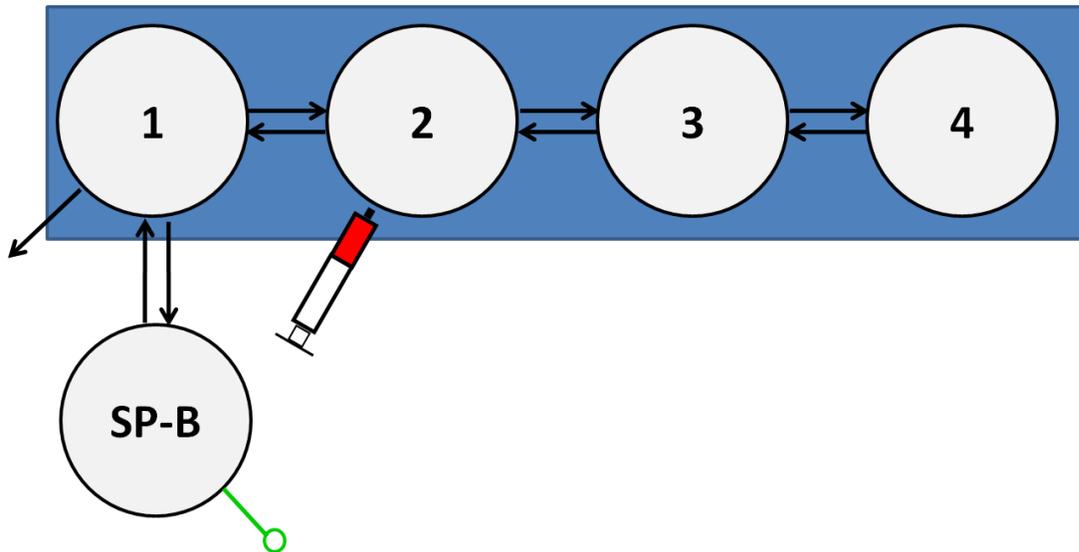
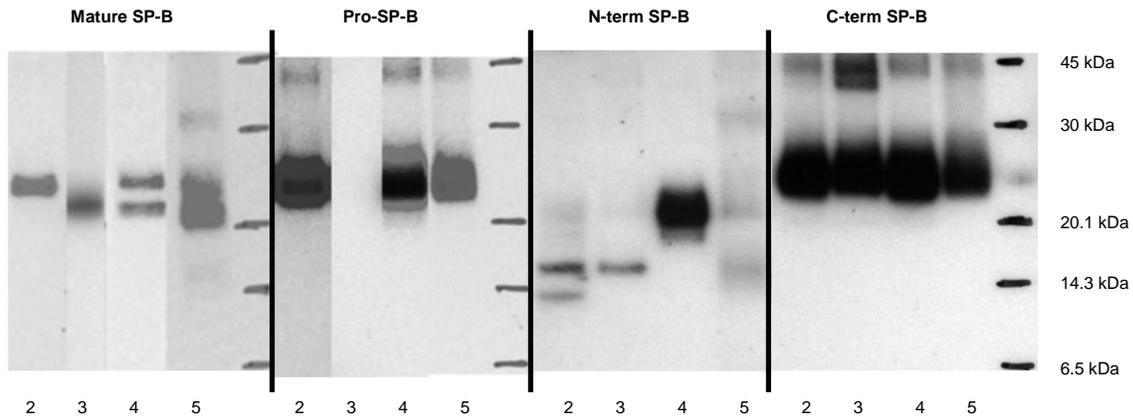


