ¹ pP0	C+98	
		Global formula	m/z
1	(1-11)	C ₃₇ H ₆₀ N ₁₂ O ₁₅ S ₁	473.2089 (2+) / 945.4100 (1+)
2	(1-14)	C ₅₂ H ₈₁ N ₁₅ O ₂₂ S ₁	650.7779 (2+) / 1300.5480 (1+)
3	(1-16)	C ₆₁ H ₉₈ N ₁₈ O ₂₅ S ₁	758.3414 (2+) / 1515.6749 (1+)
4	(1-17)	C ₆₇ H ₁₁₀ N ₂₀ O ₂₆ S ₁	822.3889 (2+) / 1643.7699 (1+)

Table S6. Identification of conjugation sites in KLH1-Cys¹pP0 conjugate using Kojak [2] and pLink2 [3] software. Type 2 peptides were identified using transcyclized and hydrolyzer linker.

#	<i>m/z</i> exp.	<i>m/z</i> theor.	<i>z</i>	Elemental composition	Assignment ¹⁾	pLink ²⁾	Kojak ³⁾	Conjugation site ⁴⁾	
1	878.4431	878.4445	2	C ₇₂ H ₁₂₂ N ₂₄ O ₂₅ S ₁	[1-6] ⁵⁾		X	Nt	
2	585.9651	585.9656	3						X
3	582.2919	582.2917	4	C ₉₆ H ₁₆₀ N ₃₀ O ₃₅ S ₁	[1-6]		X		
4	466.0280	466.0355	5						X
5	795.3931	795.4009	4	C ₁₃₃ H ₂₂₀ N ₄₀ O ₄₈ S ₁	[6-24] ⁵⁾		X	K6	
6	937.9585	937.9672	4	C ₁₅₇ H ₂₅₈ N ₄₆ O ₅₈ S ₁	[6-24]		X		
7	750.5697	750.5753	5						X
8	615.2849	615.2916	5	C ₁₂₇ H ₂₀₂ N ₄₀ O ₄₅ S ₂	[38-52] ⁶⁾	X	X	K51	
9	571.7819	571.7897	4	C ₉₄ H ₁₅₈ N ₃₀ O ₃₄ S ₁	[79-84]		X	K83	
10	571.9575	571.9615	3	C ₇₀ H ₁₂₀ N ₂₄ O ₂₄ S ₁	[79-84] ⁵⁾	X			
11	742.5496	742.5563	5	C ₁₆₃ H ₂₄₂ N ₄₆ O ₅₂ S ₁	[118-134]	X	X	K133	
12	618.9582	618.9650	6						X
13	746.1531	746.1584	5	C ₁₆₃ H ₂₄₄ N ₄₆ O ₅₃ S ₁	[118-134] ⁷⁾		H		
14	785.3786	785.3773	4	C ₁₃₉ H ₂₀₄ N ₄₀ O ₄₂ S ₁	[118-134] ⁵⁾		X		
15	628.4973	628.5034	5				X		X
16	655.0707	655.0695	4	C ₁₁₁ H ₁₇₇ N ₃₁ O ₄₀ S ₁	[126-134]	X	X		
17	682.9968	683.0025	3	C ₈₇ H ₁₃₉ N ₂₅ O ₃₀ S ₁	[126-134] ⁵⁾	X	X		
18	527.4250	527.4308	6	C ₁₃₂ H ₂₁₅ N ₄₁ O ₄₇ S ₁	[134-146]	X	X	K134	
19	632.7104	632.7156	5				X		X
20	790.6339	790.6432	4				X		X
21	648.0708	648.0770	4	C ₁₀₈ H ₁₇₇ N ₃₅ O ₃₇ S ₁	[134-146] ⁵⁾	X	X		
22	522.2539	522.2550	4	C ₈₄ H ₁₄₀ N ₂₈ O ₃₂ S ₁	[134-137]		X		
23	587.6332	587.6382	3	C ₇₇ H ₁₂₅ N ₂₁ O ₂₄ S ₁	[143-148] ⁵⁾	X	X	K146	
24	583.5411	583.5467	4	C ₁₀₁ H ₁₆₃ N ₂₇ O ₃₄ S ₁	[143-148]	X			
25	777.7212	777.7271	3				X		
26	588.0424	588.0499	4	C ₁₀₁ H ₁₆₅ N ₂₇ O ₃₅ S ₁	[143-148] ⁷⁾		H		
27	652.5242	652.5298	5	C ₁₃₇ H ₂₂₄ N ₄₂ O ₄₈ S ₁	[135-148]	X	X		
28	543.9350	543.9428	6						X
29	672.8362	672.8441	4			C ₁₁₃ H ₁₈₆ N ₃₆ O ₃₈ S ₁	[135-148] ⁵⁾		
30	538.4708	538.4768	5		X				X
31	1095.5075	1095.5154	4	C ₁₈₅ H ₂₉₂ N ₅₂ O ₆₅ S ₃	[147-170] ⁶⁾	X	X	K148	
32	876.6099	876.6139	5						X
33	1099.5063	1099.5142	4	C ₁₈₅ H ₂₉₂ N ₅₂ O ₆₆ S ₃	[147-170] ^{6,8)}		X		
34	879.8086	879.8123	5				X		X
35	880.2071	880.2160	5	C ₁₈₅ H ₂₉₄ N ₅₂ O ₆₆ S ₃	[147-170] ^{6,7)}		H		
36	952.9450	952.9492	4	C ₁₆₁ H ₂₅₄ N ₄₆ O ₅₅ S ₃	[147-170] ^{6,5)}	X	X		

37	762.5601	762.5609	5				X		
38	957.4440	957.4518	4	C ₁₆₁ H ₂₅₆ N ₄₆ O ₅₆ S ₃	[147-170] ^{5,6,7)}		H		
39	766.1576	766.1630	5					H	
40	765.7520	765.7599	5	C ₁₆₁ H ₂₅₄ N ₄₆ O ₅₆ S ₃	[147-170] ^{6,5,8)}		X		
41	956.9401	956.9479	4					X	
42	711.8349	711.8405	4	C ₁₁₉ H ₁₉₀ N ₃₆ O ₄₁ S ₂	[147-157]	X	X		
43	569.6671	569.6744	5					X	
44	572.8731	572.8734	5	C ₁₁₉ H ₁₉₀ N ₃₆ O ₄₂ S ₂	[147-157] ⁸⁾		X		
45	569.2675	569.2748	4	C ₉₅ H ₁₅₂ N ₃₀ O ₃₁ S ₂	[147-157] ⁵⁾	X			
46	573.2675	573.2730	4	C ₉₅ H ₁₅₂ N ₃₀ O ₃₂ S ₂	[147-157] ^{5,8)}	X	X		
47	766.6697	766.6766	3	C ₉₈ H ₁₄₈ N ₂₆ O ₃₄ S ₂	[168-172] ⁶⁾		X	K170	
48	804.9795	804.9852	5	C ₁₇₃ H ₂₆₆ N ₅₀ O ₅₇ S ₂	[177-198]	X	X	K189	
49	863.4078	863.4134	4	C ₁₄₉ H ₂₂₈ N ₄₄ O ₄₇ S ₂	[177-198] ⁵⁾	X			
50	690.9272	690.9327	5						X
51	694.1261	694.1317	5	C ₁₄₉ H ₂₂₈ N ₄₄ O ₄₈ S ₂	[177-198] ^{5,8)}	X			
52	723.6800	723.6796	3	C ₈₉ H ₁₄₅ N ₂₇ O ₃₄ S ₁	[228-232]		X	K229	
53	543.0110	543.0115	4						
54	838.6378	838.6405	4	C ₁₃₈ H ₂₂₃ N ₄₁ O ₅₀ S ₃	[230-244] ⁶⁾	X		K232	
55	842.6319	842.6398	4	C ₁₃₈ H ₂₂₃ N ₄₁ O ₅₁ S ₃	[230-244] ^{6,8)}		X		
56	674.3055	674.3134	5						
57	843.1373	843.1437	4	C ₁₃₈ H ₂₂₅ N ₄₁ O ₅₁ S ₃	[230-244] ^{6,7)}		H		
58	674.7166	674.7166	5						
59	927.7682	927.7639	3	C ₁₁₄ H ₁₈₅ N ₃₅ O ₄₀ S ₃	[230-244] ^{6,5)}		X		
60	700.0668	700.0735	4	C ₁₁₄ H ₁₈₅ N ₃₅ O ₄₁ S ₃	[230-244] ^{6,5,8)}		X		
61	788.1407	788.1486	5	C ₁₆₄ H ₂₅₀ N ₄₆ O ₆₁ S ₃	[331-351] ^{6,8)}		X	K337	
62	788.3730	788.3772	4	C ₁₃₆ H ₂₁₂ N ₃₆ O ₄₆ S ₂	[337-351]	X			
63	860.7427	860.7455	3	C ₁₁₂ H ₁₇₄ N ₃₀ O ₃₆ S ₂	[337-351] ⁵⁾	X			
64	866.0728	866.0772	3	C ₁₁₂ H ₁₇₄ N ₃₀ O ₃₇ S ₂	[337-351] ^{5,8)}	X			
65	964.4519	964.4559	3	C ₁₂₅ H ₁₉₁ N ₃₃ O ₄₄ S ₁	[337-349]	X			
66	1047.702	1047.6991	5	C ₂₃₇ H ₃₄₅ N ₅₉ O ₇₂ S ₂	[338-368] ⁸⁾	X	X	K351	
67	873.2523	873.2510	6						
68	876.2488	876.2527	6	C ₂₃₇ H ₃₄₇ N ₅₉ O ₇₃ S ₂	[338-368] ^{7,8)}		H		
69	870.5794	870.5846	6	C ₂₃₇ H ₃₄₅ N ₅₉ O ₇₁ S ₂	[338-368]	X	X		
70	1162.8056	1162.8076	4	C ₂₁₃ H ₃₀₇ N ₅₃ O ₆₁ S ₂	[338-368] ⁵⁾		X		
71	930.4362	930.4476	5						X
72	933.6405	933.6466	5	C ₂₁₃ H ₃₀₇ N ₅₃ O ₆₂ S ₂	[338-368] ^{5,8)}	X			
73	951.4409	951.4462	4	C ₁₇₀ H ₂₄₈ N ₄₄ O ₅₂ S ₂	[338-356]	X			
74	955.9403	955.9493	4	C ₁₇₀ H ₂₅₀ N ₄₄ O ₅₃ S ₂	[338-356] ⁷⁾		H		
75	955.4407	955.4449	4	C ₁₇₀ H ₂₄₈ N ₄₄ O ₅₃ S ₂	[338-356] ⁸⁾	X			
76	685.9835	685.9891	3	C ₉₁ H ₁₃₄ N ₂₆ O ₂₅ S ₂	[350-356] ⁵⁾	X			
77	691.3156	691.3213	3	C ₉₁ H ₁₃₄ N ₂₆ O ₂₆ S ₂	[350-356] ^{5,8)}	X			

78	846.8000	846.8035	5	C ₁₈₉ H ₂₈₁ N ₄₉ O ₅₆ S ₃	[352-372]	X	X	K368		
79	1058.2454	1058.2531	4						X	
80	850.004	850.0024	5			C ₁₈₉ H ₂₈₁ N ₄₉ O ₅₇ S ₃	[352-372] ⁸⁾		X	X
81	732.7431	732.7510	5			C ₁₆₅ H ₂₄₃ N ₄₃ O ₄₆ S ₃	[352-372] ⁵⁾			X
82	590.2802	590.2808	5	C ₁₂₄ H ₁₉₉ N ₃₅ O ₄₂ S ₃	[369-380]	X	X	K372		
83	737.5990	737.5991	4						X	X
84	983.1236	983.1303	3							X
85	593.8726	593.8834	5	C ₁₂₄ H ₂₀₁ N ₃₅ O ₄₃ S ₃	[369-380] ⁷⁾		H			
86	741.598	741.5978	4	C ₁₂₄ H ₁₉₉ N ₃₅ O ₄₃ S ₃	[369-380] ⁸⁾	X	X			
87	593.4800	593.4797	5						X	X
88	741.598	741.5978	4						X	X
89	593.4800	593.4797	5							X
90	596.6747	596.6793	5	C ₁₂₄ H ₁₉₉ N ₃₅ O ₄₄ S ₃	[369-380] ⁸⁾		X			
91	793.0405	793.0420	3	C ₁₀₀ H ₁₆₁ N ₂₉ O ₃₂ S ₃	[369-380] ⁵⁾		X			
92	595.0266	595.0334	4					X	X	
93	599.0262	599.0322	4			C ₁₀₀ H ₁₆₁ N ₂₉ O ₃₃ S ₃	[369-380] ^{5, 8)}	X	X	
94	618.7845	618.7925	4	C ₁₀₄ H ₁₆₂ N ₃₀ O ₃₈ S ₁	[375-382]		X	K380		
95	634.6263	634.6318	3	C ₈₀ H ₁₂₄ N ₂₄ O ₂₈ S ₁	[375-382] ⁵⁾	X				
96	710.6632	710.6689	3	C ₉₀ H ₁₄₀ N ₂₆ O ₃₂ S ₁	[373-382] ⁵⁾	X	X			
97	900.7518	900.7578	3	C ₁₁₄ H ₁₇₈ N ₃₂ O ₄₂ S ₁	[373-382]		X			
98	675.8142	675.8197	4					X	X	
99	852.7515	852.7520	3	C ₁₁₀ H ₁₇₄ N ₃₀ O ₃₈ S ₁	[407-415]		X	K407		
100	639.8106	639.8154	4						X	X
101	662.6586	662.6631	3			C ₈₆ H ₁₃₆ N ₂₄ O ₂₈ S ₁	[407-415] ⁵⁾		X	X
102	619.4785	619.4837	6	C ₁₆₀ H ₂₅₅ N ₄₉ O ₅₁ S ₁	[408-426]	X	X	K415		
103	531.1240	531.1299	7						X	
104	575.6627	575.6680	5	C ₁₂₂ H ₁₈₄ N ₃₆ O ₄₃ S ₁	[441-450]	X	X	K445		
105	719.3277	719.3333	4						X	X
106	579.2631	579.2707	5	C ₁₂₂ H ₁₈₆ N ₃₆ O ₄₄ S ₁	[441-450] ⁷⁾		H			
107	581.2634	581.2702	4	C ₉₈ H ₁₄₈ N ₃₀ O ₃₄ S ₁	[441-450] ^{5,7)}		H			
108	768.6823	768.6869	3	C ₉₈ H ₁₄₆ N ₃₀ O ₃₃ S ₁	[441-450] ⁵⁾	X				
109	576.7616	576.7676	4						X	
110	671.3138	671.3145	4	C ₁₀₈ H ₁₇₆ N ₃₆ O ₄₀ S ₂	[468-476] ⁶⁾	X	X	K474		
111	537.2528	537.2530	5						X	X
112	704.6560	704.6624	3	C ₈₄ H ₁₃₈ N ₃₀ O ₃₀ S ₂	[468-476] ^{6,5)}	X				
113	528.7429	528.7487	4						X	
114	568.0131	568.0183	4	C ₉₂ H ₁₄₉ N ₂₉ O ₃₆ S ₁	[471-476]	X				
115	566.9285	566.9337	3	C ₆₈ H ₁₁₁ N ₂₃ O ₂₆ S ₁	[471-476] ⁵⁾	X				
116	598.5361	598.5422	4	C ₉₈ H ₁₅₉ N ₃₃ O ₃₅ S ₁	[554-560]	X	X	K559		
117	479.0289	479.0352	5						X	X
118	455.9701	455.9760	4			C ₇₄ H ₁₂₁ N ₂₇ O ₂₅ S ₁	[554-560] ⁵⁾		X	X

119	607.6264	607.6322	3			X	X	
120	425.2052	425.2104	4	$C_{72}H_{112}N_{24}O_{22}S_1$	[651-655] ⁵⁾	X		K652
121	566.6069	566.6113	3			X		
122	567.7703	567.7767	4			$C_{96}H_{150}N_{30}O_{32}S_1$	[651-655]	
123	572.2781	572.2793	4	$C_{96}H_{152}N_{30}O_{33}S$	[651-655] ⁷⁾		H	
124	711.7656	711.7734	7	$C_{216}H_{331}N_{63}O_{67}S_3$	[653-681] ⁶⁾		X	K668
125	777.8717	777.8749	4	$C_{135}H_{206}N_{40}O_{41}S_2$	[664-681] ⁵⁾	X	X	
126	912.8345	912.8362	5	$C_{196}H_{303}N_{57}O_{63}S_3$	[656-681] ⁶⁾		X	
127	760.8655	760.8648	6			X		
128	1040.8932	1040.8925	5	$C_{229}H_{343}N_{59}O_{76}S_2$	[682-717] ⁵⁾		X	K690
129	1154.9458	1154.9454	5	$C_{253}H_{381}N_{65}O_{86}S_2$	[682-717]		X	
130	962.6211	962.6219	6			X	X	
131	1158.5446	1158.5475	5	$C_{253}H_{383}N_{65}O_{87}S_2$	[682-717] ⁷⁾		H	
132	965.6205	965.6243	6			H		
133	646.0499	646.0560	4	$C_{107}H_{173}N_{31}O_{39}S_2$	[714-721]	X	X	K717
134	861.069	861.0728	3			X		
135	650.0489	650.0552	4	$C_{107}H_{173}N_{31}O_{40}S_2$	[714-721] ⁸⁾	X	X	
136	670.9796	670.9844	3	$C_{83}H_{135}N_{25}O_{29}S_2$	[714-721] ^{5,8)}	X	X	
137	742.3402	742.3495	3	$C_{92}H_{149}N_{27}O_{33}S_2$	[717-721]		X	
138	557.0042	557.0142	4			X	X	
139	552.2560	552.2613	3	$C_{68}H_{111}N_{21}O_{23}S_2$	[717-721] ⁵⁾	X		
140	1005.3302	1005.3360	6	$C_{260}H_{413}N_{75}O_{86}S_2$	[718-756]		X	K748
141	861.8522	861.8605	7			X		
142	864.1376	864.1456	7	$C_{260}H_{413}N_{75}O_{87}S_2$	[718-756] ⁸⁾		X	
143	1001.4903	1001.4895	3	$C_{127}H_{204}N_{36}O_{46}S_1$	[739-756] ⁵⁾		X	
144	751.1180	751.1231	4			X	X	
145	715.1474	715.1530	5	$C_{151}H_{243}N_{43}O_{55}S_1$	[739-756]	X	X	
146	893.6818	893.6893	4			X		
147	718.7489	718.7551	5	$C_{151}H_{245}N_{43}O_{56}S_1$	[739-756] ⁷⁾		H	
148	1157.8936	1157.8990	3	$C_{146}H_{235}N_{43}O_{53}S_1$	[749-767]		X	K756
149	868.6730	868.6757	4			X	X	
150	695.1378	695.1420	5	X	X			
151	698.7365	698.7447	5	$C_{146}H_{237}N_{43}O_{54}S_1$	[749-767] ⁷⁾		H	
152	730.6070	730.6126	4	$C_{122}H_{199}N_{37}O_{44}S_1$	[749-767] ^{5,7)}		H	
153	584.6856	584.6917	5			H		
154	967.8090	967.8107	3	$C_{122}H_{197}N_{37}O_{43}S_1$	[749-767] ⁵⁾		X	
155	726.1093	726.1100	4			X		
156	631.7093	631.7181	5	$C_{134}H_{216}N_{40}O_{46}S_1$	[752-767]		X	
157	988.4799	988.4800	3	$C_{125}H_{203}N_{35}O_{44}S_2$	[757-770]		X	K767
158	741.612	741.6114	4			X	X	
159	593.4913	593.4905	5			X	X	

160	746.1049	746.1146	4	C ₁₂₅ H ₂₀₅ N ₃₅ O ₄₅ S ₂	[757-770] ⁷⁾		H		
161	993.8126	993.8126	3	C ₁₂₅ H ₂₀₃ N ₃₅ O ₄₅ S ₂	[757-770] ⁸⁾		X		
162	745.611	745.6101	4			X	X		
163	1197.0838	1197.0836	2	C ₁₀₁ H ₁₆₅ N ₂₉ O ₃₄ S ₂	[757-770] ⁵⁾		X		
164	798.3874	798.3917	3			X	X		
165	603.5399	603.5483	4	C ₁₀₁ H ₁₆₇ N ₂₉ O ₃₅ S ₂	[757-770] ^{5,7)}		H		
166	902.0965	902.1010	3	C ₁₁₄ H ₁₈₂ N ₃₂ O ₄₂ S ₁	[757-768]	X	X		
167	1067.5108	1067.5153	2	C ₉₀ H ₁₄₄ N ₂₆ O ₃₂ S ₁	[757-768] ⁵⁾	X			
168	712.0128	712.0132	3				X		
169	602.8903	602.8895	5	C ₁₃₂ H ₂₀₀ N ₃₆ O ₄₁ S ₂	[768-778]	X	X	K770	
170	753.3620	753.3601	4			X	X		
171	1004.1401	1004.1450	3				X		
172	757.8599	757.8633	4	C ₁₃₂ H ₂₀₂ N ₃₆ O ₄₂ S ₂	[768-778] ⁷⁾		H		
173	606.4879	606.4922	5				H		
174	1009.4764	1009.4760	3	C ₁₃₂ H ₂₀₀ N ₃₆ O ₄₂ S ₂	[768-778] ⁸⁾	X	X		
175	757.3602	757.3594	4				X		
176	606.0896	606.0891	5				X		
177	610.7901	610.7944	4			C ₁₀₈ H ₁₆₂ N ₃₀ O ₃₁ S ₂	[768-778] ⁵⁾		X
178	814.0541	814.0566	3	X	X				
179	615.2913	615.2971	4	C ₁₀₈ H ₁₆₄ N ₃₀ O ₃₂ S ₂	[768-778] ^{5,7)}		H		
180	819.3825	819.3882	3	C ₁₀₈ H ₁₆₂ N ₃₀ O ₃₂ S ₂	[768-778] ^{5,8)}	X	X		
181	614.7876	614.7932	4			X	X		
182	674.5163	674.5215	6	C ₁₇₅ H ₂₈₅ N ₅₃ O ₅₅ S ₁	[805-831]	X		K828	
183	735.3577	735.3599	3	C ₉₄ H ₁₅₀ N ₂₆ O ₃₃ S ₁	[827-831]	X	X		
184	551.7722	551.7718	4			X	X		
185	545.2670	545.2722	3	C ₇₀ H ₁₁₂ N ₂₀ O ₂₃ S ₁	[827-831] ⁵⁾	X	X		
186	583.0301	583.0370	4	C ₉₉ H ₁₅₇ N ₂₉ O ₃₄ S ₁	[829-834]		X	K831	
187	586.9534	586.9579	3	C ₇₅ H ₁₁₉ N ₂₃ O ₂₄ S ₁	[829-834] ⁵⁾	X	X		
188	771.4043	771.4114	4	C ₁₃₂ H ₂₂₀ N ₄₂ O ₄₁ S ₁	[829-840]		X		
189	617.3209	617.3307	5				X		
190	514.6029	514.6102	6				X		
191	733.6799	733.6874	3	C ₈₉ H ₁₄₇ N ₂₉ O ₃₄ S ₁	[840-844]		X	K840	
192	550.5169	550.5169	4				X		
193	668.6802	668.6787	6	C ₁₆₉ H ₂₈₄ N ₅₀ O ₅₈ S ₂	[841-861]		X	K855	
194	642.831	642.8298	4	C ₁₀₉ H ₁₈₂ N ₃₀ O ₃₉ S ₁	[856-864]	X	X	K861	
195	856.7718	856.7706	3			X	X		
196	647.3260	647.3330	4	C ₁₀₉ H ₁₈₄ N ₃₀ O ₄₀ S ₁	[856-864] ⁷⁾		H		
197	893.4140	893.4193	4	C ₁₅₂ H ₂₃₂ N ₄₄ O ₅₄ S ₁	[862-880]	X	X	K864	
198	714.9320	714.9369	5			X	X		
199	718.5317	718.5396	5	C ₁₅₂ H ₂₃₄ N ₄₄ O ₅₅ S ₁	[862-880] ⁷⁾		H		
200	750.8456	750.8536	4	C ₁₂₈ H ₁₉₄ N ₃₈ O ₄₄ S ₁	[862-880] ⁵⁾		X		

201	989.4524	989.4512	4	C ₁₇₁ H ₂₅₂ N ₄₈ O ₅₇ S ₂	[865-886] ⁶⁾		X	K880
202	1060.9844	1060.9923	4	C ₁₈₃ H ₂₇₄ N ₅₂ O ₆₁ S ₂	[865-889] ⁶⁾		X	
203	848.9908	848.9954	5			X	X	
204	852.5901	852.5975	5	C ₁₈₃ H ₂₇₆ N ₅₂ O ₆₂ S ₂	[865-889] ^{6,7)}		H	
205	710.6585	710.6659	6				H	
206	738.5376	738.5445	5	C ₁₅₉ H ₂₃₈ N ₄₆ O ₅₂ S ₂	[865-889] ^{6,5,7)}		H	
207	1224.2280	1224.2322	3	C ₁₅₉ H ₂₃₆ N ₄₆ O ₅₁ S ₂	[865-889] ^{6,5)}		X	
208	918.4220	918.4260	4			X	X	
209	690.066	690.0706	4	C ₁₁₆ H ₁₈₁ N ₃₃ O ₄₁ S ₂	[881-890]	X	X	K889
210	729.6655	729.6707	3	C ₉₂ H ₁₄₃ N ₂₇ O ₃₁ S ₂	[881-890] ⁵⁾	X	X	
211	694.5670	694.5738	4	C ₁₁₆ H ₁₈₃ N ₃₃ O ₄₂ S ₂	[881-890] ⁷⁾		H	
212	882.5682	882.5748	6	C ₂₃₆ H ₃₃₇ N ₆₁ O ₇₇ S ₁	[919-955] ⁵⁾	X		K945
213	977.6133	977.6190	6	C ₂₆₀ H ₃₇₅ N ₆₇ O ₈₇ S ₁	[919-955]	X	X	
214	1172.9337	1172.9418	5				X	
215	1066.7587	1066.7583	4	C ₁₈₁ H ₂₇₉ N ₅₅ O ₆₃ S ₁	[956-978]		X	K969
216	853.6026	853.6086	5			X		
217	711.5028	711.5079	6			X	X	
218	609.9998	610.0084	7				X	
219	714.5037	714.5103	6	C ₁₈₁ H ₂₈₁ N ₅₅ O ₆₄ S ₁	[956-978] ⁷⁾		H	
220	616.4599	616.4644	6	C ₁₅₇ H ₂₄₁ N ₄₉ O ₅₃ S ₁	[956-978] ⁵⁾	X		
221	739.5514	739.5556	5				X	
222	743.1513	743.1577	5	C ₁₅₇ H ₂₄₃ N ₄₉ O ₅₄ S ₁	[956-978] ^{5,7)}		H	
223	618.7838	618.7924	6	C ₁₅₈ H ₂₃₉ N ₄₇ O ₅₅ S ₁	[956-973]		X	
224	742.3406	742.3487	5			X	X	
225	927.6770	927.6846	4				X	
226	870.8224	870.8278	5			C ₁₈₈ H ₂₉₃ N ₅₃ O ₆₄ S ₁	[970-993]	X
227	725.8507	725.8582	6		X			
228	945.7093	945.7172	4	C ₁₆₄ H ₂₅₅ N ₄₇ O ₅₄ S ₁	[970-993] ⁵⁾		X	K978
229	778.9748	778.9789	5	C ₁₇₀ H ₂₆₀ N ₄₆ O ₅₇ S ₁	[974-993]	X	X	
230	973.4654	973.4724	4				X	
231	830.9014	830.9061	4	C ₁₄₆ H ₂₂₂ N ₄₀ O ₄₇ S ₁	[974-993] ⁵⁾	X		
232	962.7864	962.7906	3	C ₁₂₂ H ₁₉₂ N ₃₄ O ₄₅ S ₁	[974-984]	X	X	
233	772.6981	772.7029	3	C ₉₈ H ₁₅₄ N ₂₈ O ₃₅ S ₁	[974-984] ⁵⁾	X	X	
234	840.1704	840.1750	4	C ₁₄₈ H ₂₃₃ N ₃₉ O ₄₈ S ₁	[985-1000]	X		K993
235	786.6422	786.6493	4	C ₁₄₁ H ₂₁₅ N ₃₉ O ₄₁ S ₁	[1052-1063]		X	K1061
236	809.6231	809.6287	4	C ₁₃₃ H ₂₁₅ N ₄₁ O ₄₇ S ₃	[1064-1078] ⁶⁾	X	X	K1065
237	813.6219	813.6274	4	C ₁₃₃ H ₂₁₅ N ₄₁ O ₄₈ S ₃	[1064-1078] ^{6,8)}	X	X	
238	889.0747	889.0814	3	C ₁₀₉ H ₁₇₇ N ₃₅ O ₃₇ S ₃	[1064-1078] ^{6,5)}	X	X	
239	945.2021	945.2061	4	C ₁₆₁ H ₂₅₃ N ₄₅ O ₅₈ S ₁	[1079-1098]	X	X	K1097
240	922.9169	922.9217	4	C ₁₆₄ H ₂₃₄ N ₄₂ O ₅₄ S ₁	[1099-1115]	X	X	K1109
241	1040.1314	1040.1387	3	C ₁₄₀ H ₁₉₆ N ₃₆ O ₄₄ S ₁	[1099-1115] ⁵⁾	X	X	

