

Ultrasensitive single-cell proteomics workflow identifies >1000 protein groups per mammalian cell

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Supporting Information

Table S1. Protein groups identified from 0.5 ng of HeLa digest

Method	Protein groups			
	Replicate 1	Replicate 2	Replicate 3	Average
OTOT-HCD 2 FAIMS CVs	1498	1627	1633	1586
OTIT-CID 2 FAIMS CVs	1729	1844	1933	1835
OTIT-HCD 2 FAIMS CVs	2171	2009	2003	2061
OTIT-HCD 3 FAIMS CVs	1847	1937	1803	1862

Supporting Information

Table S2. Protein groups identified from single cells and blank sample analyzed by LC/MS

	Sample	Protein groups				Peptide groups			
		R1	R2	R3	Average	R1	R2	R3	Average
FAIMS	SH	1031	1156	980	1056	4064	4395	3276	3912
	SM	972	1192	1126	1097	3063	3595	3232	3297
	SI	989	1265	1322	1192	2978	3842	4195	3672
	Blank	149	200	224	191	320	443	524	429
No FAIMS	SH	532	430	418	460	2460	1814	1545	1940

Note: SH: single HeLa cell, SM: single motor neuron, SI: single interneuron; R1,2,3: replicates 1,2,3;
Blank: HeLa cell supernatant

Supporting Information

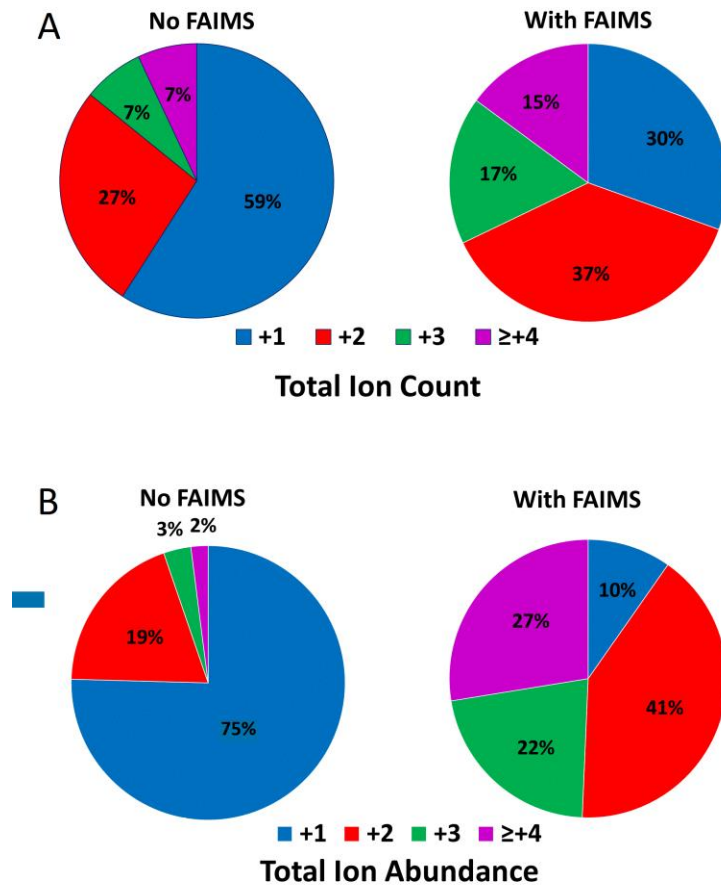


Figure S1 The percentage of detected charged ions from single HeLa cells with/without FAIMS Pro interface incorporated . A) Percentage change of charged ions count (species). B) Percentage change of charged ions abundance.

