

*Supporting Information for:*

**Biosynthesis of the sactipeptide Ruminococcin C by the human microbiome: Mechanistic insights into thioether bond formation by radical SAM enzymes**

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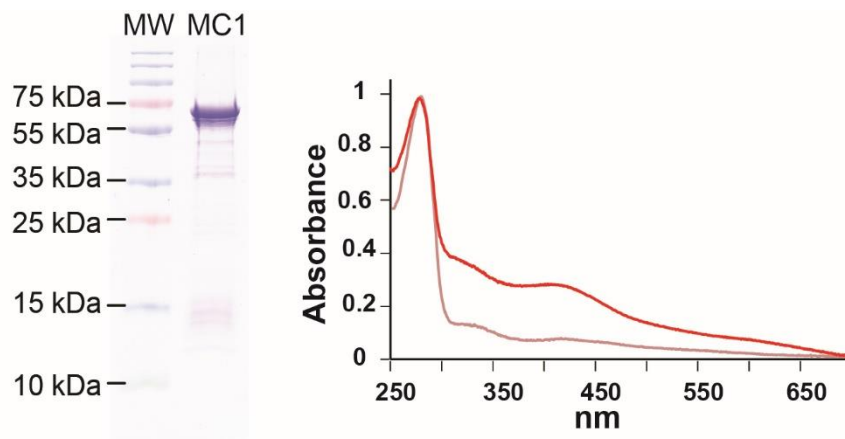
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<b>Figure S1</b> - Sequence alignment of RumMC1 & RumMC2 with other SPASM-domain radical SAM enzymes.....	2
<b>Figure S2</b> – SDS-PAGE analysis and UV-visible spectrum of RumMC1 before (pink trace) and after (red trace) iron-sulfur cluster reconstitution.....	3
<b>Figure S3</b> – LC-MS analysis of the reaction performed with the C <sub>28-63</sub> peptide.....	4
<b>Figure S4</b> - Sequence of the C1 to C5 peptides encoded by the RumC operon .....	5
<b>Figure S5</b> - MS/MS spectrum of the P2d peptide obtained after incubation with RumMC2.....	6
<b>Figure S6</b> - MS/MS spectrum of the P3d peptide obtained after incubation with RumMC2.....	7
<b>Figure S7</b> – MS/MS spectrum of the P4d peptide obtained after incubation with RumMC2 .....	7
<b>Figure S8</b> – MS/MS spectrum of the P6s peptide .....	8
<b>Figure S9</b> – MS/MS spectrum of the P6d peptide.....	8
<b>Figure S10</b> – MS/MS analysis of the C2 peptide after incubation in the presence of RumMC2ΔRRE and RRE-MC2.....	9
<b>Table S1</b> – Peptides used in this study and their theoretical and observed (obs) masses .....	10
<b>Table S2</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> [M+H] <sup>+</sup> .....	11
<b>Table S3</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> K <sup>53</sup> K <sup>61</sup> [M+H] <sup>+</sup> .....	12
<b>Table S4</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> A <sup>61</sup> [M+H] <sup>+</sup> .....	13
<b>Table S5</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> T <sup>61</sup> [M+H] <sup>+</sup> .....	14
<b>Table S6</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> G <sup>61</sup> [M+H] <sup>+</sup> .....	15
<b>Table S7</b> - Theoretical mass fragments of peptide C <sub>28-63</sub> A <sup>61</sup> d <sub>4</sub> .....	16
<b>Table S8</b> - Theoretical Mass fragments of peptide C <sub>28-63</sub> A <sup>61</sup> d <sub>3</sub> .....	17

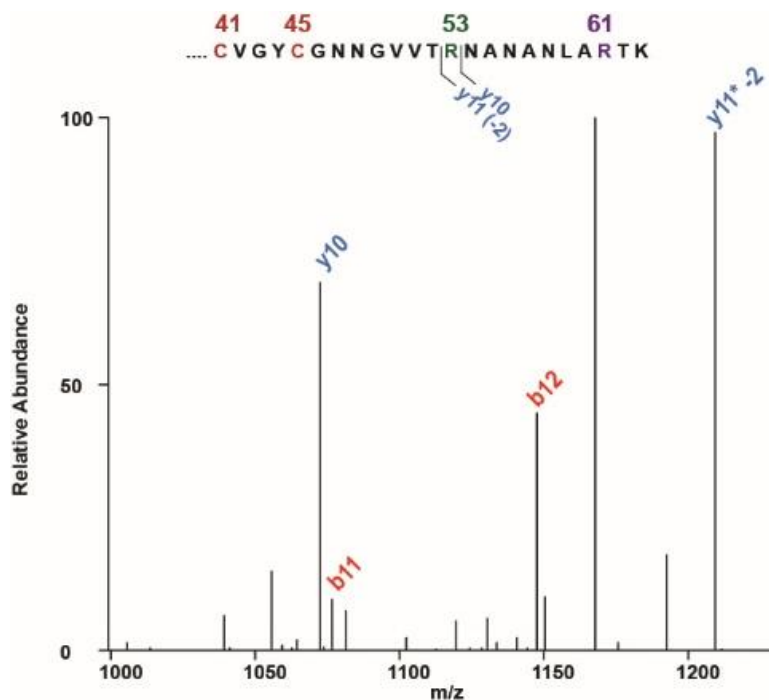
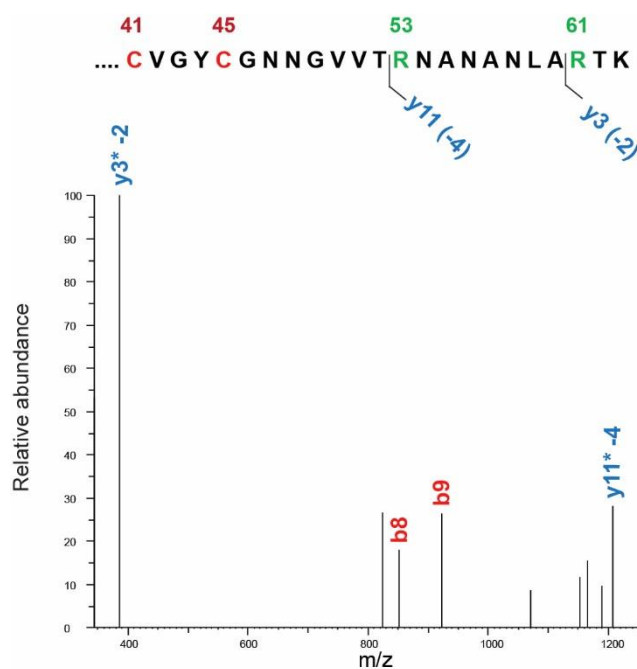