242	955.1780	955.1823	4	C ₁₆₉ H ₂₄₁ N ₄₃ O ₅₇ S ₁	[1098-1115]	X		
243	1083.1485	1083.1523	3	C ₁₄₅ H ₂₀₃ N ₃₇ O ₄₇ S ₁	[1098-1115] ⁵⁾	X		
244	1128.9239	1128.9299	5	C ₂₅₁ H ₃₆₇ N ₆₃ O ₈₄ S ₁	[1097-1130]		X	
245	1268.3416	1268.3442	4	C ₂₂₇ H ₃₂₉ N ₅₇ O ₇₄ S ₁	[1097-1131] ⁵⁾	X		
246	568.0434	568.0428	4	C ₉₆ H ₁₆₁ N ₂₇ O ₃₄ S ₁	[1132-1137]	X	X	
247	572.5384	572.5460	4	C ₉₆ H ₁₆₃ N ₂₇ O ₃₅ S ₁	[1132-1137] ⁷⁾		H	
248	458.2307	458.2383	5				H	
249	572.9630	572.9703	3	C ₇₂ H ₁₂₅ N ₂₁ O ₂₅ S ₁	[1132-1137] ^{5,7)}		H	
250	566.9621	566.9663	3	C ₇₂ H ₁₂₃ N ₂₁ O ₂₄ S ₁	[1132-1137] ⁵⁾	X	X	
251	657.5689	657.5737	4	C ₁₀₈ H ₁₇₉ N ₃₃ O ₄₁ S ₁	[1133-1141]		X	K1137
252	683.9344	683.9406	5	C ₁₅₂ H ₂₃₁ N ₄₁ O ₄₇ S ₁	[1138-1153]	X	X	
253	854.6695	854.6740	4			X	X	
254	859.1713	859.1771	4	C ₁₅₂ H ₂₃₃ N ₄₁ O ₄₈ S ₁	[1138-1153] ⁷⁾		H	
255	687.5370	687.5433	5				H	
256	573.1142	573.1207	6				H	
257	712.1029	712.1083	4	C ₁₂₈ H ₁₉₃ N ₃₅ O ₃₇ S ₁	[1138-1153] ⁵⁾	X		
258	733.1138	733.1193	4	C ₁₃₄ H ₂₀₁ N ₃₃ O ₃₉ S ₁	[1142-1153]	X	X	
259	553.6184	553.6230	3	C ₆₉ H ₁₁₉ N ₂₁ O ₂₄ S ₁	[1153-1158] ⁵⁾	X	X	
260	562.5303	562.5381	4	C ₉₃ H ₁₅₉ N ₂₇ O ₃₅ S ₁	[1153-1158] ⁷⁾		H	
261	450.2242	450.2321	5				H	
262	1233.7552	1233.7631	5	C ₂₆₈ H ₃₉₉ N ₇₁ O ₈₉ S ₄	[1154-1192] ⁶⁾		X	
263	1028.2960	1028.3038	6				X	K1158
264	981.4701	981.4788	3	C ₁₂₆ H ₂₀₀ N ₃₄ O ₄₅ S ₁	[1193-1203]		X	
265	736.361	736.3605	4			X	X	
266	589.2900	589.2899	5	C ₁₀₂ H ₁₆₄ N ₂₈ O ₃₆ S ₁	[1193-1203] ^{5,7)}	X	X	
267	598.2896	598.2974	4				H	
268	740.8566	740.8636	4				H	
269	592.8851	592.8925	5	C ₁₂₆ H ₂₀₂ N ₃₄ O ₄₆ S ₁	[1193-1203] ⁷⁾		H	
270	791.3863	791.3904	3	C ₁₀₂ H ₁₆₂ N ₂₈ O ₃₅ S ₁	[1193-1203] ⁵⁾	X	X	
271	569.9479	569.9543	3	C ₇₂ H ₁₁₈ N ₂₂ O ₂₄ S ₁	[1202-1207] ⁵⁾	X	X	
272	570.2786	570.2843	4	C ₉₆ H ₁₅₆ N ₂₈ O ₃₄ S ₁	[1202-1207]	X	X	
273	661.5712	661.5762	4	C ₁₁₂ H ₁₇₉ N ₃₃ O ₃₉ S ₁	[1199-1207]		X	
274	529.4550	529.4630	5			X	X	
275	519.0027	519.0106	4	C ₈₈ H ₁₄₁ N ₂₇ O ₂₉ S ₁	[1199-1207] ⁵⁾	X	X	
276	752.3671	752.3655	5	C ₁₇₀ H ₂₄₉ N ₄₃ O ₅₂ S ₁	[1199-1220] ⁵⁾		X	
277	866.4133	866.4185	5	C ₁₉₄ H ₂₈₇ N ₄₉ O ₆₂ S ₁	[1199-1220]		X	
278	722.1778	722.1833	6			X	X	
279	619.1523	619.1583	7			X		
280	1069.5106	1069.5145	4	C ₁₉₁ H ₂₈₄ N ₄₈ O ₆₂ S ₁	[1204-1226]		X	
281	855.8153	855.8126	5			X	X	
282	713.3400	713.3450	6			X	X	K1220

283	1235.5961	1235.5951	3	C ₁₆₇ H ₂₄₆ N ₄₂ O ₅₂ S ₁	[1204-1226] ⁵⁾		X		
284	926.9425	926.9482	4			X	X		
285	716.3395	716.3474	6	C ₁₉₁ H ₂₈₆ N ₄₈ O ₆₃ S ₁	[1204-1226] ⁷⁾		H		
286	859.4074	859.4153	5				H		
287	724.7014	724.7094	3	C ₉₅ H ₁₅₄ N ₂₆ O ₃₀ S ₁	[1216-1226] ⁵⁾		X		
288	596.0538	596.0587	4	C ₁₀₁ H ₁₆₉ N ₂₉ O ₃₅ S ₁	[1219-1226]	X	X		
289	604.3158	604.3209	3	C ₇₇ H ₁₃₁ N ₂₃ O ₂₅ S ₁	[1219-1226] ⁵⁾	X	X		
290	661.5776	661.5831	4	C ₁₁₄ H ₁₈₃ N ₃₁ O ₃₉ S ₁	[1221-1230]	X	X		K1226
291	691.6829	691.6865	3	C ₉₀ H ₁₄₅ N ₂₅ O ₂₉ S ₁	[1221-1230] ⁵⁾	X	X		
292	1045.1789	1045.1838	3	C ₁₃₅ H ₂₁₃ N ₃₉ O ₄₅ S ₁	[1227-1239]	X	X	K1230	
293	784.1342	784.1397	4			X	X		
294	627.5074	627.5132	5			X	X		
295	855.0922	855.0961	3	C ₁₁₁ H ₁₇₅ N ₃₃ O ₃₅ S ₁	[1227-1239] ⁵⁾	X	X		
296	641.5682	641.5735	4			X	X		
297	551.7740	551.7817	10	C ₂₃₇ H ₃₇₅ N ₇₃ O ₇₇ S ₁	[1227-1260]		X		
298	612.9710	612.9789	9				X		
299	689.4657	689.4752	8				X		
300	787.8270	787.8271	7			X	X		
301	918.9559	918.9644	6				X		
302	706.3595	706.3613	7	C ₂₁₃ H ₃₃₇ N ₆₇ O ₆₇ S ₁	[1227-1260] ⁵⁾		X		
303	618.1919	618.1921	8				X		
304	646.0688	646.0767	4	C ₁₁₁ H ₁₇₇ N ₃₃ O ₃₆ S	[1227-1239] ^{5,7)}		H		
305	517.0551	517.0629	5				H		
306	788.6351	788.6429	4	C ₁₃₅ H ₂₁₅ N ₃₉ O ₄₆ S	[1227-1239] ⁷⁾		H		
307	631.1081	631.1159	5				H		
308	526.0901	526.0979	6				H		
309	620.4356	620.4434	8	C ₂₁₃ H ₃₃₉ N ₆₇ O ₆₈ S ₁	[1227-1260] ^{5,7)}		H		
310	614.9726	614.9800	9	C ₂₃₇ H ₃₇₇ N ₇₃ O ₇₈ S ₁	[1227-1260] ⁷⁾		H		
311	821.4076	821.4122	4	C ₁₃₇ H ₂₂₈ N ₄₀ O ₅₁ S ₁	[1260-1274]	X	X	K1260	
312	904.7876	904.7922	3	C ₁₁₃ H ₁₉₀ N ₃₄ O ₄₁ S ₁	[1260-1274] ⁵⁾	X	X		
313	660.9260	660.9339	5	C ₁₃₇ H ₂₃₀ N ₄₀ O ₅₂ S	[1260-1274] ⁷⁾		H		
314	1019.2111	1019.2093	3	C ₁₃₅ H ₂₂₃ N ₃₅ O ₄₃ S ₁	[1322-1334]	X	X	K1333	
315	764.6542	764.6593	4				X		
316	829.1136	829.1215	3	C ₁₁₁ H ₁₈₅ N ₂₉ O ₃₃ S ₁	[1322-1334] ⁵⁾		X		
317	752.0474	752.0553	3	C ₉₅ H ₁₆₀ N ₂₈ O ₃₃ S ₁	[1329-1334]		X		
318	564.2878	564.2934	4			X	X		
319	561.9631	561.9664	3	C ₉₅ H ₁₆₀ N ₂₈ O ₃₃ S ₁	[1329-1334] ⁵⁾	X			
320	615.5233	615.5311	5	C ₁₃₅ H ₂₂₅ N ₃₅ O ₄₄ S	[1322-1334] ⁷⁾		H		
321	926.0926	926.1015	3	C ₁₂₀ H ₁₈₂ N ₃₂ O ₄₂ S ₁	[1334-1344]		X	K1334	
322	805.6307	805.6346	4	C ₁₄₁ H ₂₁₁ N ₃₉ O ₄₆ S ₁	[1335-1348]	X	X	K1347	
323	550.1286	550.1368	8	C ₁₈₇ H ₂₈₁ N ₅₉ O ₆₃ S ₁	[1366-1388]		X	K1378	

324	638.1296	638.1351	6	C ₁₆₃ H ₂₄₃ N ₅₃ O ₅₃ S ₁	[1366-1388] ⁵⁾	X		
325	665.3010	665.3088	4	C ₁₁₂ H ₁₆₈ N ₃₆ O ₃₈ S ₁	[1371-1384] ⁵⁾		X	
326	915.0206	915.0264	5	C ₂₀₁ H ₂₉₆ N ₅₄ O ₆₅ S ₂	[1435-1460]	X	X	K1436
327	762.6899	762.6899	6			X	X	
328	804.5682	804.5761	5	C ₁₇₇ H ₂₆₀ N ₄₈ O ₅₆ S ₂	[1435-1460] ^{5,7)}		H	
329	918.6212	918.6291	5	C ₂₀₁ H ₂₉₈ N ₅₄ O ₆₆ S ₂	[1435-1460] ⁷⁾		H	
330	800.9677	800.9734	5	C ₁₇₇ H ₂₅₈ N ₄₈ O ₅₅ S ₂	[1435-1460] ⁵⁾	X	X	
331	1000.9573	1000.9655	4			X	X	
332	918.2202	918.2254	5	C ₂₀₁ H ₂₉₆ N ₅₄ O ₆₆ S ₂	[1435-1460] ⁸⁾	X	X	
333	765.3635	765.3557	6			X		
334	572.5379	572.5367	4	C ₉₆ H ₁₅₅ N ₃₁ O ₃₂ S ₁	[1473-1477]	X	X	
335	458.2304	458.2308	5			X	X	
336	572.9528	572.9578	3	C ₇₂ H ₁₁₇ N ₂₅ O ₂₂ S ₁	[1473-1477] ⁵⁾	X		
337	429.9649	429.9710	4			X		
338	747.6392	747.6392	7	C ₂₁₉ H ₃₄₄ N ₆₆ O ₇₇ S ₃	[1474-1504] ⁶⁾		X	K1495
339	872.0774	872.0771	6			X	X	
340	932.2377	932.2387	5	C ₁₉₅ H ₃₀₆ N ₆₀ O ₆₇ S ₃	[1474-1504] ^{6,5)}		X	
341	777.0347	777.0336	6				X	
342	931.8216	931.8275	5	C ₁₉₃ H ₃₀₄ N ₅₆ O ₇₂ S ₃	[1478-1504] ⁶⁾	X		
343	1046.7570	1046.76	7	C ₃₁₄ H ₄₆₃ N ₈₉ O ₁₁₁ S ₂	[1496-1544]	X	X	K1504
344	814.3652	814.3718	9				X	
345	766.3659	766.3738	3	C ₉₂ H ₁₅₃ N ₃₃ O ₃₄ S	[1544-1548] ⁷⁾		H	K1544
346	823.7555	823.7605	3	C ₁₀₅ H ₁₇₇ N ₂₉ O ₃₇ S ₁	[1605-1612]		X	K1609
347	618.0715	618.0718	4			X	X	
348	950.0033	950.0044	2	C ₈₁ H ₁₃₉ N ₂₃ O ₂₇ S ₁	[1605-1612] ⁵⁾		X	
349	633.6669	633.6722	3			X	X	
350	639.6679	639.6657	3	C ₈₁ H ₁₄₁ N ₂₃ O ₂₈ S ₁	[1605-1612] ^{5,7)}		H	
351	622.5671	622.5750	4	C ₁₀₅ H ₁₇₉ N ₂₉ O ₃₈ S	[1605-1612] ⁷⁾		H	
352	653.3223	653.3270	3	C ₈₁ H ₁₃₆ N ₂₄ O ₃₀ S ₁	[1669-1676] ⁵⁾	X		K1669
353	632.8130	632.8139	4	C ₁₀₅ H ₁₇₄ N ₃₀ O ₄₀ S ₁	[1669-1676]	X	X	
354	671.3490	671.3491	5	C ₁₄₁ H ₂₃₈ N ₄₂ O ₅₀ S ₁	[1670-1684]	X	X	K1676
355	838.9298	838.9351	4				X	
356	1118.2451	1118.2437	3		X	X		
357	696.3640	696.3689	4	C ₁₁₇ H ₂₀₀ N ₃₆ O ₄₀ S ₁	[1670-1684] ⁵⁾	X	X	
358	852.7502	852.7501	3	C ₁₀₅ H ₁₇₄ N ₃₂ O ₄₀ S ₁	[1670-1677]	X	X	
359	639.8146	639.8144	4			X	X	
360	662.6572	662.6618	3	C ₈₁ H ₁₃₆ N ₂₆ O ₃₀ S ₁	[1670-1677] ⁵⁾	X		
361	588.0424	588.0499	4	C ₁₀₁ H ₁₆₅ N ₂₇ O ₃₅ S	[1670-1677] ⁷⁾		H	
362	843.4299	843.4378	4	C ₁₄₁ H ₂₄₀ N ₄₂ O ₅₁ S ₁	[1670-1684] ⁷⁾		H	
363	674.9439	674.9518	5				H	
364	800.4025	800.4122	4	C ₁₃₄ H ₂₂₈ N ₄₀ O ₄₆ S ₁	[1677-1690]	X	X	K1684

365	640.531	640.5308	5			X	X			
366	657.8470	657.8459	4	C ₁₁₀ H ₁₉₀ N ₃₄ O ₃₆ S ₁	[1677-1690] ⁵⁾		X			
367	921.4351	921.4404	5	C ₁₉₇ H ₃₀₄ N ₅₈ O ₆₆ S ₂	[1691-1719] ⁶⁾	X	X	K1718		
368	768.0292	768.0349	6			X	X			
369	807.3816	807.3879	5	C ₁₇₃ H ₂₆₆ N ₅₂ O ₅₆ S ₂	[1691-1719] ^{6, 5)}	X				
370	748.8686	748.8772	4	C ₁₃₆ H ₂₀₂ N ₃₈ O ₃₇ S ₁	[1747-1763] ⁵⁾		X	K1747		
371	713.3496	713.3566	5	C ₁₆₀ H ₂₄₀ N ₄₄ O ₄₇ S ₁	[1747-1763]	X	X			
372	891.4396	891.4444	4							X
373	753.3730	753.3808	4	C ₁₃₆ H ₂₀₄ N ₃₈ O ₃₈ S	[1747-1763] ^{5,7)}		H			
374	602.8984	602.9062	5							H
375	716.9513	716.9592	5	C ₁₆₀ H ₂₄₂ N ₄₄ O ₄₈ S	[1747-1763] ⁷⁾		X			
376	983.7432	983.7432	8	C ₃₅₁ H ₅₃₅ N ₉₃ O ₁₁₁ S ₁	[1748-1803]		X	K1792		
377	874.5323	874.5504	9							X
378	1042.5166	1042.5243	7	C ₃₂₇ H ₄₉₇ N ₈₇ O ₁₀₁ S ₁	[1748-1803] ⁵⁾	X	X			
379	788.5763	788.5831	5	C ₁₇₁ H ₂₆₄ N ₄₆ O ₅₉ S ₁	[1783-1803]	X	X			
380	985.4701	985.4777	4							X
381	842.9059	842.9108	4	C ₁₅₃ H ₂₃₈ N ₄₄ O ₅₀ S ₁	[1783-1803] ⁵⁾	X				
382	757.3567	757.3622	4	C ₁₂₇ H ₂₀₀ N ₃₈ O ₄₆ S ₁	[1793-1804]	X	X	K1803		
383	606.0854	606.0912	5						X	X
384	614.7904	614.7960	4	C ₁₀₃ H ₁₆₂ N ₃₂ O ₃₆ S ₁	[1793-1804] ⁵⁾	X	X			
385	485.2308	485.2366	4	C ₇₈ H ₁₂₈ N ₂₈ O ₂₈ S ₁	[1797-1894] ⁵⁾	X				
386	502.4345	502.4443	5	C ₁₀₂ H ₁₆₆ N ₃₄ O ₃₈ S ₁	[1797-1894]	X	X			
387	711.3338	711.3401	4	C ₁₂₆ H ₁₈₄ N ₃₆ O ₃₈ S ₁	[1811-1825] ⁵⁾	X	X		K1817	
388	853.9001	853.9058	4	C ₁₅₀ H ₂₂₂ N ₄₂ O ₄₈ S ₁	[1811-1825]	X				
389	600.9977	601.0028	7	C ₁₈₆ H ₂₇₄ N ₅₀ O ₆₀ S ₁	[1911-1933]	X	X	K1914		
390	700.9961	701.0020	6						X	X
391	840.9955	841.0008	5						X	X
392	1050.9944	1050.9990	4						X	X
393	908.4240	908.4327	4	C ₁₆₂ H ₂₃₄ N ₄₂ O ₅₂ S ₁	[1911-1933] ⁵⁾	X	X			
394	726.9420	726.9478	5						X	X
395	761.4938	761.4997	7	C ₂₃₇ H ₃₄₃ N ₆₁ O ₇₈ S ₁	[1911-1942]		X			
396	888.2435	888.2484	6						X	X
397	1065.6906	1065.6967	5						X	X
398	1331.8702	1331.8690	4						X	
399	793.2019	793.2048	6	C ₂₁₃ H ₃₀₅ N ₅₅ O ₆₈ S ₁	[1911-1942] ⁵⁾	X	X			
400	951.6411	951.6442	5					X	X	
401	730.5455	730.5499	5	C ₁₆₂ H ₂₃₈ N ₄₄ O ₅₁ S ₁	[1911-1933] ^{5,7)}		H			
402	608.9525	608.9595	6						H	
403	844.5950	844.6029	5	C ₁₈₆ H ₂₇₆ N ₅₀ O ₆₁ S ₁	[1911-1933] ⁷⁾		H			
404	703.9959	704.0037	6						H	
405	603.5682	603.5757	7						H	

406	955.2385	955.2463	5	C ₂₁₃ H ₃₀₇ N ₅₅ O ₆₉ S ₁	[1911-1942] ^{5,7)}		H			
407	796.1987	796.2066	6						H	
408	1069.2914	1069.2993	5	C ₂₃₇ H ₃₄₅ N ₆₁ O ₇₉ S ₁	[1911-1942] ⁷⁾		H			
409	891.2429	891.2507	6						H	
410	764.0661	764.0732	7						H	
411	668.6835	668.6900	8						H	
412	815.0279	815.0357	6	C ₂₁₆ H ₃₀₆ N ₅₆ O ₇₃ S ₁	[1915-1942]		X	K1933		
413	863.7853	863.7833	5	C ₁₉₂ H ₂₆₈ N ₅₀ O ₆₃ S ₁	[1915-1942] ⁵⁾	X				
414	1123.2422	1123.2466	4	C ₁₉₂ H ₂₈₅ N ₅₁ O ₆₈ S ₃	[1995-2020] ⁶⁾	X	X	K2012		
415	1307.2352	1307.2386	3	C ₁₆₈ H ₂₄₇ N ₄₅ O ₅₈ S ₃	[1995-2020] ^{6,5)}	X	X			
416	980.6818	980.6809	4							X
417	1312.5663	1312.5703	3	C ₁₆₈ H ₂₄₇ N ₄₅ O ₅₉ S ₃	[1995-2020] ^{6,5,8)}	X				
418	687.5645	687.5693	4	C ₁₁₈ H ₁₇₉ N ₃₃ O ₃₉ S ₂	[2012-2020]	X				
419	795.0286	795.0339	3	C ₁₀₂ H ₁₅₅ N ₂₇ O ₃₇ S ₁	[2001-2013] ⁵⁾	X				
420	1192.0433	1192.0471	2						X	
421	1313.2343	1313.2422	3	C ₁₆₈ H ₂₄₉ N ₄₅ O ₅₉ S ₃	[1995-2020] ^{5,7)}		H			
422	985.1757	985.1835	4						H	
423	902.3935	902.4014	5			C ₁₉₂ H ₂₈₇ N ₅₁ O ₆₉ S ₃	[1995-2020] ⁷⁾		H	
424	768.1187	768.1235	4	C ₁₃₃ H ₂₀₉ N ₃₃ O ₄₈ S ₁	[2021-2032]	X	X	K2023		
425	833.7379	833.7406	3	C ₁₀₉ H ₁₇₁ N ₂₇ O ₃₈ S ₁	[2021-2032] ⁵⁾	X				
426	795.6244	795.6301	4	C ₁₃₃ H ₂₁₅ N ₃₇ O ₄₉ S ₂	[2024-2036]	X	X	K2032		
427	1060.5025	1060.5049	3							X
428	799.6214	799.6293	4	C ₁₃₃ H ₂₁₅ N ₃₇ O ₅₀ S ₂	[2024-2036] ⁸⁾		X			
429	870.4122	870.4160	3	C ₁₀₉ H ₁₇₇ N ₃₁ O ₃₉ S ₂	[2024-2036] ⁵⁾	X	X			
430	640.3106	640.3139	4	C ₁₀₅ H ₁₇₆ N ₃₂ O ₃₈ S ₂	[2029-2036]	X	X			
431	663.3236	663.3278	3	C ₈₁ H ₁₃₈ N ₂₆ O ₂₈ S ₂	[2029-2036] ⁵⁾	X	X			
432	621.0475	621.0463	4	C ₁₀₁ H ₁₆₅ N ₃₃ O ₃₆ S ₂	[2032-2038]		X			
433	657.5592	657.5670	4	C ₁₀₉ H ₁₇₉ N ₃₁ O ₄₀ S ₂	[2024-2036] ⁵⁾		X			
434	800.1254	800.1332	4	C ₁₃₃ H ₂₁₇ N ₃₇ O ₅₀ S ₂	[2024-2036] ^{5,7)}		H			
435	640.3003	640.3082	5						H	
436	1223.3406	1223.3533	4	C ₂₁₃ H ₃₃₃ N ₅₇ O ₇₃ S ₁	[2095-2127]		X	K2100		
437	978.8777	978.8836	5						X	X
438	769.7049	769.7112	6	C ₂₀₈ H ₃₀₆ N ₅₆ O ₆₂ S ₁	[2164-2189]	X	X	K2177		
439	923.4463	923.4520	5						X	X
440	608.4424	608.4426	7	C ₁₉₀ H ₂₈₂ N ₅₂ O ₅₈ S ₁	[2178-2198]		X	K2189		
441	709.6754	709.6818	6						X	X
442	851.4121	851.4166	5						X	X
443	669.3197	669.3249	6	C ₁₈₀ H ₂₆₄ N ₄₈ O ₅₅ S ₁	[2178-2196]	X				
444	860.9156	860.9180	4	C ₁₅₆ H ₂₂₆ N ₄₂ O ₄₅ S ₁	[2178-2196] ⁵⁾		X			
445	688.9298	688.9354	5						X	X
446	640.3123	640.3174	4	C ₁₁₂ H ₁₇₂ N ₃₂ O ₃₅ S ₁	[2178-2190] ⁵⁾	X				

447	686.0792	686.0838	4	C ₁₁₆ H ₁₈₅ N ₃₅ O ₃₈ S ₂	[2197-2206]	X	X	K2198
448	690.0788	690.0831	4	C ₁₁₆ H ₁₈₅ N ₃₅ O ₃₉ S ₂	[2197-2206] ⁸⁾	X	X	
449	729.6816	729.6865	3	C ₉₂ H ₁₄₇ N ₂₉ O ₂₉ S ₂	[2197-2206] ^{5,8)}	X		
450	724.3543	724.3543	3	C ₉₂ H ₁₄₇ N ₂₉ O ₂₈ S ₂	[2197-2206] ⁵⁾	X	X	
451	548.0129	548.0208	4	C ₉₂ H ₁₄₉ N ₂₉ O ₂₉ S ₂	[2197-2206] ^{5,7)}		H	
452	690.5792	690.5870	4	C ₁₁₆ H ₁₈₇ N ₃₅ O ₃₉ S ₂	[2197-2206] ⁷⁾		H	
453	552.6633	552.6711	5				H	
454	694.5779	694.5858	4			C ₁₁₆ H ₁₈₇ N ₃₅ O ₄₀ S ₂	[2197-2206] ^{7,8)}	
455	555.8623	555.8701	5					H
456	771.1319	771.1357	4	C ₁₃₅ H ₂₀₉ N ₃₉ O ₄₂ S ₁	[2297-2308]	X	X	K2307
457	1027.8387	1027.8457	3				X	
458	837.7528	837.7569	3	C ₁₁₁ H ₁₇₁ N ₃₃ O ₃₂ S ₁	[2297-2308] ⁵⁾	X		
459	633.0648	633.0727	4	C ₁₁₁ H ₁₇₃ N ₃₃ O ₃₃ S	[2297-2308] ^{5,7)}		H	
460	775.6311	775.6389	4	C ₁₃₅ H ₂₁₁ N ₃₉ O ₄₃ S	[2297-2308] ⁷⁾		H	
461	620.7048	620.7127	5			H		
462	681.3348	681.3368	4	C ₁₁₇ H ₁₈₄ N ₃₄ O ₃₉ S ₁	[2341-2350]		X	K2341
463	545.271	545.2709	5			X	X	
464	538.7654	538.7711	4	C ₉₃ H ₁₄₆ N ₂₈ O ₂₉ S ₁	[2341-2350] ⁵⁾	X	X	
465	872.9209	872.9280	4	C ₁₅₄ H ₂₃₄ N ₄₂ O ₄₉ S ₁	[2336-2351]	X	X	K2350
466	698.5386	698.5444	5				X	
467	702.1387	702.1465	5	C ₁₅₄ H ₂₃₆ N ₄₂ O ₅₀ S	[2342-2357] ⁷⁾		H	
468	1177.5658	1177.5736	4	C ₂₀₅ H ₃₂₀ N ₅₄ O ₆₉ S ₂	[2351-2376]		X	K2373
469	942.2552	942.2605	5				X	
470	785.3799	785.3850	6				X	
471	763.3593	763.3685	5	C ₁₆₂ H ₂₅₈ N ₄₄ O ₅₈ S ₂	[2358-2376]		X	
472	795.0334	795.0388	3	C ₉₈ H ₁₅₉ N ₂₉ O ₃₆ S ₂	[2363-2374] ^{5,8)}	X		
473	735.0918	735.0984	4	C ₁₂₂ H ₁₉₇ N ₃₅ O ₄₅ S ₂	[2363-2374]	X		
474	695.9886	695.9973	3	C ₈₆ H ₁₄₀ N ₂₄ O ₃₄ S ₁	[2371-2374]		X	
475	945.8584	945.8626	5	C ₂₀₅ H ₃₂₂ N ₅₄ O ₇₀ S ₂	[2351-2376] ⁷⁾		H	
476	788.3812	788.3868	6				H	
477	831.5651	831.5729	6			C ₂₁₅ H ₃₃₉ N ₅₇ O ₇₅ S ₂	[2351-2378] ^{7,8)}	
478	550.7694	550.7692	4	C ₉₁ H ₁₅₀ N ₂₆ O ₃₅ S ₁	[2374-2378]	X		K2376
479	518.5057	518.5091	4	C ₈₆ H ₁₄₃ N ₂₅ O ₃₂ S ₁	[2375-2378]		X	
480	555.2646	555.2724	4	C ₉₁ H ₁₅₂ N ₂₆ O ₃₆ S	[2374-2378] ⁷⁾		H	
481	622.5489	622.5503	4	C ₁₀₂ H ₁₆₃ N ₃₅ O ₃₆ S ₁	[2377-2383]	X	X	K2378
482	498.2405	498.2417	5			X	X	
483	870.8114	870.8119	5	C ₁₈₉ H ₂₈₅ N ₅₅ O ₅₈ S ₃	[2406-2434] ^{5,6)}	X		K2423
484	820.8884	820.8882	6	C ₂₁₃ H ₃₂₃ N ₆₁ O ₆₈ S ₃	[2406-2434] ⁶⁾	X	X	
485	907.2075	907.2154	5	C ₁₉₂ H ₂₉₁ N ₅₇ O ₆₅ S ₃	[2406-2431] ⁶⁾		X	
486	756.17295	756.1808	6			X		
487	760.8677	760.8616	4	C ₁₃₃ H ₂₀₂ N ₃₆ O ₄₄ S ₁	[2432-2443]		X	K2434

