Synthesis of Guaianolide Analogues with a Tunable α -Methylene– γ -lactam Electrophile and Correlating Bioactivity with Thiol Reactivity

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Figure S1. Structures of Supporting Information Compounds

Supporting Information Table S1

	А549/NF-к	B Luciferase Re	porter Assay	Alamar Blue Cell Viability Assay				
Compound	20µM	10µM	5µM	20µM	10µM	5µM		
45	93.6 ± 8.2	97.3 ± 9.0	96.2 ± 12.1	102.0 ± 11.3	97.3 ± 22.0	91.1 ± 13.9		
47	100.4 ± 7.7	106.2 ± 13.4	100.5 ± 15.2	87.9 ± 26.9	96.3 ± 13.8	96.8 ± 16.9		
48	101.3 ± 4.2	95.7 ± 6.5	101.6 ± 7.4	111.6 ± 17.4	98.5 ± 10.8	99.9 ± 23.0		
49	86.0 ± 17.6	95.3 ± 15.9	105.8 ± 16.4	108.4 ± 19.6	100.6 ± 11.9	97.2 ± 31.2		
51	20.9 ± 2.7	61.9 ± 8.5	85.6 ± 7.2	88.0 ± 38.3	96.8 ± 21.8	103.7 ± 22.0		
52	80.8 ± 12.8	95.7 ± 7.2	98.9 ± 11.5	99.2 ± 7.5	99.0 ± 11.1	96.6 ± 16.2		
54	2.6 ± 1.3	8.8 ± 2.2	33.1 ± 5.2	64.1 ± 7.7	63.6 ± 12.1	80.7 ± 7.0		
60	28.3 ± 13.7	59.1 ± 6.1	97.8 ± 19.5	82.3 ± 8.2	93.3 ± 10.9	89.0 ± 5.3		

A549/NF-кB-Luciferase Activity (%)

Table S1. Relative NF-κB activities (%; left) and relative cellular viabilities (%; right) for lactams dosed at various concentrations to

A549/NF- κ B-luc cells (Figure 4). Values are shown as mean \pm S.D. for $n \ge 3$ biological replicates.

				/3-INI -KB/C					
	HE	<293-NF-кB/SE	AP Reporter A	Assay	A	Alamar Blue Cell Viability Assay			
Compound	7.5µM	5µM	2.5µM	1µM	7.5µM	5µM	2.5µM	1µM	
PTL	19.7 ± 4.6	21.1 ± 5.0	49.7 ± 10.3	85.2 ± 15.8	74.4 ± 5.4	87.1 ± 9.6	93.3 ± 14.5	104.7±17.7	
49	102.0±43.1	113.2 ± 37.3	97.4 ± 29.9	143.5 ± 29.0	99.1 ± 12.2	101.3 ± 7.11	97.9 ± 12.1	94.6 ± 7.1	
51	44.1 ± 13.8	59.7 ± 18.3	93.2 ± 20.3	104.4 ± 39.3	101.2±12.3	110.7 ± 11.9	107.4 ± 6.6	113.5 ± 6.3	
54	20.1 ± 4.1	27.6 ± 6.4	54.2 ± 13.7	85.1 ± 26.6	82.0 ± 5.2	87.8 ± 8.7	92.8 ± 8.0	101.7 ± 9.1	
60	18.9 ± 3.5	20.3 ± 4.0	34.5 ± 13.3	63.4 ± 20.2	68.7 ± 9.1	76.3 ± 5.4	90.3 ± 12.7	102.6±17.1	

HEK293-NF-ĸB/SEAP Activity (%)

Table S2. Relative NF-KB activities (%; left) and relative cellular viabilities (%; right) for lactams dosed at various concentrations to

HEK293/NF- κ B-SEAP cells (Figure 4). Values are shown as mean \pm S.D. for $n \ge 3$ biological replicates.

Reaction of α -Methylene- γ -Lactams and Lactone S17 with Cysteamine: Monitoring Reaction Progress by ¹H NMR.