**Figure S1 - Sequence alignment of RumMC1 & RumMC2 with other SPASM-domain radical SAM enzymes.**

Sequences were aligned using Jalview (1). Blue highlighting is proportional to the conservation of amino acids between protein sequences. The canonical CX<sub>3</sub>CX<sub>2</sub>C motif is indicated by red arrows and the seven cysteine residues implicated in the SPASM motif are indicated by green arrows. For AlbA, a cysteine residue tentatively proposed to be involved in the coordination of the AuxII [4Fe-4S] cluster is squared in blue. The eighth cysteine residue, involved in the complete coordination of the AuxI [4Fe-4S] cluster in AnSME, MftC, PqqE and SuiB and CteB, is highlighted in orange.



**Figure S2 – SDS-PAGE analysis and UV-visible spectrum of RumMC1 before (pink trace) and after (red trace) iron-sulfur cluster reconstitution.**

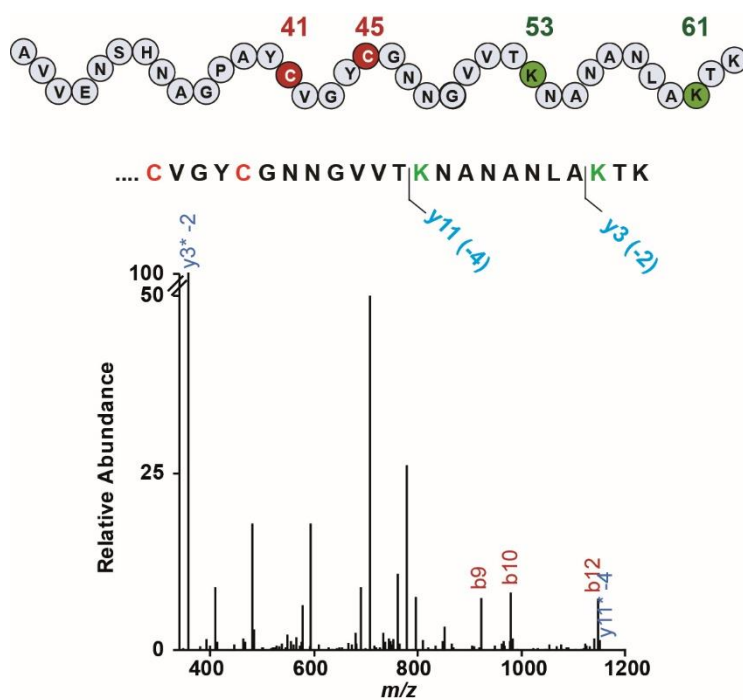
**A****B**

**Figure S3 – LC-MS analysis of the reaction performed with the C28-63 peptide**

(A) MS/MS analysis of P1s. Relevant peptide fragments are indicated showing that P1s contains a Cys<sup>45</sup>-to- Arg<sup>53</sup> thioether bond. (B) MS/MS analysis of P1d. Relevant peptide fragments are indicated showing that P1d contains a Cys<sup>41</sup>-to- Arg<sup>61</sup> thioether bond in addition to the Cys<sup>45</sup>-to- Arg<sup>53</sup> thioether bond (see table S2 for complete assignment). \* indicates loss of ammonia (-17.02 Da).