488	814.9004	814.9055	4	C ₁₃₆ H ₂₂₆ N ₃₈ O ₅₂ S ₁	[2506-2520]	X	X	K2506
489	655.7207	655.7285	5	C ₁₃₆ H ₂₂₈ N ₃₈ O ₅₃ S	[2506-2520] ⁷⁾		H	
490	901.0845	901.0895	3	C ₁₁₃ H ₁₇₇ N ₃₃ O ₄₂ S ₁	[2632-2640]		X	K2633
491	676.0688	676.0685	4			X	X	
492	541.0452	541.0563	5			X		
493	710.9965	711.0006	3	C ₈₉ H ₁₃₉ N ₂₇ O ₃₂ S ₁	[2632-2640] ⁵⁾	X	X	
494	776.8408	776.8596	4	C ₁₃₅ H ₁₉₈ N ₃₈ O ₄₅ S ₁	[2632-2648] ⁵⁾		X	
495	735.7323	735.7421	5	C ₁₅₉ H ₂₃₆ N ₄₄ O ₅₅ S ₁	[2632-2648]	X	X	
496	919.4194	919.4264	4				X	
497	537.9976	538.0045	4	C ₈₉ H ₁₄₁ N ₂₇ O ₃₃ S ₁	[2632-2640] ^{5,7)}		H	
498	680.5639	680.5717	4	C ₁₁₃ H ₁₇₉ N ₃₃ O ₄₃ S ₁	[2632-2640] ⁷⁾		H	
499	544.6511	544.6589	5				H	
500	694.9178	694.9235	5	C ₁₄₄ H ₂₂₈ N ₄₄ O ₅₀ S ₃	[2719-2734] ⁶⁾	X		K2733
501	725.8292	725.8369	4	C ₁₂₀ H ₁₉₀ N ₃₈ O ₄₀ S ₃	[2719-2734] ^{6,5)}	X	X	
502	581.9518	581.9597	3	C ₇₁ H ₁₂₂ N ₂₄ O ₂₃ S ₂	[2729-2734] ⁵⁾		X	
503	726.7763	726.7827	7	C ₂₂₁ H ₃₃₄ N ₆₆ O ₇₁ S ₁	[2734-2763]		X	K2754
504	726.6769	726.6848	6	C ₁₉₁ H ₂₈₄ N ₅₆ O ₆₀ S ₁	[2735-2763] ⁵⁾		X	
505	633.0611	633.0606	4	C ₁₀₈ H ₁₇₃ N ₃₁ O ₃₅ S ₂	[2773-2780]	X	X	K2779
506	843.7501	843.7462	3				X	
507	637.0525	637.0604	4	C ₁₀₈ H ₁₇₃ N ₃₁ O ₃₆ S ₂	[2773-2780] ⁸⁾		X	
508	653.6531	653.6573	3	C ₈₄ H ₁₃₅ N ₂₅ O ₂₅ S ₂	[2773-2780] ⁵⁾	X	X	
509	649.0696	649.0695	4	C ₁₀₉ H ₁₇₇ N ₃₁ O ₄₀ S ₁	[2780-2787]	X	X	K2780
510	741.6782	741.6836	3	C ₉₂ H ₁₄₇ N ₂₇ O ₃₅ S ₁	[2780-2784]	X	X	
511	556.5066	556.5146	4			X	X	
512	551.5897	551.5953	3	C ₆₈ H ₁₀₉ N ₂₁ O ₂₅ S ₁	[2780-2784] ⁵⁾	X		
513	570.6890	570.6868	5	C ₁₂₀ H ₁₉₇ N ₃₅ O ₄₃ S ₁	[2780-2789]	X	X	K2787
514	713.1091	713.1079	4			X	X	
515	570.5374	570.5422	4	C ₉₆ H ₁₅₉ N ₂₉ O ₃₃ S ₁	[2780-2789] ⁵⁾	X	X	
516	907.7743	907.7765	3	C ₁₁₄ H ₁₈₅ N ₃₃ O ₄₂ S ₁	[2781-2789]	X	X	
517	681.084	681.0842	4			X	X	
518	545.0617	545.0693	5				X	
519	717.6835	717.6887	3	C ₉₀ H ₁₄₇ N ₂₇ O ₃₂ S ₁	[2781-2789] ⁵⁾	X	X	
520	554.0216	554.0276	4	C ₉₂ H ₁₅₃ N ₂₇ O ₃₄ S ₁	[2785-2789]	X	X	
521	548.2741	548.2793	3	C ₆₈ H ₁₁₅ N ₂₁ O ₂₄ S ₁	[2785-2789] ⁵⁾	X	X	
522	574.2838	574.2902	5	C ₁₂₀ H ₁₉₉ N ₃₅ O ₄₄ S	[2780-2789] ⁷⁾		H	
523	685.5795	685.5873	4	C ₁₁₄ H ₁₈₇ N ₃₃ O ₄₃ S	[2781-2789] ⁷⁾		H	
524	548.6636	548.6714	5				H	
525	861.4252	861.4325	4	C ₁₅₀ H ₂₃₆ N ₄₂ O ₄₉ S ₁	[2790-2805]		X	K2804
526	689.3420	689.3470	5			X	X	
527	574.6190	574.6237	6			X	X	
528	718.8617	718.8663	4	C ₁₂₆ H ₁₉₈ N ₃₆ O ₃₉ S ₁	[2790-2805] ⁵⁾	X	X	

529	748.3428	748.3491	6	C ₁₉₅ H ₂₉₄ N ₅₂ O ₆₄ S ₃	[2845-2867]	X	X	K2852
530	897.8124	897.8175	5			X	X	
531	901.0119	901.0165	5	C ₁₉₅ H ₂₉₄ N ₅₂ O ₆₅ S ₃	[2845-2867] ⁸⁾	X	X	
532	979.4478	979.4539	4	C ₁₇₁ H ₂₅₆ N ₄₆ O ₅₄ S ₃	[2845-2867] ⁵⁾	X	X	
533	783.7573	783.7651	5			X	X	
534	872.4211	872.4246	3	C ₁₁₂ H ₁₇₉ N ₃₁ O ₃₇ S ₂	[2845-2857] ⁵⁾	X	X	
535	797.1331	797.1365	4	C ₁₃₆ H ₂₁₇ N ₃₇ O ₄₇ S ₂	[2845-2857]	X		
536	1027.4949	1027.4937	5	C ₂₂₂ H ₃₄₆ N ₆₀ O ₇₄ S ₃	[2853-2882]	X	X	K2874
537	856.4172	856.4126	6			X	X	
538	734.2055	734.2123	7				X	
539	736.4999	736.4974	7	C ₂₂₂ H ₃₄₆ N ₆₀ O ₇₅ S ₃	[2853-2882] ⁸⁾		X	
540	913.4352	913.4412	5	C ₁₉₈ H ₃₀₈ N ₅₄ O ₆₄ S ₃	[2853-2882] ⁵⁾	X	X	
541	761.3627	761.3690	6			X	X	
542	799.6550	799.6605	4	C ₁₃₆ H ₂₂₇ N ₃₇ O ₄₉ S ₁	[2868-2882]	X	X	
543	875.7863	875.7899	3	C ₁₁₂ H ₁₈₉ N ₃₁ O ₃₉ S ₁	[2868-2882] ⁵⁾	X	X	
544	661.5896	661.5975	4	C ₁₁₂ H ₁₉₁ N ₃₁ O ₄₀ S ₁	[2868-2882] ^{5,7)}		H	
545	804.1558	804.1637	4	C ₁₃₆ H ₂₂₉ N ₃₇ O ₅₀ S ₁	[2868-2882] ⁷⁾		H	
546	643.5247	643.5325	5				H	
547	721.1510	721.1589	5	C ₁₅₃ H ₂₄₉ N ₄₃ O ₅₃ S ₂	[2875-2893]		X	K2882
548	758.6255	758.6304	4	C ₁₂₉ H ₂₁₁ N ₃₇ O ₄₃ S ₂	[2875-2893] ⁵⁾	X	X	
549	810.3817	810.3806	4	C ₁₃₅ H ₂₁₂ N ₄₂ O ₄₇ S ₂	[2883-2897]	X	X	K2893
550	534.4468	534.4535	5	C ₁₁₁ H ₁₇₄ N ₃₆ O ₃₇ S ₂	[2883-2897] ⁵⁾	X		
551	667.8070	667.8149	4			X		
552	496.4720	496.4776	4	C ₈₁ H ₁₂₃ N ₂₉ O ₂₈ S ₁	[2888-2896] ⁵⁾	X	X	
553	517.2222	517.2278	4	C ₈₄ H ₁₂₄ N ₃₀ O ₃₀ S ₁	[2897-2904]	X		K2897
554	538.5373	538.5460	7	C ₁₅₈ H ₂₄₃ N ₅₃ O ₅₃ S ₁	[2894-2910]		X	
555	674.3219	674.3275	5	C ₁₄₀ H ₂₂₃ N ₄₅ O ₅₀ S ₁	[2910-2924]	X	X	K2910
556	562.1007	562.1080	6				X	
557	481.9446	481.9508	7				X	
558	700.0848	700.0919	4	C ₁₁₆ H ₁₈₅ N ₃₉ O ₄₀ S ₁	[2910-2924] ⁵⁾	X		
559	855.3878	855.3939	6	C ₂₂₂ H ₃₂₈ N ₆₂ O ₇₅ S ₂	[2925-2955] ⁶⁾	X	X	K2929
560	926.0916	926.0959	3	C ₁₁₆ H ₁₈₂ N ₃₂ O ₄₅ S ₁	[2925-2934]	X		
561	1103.5052	1103.5077	2	C ₉₂ H ₁₄₄ N ₂₆ O ₃₅ S ₁	[2925-2934] ⁵⁾	X		
562	736.0046	736.0076	3			X		
563	994.0436	994.0474	5	C ₂₁₁ H ₃₁₃ N ₆₁ O ₇₅ S ₂	[2930-2959] ⁶⁾	X	X	K2955
564	828.5431	828.5405	6			X	X	
565	710.3166	710.3215	7			X	X	
566	733.4903	733.4969	6	C ₁₈₇ H ₂₇₅ N ₅₅ O ₆₅ S ₂	[2930-2959] ^{6,5)}	X		
567	879.9897	879.9948	5			X	X	
568	1099.7371	1099.7415	4			X		
569	712.8871	712.8950	7	C ₂₁₁ H ₃₁₅ N ₆₁ O ₇₆ S ₂	[2930-2959] ^{6,7)}		H	

570	930.443	930.4433	6	C ₂₅₂ H ₃₆₂ N ₆₆ O ₇₇ S ₁	[2987-3021]	X	X	K2987
571	1116.3256	1116.3304	5			X	X	
572	1395.1649	1395.1613	4			X	X	
573	912.7641	912.7693	3	C ₁₁₈ H ₁₈₂ N ₃₂ O ₄₁ S ₁	[3038-3047]	X	X	K3044
574	684.8226	684.8293	4		X			
575	1083.5169	1083.5178	2	C ₉₄ H ₁₄₄ N ₂₆ O ₃₁ S ₁	[3038-3047] ⁵⁾	X	X	
576	722.6833	722.6815	3			X		
577	841.0538	841.0609	3	C ₁₀₉ H ₁₆₅ N ₂₉ O ₃₈ S ₁	[3038-3047] ⁹⁾	X		
578	842.7167	842.7211	3	C ₁₀₅ H ₁₆₈ N ₂₈ O ₄₀ S ₂	[3087-3094]	X	X	K3087
579	848.0482	848.0533	3	C ₁₀₅ H ₁₆₈ N ₂₈ O ₄₁ S ₂	[3087-3094] ⁸⁾	X	X	
580	978.4405	978.4455	2	C ₈₁ H ₁₃₀ N ₂₂ O ₃₀ S ₂	[3087-3094] ⁵⁾	X	X	
581	652.6338	652.6333	3			X		
582	986.4429	986.4435	2	C ₈₁ H ₁₃₀ N ₂₂ O ₃₁ S ₂	[3087-3094] ^{5,8)}		X	
583	657.9644	657.9650	3			X		
584	935.1105	935.1184	3	C ₁₁₉ H ₁₉₁ N ₃₃ O ₄₁ S ₂	[3078-3094] ⁵⁾		X	
585	940.4458	940.4500	3	C ₁₁₉ H ₁₉₁ N ₃₃ O ₄₂ S ₂	[3078-3094] ^{5,8)}	X	X	
586	848.1564	848.1557	4	C ₁₄₃ H ₂₂₉ N ₃₉ O ₅₂ S ₂	[3078-3094] ⁸⁾		X	
587	844.1477	844.1564	4	C ₁₄₃ H ₂₂₉ N ₃₉ O ₅₁ S ₂	[3078-3094]	X	X	
588	1027.5074	1027.5261	3	C ₁₃₇ H ₂₁₄ N ₃₈ O ₄₁ S ₁	[3112-3122]		X	K3121
589	770.8859	770.8965	4				X	
590	837.4347	837.4373	3	C ₁₁₃ H ₁₇₆ N ₃₂ O ₃₁ S ₁	[3112-3122] ⁵⁾	X		
591	528.7609	528.7686	4	C ₈₇ H ₁₄₆ N ₂₈ O ₃₁ S ₁	[3119-3122]		X	
592	772.1672	772.1728	5	C ₁₆₈ H ₂₅₄ N ₄₈ O ₅₅ S ₁	[3170-3192] ⁵⁾	X		K3173
593	1107.5271	1107.5309	4	C ₁₉₂ H ₂₉₂ N ₅₄ O ₆₅ S ₁	[3170-3192]		X	
594	886.2199	886.2258	5			X	X	
595	738.6832	738.6893	6			X	X	
596	780.1026	780.1085	4	C ₁₃₅ H ₁₉₇ N ₃₉ O ₄₅ S ₁	[3170-3186] ⁵⁾	X		
597	775.7676	775.7755	5	C ₁₆₈ H ₂₅₆ N ₄₈ O ₅₆ S ₁	[3170-3192] ^{5,7)}		H	
598	741.6838	741.6917	6	C ₁₉₂ H ₂₉₄ N ₅₄ O ₆₆ S ₁	[3170-3192] ⁷⁾		H	
599	691.0121	691.0128	3	C ₈₅ H ₁₄₃ N ₂₇ O ₃₁ S ₁	[3170-3192]		X	K3197
600	518.5114	518.5114	4				X	
601	418.6054	418.6132	5			C ₈₅ H ₁₄₅ N ₂₇ O ₃₂ S ₁	[3196-3199] ⁷⁾	
602	897.1288	897.1337	3	C ₁₁₇ H ₁₉₃ N ₃₁ O ₃₉ S ₁	[3211-3220]	X	X	K3216
603	673.0966	673.1021	4			X	X	
604	707.0405	707.0454	3	C ₉₃ H ₁₅₅ N ₂₅ O ₂₉ S ₁	[3211-3220] ⁵⁾	X	X	
605	713.0416	713.0494	3	C ₉₃ H ₁₅₅ N ₂₅ O ₂₉ S ₁	[3211-3220] ^{5,7)}		H	
606	677.5974	677.6053	4	C ₁₁₇ H ₁₉₃ N ₃₁ O ₃₉ S ₁	[3211-3220] ⁷⁾		H	
607	542.2779	542.2858	5				H	
608	1155.5181	1155.5214	4	C ₁₉₈ H ₂₉₆ N ₅₂ O ₇₂ S ₂	[3221-3248]		X	
609	953.9506	953.9567	4	C ₁₆₉ H ₂₅₄ N ₄₄ O ₅₃ S ₂	[3229-3248]	X		
610	957.9491	957.9554	4	C ₁₆₉ H ₂₅₄ N ₄₄ O ₅₄ S ₂	[3229-3248] ⁸⁾	X		

611	770.3733	770.3792	3	C ₁₀₁ H ₁₅₇ N ₂₇ O ₃₃ S ₁	[3229-3241] ⁵⁾	X	X		
612	1155.0667	1155.0652	2			X			
613	841.0892	841.0883	3	C ₁₁₀ H ₁₇₃ N ₃₁ O ₃₅ S ₁	[3270-3276]	X	X	K3275	
614	631.068	631.0681	4			X	X		
615	505.0491	505.0565	5		X				
616	650.9940	651.0005	3	C ₈₆ H ₁₃₅ N ₂₅ O ₂₅ S ₁	[3270-3276] ⁵⁾	X	X		
617	488.4945	488.5024	4				X		
618	555.2761	555.2757	4	C ₉₅ H ₁₅₂ N ₂₆ O ₃₃ S ₁	[3272-3276]	X	X		
619	549.9365	549.9440	3	C ₇₁ H ₁₁₄ N ₂₀ O ₂₃ S ₁	[3272-3276] ⁵⁾	X	X		
620	635.5634	635.5712	4	C ₁₁₀ H ₁₇₅ N ₃₁ O ₃₆ S ₁	[3270-3276] ⁷⁾		H		
621	508.6507	508.6586	5				H		
622	783.9791	783.9850	7	C ₂₄₃ H ₃₉₀ N ₆₆ O ₇₆ S ₁	[3276-3310]		X	K3276	
623	914.488	914.4807	6			X	X		
624	1097.1795	1097.1753	5		X	X			
625	819.4311	819.4370	6	C ₂₁₉ H ₃₅₂ N ₆₀ O ₆₆ S ₁	[3276-3310] ⁵⁾	X	X		
626	1228.6438	1228.6516	4				X		
627	926.9636	926.9707	4	C ₁₆₁ H ₂₅₈ N ₄₄ O ₅₄ S ₁	[3276-3294]		X		
628	741.7731	741.7776	5			X	X		
629	917.4802	917.4830	6	C ₂₄₃ H ₃₉₂ N ₆₆ O ₇₇ S ₁	[3276-3310] ⁷⁾		H		
630	786.5534	786.5579	7				H		
631	781.5326	781.5407	8	C ₂₇₈ H ₄₄₇ N ₇₅ O ₈₆ S ₁	[3276-3317]		X	K3310	
632	893.0420	893.0454	7			X			
633	826.2252	826.2304	5	C ₁₈₁ H ₂₉₂ N ₅₀ O ₅₈ S ₁	[3295-3317]	X	X		
634	1032.5346	1032.5363	4			X	X		
635	889.9662	889.9700	4	C ₁₅₇ H ₂₅₄ N ₄₄ O ₄₈ S ₁	[3295-3317] ⁵⁾		X		
636	1020.3621	1020.3692	6	C ₂₇₂ H ₄₃₅ N ₇₃ O ₈₅ S ₁	[3277-3317]		X		
637	874.7403	874.7456	7			X	X		
638	829.8252	829.8331	5	C ₁₈₁ H ₂₉₄ N ₅₀ O ₅₉ S ₁	[3295-3317]		X		
639	694.6140	694.6141	4	C ₁₂₀ H ₁₉₉ N ₃₃ O ₄₀ S ₁	[3311-3321]	X	X	K3317	
640	962.2581	962.2645	5	C ₂₀₇ H ₃₃₂ N ₅₄ O ₆₇ S ₅	[3322-3350]		X	K3325	
641	968.6546	968.6625	5	C ₂₀₇ H ₃₃₂ N ₅₄ O ₆₉ S ₅	[3322-3350] ⁸⁾		X		
642	965.4556	965.4635	5	C ₂₀₇ H ₃₃₂ N ₅₄ O ₆₈ S ₅	[3322-3350] ⁸⁾		X		
643	968.6537	968.6625	5	C ₂₀₇ H ₃₃₂ N ₅₄ O ₆₉ S ₅	[3322-3350] ⁸⁾	X	X		
644	942.4276	942.4312	3	C ₁₁₈ H ₁₈₉ N ₃₁ O ₄₃ S ₃	[3323-3333]	X	X		
645	969.0578	969.0656	5	C ₂₀₇ H ₃₃₄ N ₅₄ O ₆₉ S ₅	[3322-3350] ^{7,8)}		H		
646	1164.0449	1164.0432	4	C ₂₀₀ H ₃₁₈ N ₅₀ O ₆₅ S ₆	[3326-3352] ⁶⁾	X	X	K3350	
647	1168.0371	1168.0425	4	C ₂₀₀ H ₃₁₈ N ₅₀ O ₆₆ S ₆	[3326-3352] ^{6,8)}	X			
648	833.6492	833.6523	4	C ₁₄₈ H ₂₂₃ N ₃₉ O ₄₅ S ₂	[3351-3365] ⁶⁾	X	X	K3352	
649	667.1186	667.1233	5			X	X		
650	691.0803	691.0866	4	C ₁₂₄ H ₁₈₅ N ₃₃ O ₃₅ S ₂	[3351-3365] ^{6,5)}	X	X		
651	1148.6885	1148.6906	6	C ₂₉₉ H ₄₄₅ N ₇₉ O ₁₀₁ S ₄	[3353-3398]	X	X	K3365	

652	984.739	984.7358	7			X	X
653	1192.5357	1192.5397	5	C ₂₅₈ H ₃₈₅ N ₆₇ O ₈₈ S ₄	[3353-3390] ⁸⁾	X	
654	832.0572	832.0583	3	C ₁₀₅ H ₁₆₈ N ₂₈ O ₃₈ S ₂	[3363-3370]	X	X
655	837.3845	837.3900	3	C ₁₀₅ H ₁₆₈ N ₂₈ O ₃₉ S ₂	[3363-3370] ⁸⁾	X	X
656	962.4481	962.4506	2	C ₈₁ H ₁₃₀ N ₂₂ O ₂₈ S ₂	[3363-3370] ⁵⁾	X	X
657	641.9653	641.9700	3			X	X

¹⁾ Numbers inside brackets correspond to the position of the proteolytic peptide within the sequence of KLH2. All peptides from KLH2 shown in this table are linked to (C¹-K¹⁶) fragment of Cys¹pP0, except those peptides where it is specified if they are linked to (C¹-K¹¹), (C¹-K¹⁴) or (C¹-K¹⁷) fragment of Cys¹pP0.

²⁾ Identification of type 2 peptides in the KLH2-Cys¹pP0 conjugate using pLink2 software [2].

³⁾ Identification of type 2 peptides in the KLH2-Cys¹pP0 conjugate using Kojak software [3].

⁴⁾ Indicates the position where the peptide Cys¹pP0 is added by the Michael addition to the free amino groups in the Lys residues or *N*-terminus of the KLH2.

⁵⁾ Peptide from KLH1 linked to (C¹-K¹¹) fragment of Cys¹pP0

⁶⁾ Carbamidomethylation of Cys

⁷⁾ Identification of type 2 peptides with the hydrolyzed linker

⁸⁾ Met→ Met sulfoxide

⁹⁾ Peptide from KLH1 linked to (C¹-K¹⁴) fragment of Cys¹pP0

Table S7. Identification of conjugation sites in KLH2-Cys¹pP0 conjugate using Kojak [2] and pLink2 [3] software. Type 2 peptides were identified using transcyclized and hydrolyzer linker.

#	<i>m/z</i> exp.	<i>m/z</i> theor.	<i>z</i>	Elemental composition	Assignment ¹⁾	pLink ²⁾	Kojak ³⁾	Conjugation site ⁴⁾	
1	907.4576	907.4655	2	C ₇₅ H ₁₂₈ N ₂₄ O ₂₆ S	[1-7] ⁵⁾		X	Nt	
2	605.3050	605.3129	3				X		
3	795.3933	795.4012	3	C ₉₉ H ₁₆₆ N ₃₀ O ₃₆ S	[1-7]		X		
4	596.7950	596.8029	4				X		
5	477.6360	477.6439	5				X		
6	601.2979	601.3055	4	C ₉₉ H ₁₆₆ N ₃₀ O ₃₇ S	[1-7] ⁶⁾		H		
7	503.2550	503.2628	5	C ₁₀₅ H ₁₇₈ N ₃₂ O ₃₇ S	[1-7] ⁷⁾		X		
8	854.1786	854.1860	4	C ₁₄₃ H ₂₄₁ N ₄₁ O ₅₃ S	[1-22] ⁵⁾		X	K7	
9	996.7448	996.7527	4	C ₁₆₇ H ₂₇₉ N ₄₇ O ₆₃ S	[1-22]		X		
10	797.5958	797.6037	5				X		
11	801.1958	801.2058	5	C ₁₆₇ H ₂₈₁ N ₄₇ O ₆₄ S	[1-22] ⁶⁾		H		
12	1045.9611	1045.9690	2	C ₈₃ H ₁₃₅ N ₂₅ O ₃₆ S	[7-17] ⁵⁾		X		
13	887.7290	887.7369	3	C ₁₀₇ H ₁₇₃ N ₃₁ O ₄₆ S	[7-17]		X		
14	915.44443	915.4523	3	C ₁₁₄ H ₁₉₀ N ₃₂ O ₄₄ S	[7-22] ⁵⁾	X	X		
15	829.4017	829.4074	4	C ₁₃₈ H ₂₂₈ N ₃₈ O ₅₄ S	[7-22]	X	X		
16	1477.4248	1477.4322	4	C ₂₄₇ H ₃₈₉ N ₆₉ O ₉₃ S ₃	[8-52] ^{5,8)}		X	K22	
17	1176.0241	1176.0320	4	C ₁₉₆ H ₃₀₂ N ₅₈ O ₇₁ S ₃	[23-55] ^{5,8)}		X	K52	
18	941.01926	941.0272	5				X		
19	944.2197	944.2261	5	C ₁₉₆ H ₃₀₂ N ₅₈ O ₇₂ S ₃	[23-55] ^{5,8,9)}	X			
20	1318.5976	1318.5983	4	C ₂₂₀ H ₃₄₀ N ₆₄ O ₈₁ S ₃	[23-55] ⁸⁾	X	X		
21	1055.0799	1055.0801	5				X		X
22	879.4011	879.4014	6				X		X
23	756.4802	756.4896	7	C ₂₂₀ H ₃₄₂ N ₆₄ O ₈₂ S ₃	[23-55] ^{8,6)}		H		
24	1058.2797	1058.2792	5	C ₂₂₀ H ₃₄₀ N ₆₄ O ₈₂ S ₃	[23-55] ^{8,9)}	X	X		
25	882.0594	882.0673	6				X	X	
26	763.3396	763.3475	3	C ₉₆ H ₁₄₂ N ₃₂ O ₃₂ S	[117-127] ⁵⁾		X	K119	
27	572.7547	572.7625	4				X		X
28	715.32192	715.3288	4	C ₁₂₀ H ₁₈₀ N ₃₈ O ₄₂ S	[117-127]	X			
29	572.45842	572.4646	5				X		
30	1026.1641	1026.1619	3	C ₁₃₄ H ₂₀₂ N ₄₀ O ₄₂ S ₁	[120-136] ⁵⁾	X	X	K135	
31	769.8740	769.8734	4				X		X
32	616.1002	616.1002	5				X		X
33	1216.2505	1216.2501	3	C ₁₅₈ H ₂₄₀ N ₄₆ O ₅₂ S ₁	[120-136]	X	X		
34	912.4400	912.4396	4				X		X
35	730.1538	730.1533	5				X		X
36	608.6288	608.6290	6				X		X
37	619.6924	619.7023	5	C ₁₃₄ H ₂₀₄ N ₄₀ O ₄₃ S ₁	[120-136] ^{5,6)}		H		
38	916.9351	916.9422	4	C ₁₅₈ H ₂₄₂ N ₄₆ O ₅₃ S ₁	[120-136] ⁶⁾		H		
39	733.7454	733.7554	5					H	
40	611.6211	611.6307	6					H	
41	1024.0102	1024.0181	2	C ₈₈ H ₁₄₃ N ₂₅ O ₂₉ S ₁	[128-136] ⁵⁾	X			
42	683.0157	683.0146	3				X		
43	873.1030	873.1030	3	C ₁₁₂ H ₁₈₁ N ₃₁ O ₃₉ S ₁	[128-136]	X			

44	655.0790	655.0792	4			X	X	
45	515.4950	515.5028	4	C ₈₃ H ₁₃₉ N ₂₇ O ₃₂ S	[136-139]		X	K136
46	641.3182	641.3242	4	C ₁₀₇ H ₁₇₆ N ₃₄ O ₃₇ S	[136-148] ⁵⁾	X		
47	513.2546	513.2609	5			X		
48	627.3134	627.3140	5	C ₁₃₁ H ₂₁₄ N ₄₀ O ₄₇ S ₁	[136-148]	X	X	
49	522.9293	522.9296	6			X	X	
50	873.9180	873.9258	6	C ₂₂₂ H ₃₅₃ N ₆₇ O ₇₄ S ₃	[137-172] ^{5,8)}	X		K148
51	876.5838	876.5917	6	C ₂₂₂ H ₃₅₃ N ₆₇ O ₇₅ S ₃	[137-172] ^{5,8,9)}		X	
52	968.9621	968.9700	6	C ₂₄₆ H ₃₉₁ N ₇₃ O ₈₄ S ₃	[137-172] ⁸⁾		X	
53	832.9668	832.9747	7	C ₂₄₆ H ₃₉₁ N ₇₃ O ₈₅ S ₃	[137-172] ^{8,9)}	X	X	
54	596.2871	596.2949	6	C ₁₅₁ H ₂₃₈ N ₄₈ O ₅₁ S ₁	[140-156]		X	
55	612.3069	612.3071	4	C ₁₀₅ H ₁₆₄ N ₃₄ O ₃₂ S ₁	[145-156] ⁵⁾	X	X	
56	490.0476	490.0473	5			X	X	
57	754.8726	754.8734	4	C ₁₂₉ H ₂₀₂ N ₄₀ O ₄₂ S ₁	[145-156]	X	X	
58	604.0999	604.1002	5			X	X	
59	503.5841	503.5848	6			X	X	
60	526.2548	526.2612	5	C ₁₀₉ H ₁₇₅ N ₃₇ O ₃₇ S	[148-156]	X		
61	876.9180	876.9276	6	C ₂₂₂ H ₃₅₅ N ₆₇ O ₇₅ S ₃	[137-172] ^{6,5)}		H	
62	971.9621	971.9717	6	C ₂₄₆ H ₃₉₃ N ₇₃ O ₈₅ S ₃	[137-172] ⁶⁾		H	
63	616.8019	616.8098	4	C ₁₀₅ H ₁₆₆ N ₃₄ O ₃₃ S ₁	[145-156] ^{6,5)}		H	
64	493.6415	493.6494	5				H	
65	759.3677	759.3760	4	C ₁₂₉ H ₂₀₄ N ₄₀ O ₄₃ S ₁	[145-156] ⁶⁾		H	
66	607.6924	607.7023	5				H	
67	506.5770	506.5866	6				H	
68	905.6509	905.6588	4	C ₁₅₀ H ₂₃₁ N ₄₇ O ₅₂ S ₃	[230-252] ^{5,8)}		X	K231
69	724.7207	724.7286	5			X	X	
70	727.9207	727.9276	5	C ₁₅₀ H ₂₃₁ N ₄₇ O ₅₃ S ₃	[230-252] ^{5,8,9)}	X		
71	838.7754	838.7816	5	C ₁₇₄ H ₂₆₉ N ₅₃ O ₆₂ S ₃	[230-252] ⁸⁾	X	X	
72	701.8110	701.8184	6	C ₁₇₄ H ₂₆₉ N ₅₃ O ₆₃ S ₃	[230-252] ^{8,9)}	X	X	
73	775.0266	775.0275	6	C ₁₉₃ H ₃₀₂ N ₆₀ O ₆₈ S ₃	[230-261] ^{5,8)}	X	X	
74	870.0695	870.0717	6	C ₂₁₇ H ₃₄₀ N ₆₆ O ₇₈ S ₃	[230-261] ⁸⁾	X	X	
75	872.7302	872.7375	6	C ₂₁₇ H ₃₄₀ N ₆₆ O ₇₉ S ₃	[230-261] ^{8,9)}	X		
76	623.2757	623.2836	4	C ₁₀₅ H ₁₅₆ N ₃₂ O ₃₇ S	[262-274] ⁵⁾		X	K265
77	765.8486	765.8490	4	C ₁₂₉ H ₁₉₄ N ₃₈ O ₄₇ S ₁	[262-274]	X	X	
78	612.8839	612.8814	5			X	X	
79	627.7788	627.7862	4	C ₁₀₅ H ₁₅₈ N ₃₂ O ₃₈ S	[262-274] ^{5,6)}		H	
80	502.4230	502.4305	5				H	
81	770.3451	770.3525	4	C ₁₂₉ H ₁₉₆ N ₃₈ O ₄₈ S ₁	[262-274] ⁶⁾		H	
82	616.4761	616.4835	5				H	
83	513.8967	513.9042	6				H	
84	524.5907	524.5919	3	C ₆₃ H ₁₀₆ N ₂₂ O ₂₃ S ₁	[300-304] ⁵⁾		X	K302
85	720.6757	720.6837	3	C ₈₇ H ₁₄₆ N ₂₈ O ₃₄ S	[300-304] ⁶⁾		H	
86	540.7568	540.7648	4				H	
87	432.8054	432.8134	5				H	
88	1063.8282	1063.8361	3	C ₁₄₅ H ₂₀₉ N ₃₇ O ₄₁ S ₂	[340-358] ⁵⁾		X	
89	798.1291	798.1290	4			X	X	
90	1253.9211	1253.9244	3	C ₁₆₉ H ₂₄₇ N ₄₃ O ₅₁ S ₂	[340-358]	X	X	