Fraction Remaining
$$_{x} = \frac{\frac{H_{a}+H_{b}}{2}}{\frac{H_{c}+H_{d}}{2}}$$
 (eq S2)

$$\ln(Fraction Remaining_x) = -k_{pseudo1st}t + \ln(Fraction Remaining_{x_0}) \qquad (eq S3)$$

$$t_{1/2} = \frac{\ln 2}{k_{pseudo1st}} \tag{eq S4}$$

Fraction Remaining_x =
$$\frac{\frac{H_a + H_b}{2}}{\frac{H_{hexamethylbenzene}}{32}}$$
 (eq S5)

Fraction Remaining_x =
$$\frac{\frac{H_a + H_b}{2}}{\frac{H_{hexamethylbenzene}}{18}}$$
 (eq S6)

		Integration Values				
Time	Time Point	Ha	H₀	$H_{c}+H_{d}$		
mm/dd/yy					Fraction	In(Fraction
hh:mm	d:h:min	δ 5.95	δ 5.25	δ 4.8	Remaining	Remaining)
6/3/15 12:55	1:0:00	1	1.03	1.57	0.646497	-0.43619
6/4/15 10:10	1:21:15	1	1.02	2.07	0.487923	-0.7176
6/5/15 10:30	2:21:35	1	1	2.54	0.393701	-0.93216
6/8/15 11:41	5:22:46	1	1.22	8.56	0.129673	-2.04274

 Table S3.
 ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine

 Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 45







Figure S2. Fraction Remaining Values Plotted Against Time Points for Lactam 45

Figure S3. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 45





 Table S4.
 ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine

 Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 49

		Int	tegration	values		
Time	Time Point	Ha	Hb	H _c +H _d		
mm/dd/yy					Fraction	In(Fraction
hh:mm	min	δ 6.24	δ 5.46	δ 5.35	Remaining	Remaining)
5/22/19 12:05	0:0:00	1	1.06	1.07	0.962617	-0.0381
5/22/19 12:07	0:0:02	1	1.1	1.07	0.981308	-0.01887
5/22/19 12:43	0:0:38	1	1.1	1.08	0.972222	-0.02817
5/22/19 12:45	0:0:40	1	1.04	1.07	0.953271	-0.04786
5/22/19 13:36	0:1:31	1	1.06	1.11	0.927928	-0.0748
5/22/19 13:38	0:1:33	1	1.06	1.12	0.919643	-0.08377
5/22/19 14:06	0:2:01	1	1.06	1.12	0.919643	-0.08377
5/22/19 18:05	0:6:00	1	1.06	1.2	0.858333	-0.15276
5/23/19 7:34	0:19:29	1	1.12	1.53	0.69281	-0.367
5/23/19 9:39	0:21:34	1	1.19	1.65	0.663636	-0.41002
5/23/19 9:41	0:21:36	1	1.15	1.61	0.667702	-0.40391
5/23/19 13:44	1:1:39	1	1.13	1.73	0.615607	-0.48515
5/23/19 13:46	1:1:41	1	1.16	1.76	0.613636	-0.48835
5/23/19 15:31	1:0:00	1	1.17	1.82	0.596154	-0.51726
5/23/19 15:33	1:3:28	1	1.16	1.75	0.617143	-0.48265
5/23/19 16:49	1:4:44	1	1.15	1.89	0.568783	-0.56426
5/23/19 16:51	1:4:46	1	1.13	1.81	0.588398	-0.53035
5/24/19 10:32	1:22:27	1	1.18	2.41	0.452282	-0.79345
5/24/19 10:34	1:22:29	1	1.19	2.37	0.462025	-0.77214
5/24/19 13:43	2:1:38	1	1.26	2.62	0.431298	-0.84096
5/24/19 15:36	2:3:31	1	1.27	2.71	0.418819	-0.87032
5/24/19 17:56	2:5:51	1	1.27	2.77	0.409747	-0.89221
5/25/19 13:35	3:1:30	1	1.35	3.77	0.311671	-1.16581
5/25/19 22:16	3:10:11	1	1.33	4.11	0.283455	-1.2607
5/25/19 22:18	3:10:13	1	1.41	4.24	0.284198	-1.25808
5/26/19 11:16	3:23:11	1	1.57	5.16	0.249031	-1.39018
5/26/19 11:18	3:23:13	1	1.51	5.02	0.25	-1.38629
5/27/19 10:19	4:22:14	1	1.74	6.75	0.202963	-1.59473