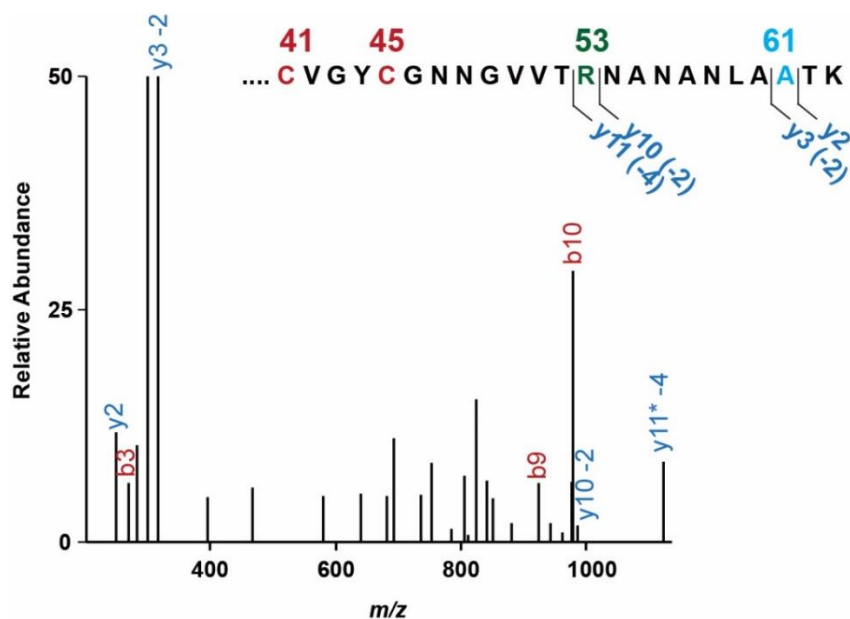
**C1** MRKIVAGKLQTFADFEKSKWGCVCSTAVANSHNAGPAYCVGYCGNNGVVTARNANANVAKTA  
**C2** MRKIVAGKLQTFADFEKSKGGCKCSGGAVVENSHNAGPAYCVGYCGNNGVVTARNANANLARTK  
**C3** MKLVETKTTKTGTNFEGNRAGCICNGTVAVANSHNAGPAYCVGYCGNSGVVTARNANANVAKTA  
**C4** MRLVQSKRIATGFNFEGSKAGCVCSGTAVANSHNAGPAYCVGYCGNNGEVTARNANYNIARRS  
**C5** MKLVTSKTMKTGTNFEGNKAGCICSGSVAVANSHNAGPAYCVGYCGNNGAVTARNANANLARTK

**Figure S4 - Sequence of the C1 to C5 peptides encoded by the RumC operon**  
 Arginine and lysine residues are highlighted in red and green respectively. Conserved cysteine residues are highlighted in blue.



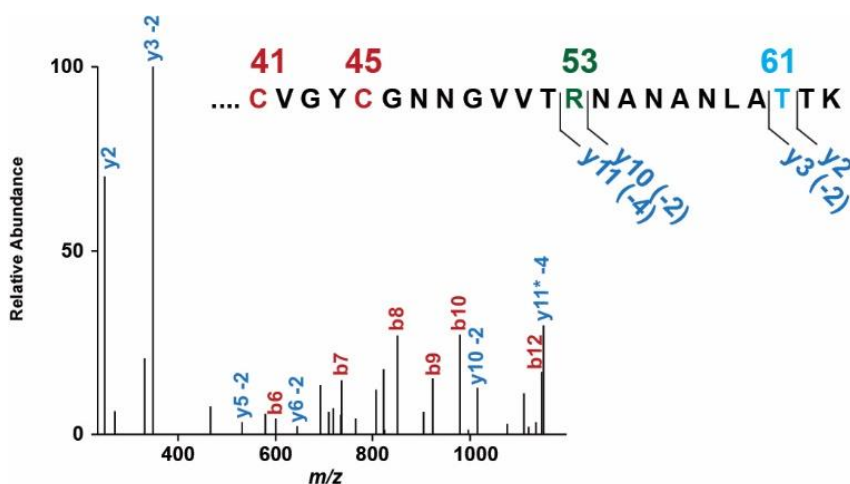
**Figure S5 - MS/MS spectrum of the P2d peptide obtained after incubation with RumMC2**

Relevant peptide fragments are indicated confirming that this peptide contains two thioether bridges (*i.e.* Lys<sup>53</sup>-Cys<sup>45</sup> and Lys<sup>61</sup>-Cys<sup>41</sup> bridges) (*see table S3* for complete assignment). \* indicates loss of ammonia (-17.02 Da).



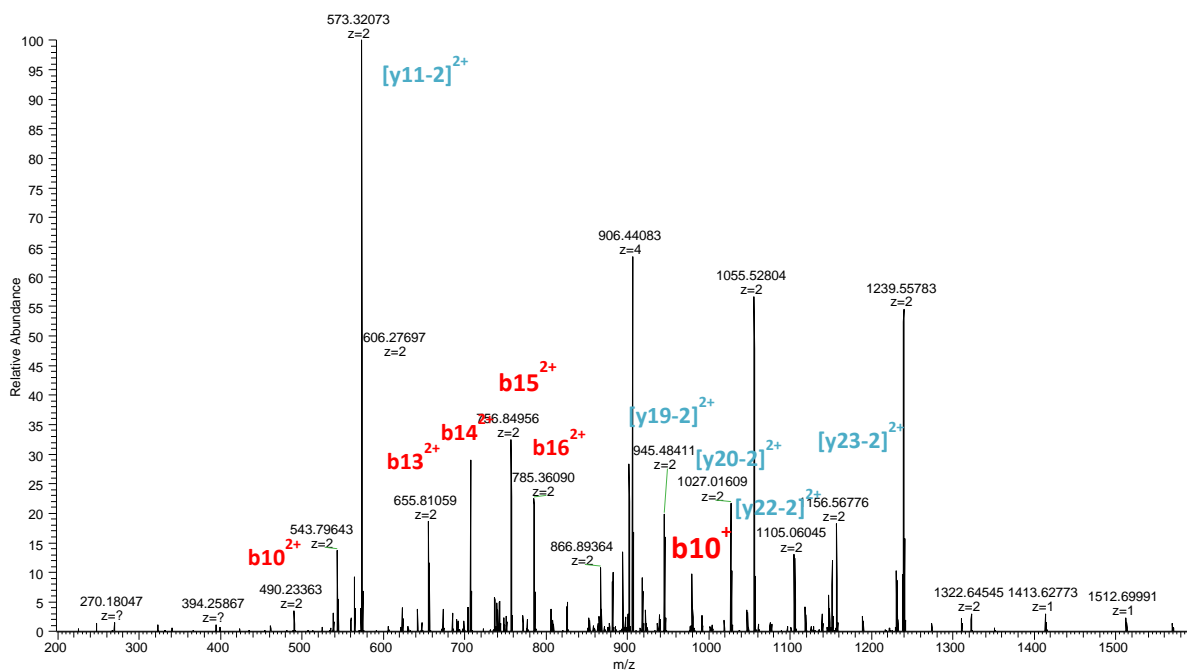
**Figure S6 - MS/MS spectrum of the P3d peptide obtained after incubation with RumMC2**