91	940.6943	940.6953	4			X	X	K353																		
92	752.7499	752.7578	5				X		K353																	
93	944.6950	944.6953	4	C ₁₆₉ H ₂₄₇ N ₄₃ O ₅₂ S ₂	[340-358] ⁹⁾	X	X			K353																
94	981.1113	981.1191	6	C ₂₆₆ H ₃₇₄ N ₆₈ O ₈₁ S ₂	[340-382] ⁵⁾	X	X				K353															
95	841.0954	841.1032	7										X	K353												
96	1076.1554	1076.1633	6	C ₂₉₀ H ₄₁₂ N ₇₄ O ₉₁ S ₂	[340-382]	X	X					K353														
97	922.5618	922.5696	7										X		X	K353										
98	807.3666	807.3745	8												X		K353									
99	1078.8212	1078.8291	6												X			K353								
100	924.8478	924.8546	7	C ₂₉₀ H ₄₁₂ N ₇₄ O ₉₂ S ₂	[340-382] ⁹⁾	X	X						K353													
101	809.3659	809.3738	8																X	K353						
102	662.3027	662.3106	3			C ₈₈ H ₁₂₉ N ₂₅ O ₂₄ S ₂	[352-358] ⁵⁾								X				X		K353					
103	667.6369	667.6423	3	C ₈₈ H ₁₂₉ N ₂₅ O ₂₅ S ₂	[352-358] ^{5,9)}	X	X								K353											
104	643.5420	643.5499	4	C ₁₁₂ H ₁₆₇ N ₃₁ O ₃₅ S ₂	[352-358] ⁹⁾	X	X												K353							
105	945.1905	945.1979	4	C ₁₆₉ H ₂₄₉ N ₄₃ O ₅₂ S ₂	[340-358] ⁶⁾		H															K353				
106	756.3524	756.3599	5																					H	K353	
107	759.5511	759.5589	5	C ₁₆₉ H ₂₄₉ N ₄₃ O ₅₃ S ₂	[340-358] ^{6,9)}		H																K353			
108	984.1128	984.1209	6	C ₂₆₆ H ₃₇₆ N ₆₈ O ₈₂ S ₂	[340-382] ^{5,6)}		H																	K353		
109	756.7852	756.7921	5	C ₁₆₅ H ₂₆₃ N ₄₉ O ₅₁ S	[407-427]	X		K409																		
110	659.6284	659.6354	7	C ₂₀₂ H ₃₂₄ N ₆₂ O ₆₀ S ₁	[400-427]		X	K414																		
111	729.1179	729.1257	4	C ₁₂₇ H ₂₀₁ N ₃₉ O ₃₈ S	[410-427] ⁵⁾		X		K414																	
112	871.6841	871.6920	4	C ₁₅₁ H ₂₃₉ N ₄₅ O ₄₈ S	[410-427]		X			K414																
113	697.5500	697.5551	5								X			X												K414
114	1014.5306	1014.5300	3	C ₁₃₃ H ₂₁₃ N ₄₁ O ₃₉ S ₁	[410-428] ⁵⁾		X				K414															
115	761.1416	761.1495	4											X		K414										
116	609.1210	609.1212	5									X		X			K414									
117	1204.6104	1204.6183	3											X				K414								
118	903.7161	903.7157	4	C ₁₅₇ H ₂₅₁ N ₄₇ O ₄₉ S ₁	[410-428]	X	X					K414														
119	723.1746	723.1741	5										X	X						K414						
120	602.8055	602.8130	6										X	X							K414					
121	516.8330	516.8409	7											X	K414											
122	765.6441	765.6521	4	C ₁₃₃ H ₂₁₅ N ₄₁ O ₄₀ S ₁	[410-428] ^{5,6)}		H						K414													
123	510.7628	510.7707	6																H			K414				
124	908.2117	908.2183	4	C ₁₅₇ H ₂₅₃ N ₄₇ O ₅₀ S ₁	[410-428] ⁶⁾		H							K414												
125	726.7694	726.7762	5																				H		K414	
126	605.8078	605.8148	6																				H	K414		
127	519.4067	519.4138	7																				H			
128	922.1135	922.1123	3			C ₁₁₅ H ₁₈₆ N ₃₄ O ₄₃ S ₁	[428-442] ⁵⁾	X											X				K428			
129	834.3945	834.4024	4	C ₁₃₉ H ₂₂₄ N ₄₀ O ₅₃ S	[428-442]		X	K428																		
130	1114.2825	1114.2806	4	C ₁₈₆ H ₂₉₇ N ₅₅ O ₆₈ S ₂	[429-452]	X	X		K443																	
131	743.1893	743.1898	6					X											K443							
132	551.2623	551.2629	4	C ₈₉ H ₁₄₈ N ₂₈ O ₃₃ S ₂	[443-447]		X	K443																		
133	555.7566	555.7656	4	C ₈₉ H ₁₅₀ N ₂₈ O ₃₄ S ₂	[443-447] ⁶⁾		H			K443																
134	444.8053	444.8140	5													H	K443									
135	555.2615	555.2617	4	C ₈₉ H ₁₄₈ N ₂₈ O ₃₄ S ₂	[443-447] ⁹⁾		X				K443															
136	961.2509	961.2588	5	C ₂₀₄ H ₃₁₇ N ₆₃ O ₆₆ S ₃	[444-476] ^{5,8)}	X	X																			
137	801.2091	801.2170	6									X				X										

138	1075.3039	1075.3118	5	C ₂₂₈ H ₃₅₅ N ₆₉ O ₇₆ S ₃	[444-476] ⁸⁾		X	K452	
139	896.2532	896.2611	6						X
140	768.3610	768.3678	7				X		X
141	898.9203	898.9269	6	C ₂₂₈ H ₃₅₅ N ₆₉ O ₇₇ S ₃	[444-476] ^{8,9)}	X	X		
142	770.6460	770.6528	7			X	X		
143	674.4393	674.4471	8				X		
144	873.4177	873.4171	4	C ₁₅₁ H ₂₂₈ N ₄₄ O ₅₀ S ₁	[448-469] ⁵⁾	X	X		
145	1015.9854	1015.9833	4	C ₁₇₅ H ₂₆₆ N ₅₀ O ₆₀ S ₁	[448-469]	X	X		
146	812.9876	812.9882	5			X	X		
147	899.2565	899.2629	6	C ₂₂₈ H ₃₅₇ N ₆₉ O ₇₇ S ₃	[444-476] ^{6,8)}		H		
148	674.6924	674.6991	8				H		
149	877.9097	877.9197	4	C ₁₅₁ H ₂₃₀ N ₄₄ O ₅₁ S ₁	[448-469] ^{5,6)}		H		
150	702.5277	702.5374	5				H		
151	816.5836	816.5903	5	C ₁₇₅ H ₂₆₈ N ₅₀ O ₆₁ S ₁	[448-469] ⁶⁾		H		
152	704.6609	704.6624	3	C ₈₄ H ₁₃₈ N ₃₀ O ₃₀ S ₂	[470-478] ^{5,8)}	X	X	K476	
153	528.7483	528.7487	4			X	X		
154	671.3071	671.3150	4	C ₁₀₈ H ₁₇₆ N ₃₆ O ₄₀ S ₂	[470-478] ⁸⁾	X	X		
155	537.2457	537.2535	5			X	X		
156	766.8545	766.8624	6	C ₂₀₃ H ₂₉₉ N ₅₅ O ₆₄ S ₂	[615-639]		X	K615	
157	659.7317	659.7396	7	C ₂₀₃ H ₂₉₉ N ₅₅ O ₆₅ S ₂	[615-639] ⁹⁾		X		
158	542.7535	542.7632	4	C ₉₀ H ₁₄₆ N ₂₆ O ₃₄ S	[652-656] ⁶⁾		H	K653	
159	906.0205	906.0284	5	C ₁₉₁ H ₂₉₇ N ₅₇ O ₆₇ S ₂	[657-682] ⁸⁾		X	K669	
160	755.1837	755.1916	6				X		
161	948.6313	948.6281	5	C ₂₀₀ H ₃₀₄ N ₅₆ O ₇₂ S ₃	[703-728]	X	X	K722	
162	790.6922	790.6913	6			X	X		
163	780.0687	780.0674	3	C ₁₀₃ H ₁₆₄ N ₂₈ O ₃₂ S ₁	[756-768] ⁵⁾	X	X	K757	
164	727.8685	727.8687	4	C ₁₂₇ H ₂₀₂ N ₃₄ O ₄₂ S ₁	[756-768]	X	X		
165	812.3994	812.4073	3	C ₁₀₄ H ₁₇₁ N ₂₉ O ₃₄ S ₂	[758-771] ⁵⁾		X	K768	
166	1002.4937	1002.4956	3	C ₁₂₈ H ₂₀₉ N ₃₅ O ₄₄ S ₂	[758-771]	X	X		
167	752.1241	752.1237	4			X	X		
168	756.1213	756.1224	4	C ₁₂₈ H ₂₀₉ N ₃₅ O ₄₅ S ₂	[758-771] ⁹⁾	X	X		
169	605.4955	605.5026	5	C ₁₂₈ H ₂₁₁ N ₃₅ O ₄₅ S ₂	[758-771] ⁶⁾		H		
170	630.7776	630.7836	4	C ₁₀₈ H ₁₆₂ N ₃₀ O ₃₄ S ₃	[769-779] ^{5,9)}	X		K771	
171	612.4893	612.4835	5	C ₁₃₂ H ₂₀₀ N ₃₆ O ₄₂ S ₃	[769-779]		X		
172	773.3420	773.3499	4	C ₁₃₂ H ₂₀₀ N ₃₆ O ₄₄ S ₃	[769-779] ⁹⁾		X		
173	618.8736	618.8815	5				X		
174	525.98605	525.9939	4	C ₈₆ H ₁₃₃ N ₂₉ O ₃₁ S	[826-835] ⁵⁾		X	K829	
175	939.9375	939.9352	4	C ₁₆₃ H ₂₄₂ N ₄₆ O ₅₃ S ₂	[864-888] ^{5,8)}	X	X	K879	
176	1082.4936	1082.5015	4	C ₁₈₇ H ₂₈₀ N ₅₂ O ₆₃ S ₂	[864-888] ⁸⁾		X		
177	866.2018	866.2028	5			X			
178	743.6663	743.6742	3	C ₉₄ H ₁₄₅ N ₂₇ O ₃₂ S ₂	[880-889] ^{5,8)}	X	X	K888	
179	700.5660	700.5738	4	C ₁₁₈ H ₁₈₃ N ₃₃ O ₄₂ S ₂	[880-889] ⁸⁾	X	X		
180	940.10267	940.1105	3	C ₁₂₇ H ₁₈₀ N ₃₆ O ₃₆ S ₁	[917-931] ⁵⁾		X	K917	
181	705.3340	705.3348	4			X	X		
182	847.9018	847.9011	4	C ₁₅₁ H ₂₁₈ N ₄₂ O ₄₆ S ₁	[917-931]	X	X		
183	678.5237	678.5225	5			X	X		
184	568.0633	568.0716	5	C ₁₂₇ H ₁₈₂ N ₃₆ O ₃₇ S ₁	[917-931] ^{5,6)}		H		

185	852.3962	852.4037	4	C ₁₅₁ H ₂₂₀ N ₄₂ O ₄₇ S ₁	[917-931] ⁶⁾		H	
186	682.1170	682.1246	5				H	
187	568.5975	568.6051	6				H	
188	633.1155	633.1231	5	C ₁₄₁ H ₂₁₇ N ₃₉ O ₄₂ S	[1051-1062] ⁶⁾		H	K1060
189	806.1214	806.1253	4	C ₁₃₂ H ₂₁₃ N ₄₁ O ₄₇ S ₃	[1063-1077] ⁸⁾	X	X	K1064
190	810.6210	810.6279	4	C ₁₃₂ H ₂₁₅ N ₄₁ O ₄₈ S ₃	[1063-1077] ⁶⁾		H	
191	648.6968	648.7039	5				H	
192	871.2720	871.2799	7	C ₂₆₆ H ₄₀₇ N ₇₅ O ₈₄ S ₃	[1065-1108] ^{5,8)}		X	K1078
193	1111.3615	1111.3694	6	C ₂₉₀ H ₄₄₅ N ₈₁ O ₉₄ S ₃	[1065-1108] ⁸⁾		X	
194	952.7395	952.7463	7			X	X	
195	1046.8442	1046.8441	3			C ₁₃₄ H ₂₁₂ N ₃₈ O ₄₇ S ₁	[1078-1096] ⁵⁾	
196	1236.9245	1236.9324	3	C ₁₅₈ H ₂₅₀ N ₄₄ O ₅₇ S ₁	[1078-1096]		X	
197	927.9522	927.9512	4			X	X	
198	742.5630	742.5626	5			X		
199	798.9866	798.9944	5	C ₁₇₃ H ₂₆₈ N ₅₀ O ₅₇ S	[1078-1103] ⁵⁾	X	X	
200	913.0473	913.0474	5	C ₁₉₇ H ₃₀₆ N ₅₆ O ₆₇ S ₁	[1078-1103]	X	X	
201	761.0397	761.0408	6			X	X	
202	789.8799	789.8876	4	C ₁₃₄ H ₂₁₄ N ₃₈ O ₄₈ S ₁	[1078-1096] ^{5,6)}		H	
203	932.4474	932.4539	4	C ₁₅₈ H ₂₅₂ N ₄₄ O ₅₈ S ₁	[1078-1096] ⁶⁾		H	
204	746.1579	746.1647	5				H	
205	668.9888	668.9884	6	C ₁₇₃ H ₂₇₀ N ₅₀ O ₅₈ S	[1078-1103] ^{5,6)}		H	
206	655.0302	655.0376	7	C ₁₉₇ H ₃₀₈ N ₅₆ O ₆₈ S ₁	[1078-1103] ⁶⁾		H	
207	1600.4044	1600.4122	3	C ₂₁₂ H ₃₁₂ N ₅₄ O ₇₂ S	[1104-1130]		X	K1108
208	960.6426	960.6505	5				X	
209	1062.4886	1062.4976	4			C ₁₈₈ H ₂₇₆ N ₄₈ O ₆₃ S	[1104-1130] ^{5,6)}	
210	918.65258	918.6605	5	C ₂₀₄ H ₃₁₄ N ₅₆ O ₆₁ S ₂	[1137-1162] ⁸⁾		X	K1157
211	837.4176	837.4231	3	C ₁₁₀ H ₁₇₆ N ₃₀ O ₃₃ S ₂	[1150-1162] ^{5,8)}	X	X	
212	955.7945	955.8024	3	C ₁₂₅ H ₁₉₇ N ₃₃ O ₄₀ S ₂	[1150-1162] ^{8,10)}	X	X	
213	770.8816	770.8855	4	C ₁₃₄ H ₂₁₄ N ₃₆ O ₄₃ S ₂	[1150-1162] ⁸⁾	X	X	
214	616.9021	616.9100	5				X	
215	922.2545	922.2626	5	C ₂₀₄ H ₃₁₆ N ₅₆ O ₆₂ S ₂	[1137-1162] ^{6,8)}		H	
216	768.7121	768.7201	6				H	
217	757.0129	757.0201	3	C ₉₈ H ₁₄₉ N ₂₅ O ₃₅ S	[1191-1195]		X	K1193
218	63.1072	763.1150	4	C ₁₃₅ H ₂₀₁ N ₃₅ O ₄₄ S	[1205-1216]	X	X	K1214
219	979.4557	979.4598	2	C ₈₈ H ₁₂₈ N ₂₂ O ₂₇ S	[1210-1216] ⁵⁾	X	X	
220	653.3034	653.3091	3			X	X	
221	843.3900	843.3975	3	C ₁₁₂ H ₁₆₆ N ₂₈ O ₃₇ S	[1210-1216]	X	X	
222	913.0583	913.0662	5	C ₁₉₆ H ₃₁₈ N ₅₆ O ₆₇ S	[1262-1292] ⁵⁾		X	K1283
223	1283.6479	1283.6470	4	C ₂₂₀ H ₃₅₆ N ₆₂ O ₇₇ S ₁	[1262-1292]		X	
224	1027.1216	1027.1191	5			X	X	
225	856.1015	856.1006	6			X	X	
226	1031.0041	1031.0077	2	C ₈₈ H ₁₄₁ N ₂₅ O ₃₀ S	[1276-1284] ⁵⁾	X	X	
227	687.6691	687.6744	3			X	X	
228	877.7574	877.7627	3	C ₁₁₂ H ₁₇₉ N ₃₁ O ₄₀ S	[1276-1284]	X	X	
229	658.5661	658.574	4				X	
230	983.8323	983.8312	3	C ₁₂₉ H ₂₀₅ N ₃₅ O ₄₂ S ₁	[1276-1292] ⁵⁾	X	X	
231	1173.9116	1173.9196	3	C ₁₅₃ H ₂₄₃ N ₄₁ O ₅₂ S ₁	[1276-1292]		X	

232	880.6933	880.6916	4			X	X			
233	916.6610	916.6683	5	C ₁₉₆ H ₃₂₀ N ₅₆ O ₆₈ S	[1276-1292] ^{6,5)}		H			
234	764.0508	764.0582	6					H		
235	885.1874	885.1943	4	C ₁₅₃ H ₂₄₅ N ₄₁ O ₅₃ S ₁	[1276-1292] ⁶⁾		H			
236	835.0730	835.0732	3	C ₁₀₉ H ₁₆₇ N ₃₁ O ₃₅ S ₁	[1284-1296] ⁵⁾	X	X	K1292		
237	626.5574	626.5569	4				X		X	
238	1025.1645	1025.1615	3	C ₁₃₃ H ₂₀₅ N ₃₇ O ₄₅ S ₁	[1284-1296]	X	X			
239	769.1233	769.1231	4				X		X	
240	615.5006	615.5006	5				X		X	
241	637.9826	637.9885	3			C ₈₂ H ₁₃₀ N ₂₆ O ₂₅ S	[1289-1296] ⁵⁾		X	X
242	478.7372	478.7433	4		X				X	
243	621.3017	621.3096	4	C ₁₀₆ H ₁₆₈ N ₃₂ O ₃₅ S	[1289-1296]		X			
244	773.6181	773.6257	4	C ₁₃₃ H ₂₀₇ N ₃₇ O ₄₆ S ₁	[1284-1296] ⁶⁾		H			
245	619.0945	619.1022	5						H	
246	516.0788	516.0864	6						H	
247	681.3048	681.3127	3	C ₈₆ H ₁₃₂ N ₂₆ O ₂₈ S ₂	[1293-1301] ^{5,8)}		X		K1296	
248	653.7967	653.8027	4	C ₁₁₀ H ₁₇₀ N ₃₂ O ₃₈ S ₂	[1293-1301] ⁸⁾	X				
249	1261.0921	1261.0956	4	C ₂₂₂ H ₃₃₀ N ₅₈ O ₇₅ S ₁	[1336-1366]	X	X	1349		
250	1009.0786	1009.0781	5				X		X	
251	914.1036	914.1085	3	C ₁₁₄ H ₁₈₆ N ₃₂ O ₄₄ S	[1349-1359]	X	X			
252	1003.8099	1003.8177	3	C ₁₂₈ H ₂₀₁ N ₃₇ O ₄₅ S	[1349-1366] ⁵⁾		X			
253	895.6736	895.6815	4	C ₁₅₂ H ₂₃₉ N ₄₃ O ₅₅ S	[1349-1366]		X			
254	1123.0256	1123.0320	4	C ₁₉₈ H ₂₉₄ N ₅₂ O ₆₆ S	[1336-1366] ^{5,6)}		H			
255	933.0779	933.0858	3	C ₁₂₂ H ₁₇₇ N ₃₁ O ₄₃ S	[1395-1403]		X	K1399		
256	1003.4851	1003.4843	4	C ₁₈₀ H ₂₆₄ N ₄₈ O ₅₅ S ₁	[1436-1461] ⁵⁾	X	X	K1437		
257	917.0454	917.0419	5	C ₂₀₄ H ₃₀₂ N ₅₄ O ₆₅ S ₁	[1436-1461]	X	X			
258	655.3101	655.3179	7							X
259	806.5812	806.5911	5	C ₁₈₀ H ₂₆₆ N ₄₈ O ₅₆ S ₁	[1436-1461] ^{5,6)}		H			
260	672.3176	672.3272	6							H
261	767.3617	767.3713	6	C ₂₀₄ H ₃₀₄ N ₅₄ O ₆₆ S ₁	[1436-1461] ⁶⁾		H			
262	657.8815	657.8909	7							H
263	739.9977	740.0081	3	C ₉₃ H ₁₄₄ N ₂₆ O ₃₅ S	[1475-1479]		X	K1475		
264	706.3223	706.3224	4	C ₁₁₅ H ₁₈₄ N ₃₆ O ₄₁ S ₃	[1475-1489] ^{5,8)}	X	X			
265	848.8807	848.8886	4	C ₁₃₉ H ₂₂₂ N ₄₂ O ₅₁ S ₃	[1475-1489] ⁸⁾		X			
266	679.3126	679.3124	5				X		X	
267	682.5050	682.5114	5	C ₁₃₉ H ₂₂₂ N ₄₂ O ₅₂ S ₃	[1475-1489] ^{8,9)}	X	X			
268	906.5835	906.5913	6	C ₂₂₈ H ₃₆₁ N ₆₅ O ₈₁ S ₄	[1475-1507] ^{8,9)}		X			
269	682.9053	682.9146	5	C ₁₃₉ H ₂₂₄ N ₄₂ O ₅₂ S ₃	[1475-1489] ^{8,6)}		H			
270	970.4482	970.4561	5	C ₂₀₄ H ₃₂₃ N ₅₉ O ₇₀ S ₄	[1475-1507] ^{5,8)}	X	X	K1489		
271	1084.5012	1084.5090	5	C ₂₂₈ H ₃₆₁ N ₆₅ O ₈₀ S ₄	[1475-1507] ⁸⁾	X	X			
272	903.92113	903.9255	6				X		X	
273	1087.7001	1087.7080	5	C ₂₂₈ H ₃₆₁ N ₆₅ O ₈₁ S ₄	[1475-1507] ^{8,9)}		X			
274	906.5835	906.5913	6							X
275	1180.7865	1180.7944	4	C ₁₉₈ H ₃₁₁ N ₅₇ O ₆₉ S ₄	[1476-1507] ^{5,8)}	X	X			
276	1058.8836	1058.8900	5	C ₂₂₂ H ₃₄₉ N ₆₃ O ₇₉ S ₄	[1476-1507] ⁸⁾	X	X			
277	1062.0812	1062.0890	5	C ₂₂₂ H ₃₄₉ N ₆₃ O ₈₀ S ₄	[1476-1507] ^{8,9)}		X			
278	917.7656	917.7735	3	C ₁₁₈ H ₁₈₃ N ₃₃ O ₄₁ S	[1489-1498]	X	X			

279	1057.5153	1057.5135	3	C ₁₃₅ H ₂₁₆ N ₃₈ O ₄₆ S ₂	[1489-1507] ⁵⁾		X	
280	1062.8396	1062.8452	3	C ₁₃₅ H ₂₁₆ N ₃₈ O ₄₇ S ₂	[1489-1507] ^{5,9)}	X		
281	935.9540	935.9534	4	C ₁₅₉ H ₂₅₄ N ₄₄ O ₅₆ S ₂	[1489-1507]	X	X	
282	939.94423	939.9521	4	C ₁₅₉ H ₂₅₄ N ₄₄ O ₅₇ S ₂	[1489-1507] ⁹⁾		X	
283	906.9176	906.9272	6	C ₂₂₈ H ₃₆₃ N ₆₅ O ₈₁ S ₄	[1475-1507] ^{6,8)}		H	
284	932.2279	932.2358	5	C ₁₉₉ H ₃₀₂ N ₅₈ O ₆₈ S ₂	[1490-1520] ⁵⁾		X	
285	1046.2809	1046.2888	5	C ₂₂₃ H ₃₄₀ N ₆₄ O ₇₈ S ₂	[1490-1520]	X	X	
286	1243.5800	1243.5879	6	C ₃₂₈ H ₄₈₄ N ₉₂ O ₁₀₅ S ₂	[1490-1537]		X	
287	1066.0686	1066.0764	7			X	X	
288	932.9350	932.9429	8				X	
289	1068.3535	1068.3615	7			C ₃₂₈ H ₄₈₄ N ₉₂ O ₁₀₆ S ₂	[1490-1537] ⁹⁾	
290	934.9344	934.9422	8		X			
291	875.0674	875.0770	6	C ₂₂₃ H ₃₄₂ N ₆₄ O ₇₉ S ₂	[1490-1520] ⁶⁾		H	
292	845.6893	845.6972	7	C ₂₆₀ H ₃₈₆ N ₇₈ O ₈₀ S	[1508-1546] ⁵⁾		X	
293	967.71083	967.7187	4	C ₁₆₉ H ₂₅₅ N ₄₉ O ₅₄ S	[1523-1545] ⁵⁾		X	
294	1110.2846	1110.2849	4	C ₁₉₃ H ₂₉₃ N ₅₅ O ₆₄ S ₁	[1523-1545]	X	X	
295	888.4303	888.4295	5			X	X	
296	892.0252	892.0316	5	C ₁₉₃ H ₂₉₅ N ₅₅ O ₆₅ S ₁	[1523-1545] ⁶⁾		H	
297	743.5210	743.5276	6				H	
298	716.3195	716.3256	3	C ₈₄ H ₁₃₉ N ₂₉ O ₃₃ S ₂	[1574-1583] ^{5,8)}	X	X	
299	680.0560	680.0624	4	C ₁₀₈ H ₁₇₇ N ₃₅ O ₄₃ S ₂	[1574-1583] ⁸⁾	X		K1581
300	863.4113	863.4192	5	C ₁₉₀ H ₂₉₀ N ₅₀ O ₆₁ S ₂	[1582-1605]		X	
301	719.6773	719.6840	6				X	
302	866.6103	866.6182	5	C ₁₉₀ H ₂₉₀ N ₅₀ O ₆₂ S ₂	[1582-1605] ⁹⁾		X	
303	722.3497	722.3498	6			X	X	
304	850.2521	850.2492	6	C ₂₂₇ H ₃₄₃ N ₅₉ O ₇₁ S ₂	[1584-1614]	X	X	
305	728.9212	728.9290	7				X	
306	609.0539	609.0618	4	C ₁₀₈ H ₁₆₉ N ₂₉ O ₃₃ S	[1603-1614] ⁵⁾		X	K1605
307	601.4961	601.504	5	C ₁₃₂ H ₂₀₇ N ₃₅ O ₄₃ S	[1603-1614]		X	
308	1126.5703	1126.5703	3	C ₁₅₀ H ₂₃₃ N ₄₁ O ₄₆ S ₁	[1606-1627] ⁵⁾	X	X	
309	845.1788	845.1797	4			X	X	
310	676.3446	676.3453	5			X	X	
311	1316.6591	1316.6586	3			C ₁₇₄ H ₂₇₁ N ₄₇ O ₅₆ S ₁	[1606-1627]	X
312	987.7473	987.7459	4	X	X			
313	790.3967	790.3983	5		X			
314	658.8326	658.8332	6	X	X			
315	849.6740	849.6823	4	C ₁₅₀ H ₂₃₅ N ₄₁ O ₄₇ S ₁	[1606-1627] ^{5,6)}		H	
316	679.9392	679.9474	5	C ₁₇₄ H ₂₇₃ N ₄₇ O ₅₇ S ₁	[1606-1627] ⁶⁾		H	
317	793.9929	794.0004	5				H	
318	661.8275	661.8350	6				H	
319	701.3513	701.3516	4			C ₁₁₇ H ₁₉₆ N ₃₂ O ₄₅ S ₁	[1671-1681]	X
320	886.44257	886.4504	3	C ₁₁₄ H ₁₈₅ N ₃₃ O ₃₈ S	[1672-1685] ⁵⁾		X	
321	665.08193	665.0898	4			X	X	
322	1076.5309	1076.5388	3	C ₁₃₈ H ₂₂₃ N ₃₉ O ₄₈ S	[1672-1685]		X	
323	807.6510	807.6560	4			X	X	
324	646.3195	646.3264	5			X	X	
325	929.1409	929.1488	3	C ₁₂₀ H ₁₉₇ N ₃₅ O ₃₉ S ₁	[1672-1686] ⁵⁾		X	