Figure S4. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 49





Figure S5. Fraction Remaining Values Plotted Against Time Points for Lactam 49

Figure S6. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 49



Table S5. ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 49 and Internal Standard

		In	tegration V	Values		
Time	Time Point	Ha	Hb	Internal standard	Fraction Remaining	In(Fraction Remaining)
m/dd/yy hh:mm	d:h:min	δ 6.18	δ 5.4	δ 2.2		
5/22/2019 12:05	0:0:00	1	1.06	32.59	1.011353	0.011289
5/22/2019 12:07	0:0:02	1	1.1	33.12	1.014493	0.014389
5/22/2019 12:43	0:0:38	1	1.1	33.79	0.994377	-0.00564
5/22/2019 12:45	0:0:40	1	1.04	33.72	0.967972	-0.03255
5/22/2019 13:36	0:1:31	1	1.06	34.53	0.954532	-0.04653
5/22/2019 13:38	0:1:33	1	1.06	35.12	0.938497	-0.06348
5/22/2019 14:06	0:2:01	1	1.06	35.04	0.940639	-0.0612
5/22/2019 18:05	0:6:00	1	1.06	38.49	0.856326	-0.1551
5/23/2019 7:34	0:19:29	1	1.12	50.04	0.677858	-0.38882
5/23/2019 9:39	0:21:34	1	1.19	53.7	0.652514	-0.42692
5/23/2019	0:21:36	1	1.15	52.37	0.656865	-0.42028
5/23/2019 13:44	1:1:39	1	1.13	56.2	0.606406	-0.50021
5/23/2019 13:46	1:1:41	1	1.16	56.51	0.611573	-0.49172
5/23/2019 15:31	1:0:00	1	1.17	57.77	0.601004	-0.50915
5/23/2019 15:33	1:3:28	1	1.16	56.65	0.610062	-0.4942
5/23/2019 16:49	1:4:44	1	1.15	59.86	0.574674	-0.55395
5/23/2019 16:51	1:4:46	1	1.13	59.34	0.574317	-0.55457
5/24/2019 10:32	1:22:27	1	1.18	78.15	0.446321	-0.80672
5/24/2019 10:34	1:22:29	1	1.19	78.49	0.446426	-0.80648
5/24/2019 13:43	2:1:38	1	1.26	83.98	0.430579	-0.84263
5/24/2019 15:36	2:3:31	1	1.27	88.54	0.41021	-0.89109
5/24/2019 17:56	2:5:51	1	1.27	95.41	0.380673	-0.96581
5/25/2019 13:35	3:1:30	1	1.35	124.4	0.302251	-1.1965
5/25/2019 22:16	3:10:11	1	1.33	144.36	0.258243	-1.35385
5/25/2019	3:10:13	1	1.41	146.97	0.262366	-1.33801
5/26/2019 11:16	3:23:11	1	1.57	184.74	0.222583	-1.50245
5/26/2019 11:18	3:23:13	1	1.51	178.01	0.225605	-1.48897
5/27/2019 10:19	4:22:14	1	1.74	248.15	0.176667	-1.73349

Figure S7. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 49 with Internal Standard





Figure S8. Fraction Remaining Values Plotted Against Time Points for Lactam 49

Figure S9. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 49





 Table S6.
 ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine

 Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 51

		Integration				
Time	Timepoint	Proton Ha	Proton H₅	Proton Hc+HD		
m/dd/yy		6.28	5.52			
hh:mm	d:h:min	ppm	ppm	5.34 ppm	%rem	ln%rem
5/14/18 13:38	0:0:02	1	1	1.15	0.869565	-0.13976
5/14/18 14:55	0:1:19	1	0.98	1.31	0.755725	-0.28008
5/14/18 15:43	0:2:07	1	1.09	1.49	0.701342	-0.35476
5/14/18 16:42	0:3:06	1	1.07	1.56	0.663462	-0.41028
5/14/18 16:51	0:3:15	1	1.06	1.62	0.635802	-0.45287
5/14/18 16:53	0:3:17	1	1	1.61	0.621118	-0.47623
5/15/18 2:25	0:12:49	1	1.09	3.2	0.326563	-1.11913
5/15/18 8:44	0:19:08	1	1.04	5.32	0.191729	-1.65167
5/15/18 9:40	0:20:04	1	1.11	5.94	0.177609	-1.72817
5/15/18 10:43	0:21:07	1	0.86	4.82	0.192946	-1.64534
5/15/18 11:32	0:21:56	1	1.18	6.19	0.17609	-1.73676
5/15/18 12:34	0:22:58	1	1.02	6.27	0.161085	-1.82583
5/15/18 14:37	1:0:59	1	0.94	6.39	0.1518	-1.88519