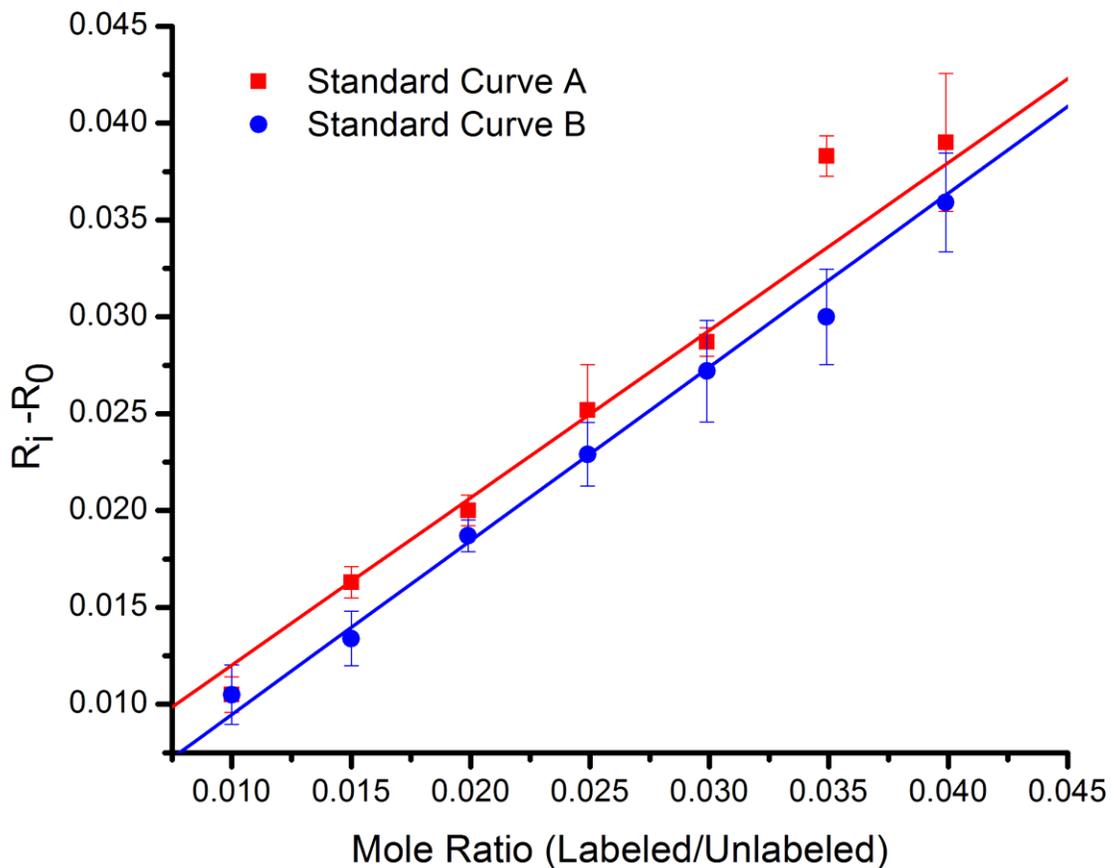
Supplementary material



Supplementary Figure 1: Compartmental model used to calculate SP-B kinetics. We used a four pool compartmental model to estimate leucine kinetics (blue box) with the addition of a fifth pool for SP-B. The measured KIC enrichment in plasma was used as a proxy of the intracellular leucine enrichment in pool 1. A population approach was used to fit the SP-B enrichments using nonlinear mixed-effects modeling²⁸. The population modeling enabled us to calculate individual rate constants even in patients with sparsely sampled data.