326	697.1057	697.1135	4				X	
327	557.8934	557.8924	5			X	X	
328	839.6788	839.6797	4			X	X	
329	671.9458	671.9454	5	C ₁₄₄ H ₂₃₅ N ₄₁ O ₄₉ S ₁	[1672-1686]	X	X	
330	560.1146	560.1224	6				X	
331	566.6155	566.6234	3	C ₇₃ H ₁₁₆ N ₂₄ O ₂₁ S	[1681-1685] ⁵⁾		X	
332	454.4223	454.4302	5	C ₉₇ H ₁₅₄ N ₃₀ O ₃₁ S	[1681-1685]		X	
333	669.5844	669.5925	4	C ₁₁₄ H ₁₈₇ N ₃₃ O ₃₉ S	[1672-1685] ^{5,6)}		H	
334	535.8675	535.8755	5				H	
335	812.1517	812.1587	4	C ₁₃₈ H ₂₂₅ N ₃₉ O ₄₉ S	[1672-1685] ⁶⁾		H	
336	561.4871	561.4945	5	C ₁₂₀ H ₁₉₉ N ₃₅ O ₄₀ S ₁	[1672-1686] ^{5,6)}		H	
337	675.5407	675.5475	5	C ₁₄₄ H ₂₃₇ N ₄₁ O ₅₀ S ₁	[1672-1686] ⁶⁾		H	
338	563.1172	563.1242	6				H	
339	832.7459	832.7450	3	C ₁₀₈ H ₁₇₀ N ₃₀ O ₃₆ S ₁	[1738-1749] ⁵⁾	X	X	
340	1022.8323	1022.8333	3			X		
341	767.3778	767.3770	4	C ₁₃₂ H ₂₀₈ N ₃₆ O ₄₆ S ₁	[1738-1749]	X	X	K1748
342	614.1044	614.1031	5			X	X	
343	656.9231	656.9305	5	C ₁₄₈ H ₂₂₂ N ₄₀ O ₄₃ S	[1779-1792]	X	X	
344	1096.2967	1096.2948	4				X	
345	877.2378	877.2374	5	C ₁₉₈ H ₂₉₇ N ₅₁ O ₆₀ S ₁	[1764-1792] ⁵⁾	X	X	
346	1238.8531	1238.861	4				X	
347	991.2921	991.2903	5	C ₂₂₂ H ₃₃₅ N ₅₇ O ₇₀ S ₁	[1764-1792]	X	X	K1784
348	826.243	826.2433	6			X	X	
349	708.3467	708.3525	7				X	
350	880.8295	880.8395	5	C ₁₉₈ H ₂₉₉ N ₅₁ O ₆₁ S ₁	[1764-1792] ^{5,6)}		H	
351	994.8839	994.8925	5				H	
352	829.2366	829.2451	6	C ₂₂₂ H ₃₃₇ N ₅₇ O ₇₁ S ₁	[1764-1792] ⁶⁾		H	
353	710.9171	710.9254	7				H	
354	1016.7317	1016.7396	4	C ₁₇₇ H ₂₇₁ N ₄₇ O ₆₁ S	[1785-1806]		X	
355	924.6497	924.6493	5	C ₂₀₄ H ₃₀₈ N ₅₄ O ₆₇ S ₁	[1785-1816] ⁵⁾		X	
356	1298.1180	1298.1260	4				X	
357	1038.6979	1038.7023	5	C ₂₂₈ H ₃₄₆ N ₆₀ O ₇₇ S ₁	[1785-1816]	X		K1792
358	865.7516	865.7516	6			X	X	
359	817.1853	817.1953	5	C ₁₇₇ H ₂₇₃ N ₄₇ O ₆₂ S	[1785-1806]		H	
360	868.7477	868.7550	6	C ₂₂₈ H ₃₄₈ N ₆₀ O ₇₈ S ₁	[1785-1816]		H	
361	794.3679	794.3757	2	C ₆₄ H ₁₀₆ N ₂₀ O ₂₅ S	[1898-1903] ⁵⁾		X	
362	720.0002	720.0081	3	C ₈₈ H ₁₄₄ N ₂₆ O ₃₅ S	[1898-1903]		X	
363	886.0734	886.0813	3				X	
364	664.8123	664.8129	4	C ₁₀₈ H ₁₇₈ N ₃₄ O ₃₈ S ₃	[1898-1912] ^{5,8)}	X	X	K1902
365	668.8051	668.8116	4	C ₁₀₈ H ₁₇₈ N ₃₄ O ₃₉ S ₃	[1898-1912] ^{5,8,9)}	X		
366	807.3787	807.3792	4			X	X	
367	646.1041	646.1049	5	C ₁₃₂ H ₂₁₆ N ₄₀ O ₄₈ S ₃	[1898-1912] ⁸⁾	X	X	
368	649.3046	649.3039	5	C ₁₃₂ H ₂₁₆ N ₄₀ O ₄₉ S ₃	[1898-1912] ^{8,9)}	X		
369	889.7416	889.7495	3				X	
370	667.5562	667.5641	4	C ₁₁₀ H ₁₇₉ N ₃₃ O ₃₈ S ₃	[1903-1916] ^{5,8)}		X	K1912
371	810.1289	810.1303	4			X	X	
372	648.3060	648.3058	5	C ₁₃₄ H ₂₁₇ N ₃₉ O ₄₈ S ₃	[1903-1916] ⁸⁾	X	X	

373	814.1230	814.1291	4	C ₁₃₄ H ₂₁₇ N ₃₉ O ₄₉ S ₃	[1903-1916] ^{8,9)}	X			
374	651.4969	651.5048	5						X
375	619.6306	619.6385	3	C ₇₇ H ₁₂₉ N ₂₃ O ₂₆ S ₂	[1910-1916] ⁵⁾		X		
376	607.5392	607.5471	4	C ₁₀₁ H ₁₆₇ N ₂₉ O ₃₆ S ₂	[1910-1916]		X		
377	611.5381	611.5458	4	C ₁₀₁ H ₁₆₇ N ₂₉ O ₃₇ S ₂	[1910-1916] ⁹⁾		X		
378	941.1236	941.1259	3	C ₁₂₆ H ₁₈₉ N ₃₃ O ₃₉ S ₁	[1913-1928] ⁵⁾		X	K1916	
379	706.0885	706.0964	4						
380	1131.2161	1131.2142	3	C ₁₅₀ H ₂₂₇ N ₃₉ O ₄₉ S ₁	[1913-1928]	X	X		
381	848.6628	848.6626	4				X		X
382	679.1238	679.1317	5						X
383	1137.2104	1137.2177	3	C ₁₅₀ H ₂₂₉ N ₃₉ O ₅₀ S ₁	[1913-1928] ⁶⁾		H		
384	853.1578	853.1652	4						H
385	682.7263	682.7338	5						H
386	842.0769	842.0733	3			C ₁₁₁ H ₁₆₆ N ₃₂ O ₃₄ S ₁	[1917-1929] ⁵⁾	X	X
387	1032.1610	1032.1616	3	C ₁₃₅ H ₂₀₄ N ₃₈ O ₄₄ S ₁	[1917-1929]	X		K1928	
388	774.3732	774.3732	4				X		X
389	1082.1802	1082.1881	3	C ₁₄₃ H ₂₁₄ N ₄₀ O ₄₅ S	[1917-1935] ⁵⁾		X		
390	954.4606	954.4592	4	C ₁₆₇ H ₂₅₂ N ₄₆ O ₅₅ S ₁	[1917-1935]	X	X		
391	763.7697	763.7690	5				X		X
392	725.5295	725.5355	5	C ₁₅₄ H ₂₃₁ N ₄₅ O ₅₅ S	[1929-1944]	X		K1935	
393	604.7730	604.7809	6						
394	899.4106	899.4185	7	C ₂₇₈ H ₄₀₆ N ₇₂ O ₉₂ S ₂	[1929-1967]		X		
395	787.1093	787.1171	8						
396	725.0758	725.0759	4	C ₁₂₄ H ₁₈₁ N ₃₅ O ₄₄ S ₁	[1930-1944] ⁵⁾	X	X		
397	867.6415	867.6421	4	C ₁₄₈ H ₂₁₉ N ₄₁ O ₅₄ S ₁	[1930-1944]	X	X		
398	694.3154	694.3152	5				X		X
399	622.6776	622.6855	5	C ₁₃₄ H ₁₉₇ N ₃₇ O ₄₇ S	[1933-1944]		X		
400	729.5714	729.5786	4	C ₁₂₄ H ₁₈₃ N ₃₅ O ₄₅ S ₁	[1930-1944] ^{5,6)}		H		
401	583.8571	583.8644	5						
402	697.9094	697.9174	5	C ₁₄₈ H ₂₂₁ N ₄₁ O ₅₅ S ₁	[1930-1944] ⁶⁾		H		
403	581.7579	581.7658	6						H
404	1022.4760	1022.4727	5	C ₂₂₁ H ₃₃₁ N ₅₉ O ₇₇ S ₂	[1945-1974]	X	X	K1967	
405	680.9995	681.0074	3	C ₉₀ H ₁₄₁ N ₂₃ O ₂₉ S	[2023-2030] ⁵⁾	X	X	K2025	
406	653.5659	653.5737	4	C ₁₁₄ H ₁₇₉ N ₂₉ O ₃₉ S	[2023-2030]		X		
407	989.9914	989.9993	2	C ₈₅ H ₁₃₉ N ₂₃ O ₂₉ S ₁	[2026-2033] ⁵⁾		X	K2030	
408	660.3355	660.3355	3				X		X
409	850.4239	850.4238	3	C ₁₀₉ H ₁₇₇ N ₂₉ O ₃₉ S ₁	[2026-2033]	X	X		
410	638.0701	638.0698	4				X		X
411	856.4190	856.4273	3	C ₁₀₉ H ₁₇₉ N ₂₉ O ₄₀ S ₁	[2026-2033] ⁶⁾		H		
412	736.1038	736.1130	4	C ₁₂₄ H ₂₀₅ N ₃₃ O ₄₅ S ₂	[2026-2036] ⁶⁾		H		
413	866.9178	866.9220	2	C ₇₁ H ₁₂₁ N ₂₁ O ₂₅ S ₂	[2031-2036] ⁵⁾	X		K2033	
414	578.2851	578.2839	3				X		
415	583.6077	583.6156	3	C ₇₁ H ₁₂₁ N ₂₁ O ₂₆ S ₂	[2031-2036] ^{5,9)}	X	X		
416	768.3729	768.3723	3	C ₉₅ H ₁₅₉ N ₂₇ O ₃₅ S ₂	[2031-2036]	X	X		
417	576.5322	576.5312	4				X		X
418	773.7045	773.7039	3	C ₉₅ H ₁₅₉ N ₂₇ O ₃₆ S ₂	[2031-2036] ⁹⁾	X			
419	580.5301	580.5299	4				X		X

420	981.1381	981.1481	3	C ₁₂₄ H ₂₀₅ N ₃₃ O ₄₅ S ₂	[2026-2036] ⁶⁾		H	
421	736.1038	736.1130	4				H	
422	589.0829	589.0920	5				H	
423	744.0252	744.0251	3	C ₉₁ H ₁₅₂ N ₂₈ O ₃₃ S ₂	[2034-2038]		X	K2036
424	558.2705	558.2708	4				X	
425	881.75134	881.7568	3	C ₁₁₃ H ₁₇₉ N ₃₁ O ₃₈ S ₂	[2034-2046] ⁵⁾	X		
426	661.5690	661.5695	4			X	X	
427	1071.8372	1071.8451	3	C ₁₃₇ H ₂₁₇ N ₃₇ O ₄₈ S ₂	[2034-2046]		X	
428	804.1359	804.1358	4			X	X	
429	643.5099	643.5102	5			X	X	
430	808.1352	808.1345	4				X	
431	646.7136	646.7092	5	C ₁₃₇ H ₂₁₇ N ₃₇ O ₄₉ S ₂	[2034-2046] ⁹⁾		X	
432	705.6749	705.6828	3				X	
433	916.1302	916.1381	3	C ₁₂₁ H ₁₉₂ N ₃₄ O ₃₇ S	[2037-2050] ⁵⁾		X	K2046
434	687.3477	687.3555	4				X	
435	550.0781	550.0860	5				X	
436	1106.2185	1106.2264	3	C ₁₄₅ H ₂₃₀ N ₄₀ O ₄₇ S ₁	[2037-2050]		X	
437	829.9225	829.9218	4			X	X	
438	664.1386	664.1390	5			X	X	
439	553.6176	553.6171	6			X	X	
440	826.4116	826.4097	3	C ₁₀₉ H ₁₆₉ N ₂₉ O ₃₅ S ₁	[2039-2050] ⁵⁾	X	X	
441	620.0587	620.0593	4			X	X	
442	762.6245	762.6255	4	C ₁₃₃ H ₂₀₇ N ₃₅ O ₄₅ S ₁	[2039-2050]	X	X	
443	610.3008	610.3019	5			X	X	
444	834.4177	834.4244	4				H	
445	667.7342	667.7411	5	C ₁₄₅ H ₂₃₂ N ₄₀ O ₄₈ S ₁	[2037-2050] ⁶⁾		H	
446	556.6118	556.6189	6				H	
447	536.9246	536.9320	3	C ₆₇ H ₁₀₉ N ₂₁ O ₂₃ S	[2047-2051] ⁵⁾		X	K2050
448	684.0007	684.0062	3	C ₈₆ H ₁₄₀ N ₂₆ O ₃₀ S	[2047-2056] ⁵⁾		X	
449	760.0366	760.0432	3	C ₉₆ H ₁₅₆ N ₂₈ O ₃₄ S	[2047-2058] ⁵⁾		X	
450	502.4957	502.5016	4	C ₈₂ H ₁₃₅ N ₂₉ O ₂₈ S	[2071-2079] ⁵⁾	X	X	K2073
451	470.2350	470.2410	4	C ₇₇ H ₁₂₈ N ₂₈ O ₂₅ S	[2072-2079] ⁵⁾	X	X	
452	490.4411	490.4473	5	C ₁₀₁ H ₁₆₆ N ₃₄ O ₃₅ S	[2072-2079]	X	X	
453	1045.0104	1045.0125	2	C ₈₅ H ₁₄₅ N ₂₇ O ₃₀ S ₂	[2097-2106] ⁵⁾		X	K2102
454	697.0102	697.0109	3			X	X	
455	523.00228	523.0101	4			X	X	
456	702.3435	702.3425	3	C ₈₅ H ₁₄₅ N ₂₇ O ₃₁ S ₂	[2097-2106] ^{5,9)}	X	X	
457	887.0999	887.0992	3	C ₁₀₉ H ₁₈₃ N ₃₃ O ₄₀ S ₂	[2097-2106]	X	X	
458	665.5765	665.5764	4			X	X	
459	532.6548	532.6627	5				X	
460	892.4281	892.4308	3				X	
461	669.5761	669.5751	4	C ₁₀₉ H ₁₈₃ N ₃₃ O ₄₁ S ₂	[2097-2106] ⁹⁾	X	X	
462	658.6636	658.6686	3				X	
463	636.8118	636.8196	4	C ₁₀₅ H ₁₇₈ N ₃₂ O ₃₇ S ₂	[2098-2106]	X	X	
464	640.8105	640.8184	4	C ₁₀₅ H ₁₇₈ N ₃₂ O ₃₈ S ₂	[2098-2106] ⁹⁾		X	
465	703.0065	703.0144	3	C ₈₅ H ₁₄₇ N ₂₇ O ₃₁ S ₂	[2097-2106] ^{5,6)}		H	
466	527.5049	527.5128	4				H	

467	893.0948	893.1027	3	C ₁₀₉ H ₁₈₅ N ₃₃ O ₄₁ S ₂	[2097-2106] ⁶⁾		H		
468	670.0711	670.0790	4				H		
469	536.2569	536.2648	5				H		
470	674.0689	674.0778	4	C ₁₀₉ H ₁₈₅ N ₃₃ O ₄₂ S ₂	[2097-2106] ^{6,9)}		H		
471	539.4551	539.4637	5				H		
472	1001.4974	1001.4947	5	C ₂₂₄ H ₃₄₄ N ₅₆ O ₆₈ S ₃	[2154-2183]	X	X	K2179	
473	828.60706	828.6136	5	C ₁₈₅ H ₂₈₄ N ₄₈ O ₅₆ S ₂	[2161-2183]	X			
474	1004.4886	1004.4943	3	C ₁₃₁ H ₂₀₇ N ₃₃ O ₄₆ S	[2180-2192]		X	K2183	
475	1107.8684	1107.8683	3	C ₁₅₁ H ₂₂₁ N ₃₇ O ₄₆ S ₁	[2180-2198] ⁵⁾		X		
476	973.7210	973.7194	4	C ₁₇₅ H ₂₅₉ N ₄₃ O ₅₆ S ₁	[2180-2198]	X	X		
477	1095.0389	1095.0381	4	C ₁₉₉ H ₂₉₄ N ₅₀ O ₆₀ S ₁	[2180-2208] ⁵⁾		X		
478	1237.6053	1237.6044	4	C ₂₂₃ H ₃₃₂ N ₅₆ O ₇₀ S ₁	[2180-2208]		X		
479	1471.7151	1471.7148	3	C ₂₀₂ H ₂₉₄ N ₅₀ O ₆₀ S ₁	[2184-2212] ⁵⁾		X	K2208	
480	1104.0413	1104.0381	4				X		
481	1246.6050	1246.6044	4	C ₂₂₆ H ₃₃₂ N ₅₆ O ₇₀ S ₁	[2184-2212]	X	X		
482	997.4853	997.4850	5				X		X
483	831.4113	831.4055	6				X		
484	1035.1693	1035.1771	3	C ₁₃₅ H ₂₁₁ N ₃₇ O ₄₅ S	[2199-2212]	X	X		
485	776.6325	776.6348	4				X		X
486	887.0268	887.0342	5	C ₂₀₂ H ₂₉₆ N ₅₀ O ₆₁ S ₁	[2184-2212] ^{5,6)}		H		
487	1251.0996	1251.1071	4	C ₂₂₆ H ₃₃₄ N ₅₆ O ₇₁ S ₁	[2184-2212] ⁶⁾		H		
488	834.3997	834.4073	6				H		
489	800.3833	800.3820	3	C ₁₀₂ H ₁₅₉ N ₂₉ O ₃₆ S ₁	[2209-2220] ⁵⁾	X	X	K2212	
490	600.5363	600.5384	4				X		
491	990.4624	990.4703	3	C ₁₂₆ H ₁₉₇ N ₃₅ O ₄₆ S ₁	[2209-2220]		X		
492	743.1045	743.1046	4				X		X
493	594.6789	594.6853	5				X		
494	605.0334	605.0411	4	C ₁₀₂ H ₁₆₁ N ₂₉ O ₃₇ S ₁	[2209-2220] ^{5,6)}		H		
495	747.5998	747.6073	4	C ₁₂₆ H ₁₉₉ N ₃₅ O ₄₇ S ₁	[2209-2220] ⁶⁾		H		
496	790.0796	790.0798	3	C ₁₀₇ H ₁₆₆ N ₃₀ O ₂₉ S ₁	[2299-2309] ⁵⁾	X	X	K2308	
497	980.1602	980.1681	3	C ₁₃₁ H ₂₀₄ N ₃₆ O ₃₉ S ₁	[2299-2309]		X		
498	735.3780	735.3780	4				X		X
499	986.1629	986.1716	3	C ₁₃₁ H ₂₀₆ N ₃₆ O ₄₀ S ₁	[2299-2309] ⁶⁾		H		
500	770.3677	770.3755	5	C ₁₅₉ H ₂₆₃ N ₅₁ O ₅₂ S ₄	[2309-2326] ^{8,9)}		X	K2311	
501	496.2516	496.2594	4	C ₈₄ H ₁₃₆ N ₃₀ O ₂₄ S	[2310-2316] ⁵⁾		X		
502	397.2013	397.2091	5				X		
503	638.8247	638.8257	4	C ₁₀₈ H ₁₇₄ N ₃₆ O ₃₄ S ₁	[2310-2316]	X	X		
504	511.2557	511.2621	5				X		
505	599.7925	599.8004	4	C ₁₀₂ H ₁₆₂ N ₃₂ O ₃₃ S	[2311-2316]	X	X		
506	480.0340	480.0419	5				X	X	
507	709.0807	709.0811	4	C ₁₁₇ H ₁₈₉ N ₃₇ O ₃₇ S ₄	[2312-2326] ^{5,8)}	X	X	K2316	
508	681.5115	681.5193	5	C ₁₄₁ H ₂₂₇ N ₄₃ O ₄₇ S ₄	[2312-2326] ⁸⁾		X		
509	568.0941	568.1008	6				X		
510	684.7105	684.7184	5	C ₁₄₁ H ₂₂₇ N ₄₃ O ₄₈ S ₄	[2312-2326] ^{8,9)}		X		
511	1012.9605	1012.9593	4	C ₁₇₂ H ₂₆₂ N ₅₂ O ₅₂ S ₅	[2317-2342] ^{5,8)}	X		K2326	
512	810.5692	810.5690	5				X		X
513	675.6343	675.6422	6						X

514	813.7601	813.7680	5	C ₁₇₂ H ₂₆₂ N ₅₂ O ₅₃ S ₅	[2317-2342] ^{5,8,9)}		X		
515	1155.5272	1155.5255	4	C ₁₉₆ H ₃₀₀ N ₅₈ O ₆₂ S ₅	[2317-2342] ⁸⁾	X	X		
516	924.6141	924.6220	5					X	
517	927.8157	927.8210	5	C ₁₉₆ H ₃₀₀ N ₅₈ O ₆₃ S ₅	[2317-2342] ^{8,9)}		X		
518	773.3455	773.3521	6					X	
519	814.1612	814.1711	5	C ₁₇₂ H ₂₆₄ N ₅₂ O ₅₃ S ₅	[2317-2342] ^{8,6,5)}		H		
520	678.6344	678.6439	6					H	
521	928.2170	928.2241	5	C ₁₉₆ H ₃₀₂ N ₅₈ O ₆₃ S ₅	[2317-2342] ^{8,6)}		H		
522	773.6809	773.6880	6					H	
523	663.2979	663.3052	7					H	
524	547.5096	547.5175	4	C ₈₈ H ₁₄₇ N ₂₉ O ₃₄ S	[2381-2385]		X	K2381	
525	894.8991	894.9070	4	C ₁₄₈ H ₂₃₄ N ₄₂ O ₅₅ S ₃	[2408-2426] ⁸⁾		X		
526	793.9553	793.9556	5	C ₁₆₈ H ₂₅₃ N ₄₉ O ₅₅ S ₄	[2408-2433] ^{5,8)}		X		
527	908.00072	908.0086	5	C ₁₉₂ H ₂₉₁ N ₅₅ O ₆₅ S ₄	[2408-2433] ⁸⁾		X		
528	756.8403	756.8418	6					X	
529	726.6587	726.6653	6	C ₁₈₇ H ₂₈₀ N ₅₄ O ₅₉ S ₄	[2408-2436] ^{5,8)}	X			
530	985.8549	985.8499	5	C ₂₁₁ H ₃₁₈ N ₆₀ O ₆₉ S ₄	[2408-2436] ⁸⁾	X			
531	821.7090	821.7095	6				X	X	
532	704.4586	704.4664	7					X	
533	989.0410	989.0488	5	C ₂₁₁ H ₃₁₈ N ₆₀ O ₇₀ S ₄	[2408-2436] ^{8,9)}		X		
534	824.3687	824.3754	6				X	X	
535	989.0410	989.0488	5					X	
536	824.3675	824.3754	6				X	X	
537	706.7446	706.7514	7			X	X		
538	707.0314	707.0394	7	C ₂₁₁ H ₃₂₀ N ₆₀ O ₇₀ S ₄	[2408-2436] ^{6,8)}		H		
539	857.3980	857.4059	4	C ₁₅₁ H ₂₂₄ N ₄₂ O ₄₆ S ₂	[2426-2444] ⁵⁾		X		
540	686.1259	686.1263	5				X	X	
541	800.1714	800.1793	5	C ₁₇₅ H ₂₆₂ N ₄₈ O ₅₆ S ₂	[2426-2444]		X		
542	803.3718	803.3782	5	C ₁₇₅ H ₂₆₂ N ₄₈ O ₅₇ S ₂	[2426-2444] ⁹⁾	X	X	2436	
543	780.0558	780.0553	3	C ₁₀₂ H ₁₆₀ N ₂₈ O ₃₃ S ₁	[2434-2444] ⁵⁾	X	X		
544	970.1357	970.1436	3	C ₁₂₆ H ₁₉₈ N ₃₄ O ₄₃ S ₁	[2434-2444]	X			
545	727.8608	727.8596	4				X	X	
546	1369.6661	1369.6655	3	C ₁₈₄ H ₂₇₆ N ₄₆ O ₅₉ S ₁	[2437-2463] ⁵⁾		X		
547	1027.5017	1027.5011	4				X	X	
548	822.1946	822.2025	5					X	
549	1559.7536	1559.7539	3	C ₂₀₈ H ₃₁₄ N ₅₂ O ₆₉ S ₁	[2437-2463]	X	X		
550	1170.0685	1170.0674	4				X	X	
551	936.2565	936.2554	5				X	X	
552	1031.9957	1032.0038	4			C ₁₈₄ H ₂₇₈ N ₄₆ O ₆₀ S ₁	[2437-2463] ^{5,6)}		H
553	825.7966	825.8046	5					H	
554	1174.5621	1174.5699	4	C ₂₀₈ H ₃₁₆ N ₅₂ O ₇₀ S ₁	[2437-2463] ⁶⁾		H		
555	939.8496	939.8575	5					H	
556	783.3747	783.3826	6					H	
557	912.09847	912.1064	3	C ₁₂₀ H ₁₈₄ N ₃₀ O ₄₁ S	[2456-2465]		X	K2463	
558	891.7765	891.7770	3	C ₁₁₀ H ₁₈₅ N ₃₃ O ₄₂ S ₁	[2508-2522] ⁵⁾	X	X		
559	1081.8637	1081.8654	3	C ₁₃₄ H ₂₂₃ N ₃₉ O ₅₂ S ₁	[2508-2522]	X	X	K2508	
560	811.6508	811.6509	4					X	X

561	946.4485	946.4562	2	C ₈₁ H ₁₃₀ N ₂₂ O ₂₆ S ₂	[2574-2580] ⁵⁾		X	K2577
562	710.14588	710.1537	5	C ₁₅₅ H ₂₄₀ N ₄₆ O ₄₆ S ₂	[2719-2739] ^{5,8)}		X	K2735
563	824.2070	824.2067	5	C ₁₇₉ H ₂₇₈ N ₅₂ O ₅₆ S ₂	[2719-2739] ⁸⁾	X	X	
564	686.99905	687.0669	6				X	
565	807.6505	807.6491	4	C ₁₄₀ H ₂₁₉ N ₄₁ O ₄₃ S ₂	[2721-2739] ^{5,8)}	X	X	
566	760.3733	760.3738	5	C ₁₆₄ H ₂₅₇ N ₄₇ O ₅₃ S ₂	[2721-2739] ⁸⁾	X	X	
567	688.7094	688.7144	3	C ₉₀ H ₁₅₄ N ₂₆ O ₂₇ S	[2731-2739] ⁵⁾	X	X	
568	878.7948	878.8027	3	C ₁₁₄ H ₁₉₂ N ₃₂ O ₃₇ S	[2731-2739]		X	
569	659.3468	659.3540	4				X	
570	838.3967	838.4046	3	C ₁₀₈ H ₁₆₉ N ₂₉ O ₃₈ S	[2736-2743]	X	X	K2739
571	729.3596	729.3607	3	C ₉₃ H ₁₄₈ N ₂₈ O ₃₁ S ₁	[2736-2745] ⁵⁾	X	X	
572	547.2690	547.2725	4				X	
573	919.4480	919.4490	3	C ₁₁₇ H ₁₈₆ N ₃₄ O ₄₁ S ₁	[2736-2745]	X	X	
574	689.8375	689.8387	4				X	
575	552.0725	552.0725	5				X	
576	422.4539	422.4617	4	C ₆₈ H ₁₁₅ N ₂₃ O ₂₅ S	[2791-2795] ⁵⁾		X	K2792
577	565.0201	565.0280	4	C ₉₂ H ₁₅₃ N ₂₉ O ₃₅ S	[2791-2795]		X	
578	710.3357	710.3363	3	C ₈₆ H ₁₄₁ N ₂₇ O ₃₄ S ₁	[2792-2795]		X	
579	533.0034	533.0042	4				X	
580	537.4983	537.5068	4	C ₈₆ H ₁₄₃ N ₂₇ O ₃₅ S ₁	[2792-2795] ⁶⁾		H	
581	430.1986	430.2070	5				H	
582	989.0093	989.0097	2	C ₈₆ H ₁₄₁ N ₂₃ O ₂₈ S ₁	[2847-2854] ⁵⁾	X	X	K2851
583	659.6765	659.6757	3				X	
584	849.7626	849.7640	3	C ₁₁₀ H ₁₇₉ N ₂₉ O ₃₈ S ₁	[2847-2854]	X	X	
585	637.5747	637.5750	4				X	
586	665.6699	665.6793	3	C ₈₆ H ₁₄₃ N ₂₃ O ₂₉ S ₁	[2847-2854] ^{5,6)}		H	
587	642.0691	642.0776	4	C ₁₁₀ H ₁₈₁ N ₂₉ O ₃₉ S ₁	[2847-2854] ⁶⁾		H	
588	513.8553	513.8636	5				H	
589	608.6244	608.6323	6	C ₁₅₄ H ₂₄₈ N ₄₂ O ₅₈ S	[2916-2934]		X	K2929
590	676.6543	676.6622	3	C ₈₄ H ₁₃₈ N ₂₄ O ₃₂ S	[2989-2992]		X	K2991
591	618.4613	618.4692	6	C ₁₆₅ H ₂₄₅ N ₄₅ O ₅₁ S	[2989-3006]		X	
592	742.5957	742.6019	4	C ₁₂₄ H ₁₉₅ N ₃₉ O ₄₂ S ₂	[3027-3038]	X	X	K3031
593	594.2767	594.2831	5				X	
594	495.3960	495.4039	6				X	
595	557.0317	557.0396	4	C ₉₃ H ₁₅₇ N ₂₉ O ₃₂ S	[3124-3128]		X	K3126
596	423.2154	423.2237	5	C ₈₈ H ₁₅₀ N ₂₈ O ₃₀ S	[3127-3130]		X	K3129
597	622.5556	622.5634	4	C ₁₀₆ H ₁₇₁ N ₃₁ O ₃₆ S	[3129-3136]		X	K3130
598	498.2444	498.2523	5				X	
599	787.0449	787.0503	3	C ₁₀₀ H ₁₅₉ N ₂₉ O ₃₅ S ₁	[3130-3136]	X		
600	590.5405	590.5397	4				X	
601	594.6797	594.6875	5	C ₁₂₅ H ₁₉₇ N ₃₇ O ₄₅ S	[3162-3174]	X	X	K3165
602	898.0106	898.0134	5	C ₁₉₃ H ₂₈₉ N ₅₃ O ₆₇ S ₂	[3227-3252]	X	X	K3244
603	772.6464	772.6543	4	C ₁₃₇ H ₂₁₉ N ₃₇ O ₄₂ S	[3303-3321] ⁵⁾		X	K3317
604	915.2127	915.2205	4	C ₁₆₁ H ₂₅₇ N ₄₃ O ₅₂ S ₁	[3303-3321]		X	
605	732.3785	732.3780	5				X	
606	708.3740	708.3790	4			C ₁₂₇ H ₂₀₄ N ₃₄ O ₃₇ S	[3305-3321] ⁵⁾	
607	680.9523	680.9578	5	C ₁₅₁ H ₂₄₂ N ₄₀ O ₄₇ S	[3305-3321]	X		

608	546.2803	546.2881	4	C ₉₂ H ₁₅₆ N ₂₈ O ₃₁ S	[3322-3326]		X	K3325
609	778.68202	778.6873	3	C ₁₀₀ H ₁₄₈ N ₂₈ O ₃₅ S	[3375-3386] ⁵⁾	X	X	K3385
610	968.77071	968.7756	3	C ₁₂₄ H ₁₈₆ N ₃₄ O ₄₅ S	[3375-3386]	X	X	
611	726.82713	726.8336	4			X	X	
612	806.1199	806.1245	4	C ₁₃₅ H ₂₀₉ N ₄₁ O ₄₇ S ₂	[3375-3394] ^{5,8)}	X		

¹⁾ Numbers inside brackets correspond to the position of the proteolytic peptide within the sequence of KLH2. All peptides from KLH2 shown in this table are linked to (C¹-K¹⁶) fragment of Cys¹pP0, except those peptides where it is specified if they are linked to (C¹-K¹¹), (C¹-K¹⁴) or (C¹-K¹⁷) fragment of Cys¹pP0.