Figure S10. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 51



Figure S11. Fraction Remaining Values Plotted Against Time Points for Lactam 51

Figure S12. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 51





 Table S7.
 ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine

 Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 52

		Int	egration	Values		
Time	Time Point	Ha	H₀	$H_{c}+H_{d}$		
mm/dd/yy					Fraction	In(Fraction
hh:mm	d:h:min	δ 6.22	δ 5.45	δ 5.31	Remaining	Remaining)
4/11/19 14:25	0:0:02	1	1.03	1.06	0.957547	-0.04338
4/11/19 14:28	0:0:05	1	1.04	1.09	0.93578	-0.06638
4/11/19 14:30	0:0:07	1	1.03	1.07	0.948598	-0.05277
4/11/19 14:45	0:0:22	1	1.03	1.09	0.931193	-0.07129
4/11/19 14:47	0:0:24	1	1.03	1.09	0.931193	-0.07129
4/11/19 15:08	0:0:45	1	1.03	1.09	0.931193	-0.07129
4/11/19 15:10	0:0:47	1	1.03	1.07	0.948598	-0.05277
4/11/19 16:39	0:2:16	1	1.06	1.1	0.936364	-0.06575
4/11/19 16:41	0:2:18	1	1.05	1.12	0.915179	-0.08864
4/12/19 8:59	0:18:36	1	1.09	1.23	0.849593	-0.163
4/12/19 10:13	0:19:50	1	1.09	1.26	0.829365	-0.18709
4/12/19 10:15	0:19:52	1	1.03	1.23	0.825203	-0.19213
4/12/19 10:17	0:19:54	1	1.05	1.2	0.854167	-0.15763
4/12/19 12:52	0:22:29	1	1.04	1.21	0.842975	-0.17082
4/12/19 12:54	0:22:31	1	1.05	1.26	0.813492	-0.20642
4/12/19 13:47	0:23:24	1	1.05	1.25	0.82	-0.19845
4/12/19 16:04	1:1:41	1	1.05	1.25	0.82	-0.19845
4/12/19 16:52	1:2:29	1	1.07	1.26	0.821429	-0.19671
4/13/19 12:01	1:21:38	1	1.07	1.32	0.784091	-0.24323
4/13/19 14:13	1:23:50	1	1.09	1.34	0.779851	-0.24865
4/13/19 17:46	2:3:23	1	1.04	1.26	0.809524	-0.21131
4/13/19 17:48	2:3:25	1	1.05	1.27	0.807087	-0.21432
4/14/19 12:29	2:22:06	1	1.06	1.38	0.746377	-0.29252
4/14/19 12:32	2:22:09	1	1.09	1.4	0.746429	-0.29246
4/15/19 10:10	3:19:47	1	1.09	1.56	0.669872	-0.40067
4/15/19 10:13	3:19:50	1	1.07	1.54	0.672078	-0.39738
4/15/19 13:54	3:23:31	1	1.08	1.6	0.65	-0.43078
4/16/19 9:37	4:19:14	1	1.08	1.64	0.634146	-0.45548

4/16/19 11:51	4:21:28	1	1.09	1.67	0.625749	-0.46881
4/16/19 17:27	5:3:04	1	1.08	1.7	0.611765	-0.49141
4/16/19 21:07	5:6:44	1	1.11	1.76	0.599432	-0.51177
4/17/19 10:14	5:19:51	1	1.11	1.84	0.57337	-0.55622

Figure S13. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 52



.0 6.9 6.8 6.7 6.6 6.5 6.4 6.3 6.2 6.1 6.0 5.9 5.8 5.7 5.6 5.5 5.4 5.3 5.2 5.1 5.0 4.9 4.8 $^{\rm f1\,(ppm)}$



Figure S14. Fraction Remaining Values Plotted Against Time Points for Lactam 52

Figure S15. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 52



Table S8. ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 52 with Internal Standard