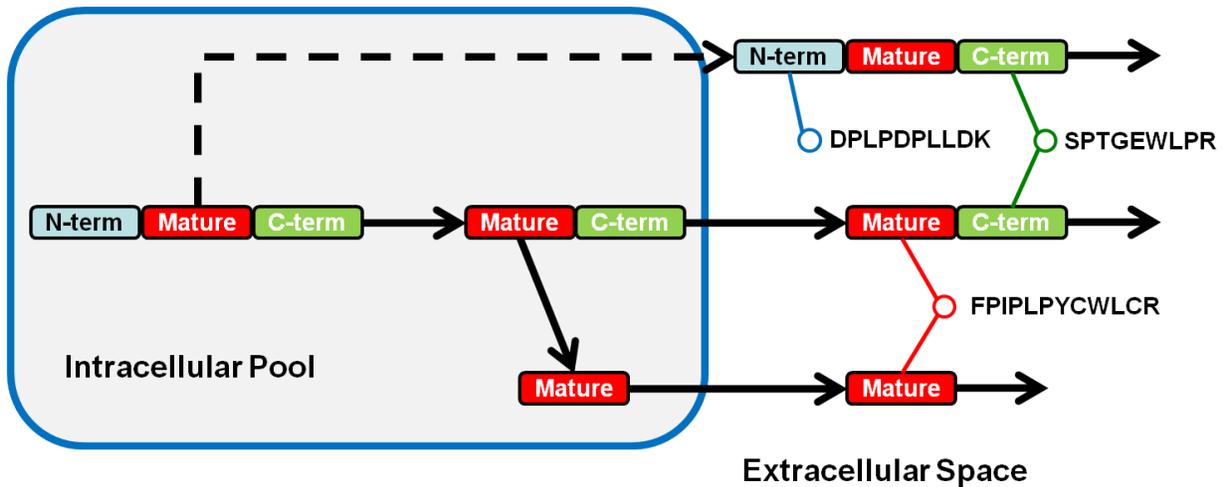


Supplementary Figure 2: Protein immunoblotting under non-reducing conditions for SP-B in unfractionated tracheal aspirates from Patients 2-5 using antisera directed against mature SP-B, proSP-B (Chemicon #AB3430), N-terminal proSP-B, and C-terminal proSP-B. In addition to mature SP-B, peptide fragments specific to the N- and C-terminal portions of proSP-B are present. Although it is difficult to determine exactly, the N- and C-terminal fragments likely correspond to the fragments represented by DPLPDPLLDK and SPTGEWLPR, respectively, identified with mass spectrometry.



Supplementary Figure 3. Measurement of the tracer:tracee ratio from synthetic peptide standards of known enrichment. The peptide SPTGEWLPR was synthesized with endogenous isotopes and enriched with $^2\text{H}_3$ -leucine. The standards were mixed in known amounts on two different weeks to create two standard enrichment curves (A and B). The tracer:tracee ratio was calculated from the mass spectrometry data as described in the methods and plotted against the expected mole ratio of the standards. Points represent mean values ($n=5$); error bars are \pm SD. Standard curve A had a regression that resulted in a linear equation of $Y = (0.8645 \pm 0.0728) * X + (0.0034 \pm 0.0022)$ $r^2 = 0.9794$. Standard curve B had a linear equation of $Y =$

$(0.8966 \pm 0.0306) * X + (0.0005 \pm 0.0009) \quad r^2 = 0.9965$. The sloped deviation from unity is probably a result of differences between the amount of peptide actually aliquoted by the peptide synthesis company and claimed. Note: these curves were only used to demonstrate a linear response and not to calibrate the instrument measurement because the source of the labeled amino acid in these peptides was different from that used as the tracer infusate.



Supplementary Figure 4: Model illustrating the forms accounted for by the measurement of individual tryptic peptides mapping to SP-B. The peptide DPLPDPLLDK sits on the N-terminus of the protein, the peptide FPIPLPYCWLCR is a partially tryptic peptide sitting on the N-terminal end of the mature SP-B peptide. The peptide SPTGEWLPR is present on the C-terminal portion of the proSP-B. The C-terminal peptide could have been derived from isoforms containing the N-terminus through to the C-terminus or following the cleavage and removal of the N-terminal portion of the protein. Given the Western blot data, the high abundance of the peptide SPTGEWLPR and the observation that the peptide DPLPDPLLDK is present at ~100x lower abundance, we assume that a majority of the SP-B in these tracheal aspirates are missing the N-terminus but contain the mature and C-terminal domains of the protein.

Supplementary Table 1: Measurements of SPTGEW(²H₃-L)PR enrichment (MPE) from tracheal aspirates.