²⁾ Identification of type 2 peptides in the KLH2-Cys¹pP0 conjugate using pLink2 software [2].

³⁾ Identification of type 2 peptides in the KLH2-Cys¹pP0 conjugate using Kojak software [3].

⁴⁾ Indicates the position where the peptide Cys¹pP0 is added by the Michael addition to the free amino groups in the Lys residues or *N*-terminus of the KLH2.

⁵⁾ Peptide from KLH2 linked to (C¹-K¹¹) fragment of Cys¹pP0

⁶⁾ Identification of type 2 peptides with the hydrolyzed linker

⁷⁾ Peptide from KLH2 linked to (C¹-K¹⁷) fragment of Cys¹pP0

⁸⁾ Carbamidomethylation of Cys

⁹⁾ Met→ Met sulfoxide

¹⁰⁾ Peptide from KLH2 linked to (C¹-K¹⁴) fragment of Cys¹pP0

Table S8. Identification of type 2 peptides derived from the proteolytic digestion of KLH2-Cys¹pP0 conjugate with trypsin using the MED-FASP method [4]. Type 2 peptides were identified using transcyclized linker.

#	Assignment ¹⁾	z	Conjugation site ²⁾	m/z theor	m/z exp	Error (ppm)	RT1 ³⁾	RT2 ⁴⁾	Ion ⁵⁾
1	[1-7]-(C1-K16)	4	Nterm	596.8019	596.8028	-1.51	15.29	23.19	3
2	[7-22]-(C1-K11)	3	K7	915.4508	915.4522	-1.57	39.50	39.75	2
3	[7-22]-(C1-K16)	4		829.4060	829.4074	-1.63	36.59	36.83	3
4	[23-55]-(C1-K11) ⁶⁾	5	K52	941.0257	941.0270	-1.45	39.02	-	-
5	[23-55]-(C1-K16) ⁶⁾	6		879.4001	879.4014	-1.42	37.23	-	3
6		5		1055.0785	1055.0801	-1.48	37.23	-	3
7	[128-136]-(C1-K11)	3	K135	683.0135	683.0146	-1.56	34.11	34.52	1,2
8		2		1024.0164	1024.0180	-1.51	34.11	-	1
9	[128-136]-(C1-K16)	4		655.0781	655.0791	-1.53	31.48	31.86	1,3
10		3		873.1015	873.1028	-1.57	31.47	31.86	1,3
11	[136-139]-(C1-K16)	4	K136	515.5020	515.5028	-1.50	12.41	12.68	1,3
12	[140-156]-(C1-K11)	5	K148	601.2984	601.2993	-1.50	29.75	-	-
13	[140-156]-(C1-K16)	6		596.2940	596.2949	-1.45	28.34	-	3
14	[145-156]-(C1-K11)	4		612.3062	612.3071	-1.39	22.63	23.05	1,2
15		5		490.0465	490.0472	-1.43	22.63	23.06	1
16	[145-156]-(C1-K16)	6		503.5840	503.5848	-1.56	21.96	22.29	1,3
17		4		754.8722	754.8733	-1.42	21.96	22.29	1,3
18		5		604.0993	604.1002	-1.46	21.98	22.30	1,2,3
19		[230-252]-(C1-K11) ⁶⁾		5	K231	724.7275	724.7285	-1.41	35.68
20	4		905.6574	905.6587		-1.44	35.68	-	2
21	[230-252]-(C1-K16) ^{6,7)}	6	701.8174	701.8184		-1.38	30.13	-	3
22	[230-252]-(C1-K16) ⁶⁾	5	838.7803	838.7815		-1.43	33.62	33.78	3

23	[230-261]-(C1-K11) ⁶⁾	6		775.0263	775.0274	-1.46	43.39	-	2
24		5		929.8300	929.8314	-1.46	43.37	-	2
25	[230-261]-(C1-K16) ⁶⁾	6		870.0703	870.0716	-1.48	41.44	-	2,3
26		7		745.9185	745.9196	-1.49	41.44	-	3
27	[300-304]-(C1-K11)	3	K302	524.5911	524.5919	-1.40	12.47	12.88	1
28	[300-304]-(C1-K16)	4		536.2612	536.2620	-1.59	13.72	14.09	1,3
29		3		714.6791	714.6801	-1.45	13.72	14.09	1,3
30	[340-358]-(C1-K11)	3	K353	1063.8345	1063.8360	-1.41	54.31	-	2
31		4		798.1278	798.1290	-1.44	54.31	-	2
32	[340-358]-(C1-K16) ⁷⁾	4		944.6925	944.6939	-1.48	52.26	53.55	3
33	[340-358]-(C1-K16)	4		940.6938	940.6952	-1.46	53.87	-	2,3
34		3		1253.9225	1253.9243	-1.44	53.86	-	3
35		5		752.7566	752.7577	-1.46	53.90	-	3
36	[400-414]-(C1-K11)	5	K409	546.4802	546.4810	-1.54	20.55	20.72	-
37	[400-427]-(C1-K16)	7		659.6343	659.6353	-1.54	31.59	31.80	3
38		6		769.4054	769.4066	-1.52	31.59	31.78	3
39	[407-427]-(C1-K11)	5		642.7381	642.7390	-1.46	33.99	34.20	-
40	[407-427]-(C1-K16)	6		630.8270	630.8280	-1.56	32.25	32.46	3
41	[410-427]-(C1-K11)	4	K414	729.1246	729.1257	-1.44	38.01	38.20	2
42	[410-427]-(C1-K16)	5		697.5540	697.5551	-1.52	35.59	35.80	3
43		4		871.6906	871.6919	-1.46	35.60	35.80	3
44	[428-442]-(C1-K11)	3	K428	922.1108	922.1122	-1.52	36.57	36.85	2
45	[428-442]-(C1-K16)	4		834.4010	834.4023	-1.59	34.01	34.32	3
46	[443-447]-(C1-K11)	3	K443	544.5922	544.5930	-1.35	15.62	16.10	1
47	[443-447]-(C1-K16) ⁷⁾	4		555.2608	555.2616	-1.49	13.45	13.67	3
48		4		551.2621	551.2629	-1.45	16.19	16.60	1,3
49	[443-452]-(C1-K16)	4		713.3335	713.3346	-1.51	28.34	28.63	3

50	[448-469]-(C1-K16)	5	K452	812.9870	812.9882	-1.45	36.41	36.58	2,3
51		4		1015.9818	1015.9833	-1.45	36.41	36.58	2,3
52	[448-469]-(C1-K11)	4		873.4158	873.4171	-1.43	38.13	38.66	2
53	[470-478]-(C1-K16) ⁶⁾	5	K476	537.2527	537.2535	-1.45	15.85	16.20	2,3
54		4		671.3140	671.3149	-1.34	15.85	16.19	3
55	[652-656]-(C1-K16)	4	K653	538.2597	538.2605	-1.49	16.06	-	1,3
56	[666-682]-(C1-K11)	3	K669	977.1421	977.1436	-1.54	44.75	44.94	2
57	[703-728]-(C1-K16)	5	K722	948.6266	948.6279	-1.43	51.88	-	3
58		6		790.6901	790.6913	-1.48	51.88	-	3
59	[769-776]-(C1-K16)	4	K771	659.8017	659.8027	-1.44	31.36	-	3
60	[769-776]-(C1-K11)	3		689.3117	689.3126	-1.31	34.22	-	-
61	[826-835]-(C1-K11)	4	K829	525.9931	525.9938	-1.43	34.11	-	-
62	[917-931]-(C1-K11)	4	K917	705.3338	705.3348	-1.42	44.32	44.50	2
63		3		940.1092	940.1105	-1.35	44.29	-	-
64	[917-931]-(C1-K16)	5		678.5214	678.5224	-1.44	41.22	41.38	3
65		4		847.8998	847.9010	-1.44	41.22	41.38	3
66	[1046-1062]-(C1-K11)	4		K1060	787.3945	787.3956	-1.46	50.13	-
67	[1063-1077]-(C1-K11) ⁶⁾	3	K1064	884.4082	884.4094	-1.39	30.62	30.92	-
68	[1063-1077]-(C1-K16) ⁶⁾	4		806.1241	806.1252	-1.43	28.60	28.89	3
69	[1078-1096]-(C1-K11)	3	K1078	1046.8424	1046.8440	-1.53	47.37	47.55	-
70		4		785.3838	785.3850	-1.46	47.39	47.55	-
71	[1078-1096]-(C1-K16)	4		927.9497	927.9512	-1.59	44.07	44.26	3
72		3		1236.9304	1236.9323	-1.54	44.08	44.26	3
73	[1078-1103]-(C1-K11)	5		798.9932	798.9944	-1.45	45.85	45.97	-
74		4		998.4895	998.4910	-1.50	45.86	45.97	-
75	[1078-1103]-(C1-K16)	6		761.0396	761.0408	-1.51	43.49	43.64	3
76		5		913.0460	913.0474	-1.47	43.49	43.64	3

77	[1261-1264]-(C1-K16)	3	K1261	700.0158	700.0169	-1.52	16.46	16.89	1,3
78	[1276-1292]-(C1-K11)	3	K1283	983.8296	983.8311	-1.56	45.72	45.85	-
79	[1276-1292]-(C1-K16)	4		880.6902	880.6915	-1.50	42.71	-	3
80		3		1173.9176	1173.9194	-1.56	42.71	-	3
81	[1284-1296]-(C1-K11)	4	K1292	626.5559	626.5568	-1.44	32.61	32.95	-
82	[1336-1366]-(C1-K16)	4	K1349	1261.0936	1261.0956	-1.55	54.08	54.35	2,3
83		5		1009.0765	1009.0780	-1.49	54.08	54.35	2,3
84	[1436-1461]-(C1-K11)	4	K1437	1003.4827	1003.4842	-1.47	53.82	-	2
85	[1436-1461]-(C1-K16)	5		917.0405	917.0419	-1.50	53.41	-	2,3
86		7		655.3169	655.3179	-1.46	53.42	-	3
87	[1472-1488]-(C1-K11) ⁶⁾	6	K1474	524.4108	524.4116	-1.40	27.59	-	-
88	[1474-1488]-(C1-K11) ⁶⁾	5		565.2586	565.2594	-1.42	27.47	-	-
89	[1475-1488]-(C1-K11) ⁶⁾	3	K1475	898.7276	898.7288	-1.37	27.70	28.00	-
90	[1489-1507]-(C1-K11)	3	K1489	1057.5118	1057.5134	-1.48	41.89	42.10	-
91	[1489-1507]-(C1-K16) ⁷⁾	4		939.9506	939.9520	-1.52	35.00	35.25	3
92	[1489-1507]-(C1-K16)	4		935.9518	935.9533	-1.60	38.94	39.14	3
93	[1490-1520]-(C1-K11)	5	K1507	932.2344	932.2357	-1.39	46.37	46.47	-
94	[1490-1520]-(C1-K16)	5		1046.2872	1046.2887	-1.43	44.37	-	3
95	[1523-1545]-(C1-K11)	3	K1537	1289.9537	1289.9556	-1.45	49.31	-	-
96		4		967.7172	967.7186	-1.47	49.31	-	-
97	[1523-1545]-(C1-K16)	5		888.4281	888.4294	-1.51	46.95	-	3
98		4		1110.2832	1110.2848	-1.49	46.96	-	3
99	[1603-1614]-(C1-K11)	4	K1605	609.0608	609.0617	-1.52	41.23	41.50	1
100	[1603-1614]-(C1-K16)	5		601.5030	601.5039	-1.53	38.38	38.60	3
101	[1606-1627]-(C1-K16)	5	K1614	790.3970	790.3982	-1.54	53.89	-	3
102		4		987.7443	987.7459	-1.54	53.88	-	3
103		6		658.8321	658.8331	-1.59	53.90	-	3

104	[1615-1631]-(C1-K11)	4	K1627	709.3612	709.3623	-1.52	46.17	46.32	-
105	[1671-1681]-(C1-K11)	3	K1671	744.7100	744.7112	-1.61	29.75	-	-
106	[1671-1681]-(C1-K16)	4		701.3504	701.3516	-1.68	27.81	28.12	3
107	[1672-1685]-(C1-K16)	5	K1681	646.3253	646.3263	-1.58	41.40	-	3
108		4		807.6547	807.6559	-1.55	41.41	-	3
109		3		1076.5370	1076.5387	-1.55	41.41	-	3
110	[1672-1685]-(C1-K11)	4		665.0887	665.0897	-1.54	40.05	-	2
111		3		886.4490	886.4504	-1.54	44.51	-	-
112	[1686-1689]-(C1-K16)	4		K1686	507.5108	507.5116	-1.58	16.15	16.56
113	[1738-1749]-(C1-K11)	3	K1748	832.7436	832.7449	-1.56	38.93	39.33	-
114	[1738-1749]-(C1-K16)	4		767.3757	767.3768	-1.50	36.02	36.27	2,3
115		3		1022.8316	1022.8332	-1.56	36.02	36.27	3
116	[1764-1792]-(C1-K16)	6	K1784	826.2419	826.2432	-1.57	53.57	-	3
117		5		991.2888	991.2903	-1.49	53.57	-	3
118	[1785-1806]-(C1-K16)	4	K1792	1016.7379	1016.7395	-1.55	49.56	-	3
119	[1898-1912]-(C1-K11) ⁶⁾	4	K1902	664.8119	664.8129	-1.43	26.51	26.75	1
120		3		886.0799	886.0812	-1.47	26.52	26.75	-
121	[1898-1912]-(C1-K16) ⁶⁾	5		646.1039	646.1048	-1.42	25.14	25.40	1,3
122		4		807.3779	807.3791	-1.46	25.14	25.40	1,3
123	[1913-1928]-(C1-K16)	4	K1916	848.6612	848.6625	-1.56	44.13	44.35	3
124	[1917-1929]-(C1-K11)	3	K1928	842.0720	842.0732	-1.46	39.34	39.60	-
125	[1917-1929]-(C1-K16)	3		1032.1600	1032.1616	-1.52	36.62	-	3
126		4		774.3720	774.3731	-1.45	36.35	36.62	3
127	[1929-1944]-(C1-K11)	5	K1935	611.4816	611.4824	-1.37	28.19	28.50	-
128		4		764.1000	764.1011	-1.44	28.16	28.50	-
129	[1929-1944]-(C1-K16)	6		604.7800	604.7808	-1.35	26.90	27.22	3
130	[1930-1944]-(C1-K11)	4		725.0748	725.0758	-1.41	31.07	31.38	2

131	[1930-1944]-(C1-K16)	4		867.6408	867.6420	-1.44	29.29	29.59	3	
132		5		694.3142	694.3152	-1.44	29.30	29.59	3	
133	[1945-1974]-(C1-K16)	5	K1967	1022.4710	1022.4725	-1.51	53.76	-	3	
134	[1945-1974]-(C1-K11)	4		1135.2709	1135.2725	-1.41	54.04	-	-	
135	[2023-2030]-(C1-K11)	3	K2025	681.0062	681.0073	-1.62	29.13	29.55	1,2	
136	[2023-2030]-(C1-K16)	4		653.5726	653.5737	-1.61	27.30	27.64	1,3	
137	[2026-2033]-(C1-K11)	3	K2030	660.3344	660.3354	-1.51	30.26	30.60	1	
138	[2026-2033]-(C1-K16)	4		638.0687	638.0698	-1.65	28.19	28.50	1,3	
139	[2031-2036]-(C1-K16)	4	K2033	576.5302	576.5311	-1.56	21.06	21.33	1,3	
140	[2034-2038]-(C1-K16)	3	K2036	744.0239	744.0250	-1.52	21.96	22.32	1,3	
141		4		558.2699	558.2708	-1.48	21.96	22.32	1,3	
142	[2037-2050]-(C1-K16)	5	K2046	664.1379	664.1389	-1.54	35.32	35.55	3	
143	[2039-2050]-(C1-K11)	4		620.0582	620.0592	-1.57	37.60	37.87	-	
144		3		826.4084	826.4096	-1.49	37.60	37.87	-	
145	[2039-2050]-(C1-K16)	4		762.6242	762.6254	-1.57	34.90	35.16	3	
146		5		610.3009	610.3019	-1.61	34.90	35.17	1,3	
147	[2097-2106]-(C1-K16) ⁷⁾	3		K2102	892.4294	892.4308	-1.53	28.32	28.68	2,3
148		4	669.5740		669.5750	-1.53	28.32	28.68	3	
149	[2097-2106]-(C1-K16)	4	665.5753		665.5763	-1.50	35.21	35.57	1,2,3	
150		3	887.0978		887.0991	-1.50	35.20	35.58	1,2,3	
151	[2097-2106]-(C1-K11) ⁷⁾	3	702.3414		702.3425	-1.52	30.32	30.75	1,2	
152	[2097-2106]-(C1-K11)	3	697.0098		697.0108	-1.48	38.37	38.75	1,2	
153		2	1045.0107		1045.0123	-1.48	38.37	38.75	2	
154	[2154-2183]-(C1-K16)	5	K2179		1001.4931	1001.4946	-1.52	54.21	-	3
155	[2180-2198]-(C1-K11)	3	K2183		1107.8665	1107.8682	-1.47	54.08	-	2
156	[2180-2198]-(C1-K16)	4			973.7178	973.7193	-1.51	53.56	-	2,3
157	[2199-2212]-(C1-K16)	4	K2208	776.6336	776.6348	-1.51	37.61	37.83	3	

158		3		1035.1755	1035.1771	-1.55	37.63	37.83	3
159	[2199-2212]-(C1-K11)	3		845.0875	845.0888	-1.50	40.90	41.15	-
160	[2209-2220]-(C1-K11)	3	K2212	800.3807	800.3819	-1.46	28.23	28.55	-
161	[2209-2220]-(C1-K16)	3		990.4687	990.4702	-1.48	26.59	26.90	1,2,3
162		4		743.1034	743.1046	-1.58	26.59	26.90	1,2,3
163	[2299-2309]-(C1-K11)	3	K2308	790.0785	790.0797	-1.48	48.42	48.60	-
164	[2299-2309]-(C1-K16)	4		735.3768	735.3779	-1.53	44.90	45.10	3
165	[2311-2316]-(C1-K11)	4	K2311	457.2334	457.2341	-1.53	10.38	10.99	-
166	[2311-2316]-(C1-K16)	5		480.0411	480.0418	-1.50	12.33	12.52	1,3
167		4		599.7994	599.8004	-1.54	12.33	12.52	1,3
168	[2381-2385]-(C1-K16)	4	K2381	547.5166	547.5174	-1.46	12.58	12.90	1,3
169	[2408-2433]-(C1-K11) 6,7)	5	K2425	797.1534	797.1545	-1.41	45.03	-	-
170	[2408-2433]-(C1-K11) ⁶⁾	5		793.9544	793.9556	-1.44	48.44	-	-
171		4		992.1911	992.1925	-1.39	48.45	-	-
172	[2408-2433]-(C1-K16) ⁶⁾	5		908.0072	908.0085	-1.45	46.00	-	3
173		6		756.8407	756.8417	-1.37	46.04	-	3
174	[2434-2444]-(C1-K11)	3	K2436	780.0540	780.0552	-1.54	42.71	42.93	2
175	[2434-2444]-(C1-K16)	4		727.8585	727.8596	-1.48	39.39	39.60	1,2,3
176		3		970.1420	970.1435	-1.55	39.38	39.60	3
177	[2437-2463]-(C1-K16)	4	K2444	1170.0654	1170.0673	-1.58	53.83	-	2,3
178	[2508-2522]-(C1-K11)	3	K2508	891.7755	891.7769	-1.61	41.67	41.93	2
179	[2508-2522]-(C1-K16)	4		811.6496	811.6509	-1.57	38.76	39.00	2,3
180		3		1081.8635	1081.8652	-1.60	38.78	-	3
181	[262-274]-(C1-K16)	5	K265	612.8805	612.8814	-1.37	23.32	23.58	3
182		4		765.8486	765.8497	-1.47	23.32	23.58	3
183	[262-274]-(C1-K11)	4		623.2826	623.2835	-1.44	24.35	24.73	1,2
184		3		830.7076	830.7087	-1.36	24.35	24.73	-

185	[2721-2739]-(C1-K16) ⁶⁾	5	K2735	760.3726	760.3738	-1.55	36.80	36.95	3
186	[2721-2739]-(C1-K11) ⁶⁾	4		807.6478	807.6491	-1.55	39.06	39.23	-
187	[2736-2745]-(C1-K11)	3	K2739	729.3595	729.3607	-1.55	29.25	29.65	2
188	[2736-2745]-(C1-K16)	4		689.8376	689.8386	-1.52	27.30	27.60	2,3
189		3	919.4475	919.4489	-1.56	27.30	27.60	3	
190	[2791-2795]-(C1-K11)	4	K2792	422.4610	422.4616	-1.54	9.62	9.80	1
191	[2791-2795]-(C1-K16)	4		565.0270	565.0279	-1.55	11.34	11.59	1,3
192		4		533.0034	533.0042	-1.41	13.16	13.55	1,3
193	[2792-2795]-(C1-K16)	3		710.3352	710.3362	-1.50	13.16	13.55	1,3
194	[2847-2854]-(C1-K11)	3	K2851	659.6746	659.6757	-1.62	36.71	37.02	1,2
195		2		989.0080	989.0096	-1.62	36.70	37.02	-
196	[2847-2854]-(C1-K16)	3		849.7626	849.7640	-1.61	33.64	33.96	3
197		4		637.5739	637.5750	-1.61	33.65	33.95	1,3
198	[2989-3006]-(C1-K16)	6	K2991	618.4682	618.4691	-1.51	36.38	-	3
199	[3027-3038]-(C1-K16)	5	K3031	594.2822	594.2830	-1.38	19.74	19.94	1,3
200		6		495.4031	495.4038	-1.45	19.75	19.94	3
201		4		742.6008	742.6018	-1.38	19.74	19.94	3
202	[3129-3136]-(C1-K16)	5	K3130	498.2515	498.2522	-1.49	14.84	15.20	3
203		4		622.5624	622.5634	-1.53	14.85	15.20	1,3
204	[3162-3174]-(C1-K16)	5	K3165	594.6866	594.6874	-1.45	23.00	23.35	3
205	[3227-3252]-(C1-K16)	5	K3244	898.0120	898.0133	-1.45	47.41	-	3
206	[3322-3326]-(C1-K16)	4	K3325	546.2872	546.2881	-1.60	20.03	20.35	1,3
207	[3375-3394]-(C1-K16) ⁶⁾	4	K3385	948.6892	948.6906	-1.50	31.30	31.50	3
208	[3375-3394]-(C1-K11) ⁶⁾	4		806.1232	806.1244	-1.49	33.38	33.60	2

¹⁾ Numbers inside brackets and parentheses correspond to the position of the proteolytic peptide within the sequences of KLH2 and Cys¹pP0, respectively.

- 2) Indicates the position where the peptide Cys¹pP0 is added by the Michael addition to the free amino groups in the Lys residues or N-terminus of the KLH2.
- 3) Retention time of the first peak of type 2 peptides
- 4) Retention time of the second peak of type 2 peptides
- 5) Type 2 peptides containing the P+71 ion fragment (1), C+80 ion fragment (2), y5 α/β fragment ion ($m/z = 589.282, 1+$) (3) or not containing any ion fragments (-).
- 6) Carbamidomethylation of Cys
- 7) Met \rightarrow Met sulfoxide

Table S9. Identification of type 2 peptides derived from the proteolytic digestion of KLH2-Cys¹pP0 conjugate with trypsin using the MED-FASP method [4]. Type 2 peptides were identified using hydrolyzed thiosuccinimide linker.