Relevant peptide fragments are indicated showing that this peptide contains two thioether bridges (*i.e.* Arg<sup>53</sup>-Cys<sup>45</sup> and Ala<sup>61</sup>-Cys<sup>41</sup> bridges) (*see table S4* for complete assignment). \* indicates loss of ammonia (-17.02 Da).



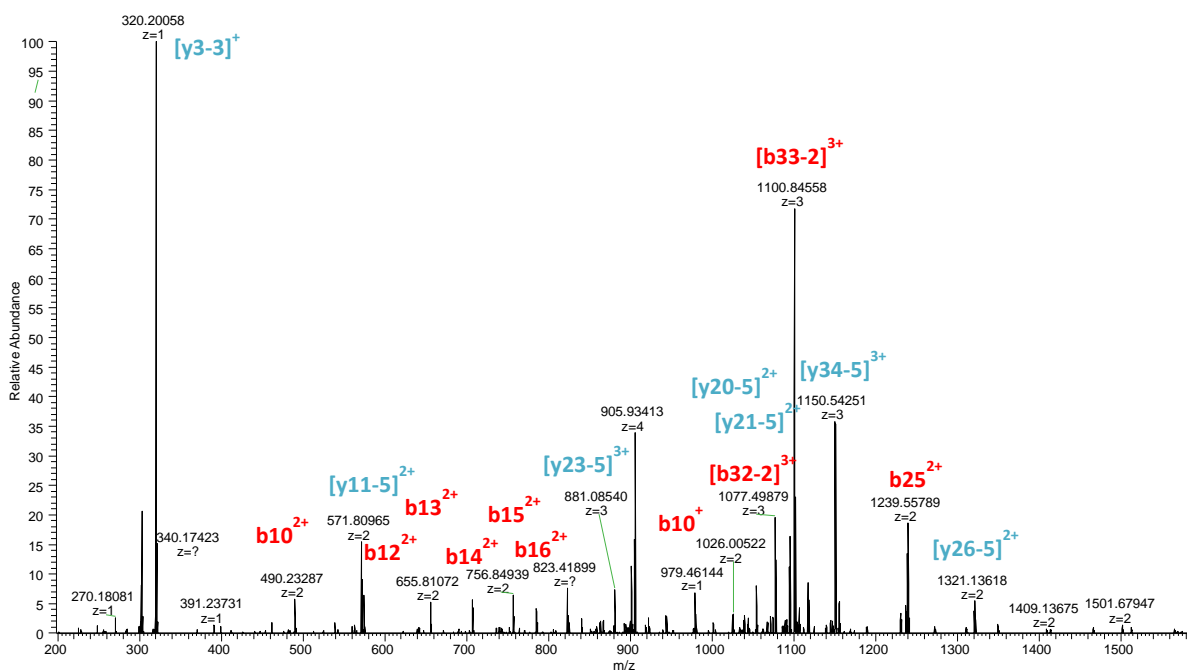
**Figure S7 – MS/MS spectrum of the P4d peptide obtained after incubation with RumMC2**

Relevant peptide fragments are indicated showing that this peptide contains two thioether bridges (*i.e.* Arg<sup>53</sup>-Cys<sup>45</sup> and Thr<sup>61</sup>-Cys<sup>41</sup> bridges) (*see table S5* for complete assignment). \* indicates loss of ammonia (-17.02 Da).