		Int	egration `	Values		
Time	Time Point	Ha	H _b	Internal standard	Fraction Remaining	ln(Fraction Remaining)
m/dd/yy hh:mm	d:h:min	δ 6.22	δ 5.45	δ 2.20		
5/3/2019 16:19	0:0:00	1	1.02	17.83	1.01963	0.01944
5/4/2019 12:07	0:19:48	1	1.07	20.23	0.92091	-0.08239
5/4/2019 15:05	0:22:46	1	1.13	20.6	0.930583	-0.07194
5/4/2019 15:07	0:22:48	1	1.05	20.63	0.894329	-0.11168
5/6/2019 9:51	2:17:32	1	1.12	25.26	0.755344	-0.28058
5/7/2019 13:30	3:21:11	1	1.19	27.59	0.714389	-0.33633
5/8/2019 13:42	4:21:23	1	1.03	28.02	0.652034	-0.42766
5/9/2019 8:14	5:15:55	1	1.12	32.13	0.593838	-0.52115
5/9/2019 14:34	5:22:15	1	1.22	34.19	0.584381	-0.5372
5/10/2019 4:08	6:11:49	1	1.31	36.35	0.571939	-0.55872

Figure S16. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 52 with Internal Standard



Figure S17. Fraction Remaining Values Plotted Against Time Points for Lactam 52 with Internal Standard



Figure S18. ln(Fraction Remaining) Values Plotted Against Time Points for Lactam 52 with Internal Standard





 Table S9.
 ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine

 Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactam 60

		Int	tegration	/alues		
	Time				Fraction	In(Fraction
Time	Point	Ha	Hb	$H_{c}+H_{d}$	Remaining	Remaining)
m/dd/yy						
hh:mm	min	δ 6.28	δ 5.50	δ 5.06		
5/17/18 15:12	4	1	0.82	2.17	0.419354839	-0.869037847
5/17/18 15:14	6	1	1.07	3.5	0.295714286	-1.218361542
5/17/18 15:16	8	1	0.83	3.81	0.24015748	-1.426460403

Figure S19. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactam 60





Figure S20. Fraction Remaining Values Plotted Against Time Points for Lactam 60

Figure S21. In(Fraction Remaining) Values Plotted Against Time Points for Lactam 60





Table S10. ¹H NMR Integration and In(Fraction Remaining) Values Used to Determine Pseudo-first Order Rate Constant for Reaction of Cysteamine with Lactone S17

	Int	tegration	/alues		
Time				Fraction	In(Fraction
Point	Ha	Hb	$H_{c}+H_{d}$	Remaining	Remaining)
min	δ 6.22	δ 5.53	δ 5.45		
3	1	1	1.15	0.869565	-0.13976
6	1	0.98	1.31	0.755725	-0.28008
8	1	1.09	1.49	0.701342	-0.35476
10	1	1.07	1.56	0.663462	-0.41028
12	1	1.06	1.62	0.635802	-0.45287
14	1	1	1.61	0.621118	-0.47623
17	1	1.09	3.2	0.326563	-1.11913
19	1	1.04	5.32	0.191729	-1.65167
21	1	1.11	5.94	0.177609	-1.72817
24	1	0.86	4.82	0.192946	-1.64534
26	1	1.18	6.19	0.17609	-1.73676
28	1	1.02	6.27	0.161085	-1.82583
30	1	0.94	6.39	0.1518	-1.88519



Figure 22. ¹H NMR Spectra at Reaction Time Points for Cysteamine and Lactone S17



Figure S23. Fraction Remaining Values Plotted Against Time Points for Lactone S17

Figure S24. In(Fraction Remaining) Values Plotted Against Time Points for Lactone S17



Biological Materials and Methods

Compound	Retention Time (min)	Purity (%)
45	14.8	>99
47	11.0	98.4
48	15.2	95.8
49	20.1	95.6
51	22.5	93.6
52	19.1	92.1
54	20.2	91.0
60	13.3	98.6

 Table S11. HPLC purities of compounds.

Figure S25. HPLC trace of **45**, > 99% pure. RT = 14.8 min. Peak at 1.5-3 min is DMSO.



Figure S26. HPLC trace of **47**, 98.4% pure. RT = 11.0 min. Peak at 1.5-3 min is DMSO. The peak at 18.0 min appears in the blank run, and was excluded from purity analysis.