Time points	Replicate 1	Replicate 2	Replicate 3	Replicate 4	Replicate 5	Average	Stdev
<i>Patient 1</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
3.00	0.9808	0.9997	0.9549	1.1328	0.8221	0.9781	0.1110
6.00	2.3979	2.5389	1.9522	2.6381	2.1924	2.3439	0.2756
18.00	2.8447	2.4048	2.9990	3.0429	2.5727	2.7728	0.2760
34.00	1.0186	0.5236	0.9452	0.8877	1.1714	0.9093	0.2404
48.00	1.2920	1.1658	1.2514	1.2543	1.3541	1.2635	0.0685
62.00	0.9936	0.8598	0.9814	0.8081	0.8167	0.8919	0.0895
75.00	0.8119	0.9616	0.9241	1.0353	1.0548	0.9575	0.0973
96.00	0.9129	1.0655	0.8451	1.0860	0.9148	0.9649	0.1053
<i>Patient 2</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
22.67	1.6405	1.7925	2.0718	1.6721	1.6248	1.7604	0.1861
46.41	1.4410	1.3411	1.1796	1.2926	1.6118	1.3732	0.1632
59.50	-0.0540	0.1958	-0.1370	0.2116	0.3542	0.1141	0.2032
74.50	-0.0742	0.0284	-0.1181	-0.1034	-0.2546	-0.1044	0.1016
80.50	0.1448	0.4845	0.3899	0.5022		0.3804	0.1646
<i>Patient 3</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
5.00	2.9273	2.5827	3.2303	3.1076		2.9620	0.2818
20.16	1.8593	1.9837	1.9189	1.4808	2.0403	1.8566	0.2208
26.00	1.1630	0.9069	1.4093	1.1087	0.7614	1.0699	0.2485
47.00	1.0472	1.4619	1.1338	0.8738	0.6072	1.0248	0.3165
56.00	0.3933	0.6851	0.6536	0.6321	0.2328	0.5194	0.1976
65.00	1.4180	1.3631	1.3179	1.6023	1.3142	1.4031	0.1190
<i>Patient 4</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
2.25	-0.1193	-0.0769	-0.3065	-0.2361	-0.3426	-0.2163	0.1154
10.25	2.7841	2.7953	2.3864	2.6503	2.4448	2.6122	0.1894
26.00	1.0613	1.0336	1.0848	0.8987	0.9539	1.0064	0.0779
34.16	0.7612	0.9048	0.8753	0.8219	0.8717	0.8470	0.0565
<i>Patient 5</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
0.25	1.3437	1.2696	1.0705	0.7560	0.8867	1.0653	0.2484
2.41	1.0560	0.8883	0.8768	0.8763	0.6616	0.8718	0.1400
4.25	1.4183	1.0619	1.0810	1.3321	1.0805	1.1948	0.1677
10.50	2.2442	2.4026	2.0539	2.4736	1.7895	2.1928	0.2773
21.71	1.7388	1.9192	1.8910	2.1019	2.3195	1.9941	0.2230
22.00	2.2496	2.3441	2.6114	1.8388	2.1507	2.2389	0.2819
25.25	1.9701	2.0194	1.7640	1.6456	1.7687	1.8335	0.1561
<i>Patient 6</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
5.00	0.6934		0.6287	0.9227		0.7483	0.1545
13.00	2.1687	2.2543	2.3203	2.1814	2.1097	2.2069	0.0816
17.25	2.5211	2.3957	2.4627	2.1155	2.5770	2.4144	0.1802
21.00	1.9733	2.1946	2.0217	1.9704	2.0662	2.0453	0.0923
25.00	0.5565	0.4092	0.4734			0.4797	0.0738
33.02	-0.0567	0.4202	-0.2338	0.0706	0.0120	0.0425	0.2401
41.00	0.7698	0.1794	0.4413	0.0656	0.2001	0.3312	0.2808
49.00	0.3703	0.1702	0.5504	0.4044	0.3741	0.3739	0.1356
<i>Patient 7</i>							