**Figure S8 – MS/MS spectrum of the P6s peptide**

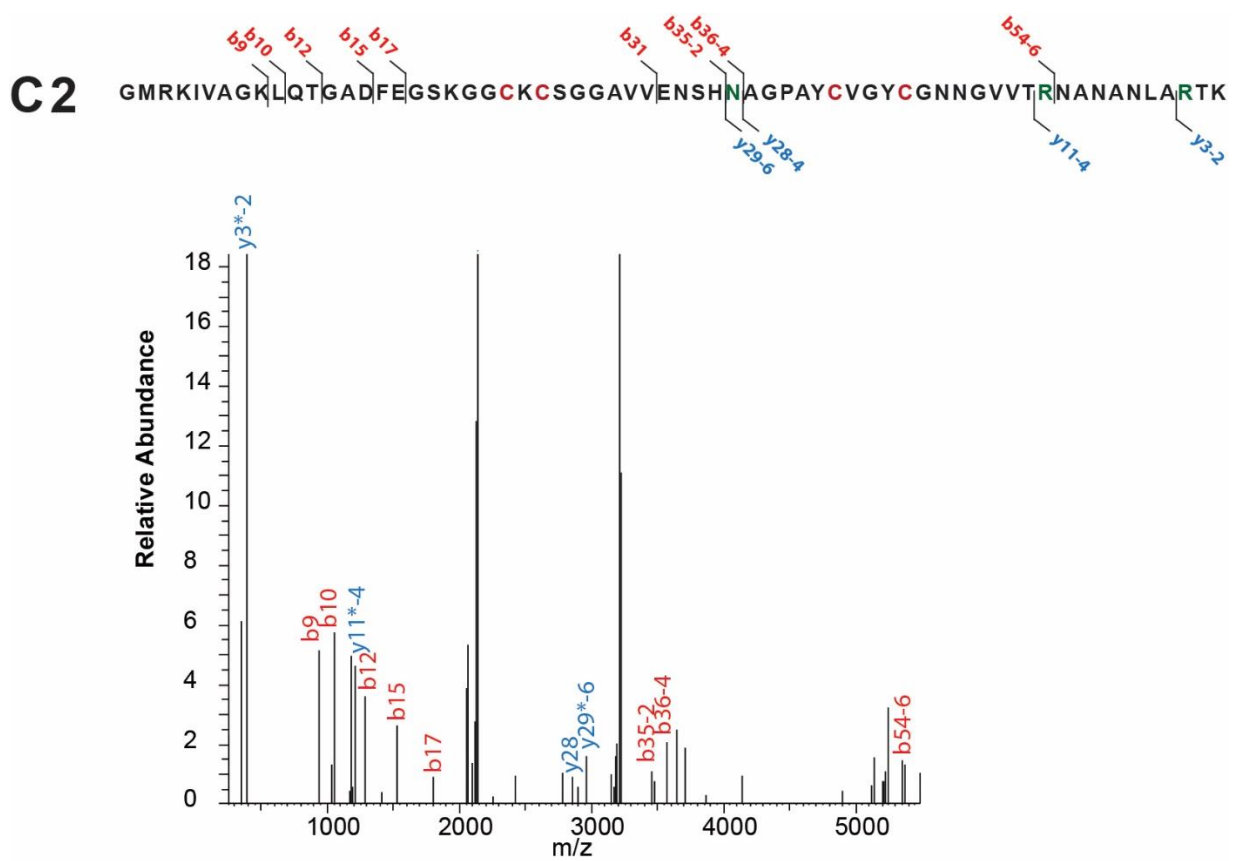
See table S7 for complete assignment



**Figure S9 – MS/MS spectrum of the P6d peptide**

See table S7 for complete assignment





**Figure S10 – MS/MS analysis of the C2 peptide after incubation in the presence of RumMC2 $\Delta$ RRE and RRE-MC2.**

C2 (250  $\mu$ M) was incubated for 4 hours with RumMC2 $\Delta$ RRE (250  $\mu$ M), RRE-MC2 (250  $\mu$ M), SAM (2 mM) and sodium dithionite (3 mM). \* indicates loss of ammonia (-17.02 Da).

**Table S1 – Peptides used in this study and their theoretical and observed (obs) masses**

Peptide	Formula	[M+4H] <sup>4+</sup>	[M+4H] <sup>4+</sup> <sub>obs</sub>
<b>C2<sub>28-63</sub></b>	C154 H246 N52 O51 S2	926.9497	926.9533
<b>C2<sub>28-63</sub>-2H</b>	C154 H244 N52 O51 S2	926.4457	926.4463
<b>C2<sub>28-63</sub>-4H</b>	C154 H242 N52 O51 S2	925.9419	925.9443
<b>C2<sub>28-63</sub>K<sup>53</sup>K<sup>61</sup></b>	C154 H246 N48 O51 S2	912.9466	912.9545
<b>C2<sub>28-63</sub>K<sup>53</sup>K<sup>61</sup>-2H</b>	C154 H244 N48 O51 S2	912.4427	912.4449
<b>C2<sub>28-63</sub>K<sup>53</sup>K<sup>61</sup>-4H</b>	C154 H242 N48 O51 S2	911.9388	91.9415
<b>C2<sub>28-63</sub>A61</b>	C151 H239 N49 O51 S2	905.6837	905.6848
<b>C2<sub>28-63</sub>A61-2H</b>	C151 H237 N49 O51 S2	905.1797	905.1831
<b>C2<sub>28-63</sub>A61-4H</b>	C151 H235 N49 O51 S2	904.6759	904.6893
<b>C2<sub>28-63</sub>T<sup>61</sup></b>	C152 H241 N49 O52 S2	913.1863	913.1920
<b>C2<sub>28-63</sub>T<sup>61</sup>-2H</b>	C152 H239 N49 O52 S2	912.6824	912.6920
<b>C2<sub>28-63</sub>T<sup>61</sup>-4H</b>	C152 H237 N49 O52 S2	912.1785	912.1822
<b>C2<sub>28-63</sub>G61</b>	C150 H237 N49 O51 S2	902.1798	902.1845
<b>C2<sub>28-63</sub>G61-2H</b>	C150 H237 N49 O51 S2	901.6758	901.6862
<b>C2<sub>28-63</sub>G61-4H</b>	C150 H237 N49 O51 S2	901.1719	901.1770

