

Table S3: Average number of deuterons incorporated in Sse2 + ATP and in Sse2-Strep-Tag II in complex with Ssa1-His₁₀

PEPTIDE			AVERAGE NUMBER OF DEUTERONS INCORPORATED							
Start	End	Mass (exp) [Da]	HX time	10 s	30 s	2 min	10 min	2 h	100%	
26	39	1499.8	Sse2 + ATP	4.6	6.6	8.0	8.6	8.4	8.8	
			Sse2-Ssa1 complex	2.9	2.7	3.9	5.7	8.0		
31	39	959.5		2.4	3.5	4.0	4.1	4.1	4.4	
				1.0	1.4	2.4	2.5	3.7		
40	68	3192.7		13.4	15.2	16.6	18.1	18.7	19.1	
				8.8	10.2	11.4	13.8	16.9		
69	81	1589.9		2.6	2.9	3.6	5.1	6.2	7.8	
				2.3	2.8	3.5	4.8	6.0		
74	81	1022.5		1.8	2.0	2.3	3.1	3.5	3.4	
				1.8	2.2	2.3	3.3	3.5		
75	81	909.5		1.7	1.8	1.9	2.2	2.5	2.6	
				1.6	1.6	1.7	2.1	2.4		
93	103	1169.7		2.3	2.6	2.8	3.3	4.3	7.1	
				2.2	2.4	2.6	3.2	4.1		
104	118	1619.8		4.5	6.0	6.5	6.9	7.3	9.2	
				4.3	5.6	6.1	6.7	7.0		
106	118	1391.7		4.1	5.4	5.8	6.2	5.9	7.6	
				3.8	5.1	5.6	5.8	5.9		
122	140	2204.2		4.6	6.0	7.0	8.0	8.6	11.0	
				4.0	4.9	6.0	7.4	8.3		
123	142	2259.1		4.6	5.8	6.7	7.8	8.6	11.6	
				3.9	4.5	5.9	7.2	8.4		
144/145	159/160	1938.9		0.6	1.1	1.5	1.7	2.7	10.9	
				0.6	1.1	1.3	1.7	2.6		
156/158	165/167	970.5		3.5	3.9	4.3	5.2	5.4	5.6	
				2.2	2.8	3.2	3.4	4.1		
160	177	1849.0		0.3	0.5	0.9	1.9	2.6	12.6	
				0.3	0.5	0.7	1.7	2.5		
160	179	1991.1		0.3	0.4	0.8	1.8	2.6	13.8	
				0.3	0.3	0.6	1.9	2.4		
168	177	1082.6		0.3	0.4	0.9	1.6	1.8	8.7	
				0.2	0.3	N.D.	1.4	1.8		
186	201	1775.0		2.6	3.1	3.4	3.6	3.5	7.3	
				2.4	2.8	3.3	3.5	3.5		
186	203	1989.1		2.6	3.1	3.5	3.5	3.4	8.7	
				2.5	2.9	3.3	3.3	3.3		
202	211	1094.5		0.2	0.2	0.2	0.4	0.7	4.9	
				N.D.	0.2	0.3	0.2	0.4		
202	212	1181.5		0.2	0.2	0.2	0.4	0.7	5.8	
				0.1	0.2	N.D.	0.1	0.2		

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202	214	1425.6		0.3 0.2	0.3 0.3	0.3 0.3	0.4 0.2	0.8 0.3	7.4
215	221	837.4		1.0 0.8	1.2 1.2	1.5 1.3	1.7 1.4	1.9 1.6	4.0
215	236	2482.3		0.8 0.8	1.2 1.1	1.9 1.8	3.3 3.1	4.7 4.6	12.7
216	227	1335.7		0.6 0.5	0.8 0.8	1.3 0.9	2.0 1.8	3.1 3.2	7.5
222	236	1662.8		0.5 0.7	0.9 1.1	1.3 1.5	2.3 2.2	2.5 2.5	7.8
222	240	2152.1		0.6 0.6	1.0 1.1	1.3 1.4	2.1 1.9	2.7 2.6	9.9
237	243	850.4		0.2 0.3	0.3 0.3	0.3 0.5	0.3 0.3	0.6 0.6	3.9
269	289	2216.2		4.4 3.5	4.9 3.9	5.7 4.5	6.4 5.9	7.6 7.7	14.8
271	289	2032.1		4.2 3.1	4.6 3.4	5.4 4.2	7.6 5.7	10.0 7.8	12.5
272	289	1961.1		4.1 3.1	4.7 3.4	5.3 4.2	7.5 5.8	9.6 7.3	11.7
293	301	990.4		1.4 1.4	1.6 1.8	2.2 2.3	3.4 3.3	4.1 4.1	4.4
309	314	682.4		0.1 N.D.	N.D. N.D.	0.5 0.5	0.9 1.2	2.1 2.1	2.4
309	325	1939.1		2.7 2.6	3.6 3.6	4.9 4.9	6.6 6.6	8.3 8.1	11.2
309	330	2450.4		2.6 2.5	3.6 3.4	5.1 4.9	7.3 7.1	10.6 9.8	15.8
315	330	1786.0		1.7 1.6	2.2 1.8	2.6 2.4	3.9 4.0	6.0 6.1	11.2
315	337	2590.4		3.3 3.5	3.7 4.0	4.6 5.0	6.7 7.0	10.7 11.1	17.1
331	337	822.4		0.7 0.7	0.8 0.8	0.9 0.9	1.2 1.2	2.4 2.4	3.2
337	350	1513.9		0.7 N.D.	1.5 N.D.	1.7 1.6	3.3 1.6	5.5 N.D.	8.3
337	357	2271.3		0.7 0.6	1.5 N.D.	1.7 1.4	3.2 N.D.	7.2 N.D.	15.3
338	357	2124.2		0.8 0.6	1.1 0.9	1.6 1.2	2.7 1.4	7.2 3.4	13.8
338	366	3054.8		3.2 1.1	4.8 1.8	5.8 3.1	7.7 4.9	12.7 8.3	19.9
351	366	1706.0		2.0 1.4	3.3 1.2	4.1 2.5	4.7 3.8	5.5 5.4	9.5
358	366	948.5		2.4 0.8	3.5 1.2	4.0 2.0	5.2 3.5	5.3 4.9	5.0
358	378	2150.1		4.8 2.6	6.7 3.8	7.2 5.5	7.8 7.0	9.5 8.6	13.2
367	378	1219.6		1.0 0.7	1.3 1.4	1.7 1.7	2.0 1.9	2.7 2.5	6.9
379	387	953.5		2.7 2.0	2.8 2.3	2.8 2.6	3.1 2.8	3.8 3.0	4.2