0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
4.00	2.0529	1.6870	1.7796	2.9297	3.8890	2.4676	0.9339
9.00	2.9475	3.0426	2.8564	2.6300	2.5801	2.8113	0.2002
11.00	3.2057	2.2584	2.5355	3.1022	3.0674	2.8338	0.4138
18.00	2.6720	2.6406	2.5080	2.3227	2.5668	2.5420	0.1383
<i>Patient 8</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
1.00	0.1841	0.1200	0.2617	0.2168		0.1957	0.0596
4.00	0.6378	0.5849	0.6542	0.5839		0.6152	0.0362
7.00	1.0182	1.0057	0.9729	0.8872		0.9710	0.0590
10.02	1.3461	1.4360	1.3666	1.4297		1.3946	0.0450
16.00	1.1910	1.1434	1.1393	1.1964		1.1675	0.0304
22.00	1.0116	1.0031	0.9582	0.9893		0.9906	0.0235
25.00	0.9063	0.8212	0.8125	0.8749	0.8371	0.8504	0.0393
28.00	0.6689	0.7062	0.6395	0.7861	0.7583	0.7118	0.0608
31.00	0.8077	0.6292	0.6888	0.8155	0.7517	0.7386	0.0795
34.50	0.6189	0.5106	0.6707	0.5849		0.5963	0.0671
43.00	0.4298	0.4658	0.4872	0.5453	0.4937	0.4843	0.0422
<i>Patient 9</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
1.00	0.0959	0.0564	-0.0262	0.0410	-0.2082	-0.0082	0.1201
4.00	0.4396	0.4035	0.2484	0.2067	0.3418	0.3280	0.0993
6.00	1.1307	1.1496	0.7979	0.5900		0.9170	0.2713
13.00	1.5349	1.8093	1.6798	1.6461	1.8273	1.6995	0.1212
16.00	1.4609	1.3110	1.3151	1.4247		1.3779	0.0764
18.00	1.4103	1.0983	0.8920	1.0708	0.9579	1.0859	0.1997
24.00	1.1046	1.1774	1.2278	1.2473	1.1531	1.1820	0.0575
27.00	0.7242	0.7041	0.9066	0.7955	0.8735	0.8008	0.0891
<i>Patient 10</i>							
0.00	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
2.08	1.1385	0.7458	1.0510	0.7952	0.5880	0.8637	0.2265
3.50	1.6143	1.2535	1.4958	1.7270	0.8394	1.3860	0.3524
10.50	3.5804	4.0821	2.7929	3.5704		3.5064	0.5323
15.00	2.6551	2.6935	2.5427	2.8746	2.4695	2.6471	0.1553
16.08	2.8007	2.6093	2.6102	2.3610		2.5953	0.1803
17.75	2.1434	2.0672	2.1659	2.2845	2.2505	2.1823	0.0868
18.75	1.9054	1.8123	1.6647	2.0308	1.6830	1.8192	0.1539
23.92	1.6655	1.5337	1.4117	1.4390	0.4938	1.3087	0.4663
28.50	1.1952	1.0741	0.9095			1.0596	0.1434
40.00	0.9843	0.9125	0.9732	0.9788	1.0152	0.9728	0.0374
46.00	1.0107	0.9103	0.9095	0.7807		0.9028	0.0942

Supplementary Table 2: Measurement of plasma $^2\text{H}_3\text{-KIC}$ enrichment from a subset of the patients in this study.

	Hours	$^2\text{H}_3\text{-KIC}$ TTR	$^2\text{H}_3\text{-KIC}$ MPE
<i>Patient 2</i>			
	0	0.0000	0.00
	4.5	0.1114	10.02
<i>Patient 3</i>			
	0	0.0000	0.00
	3	0.0634	5.96
<i>Patient 4</i>			
	0	0.0000	0.00
	2.167	0.0772	7.17
<i>Patient 5</i>			
	0	0.0000	0.00
	2	0.0559	5.29
<i>Patient 8</i>			
	0	0.0000	0.00
	2.5	0.0510	4.85