Peptide	Formula	[M+4H] <sup>4+</sup>	[M+4H] <sup>4+</sup> <sub>obs</sub>
<b>C2<sub>28-63</sub>d3</b>	C151 H236 N49 O51 S2 D3	906.4384	906.4409
<b>C2<sub>28-63</sub>d3-2H</b>	C151 H234 N49 O51 S2 D3	905.9344	905.9334
<b>C2<sub>28-63</sub>d3-4H</b>	C151 H232 N49 O51 S2 D3	905.4306	905.4329

Peptide	Formula	[M+4H] <sup>4+</sup>	[M+4H] <sup>4+</sup> <sub>obs</sub>
<b>C2<sub>28-63</sub>d4</b>	C151 H235 N49 O51 S2 D4	906.6900	906.6928
<b>C2<sub>28-63</sub>d4-2H</b>	C151 H233 N49 O51 S2 D4	906.1860	906.1890
<b>C2<sub>28-63</sub>d4-4H</b>	C151 H231 N49 O51 S2 D4	905.4306	905.4344

**Table S2 - Theoretical mass fragments of peptide C2<sub>28-63</sub> [M+H]<sup>+</sup>**

			<i>b</i>	<i>y</i>	
A	28	1	72.04444	3704.77691	36
V	29	2	171.11285	3633.73979	35
V	30	3	270.18126	3534.67138	34
E	31	4	399.22386	3435.60297	33
N	32	5	513.26678	3306.56037	32
S	33	6	600.29881	3192.51745	31
H	34	7	737.35772	3105.48542	30
N	35	8	851.40065	2968.42651	29
A	36	9	922.43776	2854.38358	28
G	37	10	979.45923	2783.34647	27
P	38	11	1076.51199	2726.325	26
A	39	12	1147.5491	2629.27224	25
Y	40	13	1310.61243	2558.23513	24
C	41	14	1413.62162	2395.1718	23
V	42	15	1512.69003	2292.16261	22
G	43	16	1569.71149	2193.0942	21
Y	44	17	1732.77482	2136.07273	20
C	45	18	1835.78401	1973.00941	19
G	46	19	1892.80547	1870.00022	18
N	47	20	2006.8484	1812.97876	17
N	48	21	2120.89133	1698.93583	16
G	49	22	2177.91279	1584.8929	15
V	50	23	2276.9812	1527.87144	14
V	51	24	2376.04962	1428.80303	13
T	52	25	2477.0973	1329.73461	12
R	53	26	2633.19841	1228.68693	11
N	54	27	2747.24133	1072.58582	10
A	55	28	2818.27845	958.5429	9
N	56	29	2932.32137	887.50578	8
A	57	30	3003.35849	773.46286	7
N	58	31	3117.40141	702.42574	6
L	59	32	3230.48548	588.38281	5
A	60	33	3301.52259	475.29875	4
R	61	34	3457.6237	404.26164	3
T	62	35	3558.67138	248.16053	2
K	63	36	3686.76634	147.11285	1

**Table S3 - Theoretical mass fragments of peptide C2<sub>28-63</sub>K<sup>53</sup>K<sup>61</sup> [M+H]<sup>+</sup>**

			<i>b</i>	<i>y</i>	
A	28	1	72.04444	3648.76461	36
V	29	2	171.11285	3577.7275	35
V	30	3	270.18126	3478.65908	34
E	31	4	399.22386	3379.59067	33
N	32	5	513.26678	3250.54808	32
S	33	6	600.29881	3136.50515	31
H	34	7	737.35772	3049.47312	30
N	35	8	851.40065	2912.41421	29
A	36	9	922.43776	2798.37128	28
G	37	10	979.45923	2727.33417	27
P	38	11	1076.51199	2670.31271	26
A	39	12	1147.5491	2573.25994	25
Y	40	13	1310.61243	2502.22283	24
C	41	14	1413.62162	2339.1595	23
V	42	15	1512.69003	2236.15032	22
G	43	16	1569.71149	2137.0819	21
Y	44	17	1732.77482	2080.06044	20
C	45	18	1835.78401	1916.99711	19
G	46	19	1892.80547	1813.98793	18
N	47	20	2006.8484	1756.96646	17
N	48	21	2120.89133	1642.92353	16
G	49	22	2177.91279	1528.88061	15
V	50	23	2276.9812	1471.85914	14
V	51	24	2376.04962	1372.79073	13
T	52	25	2477.0973	1273.72232	12
K	53	26	2605.19226	1172.67464	11
N	54	27	2719.23518	1044.57968	10
A	55	28	2790.2723	930.53675	9
N	56	29	2904.31523	859.49963	8
A	57	30	2975.35234	745.45671	7
N	58	31	3089.39527	674.41959	6
L	59	32	3202.47933	560.37667	5
A	60	33	3273.51644	447.2926	4
K	61	34	3401.61141	376.25549	3
T	62	35	3502.65908	248.16053	2
K	63	36	3630.75405	147.11285	1