Figure S27. HPLC trace of **48**, 95.8% pure. RT = 15.2 min. Peak at 1.5-3 min is DMSO. The peak at 18.0 min appears in the blank run, and was excluded from purity analysis.



Figure S28. HPLC trace of **49**, 95.6% pure. RT = 20.1 min. The peak at 18.0 min appears in the blank run, and was excluded from purity analysis.



Figure S29. HPLC trace of **51**, 93.6% pure. RT = 22.5 min.





Figure S30. HPLC trace of **52**, 92.1% pure, RT = 19.1 min. Peak at 1.5-3 min is DMSO.

Figure S31. HPLC trace of **54**, 91.0% pure. RT = 20.2 min.





Figure S32. HPLC trace of **60**, 98.6% pure. RT = 13.3 min.

Figure S33. 4-15% SDS-PAGE analysis of La antigen protein expression, stained with Coomassie blue.



Table S12. Observed adducts from La Antigen mass spectrometry MS/MS analysis. Listed are the cysteine adducts identified for the three peptides searched. IAD = carbamidomethylation, D2-IAD = d_2 -carbamidomethylation. Fluconazole and CPM are the same negative and positive controls, respectively, used in the ALARM NMR assay. N/A indicated the peptide was not identified. Samples included three biological replicates analyzed each in technical triplicate.

Sample	Compound	SLEEKIG <u>C</u> ²³² LLK	IG <u>C</u> ²³² LLK	FSGDLDDQT <u>C²⁴⁵R</u>
S1_1		IAD/D2-IAD	D2-IAD	N/A
S1_2	DMSO	IAD/D2-IAD	D2-IAD	IAD
S1_3		IAD/D2-IAD	N/A	IAD
S2_1		IAD/D2-IAD	IAD/D2-IAD	IAD
S2_2	DMSO	IAD/D2-IAD	IAD/D2-IAD	IAD
S2_3		IAD/D2-IAD	IAD/D2-IAD	IAD
S3_1		D2-IAD	IAD/D2-IAD	IAD
S3_2	DMSO	IAD/D2-IAD	IAD/D2-IAD	IAD
S3_3		IAD/D2-IAD	IAD/D2-IAD	IAD

S4_1		IAD/D2-IAD	D2-IAD	IAD
S4_2	Fluconazole	IAD/D2-IAD	D2-IAD	IAD
S4_3		IAD/D2-IAD	D2-IAD	IAD
S5_1		IAD/D2-IAD	IAD/D2-IAD	IAD
S5_2	Fluconazole	IAD/D2-IAD	D2-IAD	IAD
S5_3		IAD/D2-IAD	D2-IAD	IAD
S6_1		IAD/D2-IAD	D2-IAD	IAD
S6 2	Fluconazole	IAD/D2-IAD	IAD/D2-IAD	IAD
S6 3		IAD/D2-IAD	D2-IAD	IAD
S7 1		D2-IAD	D2-IAD/CPM	D2-IAD
S7 2	СРМ	D2-IAD/CPM	D2-IAD/CPM	D2-IAD
S7 3		D2-IAD/CPM	D2-IAD/CPM	D2-IAD
S8 1		D2-IAD/CPM	D2-IAD/CPM	D2-IAD
S8 2	СРМ	D2-IAD/CPM	D2-IAD/CPM	D2-IAD
S8 3		D2-IAD/CPM	D2-IAD/CPM	D2-IAD
S9 1		D2-IAD/CPM	D2-IAD/CPM	N/A
S9 2	CPM	D2-IAD/CPM	D2-IAD/CPM	N/A
S9 3		D2-IAD/CPM	D2-IAD/CPM	N/A
S10_1		D2-IAD	D2-IAD/ S17	D2-IAD
S10_2	S17	D2-IAD/S17	D2-IAD/ S17	D2-IAD
S10_3		D2-IAD/S17	D2-IAD/ S17	D2-IAD
S11_1		D2-IAD/S17	D2-IAD/ S17	D2-IAD
S11_2	S17	D2-IAD/S17	D2-IAD/ S17	D2-IAD
S11_3		D2-IAD/S17	D2-IAD/ S17	D2-IAD
S12 1		D2-IAD/S17	D2-IAD/ S17	D2-IAD
S12_2	S17	D2-IAD/S17	D2-IAD/ S17	D2-IAD
S12_3		D2-IAD/S17	D2-IAD/ S17	D2-IAD
S13_1		IAD	IAD/D2-IAD	IAD
S13_2	51	IAD/D2-IAD	IAD/D2-IAD	IAD
S13_3		IAD	IAD/D2-IAD	IAD
S14 1		IAD/D2-IAD	D2-IAD/ 51	IAD
S14_2	51	IAD/D2-IAD	D2-IAD	IAD
S14_3		IAD/D2-IAD	D2-IAD/51	IAD
S15_1		IAD/D2-IAD	IAD/D2-IAD/51	IAD
S15 2	51	IAD/D2-IAD	IAD/D2-IAD/51	IAD
S15 3		IAD/D2-IAD	IAD/D2-IAD/51	IAD