Supporting Information

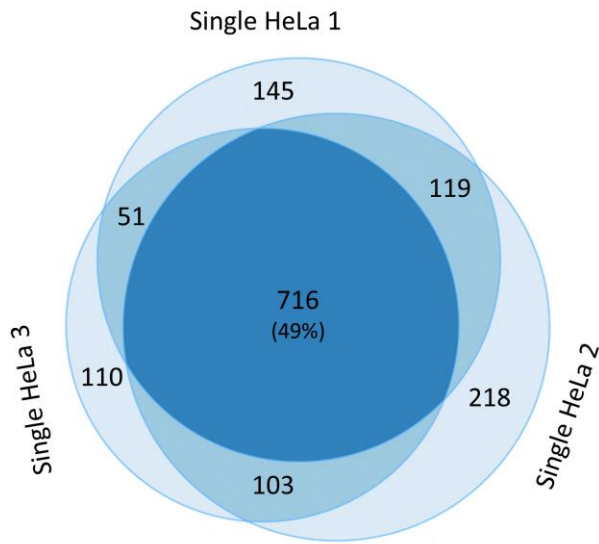


Figure S2 Venn diagram indicating overlap of identified protein groups among three replicate analyses of single HeLa cells.

Supporting Information

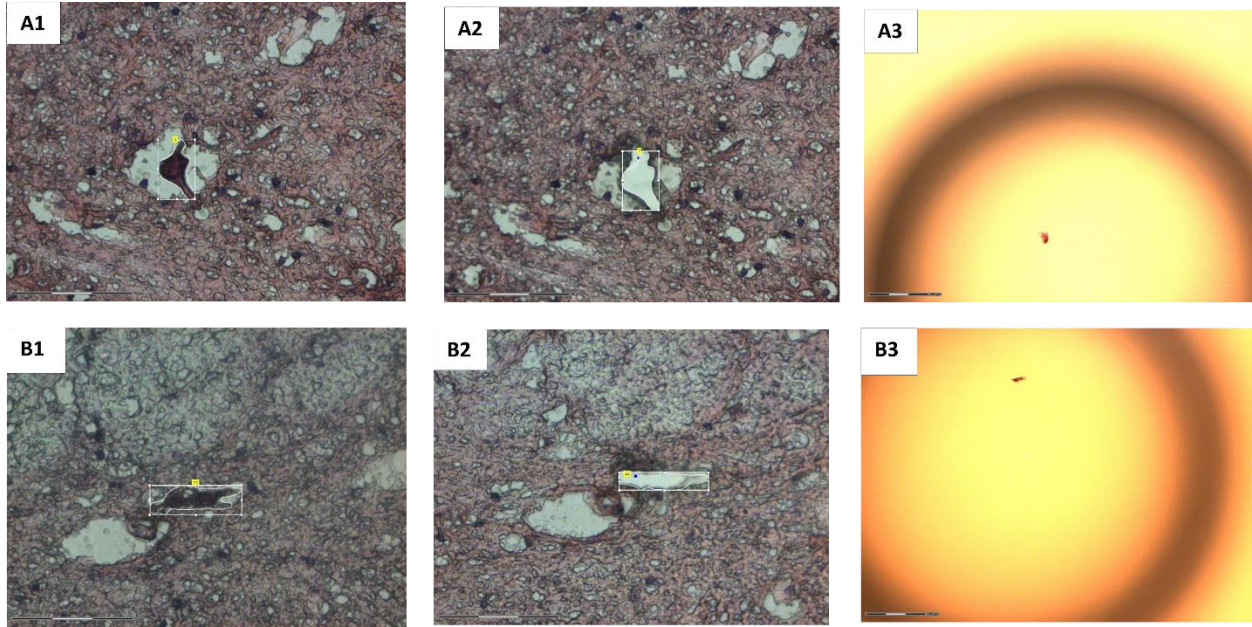


Figure S4. Representative images showing laser capture microdissection of single motor neurons and interneurons. A1 and A2 show H&E stained tissue before and after microdissection of a motor neuron. A3 shows the same motor neuron transferred to a nanowell. B1–B3 show corresponding images for an interneuron.