**Table S4 - Theoretical mass fragments of peptide C2<sub>28-63</sub>A<sup>61</sup> [M+H]<sup>+</sup>**

			<i>b</i>	<i>y</i>	
A	28	1	72.04444	3619.71291	36
V	29	2	171.11285	3548.6758	35
V	30	3	270.18126	3449.60738	34
E	31	4	399.22386	3350.53897	33
N	32	5	513.26678	3221.49638	32
S	33	6	600.29881	3107.45345	31
H	34	7	737.35772	3020.42142	30
N	35	8	851.40065	2883.36251	29
A	36	9	922.43776	2769.31958	28
G	37	10	979.45923	2698.28247	27
P	38	11	1076.51199	2641.26101	26
A	39	12	1147.5491	2544.20824	25
Y	40	13	1310.61243	2473.17113	24
C	41	14	1413.62162	2310.1078	23
V	42	15	1512.69003	2207.09861	22
G	43	16	1569.71149	2108.0302	21
Y	44	17	1732.77482	2051.00874	20
C	45	18	1835.78401	1887.94541	19
G	46	19	1892.80547	1784.93622	18
N	47	20	2006.8484	1727.91476	17
N	48	21	2120.89133	1613.87183	16
G	49	22	2177.91279	1499.82891	15
V	50	23	2276.9812	1442.80744	14
V	51	24	2376.04962	1343.73903	13
T	52	25	2477.0973	1244.67062	12
R	53	26	2633.19841	1143.62294	11
N	54	27	2747.24133	987.52183	10
A	55	28	2818.27845	873.4789	9
N	56	29	2932.32137	802.44179	8
A	57	30	3003.35849	688.39886	7
N	58	31	3117.40141	617.36174	6
L	59	32	3230.48548	503.31882	5
A	60	33	3301.52259	390.23475	4
A	61	34	3372.55971	319.19764	3
T	62	35	3473.60738	248.16053	2
K	63	36	3601.70235	147.11285	1

**Table S5 - Theoretical mass fragments of peptide C2<sub>28-63</sub>T<sup>61</sup> [M+H]<sup>+</sup>**

			<i>b</i>	<i>y</i>	
A	28	1	72.04444	3649.72348	36
V	29	2	171.11285	3578.68636	35
V	30	3	270.18126	3479.61795	34
E	31	4	399.22386	3380.54953	33
N	32	5	513.26678	3251.50694	32
S	33	6	600.29881	3137.46401	31
H	34	7	737.35772	3050.43199	30
N	35	8	851.40065	2913.37307	29
A	36	9	922.43776	2799.33015	28
G	37	10	979.45923	2728.29303	27
P	38	11	1076.51199	2671.27157	26
A	39	12	1147.5491	2574.21881	25
Y	40	13	1310.61243	2503.18169	24
C	41	14	1413.62162	2340.11836	23
V	42	15	1512.69003	2237.10918	22
G	43	16	1569.71149	2138.04077	21
Y	44	17	1732.77482	2081.0193	20
C	45	18	1835.78401	1917.95597	19
G	46	19	1892.80547	1814.94679	18
N	47	20	2006.8484	1757.92533	17
N	48	21	2120.89133	1643.8824	16
G	49	22	2177.91279	1529.83947	15
V	50	23	2276.9812	1472.81801	14
V	51	24	2376.04962	1373.74959	13
T	52	25	2477.0973	1274.68118	12
R	53	26	2633.19841	1173.6335	11
N	54	27	2747.24133	1017.53239	10
A	55	28	2818.27845	903.48946	9
N	56	29	2932.32137	832.45235	8
A	57	30	3003.35849	718.40942	7
N	58	31	3117.40141	647.37231	6
L	59	32	3230.48548	533.32938	5
A	60	33	3301.52259	420.24532	4
T	61	34	3402.57027	349.20821	3
T	62	35	3503.61795	248.16053	2
K	63	36	3631.71291	147.11285	1

**Table S6 - Theoretical mass fragments of peptide C228-63G<sup>61</sup> [M+H]<sup>+</sup>**

			<i>b</i>	<i>y</i>	
A	28	1	72.04444	3605.69726	36
V	29	2	171.11285	3534.66015	35
V	30	3	270.18126	3435.59173	34
E	31	4	399.22386	3336.52332	33
N	32	5	513.26678	3207.48073	32
S	33	6	600.29881	3093.4378	31
H	34	7	737.35772	3006.40577	30
N	35	8	851.40065	2869.34686	29
A	36	9	922.43776	2755.30393	28
G	37	10	979.45923	2684.26682	27
P	38	11	1076.51199	2627.24536	26
A	39	12	1147.5491	2530.19259	25
Y	40	13	1310.61243	2459.15548	24
C	41	14	1413.62162	2296.09215	23
V	42	15	1512.69003	2193.08296	22
G	43	16	1569.71149	2094.01455	21
Y	44	17	1732.77482	2036.99309	20
C	45	18	1835.78401	1873.92976	19
G	46	19	1892.80547	1770.92057	18
N	47	20	2006.8484	1713.89911	17
N	48	21	2120.89133	1599.85618	16
G	49	22	2177.91279	1485.81326	15
V	50	23	2276.9812	1428.79179	14
V	51	24	2376.04962	1329.72338	13
T	52	25	2477.0973	1230.65497	12
R	53	26	2633.19841	1129.60729	11
N	54	27	2747.24133	973.50618	10
A	55	28	2818.27845	859.46325	9
N	56	29	2932.32137	788.42614	8
A	57	30	3003.35849	674.38321	7
N	58	31	3117.40141	603.34609	6
L	59	32	3230.48548	489.30317	5
A	60	33	3301.52259	376.2191	4
G	61	34	3358.54406	305.18199	3
T	62	35	3459.59173	248.16053	2
K	63	36	3587.6867	147.11285	1