	Assignment ¹⁾	z	Conjugation site ²⁾	m/z theor.	m/z exp.	Error (ppm)	RT1 ³⁾	RT2 ⁴⁾	RT3 ⁵⁾	RT4 ⁶⁾	Ion ⁷⁾
1	[128 - 136]-(C1-K11)	3	(K135)	689.0140	689.0181	-1.60	30.08	-	-	-	-
2	[128 - 136]-(C1-K16)	4		659.5804	659.5814	-1.59	24.03	24.28	-	-	5
3	[145 - 156]-(C1-K11)	4	(K148)	616.8088	616.8094	-1.46	19.64	19.48	20.35	20.55	1,4
4		5		493.6486	493.6493	-1.46	19.64	19.48	20.36	20.55	-
5	[145 - 156]-(C1-K16)	5		604.4014	604.4023	-1.48	19.60	19.43	20.12	20.32	1,3,6,5
6		6		506.5858	506.5866	-1.48	19.59	19.43	20.12	20.32	1,3,5
4		4		459.3448	459.3459	-1.48	19.60	19.43	20.11	20.31	1,5
8	[262 - 244]-(C1-K11)	4		(K265)	624.4853	624.4861	-1.35	20.88	20.98	21.65	21.8
9		5	502.4298		502.4305	-1.35	20.88	20.98	21.65	21.8	-
10	[262 - 244]-(C1-K16)	5	616.4826		616.4835	-1.40	20.53	20.65	21.09	21.20	1, 5
11		6	513.9034		513.9042	-1.52	20.53	20.65	21.09	21.20	5
12	[300 - 304]-(C1-K16)	4	(K302)	540.4639	540.4644	-1.48	9.68	9.80	10.60	11.04	1,3, 5
13		5		432.8124	432.8133	-1.43	9.68	9.82	10.60	11.06	1,3,4,5
14	[340 - 358]-(C1-K16) ⁸⁾	5	(K353)	459.5544	459.5588	-1.45	50.11	-	-	-	5
15	[340 - 358]-(C1-K16)	5		456.3584	456.3598	-1.48	52.26	53.50	-	-	2,5
16		4		945.1964	945.1948	-1.51	52.25	-	-	-	5
14	[443 - 444]-(C1-K16)	4	(K443)	555.4644	555.4656	-1.53	12.49	12.63	-	-	1,3,5
18		5		444.8133	444.8140	-1.54	12.49	12.63	13.45	13.65	5
19	[448 - 469]-(C1-K11)	5	(K452)	402.5363	402.5343	-1.45	35.24	36.52	-	-	2
20	[448 - 469]-(C1-K16)	5		816.5891	816.5903	-1.44	33.32	34.35	-	-	2,5
21		6		680.6589	680.6599	-1.44	33.33	34.35	-	-	5
22	[652 - 656]-(C1-K11) ⁹⁾	3	(K653)	533.2592	533.2600	-1.44	13.15	13.55	-	-	-
23	[652 - 656]-(C1-K16) ⁹⁾	4		542.4623	542.4631	-1.54	14.24	14.68	-	-	5
24	[652 - 656]-(C1-K16)	3		423.3442	423.3483	-1.44	14.24	14.68	-	-	5
25	[914 - 931]-(C1-K11)	4	(K914)	409.8364	409.8345	-1.48	41.18	42.58	-	-	1

26	[914 - 931]-(C1-K16)	5		682.1235	682.1245	-1.44	38.43	39.64	-	-	6,5
24		4		852.4024	852.4034	-1.50	38.42	-	-	-	6,5
28	[1063 - 1044]-(C1-K11)	4	(K1064)	668.0604	668.0614	-1.46	24.45	24.90	-	-	-
29	[1063 - 1044]-(C1-K16)	4		810.6264	810.6249	-1.48	24.43	24.89	26.14	26.30	5
30	[1063 - 1044]-(C1-K16) ⁹⁾	5		648.4029	648.4039	-1.51	24.43	24.89	26.14	26.30	5
31	[1048 - 1096]-(C1-K11)	4		489.8864	489.8846	-1.52	45.15	-	-	-	2
32	[1048 - 1103]-(C1-K11)	5	(K1048)	802.5953	802.5965	-1.44	44.34	-	-	-	2
33	[1048 - 1096]-(C1-K16)	4		932.4524	932.4538	-1.53	40.53	42.03	-	-	5
34		5		446.1635	446.1646	-1.50	40.53	42.03	-	-	5
35		4		655.0366	655.0345	-1.46	40.98	42.11	-	-	5
36		6		464.0414	464.0425	-1.46	40.98	42.12	-	-	5
34	[1104 - 1130]-(C1-K11)	4	(K1108)	1062.4958	1062.4946	-1.51	54.09	-	-	-	2
38	[1261 - 1264]-(C1-K11)	4	(K1261)	384.2004	384.2010	-1.55	12.42	12.40	-	-	1, 3
39	[1261 - 1264]-(C1-K16)	4		529.4664	529.4643	-1.60	12.63	12.9	13.61	13.90	1,3,5
40		5		424.0144	424.0154	-1.56	12.63	12.9	13.61	13.90	1,3,5
41	[1246 - 1292]-(C1-K16)	4	(K1283)	885.1928	885.1942	-1.55	40.19	-	-	-	5
42	[1284 - 1296]-(C1-K16)	5	(K1292)	619.1012	619.1021	-1.45	24.46	24.58	28.35	28.50	6, 5
43	[1336 - 1366]-(C1-K11) ⁹⁾	4	(K1349)	1123.0303	1123.0320	-1.44	54.31	54.45	54.68	-	2
44	[1336 - 1366]-(C1-K16) ⁹⁾	5		1012.6486	1012.6801	-1.50	54.00	-	-	-	2, 5
45	[1436 - 1461]-(C1-K11)	5	(K1434)	806.5898	806.5910	-1.51	53.42	53.65	53.83	-	2, 6
46	[1436 - 1461]-(C1-K16)	5		920.6426	920.6440	-1.52	51.22	52.44	-	-	5
44		6		464.3401	464.3413	-1.56	51.26	52.46	-	-	5
48		4		654.8898	654.8908	-1.52	51.24	52.46	-	-	5
49	[1490 - 1520]-(C1-K16)	6	(K1504)	845.0454	845.0440	-1.44	41.85	42.45	42.92	-	2, 5
50	[1523 - 1545]-(C1-K16)	6	(K1534)	443.5265	443.5246	-1.48	44.43	45.64	-	-	5
51		5		892.0302	892.0316	-1.52	44.42	45.64	44.44	-	5
52	[1546 - 1550]-(C1-K16)	4	(K1546)	568.0208	568.0216	-1.41	15.68	-	-	-	5
53	[1642 - 1685]-(C1-K11)	4	(K1681)	669.5913	669.5924	-1.61	41.33	43.04	-	-	2, 6, 4
54		5		535.8446	535.8455	-1.60	41.36	43.04	-	-	6, 4
55	[1642 - 1685]-(C1-K16)	4		812.1543	812.1586	-1.60	38.64	40.13	-	-	1, 2, 5

56		5		649.9244	649.9285	-1.60	38.64	40.13	-	-	2, 5
54	[1438 - 1449]-(C1-K11)	4	(K1448)	629.3123	629.3133	-1.55	34.69	35.95	-	-	2, 6
58	[1438 - 1449]-(C1-K16)	4		441.8483	441.8495	-1.55	32.25	32.38	33.36	-	5
59	[1485 - 1806]-(C1-K16)	5	(K1492)	814.1940	814.1953	-1.54	44.50	-	-	-	2, 5
60	[1898 - 1912]-(C1-K11)	4	(K1902)	669.3145	669.3155	-1.49	23.01	23.15	24.20	24.35	1, 4
61	[1898 - 1912]-(C1-K16)	4		811.8805	811.8814	-1.51	21.25	21.45	22.09	22.20	5
62	[1898 - 1912]-(C1-K16) ⁹⁾	6		541.5896	541.5904	-1.51	23.14	-	-	-	5
63	[1930 - 1944]-(C1-K11)	5	(K1935)	583.8635	583.8643	-1.44	24.24	24.35	28.25	28.42	-
64		4		429.5444	429.5485	-1.44	24.24	24.35	28.28	28.42	-
65	[1930 - 1944]-(C1-K16)	6		581.4649	581.4658	-1.43	26.00	26.88	24.08	-	5
66		5		694.9163	694.9143	-1.46	26.00	26.15	26.88	24.08	5
64	[2023 - 2030]-(C1-K11)	4	(K2025)	515.5092	515.5101	-1.40	23.85	24.10	-	-	-
68	[2094 - 2106]-(C1-K11) ⁸⁾	3	(K2102)	403.0133	403.0143	-1.52	32.25	32.4	34.03	34.24	1, 2, 6, 4
69	[2094 - 2106]-(C1-K11)	3		408.3449	408.3460	-1.55	25.00	25.25	26.24	26.49	2,
40		4		524.5119	524.5124	-1.56	32.21	32.41	34.03	34.24	6
41	[2094 - 2106]-(C1-K16) ⁸⁾	4		644.0466	644.0444	-1.59	23.45	23.95	24.80	25.05	5
42		5		539.4629	539.4634	-1.48	23.49	23.95	24.80	25.10	6, 5
43		4		640.0449	640.0490	-1.54	29.96	30.15	31.483	31.40	1, 2, 5
44	[2094 - 2106]-(C1-K16)	3		893.1013	893.1024	-1.53	29.96	30.15	31.483	31.40	1, 2, 5
45		5		536.2639	536.2644	-1.53	29.96	30.15	31.483	31.40	5
46	[2209 - 2220]-(C1-K11)	4	(K2212)	605.0401	605.0410	-1.49	23.45	23.95	25.14	25.30	5
44	[2209 - 2220]-(C1-K16)	5		598.2864	598.2844	-1.54	22.44	22.9	23.48	23.94	6,5
48		4		444.6061	444.6042	-1.50	22.44	22.9	23.48	23.94	5
49	[2381 - 2385]-(C1-K16)	4	(K2381)	552.0192	552.0200	-1.54	9.39	9.48	9.65	9.45	1, 3, 5
80	[2434 - 2444]-(C1-K16)	4	(K2436)	432.3611	432.3622	-1.54	36.43	-	-	-	5
81		5		586.0904	586.0914	-1.60	36.45	-	-	-	5
82	[2508 - 2522]-(C1-K16)	5	(K2508)	653.1233	653.1244	-1.65	36.55	-	-	-	5
83	[2436 - 2445]-(C1-K11)	4	(K2439)	551.4442	551.4451	-1.59	23.46	23.95	25.43	25.65	5
84	[2492 - 2495]-(C1-K16)	5	(K2492)	430.2063	430.2040	-1.63	9.58	9.64	10.00	10.30	1, 3, 6,5
85	[2844 - 2854]-(C1-K11)	3	(K2851)	665.6481	665.6492	-1.65	32.21	34.03	-	-	1

86		4	499.5105	499.5114	-1.40	32.24	34.02	-	-	-
84	[2844 - 2854]-(C1-K16)	5	513.8628	513.8634	-1.60	29.92	31.40	-	-	5
88		4	642.0465	642.0446	-1.64	29.89	31.39	-	-	1, 5

- 1) Numbers inside brackets and parentheses correspond to the position of the proteolytic peptide within the sequences of KLH2 and Cys¹pP0, respectively.
- 2) Indicates the position where the peptide Cys¹pP0 is added by the Michael addition to the free amino groups in the Lys residues or N-terminus of the KLH2.
- 3) Retention time of the first peak of type 2 peptides
- 4) Retention time of the second peak of type 2 peptides
- 5) Retention time of the third peak of type 2 peptides
- 6) Retention time of the fourth peak of type 2 peptides
- 7) Type 2 peptides containing the P+71 ion fragment (1), C+98 ion fragment (2), P+203 ion fragment (3), backbone fragment ions $*(\alpha/\beta)b_n/*(\alpha/\beta)y_n$ generated from P+71 ion (4), backbone fragment ions $*(\alpha/\beta)b_n/*(\alpha/\beta)y_n$ generated from C+98 ion (5), y5 α/β fragment ion (m/z = 589.282, 1+) (6) or not containing any ion fragments (-).
- 8) Met → Met sulfoxide
- 9) Carbamidomethylation of Cys

Table S10. Amino acid changes detected in KLH1 using the Peaks software [5].

#	Assignment ⁱ⁾	Amino acid change	m/z exp	z	error (ppm)
1	[84-117]	Leu ⁸⁹ →Met	1279.2992	3	2.2
1	[84-117] ⁱ⁾	Leu ⁸⁹ →Met	963.7238	4	1.0
1	[86-117]	Leu ⁸⁹ →Met	1175.2340	3	4.2
2	[190-215]	Glu ¹⁹⁸ →Gln	789.3654	4	-2.5
3	[190-215]	Asp ¹⁹¹ →Val	1047.1602	3	-2.3
4 y 5	[451-462]	Ala ⁴⁵⁹ →Val, Tyr ⁴⁶¹ →Phe	676.8098	2	0.4
6	[760-767]	Ser ⁷⁶⁶ →Ala	381.7296	2	1.4
7	[907-916]	Leu ⁹¹⁶ →Arg	565.8121	2	0.4
8	[956-969]	Trp ⁹⁵⁹ →Ser	527.9172	3	5.0
9	[1079-1097]	Glu ¹⁰⁷⁹ →Gln	694.0261	3	-8.9
10, 11	[1605-1612]	Leu ¹⁶⁰⁵ →Tyr, Lys ¹⁶⁰⁹ →Ser	455.7479	2	0.5
12	[1764-1774]	Thr ¹⁷⁷⁴ →Arg	622.3472	2	-1.6
13	[1764-1782]	Glu ¹⁷⁶⁸ →Gln	698.3643	3	-4.4
14	[1845-1857]	Ile ¹⁸⁴⁵ →Gln	484.2420	3	7.2
15	[1858-1883]	Glu ¹⁸⁵⁸ →Gln	779.6208	4	-6.2
16	[1915-1942]	Ser ¹⁹¹⁷ →Val	833.1284	4	-7.6
17	[2101-2127]	Ala ²¹¹⁷ →Val	913.1328	3	0.3
18	[2101-2127]	Asp ²¹⁰⁹ →Asn	903.4582	3	-2.9
19	[2105-2127]	Ala ²¹¹⁷ →Val	768.3929	3	-1.0
20	[2152-2162]	Arg ²¹⁶² →Lys	677.3751	2	1.2
21	[2178-2189]	His ²¹⁸⁴ →Asp	470.5863	3	1.5
22, 23	[2271-2290]	Ser ²²⁷⁴ →Ala, Ile ²²⁸⁶ →Ala	777.0494	3	2.1
22, 23	[2271-2291]	Ser ²²⁷⁴ →Ala, Ile ²²⁸⁶ →Ala	822.7377	3	4.4
24	[2274-2296]	Asp ²²⁸⁴ →Asn	542.6719	5	-5.8
25	[2276-2296]	Ile ²²⁸⁶ →Ala	617.7957	4	-2.5
25	[2276-2291]	Ile ²²⁸⁶ →Ala	631.2957	3	-1.9
25	[2276-2296]	Ile ²²⁸⁶ →Ala	617.7974	4	0.4
25	[2277-2296]	Ile ²²⁸⁶ →Ala	596.0397	4	0.9
25	[2277-2291]	Ile ²²⁸⁶ →Ala	902.9265	2	0.9
25	[2277-2296]	Ile ²²⁸⁶ →Ala	1191.0726	2	1.3
25	[2278-2291]	Ile ²²⁸⁶ →Ala	430.2085	4	0.1
25	[2278-2296]	Ile ²²⁸⁶ →Ala	459.6284	5	4.5
26	[2318-2340]	Tyr ²³³¹ →His	695.8424	4	-1.1
26	[2321-2340]	Tyr ²³³¹ →His	631.5591	4	0.5
26	[2325-2340]	Tyr ²³³¹ →His	513.2506	4	-0.9
26	[2326-2340]	Tyr ²³³¹ →His	645.9855	3	1.2

26	[2326-2340]	Tyr ²³³¹ →His	645.9855	3	1.2
26	[2328-2340]	Tyr ²³³¹ →His	412.9547	4	-0.7
26	[2330-2340]	Tyr ²³³¹ →His	468.8961	3	-2.1
26	[2330-2340]	Tyr ²³³¹ →His	468.8969	3	-0.5
26	[2331-2340]	Tyr ²³³¹ →His	634.3090	2	-5.5
27	[2576-2587]	Gln ²⁵⁷⁶ →Lys	650.3290	2	2.6
28	[2576-2633] ⁱ⁾	Gly ²⁵⁹⁵ →Trp	827.6458	8	-0.2
29, 30, 31, 32	[2790-2804]	Glu ²⁷⁹⁰ →Ser, Glu ²⁷⁹¹ →His, Thr ²⁷⁹⁴ →Val, Ala ²⁷⁹⁷ →Gly	424.9880	4	1.3
33	[2843-2852]	Arg ²⁸⁴⁴ →His	632.3389	2	-2.1

¹⁾ Numbers inside brackets correspond to the position of the proteolytic peptide of the sequence of KLH1.

i) Methionine sulfoxide

Table S11. Amino acid changes detected in KLH2 using the Peaks software [5].

#	Assignment ¹⁾	Modification site	m/z	z	error (ppm)
1	[8-22]	Ser ¹¹ →His	834.9264	2	1.0
2	[8-22]	Ser ¹³ →Asn	823.4216	2	5.0
3	[73-80]	Leu ⁷⁸ →Met	519.2764	2	0.7
4	[73-80]	Leu ⁷⁸ →Met, Leu ⁷³ →Asn	519.7565	2	1.9
5	[87-119]	Gly ¹¹⁴ →Asp	1246.9608	3	-0.3
5	[92-119]	Gly ¹¹⁴ →Asp	1085.2017	3	2.0
5	[94-119]	Gly ¹¹⁴ →Asp	749.1242	4	2.3
5	[97-119]	Gly ¹¹⁴ →Asp	627.3265	4	0.7
5	[99-119]	Gly ¹¹⁴ →Asp	1159.6086	2	3.2
5	[99-119]	Gly ¹¹⁴ →Asp	580.3090	4	4.9
5	[100-119]	Gly ¹¹⁴ →Asp	548.0449	4	-1.1
5	[101-119]	Gly ¹¹⁴ →Asp	692.6992	3	3.4
5	[102-119]	Gly ¹¹⁴ →Asp	494.5109	4	-3.3
5	[102-119]	Gly ¹¹⁴ →Asp	659.0153	3	1.6
5	[106-119]	Gly ¹¹⁴ →Asp	500.2685	3	0.6
5	[106-116]	Gly ¹¹⁴ →Asp	628.8299	2	0.4
5	[108-119]	Gly ¹¹⁴ →Asp	643.8235	2	1.7
5	[109-119]	Gly ¹¹⁴ →Asp	600.3060	2	-0.6
5	[110-119]	Gly ¹¹⁴ →Asp	531.7769	2	0
5	[113-119]	Gly ¹¹⁴ →Asp	353.1743	2	0.1
6	[120-135]	Asp ¹²² →His	494.0011	4	2.5
7	[192-217]	Glu ²⁰⁰ →Gln	800.1133	4	-6.9
8	[202-217]	Ser ²⁰³ →Thr	655.3145	3	-0.2
8	[203-217]	Ser ²⁰³ →Thr	621.6306	3	-2.5
9	[232-265]	Asp ²³⁷ →Asn	786.1641	5	-9.8
10	[266-274]	Ala ²⁶⁷ →Leu/Ile	535.2516	2	1.1
11	[266-299] ⁱⁱ⁾	Ser ²⁶⁹ →Val	1299.2634	3	-9.3
11	[266-302] ⁱ⁾	Ser ²⁶⁹ →Val	1046.2483	4	-1.1
12	[318-339] ⁱⁱ⁾	Ser ³³³ →Ala	643.8136	4	1.7
12	[324-339] ⁱⁱ⁾	Ser ³³³ →Ala	642.9651	3	0.7
12	[326-339] ⁱⁱ⁾	Ser ³³³ →Ala	556.5994	3	1.9
12	[327-339] ⁱⁱ⁾	Ser ³³³ →Ala	504.5646	3	-0.1
12	[329-339] ⁱⁱ⁾	Ser ³³³ →Ala	672.2985	2	0.1
12	[332-339] ⁱⁱ⁾	Ser ³³³ →Ala	481.7346	2	0.9
13	[331-353] ⁱⁱ⁾	Glu ³³¹ →His	860.4118	3	-2.3
14	[340-353] ⁱ⁾	Leu ³⁴⁶ →Met oxidated	758.8500	2	7.8

15	[359-382]	Asp ³⁷³ →Asn	542.8453	5	-7.2
16	[410-422]	Ala ⁴²⁰ →Arg	483.2612	3	-7.9
17	[410-425]	Ala ⁴²⁵ →Arg	582.6499	3	-5.4
18	[429-443]	Asp ⁴³² →Asn	883.4516	2	-6.8
19	[453-476] ⁱⁱ⁾	Asp ⁴⁵⁵ →Asn	671.8336	4	-8.8
20	[453-469]	Ala ⁴⁶⁵ →Gly	616.3083	3	1.2
21,22	[459-476] ⁱⁱ⁾	Ala ⁴⁵⁹ →Trp, Thr ⁴⁶⁰ →Pro	537.5308	4	2.1
23	[495-506]	Arg ⁵⁰⁶ →Lys	688.3949	2	-0.5
24	[603-615]	Ala ⁶⁰⁵ →Asn	483.5718	3	-1.5
24	[605-615]	Ala ⁶⁰⁵ →Asn	399.8722	3	0.8
25	[606-615]	Ile ⁶⁰⁶ →His	554.2694	2	-1.3
26	[657-669] ⁱⁱ⁾	Ser ⁶⁶¹ →Trp	806.3871	2	-1.7
26	[659-669] ⁱⁱ⁾	Ser ⁶⁶¹ →Trp	702.3370	2	-4.8
27	[683-702]	Val ⁶⁹⁶ →Glu	607.5411	4	2.5
27,28,29	[683-702]	Val ⁶⁹⁶ →Glu, His ⁷⁰⁰ →Asp, His ⁶⁸⁹ →Glu	1198.5563	2	0.6
27,28	[687-702]	Val ⁶⁹⁶ →Glu, His ⁷⁰⁰ →Asp	630.6349	3	-0.8
30	[772-779]	Gly ⁷⁷³ →Ala	558.7695	2	2.1
30	[772-779] ⁱ⁾	Gly ⁷⁷³ →Ala	566.7664	2	0.9
30	[773-779]	Gly ⁷⁷³ →Ala	465.7294	2	1.5
31,32	[780-787]	Glu ⁷⁸⁴ →Asp, Leu ⁷⁸⁶ →Ala	455.7317	2	4.9
33,34	[845-863]	Val ⁸⁴⁶ →Ala, Thr ⁸⁵⁰ →Asn	731.0200	3	2.1
33,34	[846-863]	Val ⁸⁴⁶ →Ala, Thr ⁸⁵⁰ →Asn	697.3355	3	-0.5
35	[918-933]	Ser ⁹³² →Pro	640.9664	3	-0.2
35	[918-934]	Ser ⁹³² →Pro	669.9777	3	0.8
35	[918-933]	Ser ⁹³² →Pro	640.9673	3	1.2
36	[955-972]	His ⁹⁵⁹ →Tyr	1053.9719	2	-0.5
36	[958-972]	His ⁹⁵⁹ →Tyr	594.2758	3	-1.2
36	[959-972]	His ⁹⁵⁹ →Tyr	532.2478	3	-4.3
36	[963-972]	His ⁹⁵⁹ →Tyr	529.7503	2	6.8
37	[973-992]	Asp ⁹⁷⁷ →Ala	564.0306	4	3.1
38	[1006-1014]	Val ¹⁰⁰⁸ →Leu/Ile	567.2858	2	1.6
39	[1019-1033]	Val ¹⁰²¹ →Leu/Ile	797.8972	2	-1.5
40	[1019-1045]	Arg ¹⁰⁴⁵ →His	779.6302	4	0.2
40	[1030-1045]	Arg ¹⁰⁴⁵ →His	500.0043	4	2.0
40	[1031-1045]	Arg ¹⁰⁴⁵ →His	637.3237	3	-1.5
40	[1034-1045]	Arg ¹⁰⁴⁵ →His	518.5854	3	-0.4
40	[1034-1050]	Arg ¹⁰⁴⁵ →His	532.5050	4	2.4
40	[1034-1046]	Arg ¹⁰⁴⁵ →His	547.5972	3	1.6
40	[1035-1045]	Arg ¹⁰⁴⁵ →His	695.8437	2	0.8

41	[1065-1078] ⁱⁱ⁾	Thr ¹⁰⁶⁸ →Ser	528.2567	3	5.8
42	[1074-1098] ⁱ⁾	Ala ¹⁰⁷⁴ →His	734.3569	4	-3.6
42	[1074-1108] ⁱ⁾	Ala ¹⁰⁷⁴ →His	696.3461	6	-4.9
43	[1079-1098]	Asp ¹⁰⁸⁶ →Asn	760.0382	3	-4.3
44	[1109-1136]	Glu ¹¹¹⁴ →Gln	847.9255	4	-6.7
45	[1109-1136]	Glu ¹¹¹⁶ →Gln	678.5436	5	-4.2
46	[1137-1149]	Glu ¹¹⁴⁹ →Gln	509.6014	3	2.2
47	[1141-1157]	Leu ¹¹⁴⁷ →Tyr, His ¹¹⁴⁹ insertion	1008.5466	2	7.8
48	[1194-1214]	Glu ¹¹⁹⁵ →Gln	636.8230	4	-7.0
49	[1196-1210]	Asn ¹²¹⁰ →Arg	440.7334	4	1.9
50	[1276-1283]	Leu ¹²⁷⁹ →Ser	454.7443	2	-2.5
51	[1323-1335]	Leu ¹³²³ →Arg	801.9478	2	-4.9
52	[1385-1399]	Trp ¹³⁹⁶ →Ser	552.9456	3	2.1
53	[1438-1461]	Asp ¹⁴⁴⁴ →Asn	704.0975	4	0.1
54	[1439-1461]	Thr ¹⁴⁴⁶ →Ala	656.0749	4	-0.2
54	[1442-1461]	Thr ¹⁴⁴⁶ →Ala	584.2875	4	-0.9
55,56	[1476-1489] ⁱⁱ⁾	Glu ¹⁴⁷⁹ → Asp, Ala ¹⁴⁸⁰ → Pro	854.8750	2	0.6
57	[1476-1489] ⁱⁱ⁾	Asp ¹⁴⁸¹ → Asn	565.9194	3	-8.3
58	[1475-1489] ⁱⁱ⁾	Val ¹⁴⁸⁴ → Leu/Ile	613.6226	3	0.4
58	[1476-1489] ⁱⁱ⁾	Val ¹⁴⁸⁴ → Leu/Ile	570.9248	3	1.4
59	[1476-1489] ⁱⁱ⁾	Arg ¹⁴⁸⁸ → Leu/Ile	827.3665	2	0.6
59	[1475-1489] ⁱⁱ⁾	Arg ¹⁴⁸⁸ → Leu/Ile	594.6121	3	1.2
60	[1497-1507]	Pro ¹⁴⁹⁷ →Thr, Glu ¹⁴⁹⁸ →His	623.8043	2	-6.6
61	[1582-1605]	Ile ¹⁵⁸⁹ →Met	691.5926	4	7.8
62	[1584-1605]	Met ¹⁵⁹⁶ →Thr	622.3151	4	1.4
63,64,65	[1584-1605]	Ser ¹⁵⁸⁶ →Glu,Ile ¹⁵⁸⁹ →Val, His ¹⁵⁹⁹ →Leu	840.7545	3	0.5
63,64,65	[1584-1602]	Ser ¹⁵⁸⁶ →Glu,Ile ¹⁵⁸⁹ →Val, His ¹⁵⁹⁹ →Leu	1058.5076	2	1.1
63,64,65	[1584-1605] ⁱ⁾	Ser ¹⁵⁸⁶ →Glu,Ile ¹⁵⁸⁹ →Val, His ¹⁵⁹⁹ →Leu	846.0864	3	0.7
66	[1584-1605]	His ¹⁵⁹⁹ →Asp	832.0718	3	2.0
67	[1590-1605]	Phe ¹⁵⁹⁸ →Val	631.3157	3	-2.3
68	[1606-1614]	Ser ¹⁶¹⁰ →Glu	358.5264	3	1.9
69	[1606-1614]	His ¹⁶¹³ →Asn	504.7718	2	0.3
70	[1614-1627]	Lys ¹⁶¹⁴ →Arg	508.6140	3	-3.6
71	[1642-1655]	Arg ¹⁶⁵⁵ →His	782.9209	2	-1.5
72	[1672-1681]	Thr ¹⁶⁷⁷ →Asp	560.7916	2	2.4
73	[1690-1707]	Ala ¹⁶⁹³ →Asp	1012.9195	2	1.3
73	[1690-1707] ⁱ⁾	Ala ¹⁶⁹³ →Asp	1020.9155	2	-0.1
74	[1764-1784]	Asp ¹⁷⁷⁰ →Asn	610.5617	4	-2.1
75	[1764-1784]	His ¹⁷⁷⁹ →Asp	806.7402	3	9.9

76	[1792-1816]	Ala ¹⁷⁹³ →Asp	713.1020	4	9.6
77	[1793-1816]	Glu ¹⁷⁹⁶ →Gln	892.7620	3	-6.9
78	[1793-1816]	Glu ¹⁷⁹⁸ →Gln	669.8214	4	-9.9
79	[1793-1816]	Glu ¹⁸⁰⁰ →Gln	892.7606	3	-8.5
80	[1913-1928]	Lys ¹⁹¹⁶ →Asn	905.4488	2	5.4
81	[1936-1967]	Thr ¹⁹³⁹ →Leu/Ile	965.1912	4	-9.3
82	[1930-1944]	Asp ¹⁹⁴¹ →Asn	475.4724	4	0.4
83	[1936-1944]	Asp ¹⁹⁴¹ →Asn	393.5124	3	0.2
84	[1929-1967]	Glu ¹⁹⁵³ →Gln	787.7031	6	-4.6
85	[1929-1967]	Val ¹⁹⁵⁵ →Thr	1181.8087	4	9.7
86	[2004-2022]	Ala ²⁰⁰⁴ →Thr	721.9979	3	-0.6
87	[2037-2046]	Leu ²⁰³⁷ →Gln	429.5534	3	6.4
88	[2051-2073]	Leu ²⁰⁶⁹ →Phe	1208.6168	2	5.6
88	[2052-2073]	Leu ²⁰⁶⁹ →Phe	1151.0962	2	-0.4
88	[2057-2070]	Leu ²⁰⁶⁹ →Phe	773.4067	2	1.0
88	[2059-2070]	Leu ²⁰⁶⁹ →Phe	659.3520	2	2.4
88	[2061-2073]	Leu ²⁰⁶⁹ →Phe	715.4009	2	0.7
88	[2063-2073]	Leu ²⁰⁶⁹ →Phe	602.3154	2	-1.7
88	[2066-2073]	Leu ²⁰⁶⁹ →Phe	454.7404	2	1.2
88	[2067-2073]	Leu ²⁰⁶⁹ →Phe	411.2245	2	1.6
89	[2120-2129]	Ser ²¹²⁹ →Gln	520.7793	2	-1.0
90	[2154-2179]	Ala ²¹⁶² →Asn	1018.5049	3	3.3
91	[2213-2243]	Leu ²²⁴³ →Gln	888.4324	4	7.3
92,93	[2213-2248]	Leu ²²⁴³ →Ala, Glu ²²⁴⁷ →Leu	1008.7646	4	7.0
94,95, 96	[2236-2249]	Leu ²²³⁹ →Tyr, Thr ²²⁴² →Ala, Thr ²²⁴⁹ →Arg	809.9427	2	-1.1
97	[2252-2272]	Cys ²²⁵² →Ser	806.0796	3	8.2
98	[2250-2272] ⁱⁱ⁾	Val ²²⁶¹ →Met, Ala ²²⁶⁴ →Thr	704.0779	4	3.1
99	[2280-2298]	Tyr ²²⁸² →Phe	574.0240	4	-3.6
100	[2276-2298]	Ala ²²⁸⁸ →Leu/Ile	678.0875	4	6.9
101	[2343-2381] ⁱ⁾	Val ²³⁴⁶ →Pro	929.6542	5	2.9
102	[2386-2397]	His ²³⁹³ →Asp	632.8556	2	-0.7
103	[2426-2436]	His ²⁴²⁸ →Asp	492.2195	3	1.3
104	[2445-2463]	His ²⁴⁴⁹ →Asp	1078.0146	2	-0.4
105	[2480-2508]	Ile ²⁴⁸⁴ →Asn, Glu ²⁴⁹⁰ →Gln	737.1091	4	8.9
106	[2634-2650] ⁱ⁾	His ²⁶⁴² →Gln	1049.4475	2	1.0
107	[2719-2735] ⁱⁱ⁾	Gln ²⁷²³ →Pro	513.7607	4	4.5
108	[2721-2735] ⁱⁱ⁾	Gln ²⁷²³ →Pro	578.2885	3	2.0
108	[2723-2735] ⁱⁱ⁾	Gln ²⁷²³ →Pro	510.2554	3	-3.9
108	[2723-2735] ⁱⁱ⁾	Gln ²⁷²³ →Pro	764.8828	2	0.5