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379	394	1884.0		6.5	7.0	7.3	8.1	8.8	9.2
				4.8	6.2	6.8	7.4	8.0	
388	394	948.6		2.3	2.8	3.1	3.5	3.5	3.6
				1.5	2.0	2.6	3.0	3.5	
388	403	1954.0		3.0	3.5	4.0	4.7	6.0	8.9
				2.4	3.3	3.7	4.5	6.3	
388	405	2218.1		2.9	3.3	3.7	4.6	5.9	10.2
				2.3	2.8	3.4	4.0	6.0	
404	413	1297.5		3.4	3.7	3.7	4.0	4.3	5.0
				3.1	3.4	N.D.	3.8	4.3	
404	416	1681.8		4.0	4.7	5.0	5.7	6.4	6.6
				3.9	4.4	4.9	5.5	6.2	
406	416	1417.6		3.6	4.1	4.4	4.8	5.0	5.3
				3.5	4.0	4.3	4.7	4.9	
417	430	1538.8		3.0	3.0	3.1	4.3	5.7	7.0
				3.0	3.0	3.3	4.2	5.7	
431	439	1058.5		1.5	1.9	2.2	2.4	2.3	4.1
				1.5	1.8	2.4	2.2	2.2	
431	441	1276.6		1.6	2.4	2.9	3.3	3.2	5.5
				1.9	2.4	2.8	3.0	3.2	
434	441	949.4		1.2	1.8	2.2	2.4	2.4	3.8
				1.1	1.8	2.1	2.2	2.3	
442	458	1815.0		5.9	6.4	6.4	7.5	7.5	9.4
				5.7	6.3	6.4	7.0	7.8	
467	482	1852.1		3.2	3.4	3.9	N.D.	N.D.	N.D.
				3.3	3.3	3.9	N.D.	N.D.	
490	495	695.4		0.9	1.2	1.8	2.4	2.7	2.8
				1.0	1.3	1.9	2.2	2.8	
501	515	1632.8		2.3	2.7	3.1	4.7	6.7	8.9
				2.4	2.6	3.2	4.8	6.6	
501	521	2320.1		7.6	8.6	8.6	9.0	8.8	8.8
				7.0	8.4	8.5	8.8	8.7	
522	535	1588.9		7.2	7.8	8.3	8.9	8.7	8.6
				6.8	7.4	8.3	8.5	8.6	
522	539	1990.2		9.6	10.4	11.0	12.4	12.4	12.2
				9.3	10.1	11.0	11.9	12.3	
540	546	788.4		1.6	2.0	2.3	2.5	2.4	3.3
				1.6	2.0	2.6	2.6	2.8	
540	547	901.5		1.7	2.0	2.3	2.4	2.8	3.9
				1.6	2.1	2.1	2.3	2.6	
540	549	1130.6		1.9	2.3	2.9	3.5	3.8	4.9
				2.1	2.5	3.0	3.5	3.7	
540	550	1243.6		1.9	2.4	3.0	3.5	4.4	6.0
				2.0	2.4	2.9	3.5	4.3	
550	563	1740.9		1.3	2.4	4.6	6.0	7.9	8.3
				0.7	0.8	0.9	2.7	5.5	
551	563	1627.8		1.2	2.5	4.3	5.9	7.5	7.7
				0.6	1.0	1.4	2.8	5.1	
593	605	1475.8		1.0	1.4	3.0	5.6	7.2	7.8
				1.0	1.5	2.7	4.1	6.0	
611	624	1617.7		5.1	6.0	6.5	6.4	6.2	6.6
				3.9	4.8	5.4	5.8	6.4	

625	631	864.4		0.2	0.5	1.8	3.5	3.8	4.0
				0.1	0.2	0.5	1.5	2.9	
632	642	1218.7		2.3	4.5	6.7	7.7	7.6	8.2
				2.2	3.1	5.0	7.0	7.9	
632	655	2741.6		11.4	14.5	17.1	18.3	18.2	18.3
				10.1	12.2	15.2	17.4	18.2	
654	664	1292.6		6.0	6.4	6.0	6.2	6.0	6.8
				5.8	6.2	6.1	6.0	5.7	
670	690	2379.0		10.1	11.2	10.5	10.6	10.1	10.4
				9.8	10.6	10.5	10.2	10.1	