Figure S34. Heat map analysis of the observed adducts from La Antigen mass spectrometry MS/MS analysis (Table 12). Fluconazole is the negative control (non-thiol reactive) and CPM is the positive control (thiol-reactive). Asterisk indicates three samples overlap between these two categories (i.e., in three samples, $IG\underline{C}^{232}LLK$ is found adducted with IAD, d₂-IAD, and compound). Boxes in the middle of two possible adducts indicates the number of samples in which
the peptide is found with both adducts. Samples included three biological replicates analyzed each in technical triplicate.



¹H NMR, ¹³C NMR, and Peptide Mass Spectra






















































































































































































































































Figure S27. Example MS/MS spectrum from precursor ion m/z = 657.274 that is positively identified as a target peptide. The y- and b-ions of the carbamidomethylated FSGDLDDQT<u>C²⁴⁵R</u> are highlighted in blue and orange, respectively.



Figure S28. Example MS/MS spectrum from precursor ion m/z = 645.360 that is positively identified as a target peptide. The y- and b-ions of the carbamidomethylated SLEEKIG<u>C²³²LLK</u> are highlighted in blue and orange, respectively.



Figure S29. Example MS/MS spectrum from precursor ion m/z = 352.212 that is positively identified as a target peptide. The y- and b-ions of the carbamidomethylated IG<u>C</u>²³²LLK are highlighted in blue and orange, respectively.



Figure S30. Example MS/MS spectrum from precursor ion m/z = 658.281 that is positively identified as a target peptide. The y- and b-ions of the d₂-carbamidomethylated FSGDLDDQT<u>C²⁴⁵R</u> are highlighted in blue and orange, respectively.

Figure S31. Example MS/MS spectrum from precursor ion m/z = 646.366 that is positively identified as a target peptide. The y- and b-ions of the d₂-carbamidomethylated SLEEKIG<u>C²³²LLK</u> are highlighted in blue and orange, respectively.

Figure S32. Example MS/MS spectrum from precursor ion m/z = 353.218 that is positively identified as a target peptide. The y- and b-ions of the d₂-carbamidomethylated IG<u>C</u>²³²LLK are highlighted in blue and orange, respectively.

Figure S33. Example MS/MS spectrum from precursor ion m/z = 524.780 that is positively identified as a target peptide. The y- and b-ions of the CPM-adducted IGC²³²LLK are highlighted in blue and orange, respectively.

Figure S34. Example MS/MS spectrum from precursor ion m/z = 817.928 that is positively identified as a target peptide. The y- and b-ions of the CPM-adducted SLEEKIG<u>C</u>²³²LLK are highlighted in blue and orange, respectively.

Figure S35. Example MS/MS spectrum from precursor ion m/z = 476.764 that is positively identified as a target peptide. The y- and b-ions of the S17-adducted IG<u>C</u>²³²LLK are highlighted in blue and orange, respectively.

Figure S36. Example MS/MS spectrum from precursor ion m/z = 817.928 that is positively identified as a target peptide. The y- and b-ions of the S17-adducted SLEEKIG<u>C</u>²³²LLK are highlighted in blue and orange, respectively.

Figure S37. Example MS/MS spectrum from precursor ion m/z = 548.281 that is positively identified as a target peptide. The y- and b-ions of 51-adducted IGC²³²LLK are highlighted in blue and orange, respectively.