109	[2736-2754]	Lys ²⁷³⁹ →Arg	727.3795	3	-3.7
110	[2757-2765]	Arg ²⁷⁶⁵ →Lys	537.2428	2	-0.3
111,112	[2855-2869]	His ²⁸⁵⁹ →Phe, Glu ²⁸⁶⁷ →Ala	630.6305	3	-6.4
111,112	[2855-2869] ⁱ⁾	His ²⁸⁵⁹ →Phe, Glu ²⁸⁶⁷ →Ala	641.2984	3	0.9
111,112	[2855-2869]	His ²⁸⁵⁹ →Phe, Glu ²⁸⁶⁷ →Ala	635.9661	3	-0.2
111,112	[2857-2869]	His ²⁸⁵⁹ →Phe, Glu ²⁸⁶⁷ →Ala	549.5999	3	0.2
112	[2861-2869]	Glu ²⁸⁶⁷ →Ala	547.2700	2	1.3
112	[2862-2869]	Glu ²⁸⁶⁷ →Ala	489.7565	2	1.4
113,114	[2870-2884]	Ile ²⁸⁷⁰ →Val, Arg ²⁸⁸² →Ser	759.3889	2	0.8
114	[2875-2884]	Arg ²⁸⁸² →Ala	523.7724	2	1.0
115, 116	[2885-2895]	Gly ²⁸⁹² →Gln, Gly ²⁸⁹³ →Ala	424.5294	3	-2.1
117	[2916-2934]	Gly ²⁹¹⁸ →Asp	713.0278	3	1.4
117	[2916-2929]	Gly ²⁹¹⁸ →Asp	774.3738	2	-2.1
118, 119	[2989-3006]	Lys ²⁹⁹¹ →Arg, Ile ²⁹⁹⁹ →Val	718.0257	3	-4.5
119	[2992-3006]	Ile ²⁹⁹⁹ →Val	604.6218	3	0.5
119	[2992-3006]	Ile ²⁹⁹⁹ →Val	906.4286	2	0.1
119	[2992-3006]	Ile ²⁹⁹⁹ →Val	609.9546	3	-7.8
119	[2993-3006]	Ile ²⁹⁹⁹ →Val	561.6071	3	-0.4
119	[2994-3006]	Ile ²⁹⁹⁹ →Val	773.3787	2	1.0
119	[2996-3006]	Ile ²⁹⁹⁹ →Val	467.9039	3	1.6
120	[2992-3006]	Ile ²⁹⁹⁹ →Val	467.9734	4	-5.8
121, 122	[3007-3026]	Ile ³⁰⁰⁹ →Thr, Ser ³⁰²¹ →Asn	741.3451	3	-0.6
121, 122	[3007-3026]	Ile ³⁰⁰⁹ →Thr, Ser ³⁰²¹ →Asn	1111.5149	2	0.2
121, 122	[3007-3026] ⁱ⁾	Ile ³⁰⁰⁹ →Thr, Ser ³⁰²¹ →Asn	1119.5160	2	3.4
121, 122	[3009-3026]	Ile ³⁰⁰⁹ →Thr, Ser ³⁰²¹ →Asn	668.6551	3	0.5
122	[3010-3026]	Ser ³⁰²¹ →Asn	951.9561	2	1.5
122	[3014-3026]	Ser ³⁰²¹ →Asn	775.8482	2	-1.2
122	[3015-3026]	Ser ³⁰²¹ →Asn	702.3167	2	2.4
122	[3016-3026]	Ser ³⁰²¹ →Asn	628.7814	2	1.1
122	[3017-3026]	Ser ³⁰²¹ →Asn	593.2618	2	-0.5
122	[3019-3026]	Ser ³⁰²¹ →Asn	492.2336	2	1.8
123, 124	[3094-3116] ⁱ⁾	Ser ³⁰⁹⁷ →Thr, Ala ³⁰⁹⁹ →Glu	878.0879	3	2.7
123, 124	[3097-3116] ⁱ⁾	Ser ³⁰⁹⁷ →Thr, Ala ³⁰⁹⁹ →Glu	776.3851	3	1.3
125	[3101-3116]	Tyr ³¹⁰³ →Phe	902.9581	2	2.2
125	[3101-3116]	Tyr ³¹⁰³ →Phe	602.3057	3	-1.4
126, 127	[3188-3204]	Ser ³¹⁹³ →Asn, Asn ³²⁰¹ →Thr	501.7804	4	0.7
126	[3188-3200]	Ser ³¹⁹³ →Asp	508.2812	3	4.4
126	[3188-3200]	Ser ³¹⁹³ →Asn	761.4232	2	0.5
126	[3189-3200]	Ser ³¹⁹³ →Asn	478.9401	3	-0.5

126	[3190-3200]	Ser ³¹⁹³ →Asn	661.3636	2	-1.7
126	[3191-3200]	Ser ³¹⁹³ →Asn	597.3340	2	-2.5
126	[3192-3200]	Ser ³¹⁹³ →Asn	355.5460	3	2.2
126	[3192-3200]	Ser ³¹⁹³ →Asn	532.8128	2	-2.5
127, 128, 129, 130	[3222-3244]	Asn ³²³³ →Tyr, Ser ³²³⁶ →Thr, Val ³²³⁸ →Ala, Pro ³²⁴² →Ala	1255.1392	2	-0.4
131	[3267-3280]	Val ³²⁷⁹ →L/I	901.9203	2	3.0
132	[3267-3280] ⁱ⁾	Val ³²⁷⁹ →L/I	606.9470	3	1.9
133	[3356-3374]	Asn ³³⁶³ →His	566.7847	4	1.5
134	[3358-3374]	Asn ³³⁶³ →His	517.2491	4	-0.9
135	[3386-3402] ⁱⁱ⁾	Thr ³³⁸⁹ →Ala	659.9877	3	1.9

1) Numbers inside brackets correspond to the position of the proteolytic peptide of the sequence of KLH2.

i) Methionine sulfoxide

ii) Carbamidomethylation of Cys

Table S12. Post-translational modifications found in KLH proteins using the Peaks software [5].

#	Protein	Residue	Assignment ¹⁾	m/z exp.	m/z theor.	Z	Error (ppm)
Oxidation to nitro							
1	KLH1	Tyr ⁴⁶¹	[451-467]	627.2889	627.2901	3	-1.9
2		Tyr ⁹⁴⁶	[946-955]	598.2608	598.2589	2	3.2
3		Tyr ¹³⁶¹	[1349-1365]	519.2578	519.2549	4	5.6
4		Tyr ¹⁴⁴⁴	[1437-1460]	716.5759	716.5744	4	2.1
5		Tyr ¹⁷³⁷	[1736-1746]	707.8544	707.8511	2	4.7
6		Tyr ¹⁹⁴⁰	[1934-1942]	594.2476	594.2458	2	3.0
7		Tyr ¹⁹⁴⁵	[1943-1958]	991.9680	991.9668	2	1.2
8		Tyr ²⁶⁴⁷	[2641-2648]	519.2214	519.2196	2	3.5
9		Tyr ²⁷⁶⁸	[2764-2772]	631.2837	631.2831	2	1.0
1	KLH2	Tyr ⁷⁴	[73-80]	532.7890	532.7900	2	-1.9
2		Tyr ⁴³⁹	[429-443]	604.6310	604.6301	3	1.5
3		Tyr ¹²⁸⁷	[1284-1292]	541.2609	541.2613	2	-0.7
4		Tyr ¹³²⁴	[1323-1335]	802.9360	802.9349	2	1.4
5		Tyr ¹³⁶²	[1350-1366]	964.9530	964.9519	2	1.1
6		Tyr ¹⁴⁷⁷	[1476-1489]	581.2477	581.2468	3	1.5
7		Tyr ¹⁶⁰⁴	[1584-1605]	641.0587	641.0582	4	0.8
8		Tyr ¹⁷³⁹	[1738-1748]	694.3439	694.3400	2	5.6
9		Tyr ¹⁹⁴²	[1930-1944]	486.9653	486.9642	4	2.3
10		Tyr ²⁰³⁹	[2037-2046]	439.5552	439.5540	3	2.7
11		Tyr ²¹⁹³	[2184-2208]	1000.4786	1000.4775	3	1.1
12		Tyr ²⁴³¹	[2426-2436]	514.5580	514.5577	3	0.6
13		Tyr ²⁴³⁷	[2437-2444]	498.7544	498.7531	2	2.6
14		Tyr ³³⁷⁹	[3375-3385]	422.8631	422.8631	3	0.0
Hydroxylation (Tyr)							
1	KLH1	Tyr ⁹⁴⁶	[946-955]	583.7651	583.7638	2	2.2
2		Tyr ¹⁰⁴³	[1035-1051]	529.0002	528.9996	4	1.1
3		Tyr ¹¹⁰⁸	[1098-1109]	499.9062	499.9049	3	2.6
4		Tyr ¹¹⁹³	[1193-1198]	392.6902	392.6896	2	1.5
5		Tyr ¹³⁶¹	[1349-1365]	512.0078	512.0074	4	0.8
6		Tyr ¹⁷³⁷	[1736-1746]	693.3585	693.3560	2	3.6
7		Tyr ²¹⁵³	[2152-2162]	699.3745	699.3742	2	0.4
8		Tyr ²¹⁹²	[2190-2196]	523.7327	523.7323	2	0.8
9		Tyr ²²⁰⁸	[2207-2218]	470.8937	470.8929	3	1.7
10		Tyr ²²¹⁵	[2207-2218]	470.8937	470.8929	3	1.7
11		Tyr ²³⁵¹	[2351-2357]	465.2405	465.2398	2	1.5
12		Tyr ²⁴³⁵	[2435-2461]	798.8958	798.8904	4	6.8
13		Tyr ²⁷⁶⁸	[2764-2772]	616.7875	616.7880	2	-0.8
14		Tyr ³²²⁹	[3221-3240]	1002.956	1002.9542	2	1.8

1	KLH2	Tyr ⁷⁴	[73-80]	518.2947	518.2949	2	-0.4
2		Tyr ⁴³⁹	[429-442]	827.9004	827.8987	2	2.1
2		Tyr ⁴³⁹	[429-443]	891.9482	891.9461	2	2.4
3		Tyr ⁴⁵⁷	[453-469]	626.3145	626.3107	3	6.1
4		Tyr ⁶³¹	[616-639] ⁱⁱ⁾	733.8403	733.8389	4	1.9
5		Tyr ⁹⁸⁷	[976-992]	648.298	648.2983	3	-0.5
6		Tyr ¹¹¹⁵	[1109-1136]	681.9437	681.9417	5	2.9
7		Tyr ¹¹⁵⁹	[1158-1162] ⁱ⁾	366.6730	366.6728	2	0.5
8		Tyr ¹¹⁹⁴	[1194-1214]	854.4316	854.4264	3	6.1
9		Tyr ¹²⁸⁷	[1284-1292]	526.7675	526.7662	2	2.5
10		Tyr ¹⁴⁷⁷	[1475-1489] ^{i,iii)}	619.6128	619.6133	3	-0.8
11		Tyr ¹⁷⁷⁷	[1764-1784]	614.8095	614.8073	4	3.6
12		Tyr ¹⁹⁴²	[1929-1944]	518.7431	518.7419	4	2.3
13		Tyr ²⁰²⁶	[2026-2030]	328.1690	328.1684	2	1.8
13		Tyr ²⁰²⁶	[2026-2033] ⁱⁱⁱ⁾	539.2642	539.2646	2	-0.7
14		Tyr ²⁰³⁹	[2039-2046]	509.7401	509.7397	2	0.8
14		Tyr ²⁰³⁹	[2037-2046]	429.8908	429.8906	3	0.5
15		Tyr ²⁴³⁷	[2437-2444]	484.2587	484.2580	2	1.4
16		Tyr ²⁶⁷⁰	[2692-2708]	532.9986	532.9983	4	0.6
17	Tyr ³²⁵⁶	[3256-3264]	548.7690	548.7667	2	4.2	
18	Tyr ³³⁷¹	[3367-3374]	330.8383	330.8383	3	0.0	
19	Tyr ³³⁷⁸	[3375-3385]	413.1996	413.1997	3	-0.2	
Hydroxylation (Trp)							
1	KLH1	Trp ⁷⁷¹	[771-778]	535.7764	535.7742	2	4.1
2		Trp ⁹⁵⁹	[956-969]	566.2628	566.2617	3	1.9
3		Trp ¹⁷⁵⁹	[1748-1763]	628.3254	628.3244	3	1.6
4		Trp ²⁰¹⁶	[2013-2023]	491.2399	491.2392	3	1.4
4		Trp ²⁰¹⁶	[2013-2023] ⁱⁱ⁾	496.5715	496.5709	3	1.2
5		Trp ²¹⁹⁵	[2190-2198]	430.2043	430.2033	3	2.3
5		Trp ²¹⁹⁵	[2190-2196]	523.7335	523.7323	2	2.3
6		Trp ²⁴²⁷	[2424-2434]	482.2403	482.2394	3	1.9
6		Trp ²⁴²⁷	[2424-2434] ⁱⁱ⁾	487.5719	487.5710	3	1.8
6		Trp ²⁴²⁷	[2424-2431]	528.7314	528.7318	2	-0.8
7		Trp ²⁷¹¹	[2707-2718]	796.9163	796.9139	2	3.0
1	KLH2	Trp ⁶⁴¹	[640-650]	707.4044	707.4030	2	2.0
2		Trp ⁹²⁷	[918-931]	570.2665	570.2668	3	-0.5
3		Trp ¹³⁹⁶	[1390-1399]	629.291	629.2905	2	0.8
4		Trp ¹⁷⁵⁹	[1749-1763]	614.9935	614.9927	3	1.3
4		Trp ¹⁷⁵⁹	[1749-1763]	614.9935	614.9927	3	1.3
5		Trp ¹⁷⁶¹	[1749-1763]	614.9951	614.9927	3	3.9
6		Trp ¹⁸⁸⁷	[1886-1897]	522.945	522.9438	3	2.3
7		Trp ¹⁸⁹⁰	[1886-1897]	522.9448	522.9438	3	1.9
8	Trp ²⁰¹⁸	[2016-2022]	498.2248	498.2236	2	2.4	

9		Trp ²¹⁷⁵	[2161-2179]	721.3622	721.3622	3	0.0
10		Trp ²³⁰⁰	[2299-2308]	629.8516	629.8502	2	2.2
11		Trp ²³⁰³	[2299-2308]	629.8564	629.8502	2	9.8
12		Trp ²⁴⁰⁴	[2403-2407] ⁱ⁾	385.1884	385.1886	2	-0.5
12		Trp ²⁴⁰⁴	[2403-2407] ⁱ⁾	385.1882	385.1886	2	-1.0
13		Trp ²⁴²⁹	[2426-2436]	378.9226	378.9227	4	-0.3
13		Trp ²⁴²⁹	[2426-2436] ⁱⁱ⁾	382.9218	382.9214	4	1.0
14		Trp ³⁰⁸⁶	[3078-3091]	525.9484	525.9471	3	2.5
Hydroxylation (His)							
1	KLH1	His ⁷⁸³	[779-786]	494.7782	494.7764	2	3.6
2		His ¹³⁷⁶	[1366-1378]	412.6909	412.6907	4	0.5
3		His ²⁴³⁹	[2435-2461]	798.8916	798.8904	4	1.5
1	KLH2	His ¹²⁰¹	[1194-1214]	854.4259	854.4264	3	-0.6
2		His ¹²⁰⁶	[1194-1214]	854.4294	854.4264	3	3.5
3		His ¹⁹⁰⁵	[1903-1912] ⁱ⁾	616.7755	616.7731	2	3.9
4		His ²⁷²⁸	[2719-2735] ⁱ⁾	700.3460	700.3419	3	5.9
Di-hydroxylation (Trp)							
1	KLH1	Trp ³⁵²	[352-368]	562.5280	562.5268	4	2.1
1		Trp ³⁵²	[352-368]	562.5289	562.5268	4	3.7
2		Trp ⁶⁴³	[639-649] ⁱⁱ⁾	725.3702	725.3684	2	2.5
3		Trp ⁸⁸¹	[881-889] ⁱ⁾	547.2358	547.2343	2	2.7
4		Trp ⁹⁵⁹	[956-969]	571.5930	571.5933	3	-0.5
5		Trp ¹⁷⁵⁷	[1748-1763]	633.6568	633.6560	3	1.3
6		Trp ¹⁷⁵⁹	[1748-1763]	633.6569	633.6560	3	1.4
7		Trp ¹⁷⁸⁸	[1884-1893]	666.8570	666.8560	2	1.5
8		Trp ¹⁸⁸⁵	[1884-1893]	666.8566	666.8560	2	0.9
9		Trp ¹⁹⁵⁵	[1943-1958]	985.4708	985.4691	2	1.7
10		Trp ²⁰¹⁶	[2013-2023]	496.5715	496.5709	3	1.2
10		Trp ²⁰¹⁶	[2013-2023] ⁱⁱ⁾	501.9026	501.9025	3	0.2
11		Trp ²¹⁹⁵	[2190-2196]	531.7299	531.7298	2	0.2
12		Trp ²³⁰¹	[2297-2306]	631.3130	631.3138	2	-1.3
13		Trp ²⁴²⁷	[2424-2434]	730.8538	730.8526	2	1.6
13	Trp ²⁴²⁷	[2424-2434] ⁱⁱ⁾	738.8523	738.8500	2	3.1	
14	Trp ²⁷¹¹	[2707-2718]	804.9124	804.9113	2	1.4	
15	Trp ²⁹⁹⁹	[2988-3021]	979.2117	979.2106	4	1.1	
1	KLH2	Trp ⁹⁶	[87-99]	800.8608	800.8612	2	-0.5
2		Trp ³⁵⁴	[354-358]	380.6744	380.6743	2	0.3
3		Trp ⁶⁴⁴	[640-650]	715.4017	715.4005	2	1.7
3		Trp ⁶⁴⁴	[640-650]	715.4017	715.4005	2	1.7
4		Trp ⁸⁸⁰	[880-888] ⁱ⁾	568.2405	568.2395	2	1.8
5		Trp ⁹²⁷	[918-931]	575.6006	575.5984	3	3.8
6		Trp ¹³⁴⁶	[1336-1349]	820.8842	820.8824	2	2.2
7	Trp ¹³⁹⁶	[1390-1399]	637.2866	637.2880	2	-2.2	

8		Trp ¹⁷⁵⁹	[1749-1763]	465.4952	465.4952	4	0.0
9		Trp ¹⁸⁵⁵	[1847-1860]	512.9356	512.9352	3	0.8
10		Trp ¹⁸⁹⁰	[1886-1897]	791.9107	791.9092	2	1.9
11		Trp ²¹⁷⁵	[2161-2179] ⁱⁱ⁾	732.0275	732.0255	3	2.7
12		Trp ²³⁰⁰	[2299-2308]	637.8479	637.8477	2	0.3
13		Trp ²³⁰³	[2299-2308]	637.8482	637.8477	2	0.8
14		Trp ²⁴⁰⁴	[2403-2407] ⁱ⁾	393.1863	393.1861	2	0.5
15		Trp ²⁴²⁹	[2426-2436]	510.2263	510.2259	3	0.8
15		Trp ²⁴²⁹	[2427-2433]	337.4764	337.4766	3	-0.6
16		Trp ²⁸³⁹	[2835-2843]	613.7801	613.7787	2	2.3
Di-hydroxylation (Tyr)							
1	KLH1	Tyr ⁹⁴⁶	[946-955]	591.7629	591.7612	2	2.9
2		Tyr ²³⁵¹	[2351-2357]	473.2379	473.2372	2	1.5
1	KLH2	Tyr ⁷⁴	[73-80]	526.2939	526.2923	2	3.0
2		Tyr ²⁰²⁶	[2026-2033] ⁱⁱⁱ⁾	547.2623	547.2620	2	0.5
3		Tyr ²⁴³⁷	[2437-2444]	492.2556	492.2555	2	0.2
Di-hydroxylation (Phe)							
1	KLH1	Phe ³⁰⁴	[303-308]	365.6794	365.6795	2	-0.3
2		Phe ²⁶⁴⁶	[2641-2648]	512.7219	512.7220	2	-0.2
Quinone							
1	KLH1	Tyr ⁷⁴	[73-80]	525.2853	525.2846	2	1.3
1	KLH2	Trp ¹³⁹⁶	[1390-1399]	636.2809	636.2811	2	-0.3
2		Trp ¹⁸⁵⁵	[1847-1860]	384.4499	384.45	4	-0.3
Trp oxidation to kynurenine							
1	KLH1	Trp ³⁵²	[352-368]	555.5302	555.5281	4	3.8
2		Trp ⁹³⁰	[919-945]	792.1234	792.1223	4	1.4
3		Trp ⁹⁵⁹	[956-969]	421.9475	421.9482	4	-1.7
4		Trp ¹⁷⁵⁹	[1748-1763]	624.3244	624.3244	3	0.0
5		Trp ¹⁹⁵⁵	[1943-1958]	971.4746	971.4717	2	3.0
6		Trp ²⁰¹⁶	[2013-2023]	487.2401	487.2392	3	1.8
7		Trp ²¹⁹⁵	[2190-2198]	426.2041	426.2033	3	1.9
7		Trp ²¹⁹⁵	[2190-2196]	517.7333	517.7323	2	1.9
1	KLH2	Trp ⁶⁴⁴	[640-650]	701.4030	701.4030	2	0.0
2		Trp ¹⁷⁵⁹	[1749-1763]	458.4977	458.4965	4	2.6
3		Trp ¹⁷⁶¹	[1749-1763]	610.9937	610.9927	3	1.6
4		Trp ¹⁸⁵⁵	[1847-1860]	377.9547	377.9546	4	0.3
5		Trp ¹⁸⁸⁷	[1886-1897]	518.9461	518.9438	3	4.4
6		Trp ¹⁸⁹⁰	[1886-1897]	777.9177	777.9117	2	7.7
7		Trp ²¹⁹⁷	[2184-2198]	634.9679	634.9680	3	-0.2
8		Trp ²³⁰⁰	[2299-2308]	623.8522	623.8502	2	3.2
9		Trp ²³⁰³	[2299-2308]	623.8522	623.8502	2	3.2
Trp oxidation to oxolactone							
1	KLH1	Trp ⁹⁵⁹	[956-969]	565.5909	565.5898	3	1.9

2		Trp ²⁷¹¹	[2707-2718]	795.9076	795.9061	2	1.9
1	KLH2	Trp ³⁰⁸⁶	[3078-3091]	525.2755	525.2753	3	0.4
Formylation							
1	KLH1	Lys ²³²	[230-244] ⁱ⁾	604.6194	604.6183	3	1.8
2		Lys ³⁷²	[369-380]	469.8974	469.8966	3	1.7
3		Lys ¹¹³²	[1132-1137]	365.2291	365.2287	2	1.1
4		Lys ¹¹⁹⁸	[1193-1203]	468.2458	468.2450	3	1.7
5		Lys ¹³³³	[1322-1334]	758.4608	758.4599	2	1.2
6		Lys ¹⁶⁰⁹	[1605-1612]	465.2879	465.2866	2	2.8
7		Lys ²⁰³²	[2024-2036]	820.4003	820.4026	2	-2.8
8		Lys ²¹⁹⁸	[2197-2206]	601.3127	601.3105	2	3.7
9		Lys ²³⁴¹	[2341-2350]	591.8182	591.8165	2	2.9
10		Lys ²³⁵⁰	[2342-2357]	650.3360	650.3349	3	1.7
11		Lys ³¹⁷³	[3170-3192]	722.6067	722.6051	4	2.2
12		Lys ³²⁷⁵	[3270-3276]	327.8551	327.8554	3	-0.9
1	KLH2	Lys ⁵²	[23-55] ⁱ⁾	933.6757	933.6721	4	3.9
2		Lys ¹³⁵	[120-136]	527.5165	527.5141	4	4.5
3		Lys ¹³⁶	[136-148]	398.9657	398.9651	4	1.5
4		Lys ⁴¹⁴	[410-428]	518.7919	518.7901	4	3.5
5		Lys ¹²⁹²	[1284-1296]	511.9292	511.9278	3	2.7
6		Lys ¹⁴⁷⁵	[1475-1489] ⁱ⁾	618.2834	618.2817	3	2.7
7		Lys ¹⁵³⁷	[1523-1545]	966.8157	966.8095	3	6.4
8		Lys ¹⁶¹⁴	[1606-1627]	803.4274	803.4243	3	3.9
9		Lys ¹⁶⁸¹	[1672-1685]	563.3062	563.3049	3	2.3
10		Lys ¹⁷⁸⁴	[1764-1792]	853.9375	853.9349	4	3.0
11		Lys ¹⁹¹²	[1903-1916] ⁱ⁾	849.4074	849.4021	2	6.2
12		Lys ²⁰³⁰	[2026-2033]	505.2821	505.2814	2	1.4
13		Lys ²¹⁰²	[2097-2106]	560.2957	560.2946	2	2.0
14		Lys ²²¹²	[2209-2220]	715.3517	715.3509	2	1.1
15		Lys ²³⁰⁸	[2299-2309]	699.9006	699.8976	2	4.3
16		Lys ²⁴⁴⁴	[2437-2463]	1046.5217	1046.5192	3	2.4
17		Lys ²⁵⁰⁸	[2508-2522]	852.4446	852.4432	2	1.6
18		Lys ³³²¹	[3318-3324]	461.3100	461.3098	2	0.4
Phosphorylation (Thr)							
1	KLH2	Thr ²⁰³¹	[2026-2033]	531.2670	531.2671	2	-0.2
2		Thr ¹⁵⁸⁸	[1584-1605]	866.0659	866.0688	3	-3.3
3		Thr ²⁰³¹	[2026-2033] ^{iv)}	547.2623	547.2620	2	0.5
3		Thr ²⁰³¹	[2026-2033] ^{v)}	539.2642	539.2646	2	-0.7
3		Thr ²⁰³¹	[2026-2033] ^{vi)}	522.2645	522.2628	2	3.3
3		Thr ²⁰³¹	[2026-2033]	531.2677	531.2671	2	1.1
N-glycosylation							
1	KLH1	Asn ³⁸⁷	[381-407] ^{vii)}	733.1655	733.1652	5	0.4
1		Asn ³⁸⁷	[381-407] ^{viii)}	952.7232	952.7206	4	2.7

1		Asn ³⁸⁷	[381-407] ^{ix)}	820.9286	820.9244	4	5.1
1		Asn ³⁸⁷	[381-407] ^{x)}	784.4142	784.4086	4	7.1
1		Asn ³⁸⁷	[383-406] ^{viii)}	870.6787	870.6770	4	2.0
1		Asn ³⁸⁷	[383-406] ^{x)}	936.1598	936.1508	3	9.6
1	KLH2	Asn ³⁸⁹	[383-399] ^{vii)}	1276.6051	1276.6012	2	3.1
1		Asn ³⁸⁹	[383-399] ^{viii)}	1349.6321	1349.6324	2	-0.2
1		Asn ³⁸⁹	[383-399] ^{viii)}	900.0952	900.0909	3	4.8
1		Asn ³⁸⁹	[383-399] ^{xi)}	905.4226	905.4225	3	0.1
1		Asn ³⁸⁹	[383-409] ^{x)}	788.9299	788.9221	4	9.9
1		Asn ³⁸⁹	[385-399] ^{viii)}	828.0652	828.0660	3	-1.0

¹⁾ Numbers inside brackets correspond to the position of the proteolytic peptide of the sequence of KLH1.

- ⁱ⁾ Carbamidomethylation of Cys
- ⁱⁱ⁾ Methionine sulfoxide
- ⁱⁱⁱ⁾ Phosphorylation of Thr
- ^{iv)} Dihydroxylation of Tyr
- ^{v)} Hydroxylation of Tyr
- ^{vi)} Dehydration of Asp
- ^{vii)} N-glycosylation (Hex₂HexNAc₂)
- ^{viii)} N-glycosylation (Hex₂HexNAc₂dHex₁)
- ^{ix)} N-glycosylation (HexNAc₁dHex₁)
- ^{x)} N-glycosylation (HexNAcylation)
- ^{xi)} N-glycosylation (Hex₃HexNAc₂)

Table S13. Summary of diagnostic ions corresponding to PTMs found in KLH proteins using the Peaks software [5].

PTM	Δm	Immonium ions or other diagnostic ions (Th)	Reference
Tyr nitration	(+44.99)Da	181.06	[6]
Tyr hydroxylation	(+15.99)Da	152.07	[7]
Tyr di-hydroxylation	(+31.99)Da	168.07	
Trp hydroxylation	(+15.99)Da	175.09, 146.09	[7]
Trp di-hydroxylation	(+31.99)Da	191.08	
Kynurenine	(+3.99)Da	163.09	[8]
Lys formylation	(+27.99)Da	112.08	[9]
Phosphorylation	(+79.97)Da	Neutral loss -97.98 Da	[10]
N-glycosylation	(+203.08)Da (+349.14)Da (+730.26)Da (+876.32)Da (+892.32)Da	N-acetylhexosamines (HexNAc, 204.08) HexNAc – H ₂ O (186.08) HexNAc – 2H ₂ O (168.07) HexNAc – C ₂ H ₄ O ₂ (144.07) HexNAc – CH ₆ O ₃ (138.06) HexNAc – C ₂ H ₆ O ₃ (126.06) Hexosa (Hex. 163.06) HexNAc + Hex (366.14)	[11-13]

Table S14. Identification of the thioether bond in KLH1 and KLH2 using Kojak [2].

	Protein	Assignment ¹⁾	Thioether bond ²⁾	m/z exp.	m/z theor.	z	Error (ppm)
1	KLH1	[477-492]	C ⁴⁸⁰ -H ⁴⁸²	502.9702	502.9780	4	-15.5
2		[891-906] ⁱ⁾	C ⁸⁹⁴ -H ⁸⁹⁶	496.9611	496.9689	4	-15.7
3		[1292-1321] ⁱⁱ⁾	C ¹³⁰⁹ -H ¹³¹¹	574.5854	574.5939	6	-14.8
4		[1720-1735] ⁱ⁾	C ¹⁷²³ -H ¹⁷²⁵	491.2071	491.2149	4	-15.9
1	KLH2	[56-72]	C ⁶⁰ -H ⁶²	674.9725	674.9803	3	-11.6
		[56-72] ⁱ⁾		510.4781	510.4859	4	-15.3
2		[477-494]	C ⁴⁸² -H ⁴⁸⁴	742.6539	742.6617	3	-10.5
				445.9923	446.0002	5	-17.7
3		[890-905] ⁱ⁾	C ⁸⁹³ -H ⁸⁹⁵	492.9623	492.9702	4	-16.0
4		[1297-1322]	C ¹³¹⁰ -H ¹³¹²	758.8250	758.8328	4	-10.3
5		[1722-1737] ⁱ⁾	C ¹⁷²⁵ -H ¹⁷²⁷	492.2086	492.2165	4	-16.0
6	[2107-2153]	C ²¹⁴¹ -H ²¹⁴³	749.0424	749.0503	7	-10.5	
7	[2557-2573]	C ²⁵⁶¹ -H ²⁵⁶³	503.2061	503.2139	4	-15.5	

¹⁾ Numbers inside brackets correspond to the position of the proteolytic peptide of the sequence of KLH1 or KLH2.

²⁾ Residue in the sequence of KLH1 or KLH2 that is forming the thioether bond (Cys-His)

i) Methionine sulfoxide

ii) Deamidation of Asn

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