**Table S7 - Theoretical mass fragments of peptide C228-63A<sup>61</sup>d4**

		<i>b</i>	<i>y</i>	
A	1	72.04444	3623.73802	36
V	2	171.11285	3552.70091	35
V	3	270.18126	3453.63249	34
E	4	399.22386	3354.56408	33
N	5	513.26678	3225.52149	32
S	6	600.29881	3111.47856	31
H	7	737.35772	3024.44653	30
N	8	851.40065	2887.38762	29
A	9	922.43776	2773.34469	28
G	10	979.45923	2702.30758	27
P	11	1076.51199	2645.28612	26
A	12	1147.5491	2548.23335	25
Y	13	1310.61243	2477.19624	24
C	14	1413.62162	2314.13291	23
V	15	1512.69003	2211.12372	22
G	16	1569.71149	2112.05531	21
Y	17	1732.77482	2055.03385	20
C	18	1835.78401	1891.97052	19
G	19	1892.80547	1788.96133	18
N	20	2006.8484	1731.93987	17
N	21	2120.89133	1617.89694	16
G	22	2177.91279	1503.85402	15
V	23	2276.9812	1446.83255	14
V	24	2376.04962	1347.76414	13
T	25	2477.0973	1248.69573	12
R	26	2633.19841	1147.64805	11
N	27	2747.24133	991.54694	10
A	28	2818.27845	877.50401	9
N	29	2932.32137	806.4669	8
A	30	3003.35849	692.42397	7
N	31	3117.40141	621.38685	6
L	32	3230.48548	507.34393	5
A	33	3301.52259	394.25986	4
A	34	3376.58482	323.22275	3
T	35	3477.63249	248.16053	2
K	36	3605.72746	147.11285	1



**Table S8 - Theoretical Mass fragments of peptide C228-63A<sup>61</sup>d3**

		<i>b</i>	<i>y</i>	
<b>A</b>	<b>1</b>	72.04444	3622.73174	36
<b>V</b>	<b>2</b>	171.11285	3551.69463	35
<b>V</b>	<b>3</b>	270.18126	3452.62621	34
<b>E</b>	<b>4</b>	399.22386	3353.5578	33
<b>N</b>	<b>5</b>	513.26678	3224.51521	32
<b>S</b>	<b>6</b>	600.29881	3110.47228	31
<b>H</b>	<b>7</b>	737.35772	3023.44025	30
<b>N</b>	<b>8</b>	851.40065	2886.38134	29
<b>A</b>	<b>9</b>	922.43776	2772.33841	28
<b>G</b>	<b>10</b>	979.45923	2701.3013	27
<b>P</b>	<b>11</b>	1076.51199	2644.27984	26
<b>A</b>	<b>12</b>	1147.5491	2547.22707	25
<b>Y</b>	<b>13</b>	1310.61243	2476.18996	24
<b>C</b>	<b>14</b>	1413.62162	2313.12663	23
<b>V</b>	<b>15</b>	1512.69003	2210.11744	22
<b>G</b>	<b>16</b>	1569.71149	2111.04903	21
<b>Y</b>	<b>17</b>	1732.77482	2054.02757	20
<b>C</b>	<b>18</b>	1835.78401	1890.96424	19
<b>G</b>	<b>19</b>	1892.80547	1787.95505	18
<b>N</b>	<b>20</b>	2006.8484	1730.93359	17
<b>N</b>	<b>21</b>	2120.89133	1616.89066	16
<b>G</b>	<b>22</b>	2177.91279	1502.84774	15
<b>V</b>	<b>23</b>	2276.9812	1445.82627	14
<b>V</b>	<b>24</b>	2376.04962	1346.75786	13
<b>T</b>	<b>25</b>	2477.0973	1247.68945	12
<b>R</b>	<b>26</b>	2633.19841	1146.64177	11
<b>N</b>	<b>27</b>	2747.24133	990.54066	10
<b>A</b>	<b>28</b>	2818.27845	876.49773	9
<b>N</b>	<b>29</b>	2932.32137	805.46062	8
<b>A</b>	<b>30</b>	3003.35849	691.41769	7
<b>N</b>	<b>31</b>	3117.40141	620.38057	6
<b>L</b>	<b>32</b>	3230.48548	506.33765	5
<b>A</b>	<b>33</b>	3301.52259	393.25358	4
<b>A</b>	<b>34</b>	3375.57854	322.21647	3
<b>T</b>	<b>35</b>	3476.62621	248.16053	2
<b>K</b>	<b>36</b>	3604.72118	147.11285	1

## Supplementary references

1. Waterhouse, A. M., Procter, J. B., Martin, D. M., Clamp, M., and Barton, G. J. (2009) Jalview Version 2--a multiple sequence alignment editor and analysis workbench. *Bioinformatics*. **25**, 1189-1191