## **Supplemental Information**

## Spatial-CITE-seq: spatially resolved high-plex protein and whole transcriptome co-mapping

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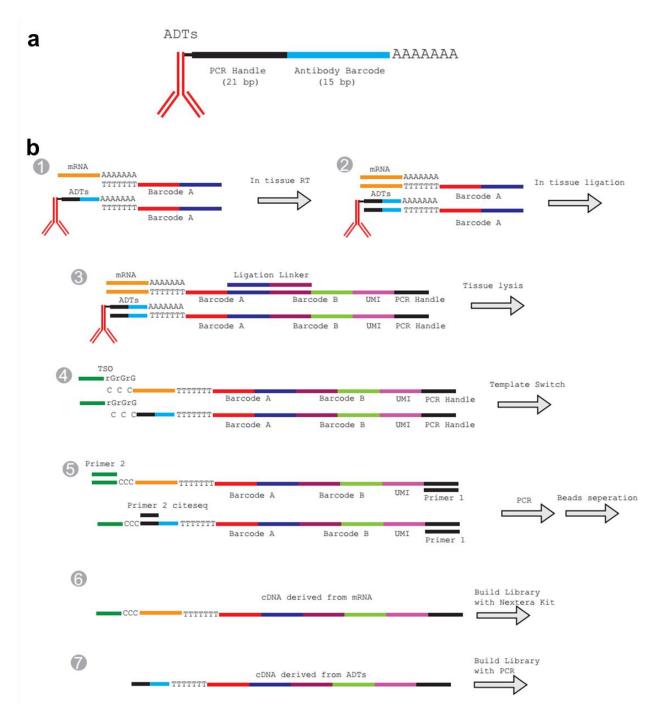
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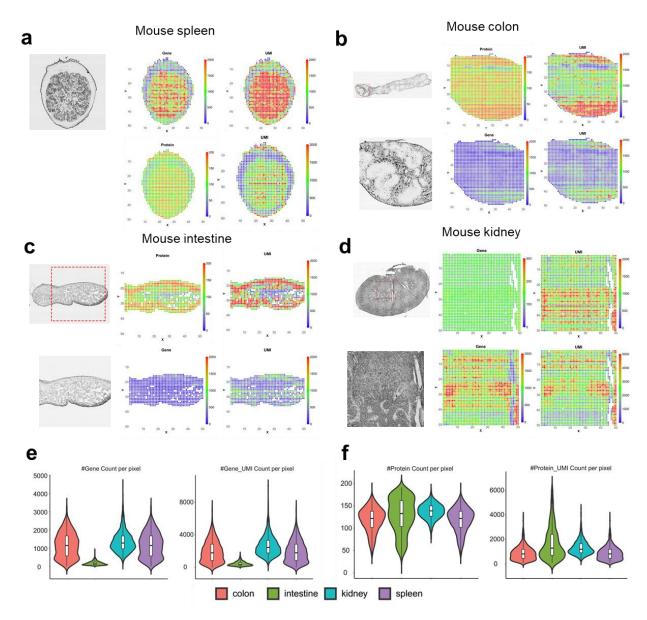
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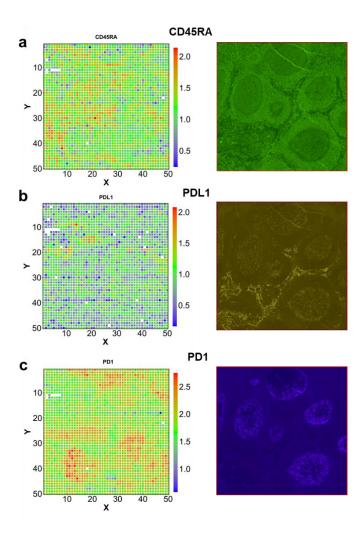


**Figure S1. Spatial-CITE-seq design and detailed workflow.** (a) ADT structure. The oligo labelled to the antibody has three functional regions: PCR handle (21 bp), antibody barcode (15 bp) and poly-A region (32 bp). (b) ADTs and mRNA with Poly-A region at the 3' end can be reverse transcribed into cDNA using Barcode A as the RT primer. Barcode A consists of three functional regions, the poly-T region, spatial barcode region and the ligation region. During the first flow, 50 Barcode As were loaded into 50 parallel channels and the RT reaction was carried out inside each isolated channel (Step 1&2). After peeling off the 1<sup>st</sup> PDMS, a 2<sup>nd</sup> PDMS was attached. The in-channel ligation was performed with injecting 50 Barcode Bs into each of the 50 channels which

are perpendicular to the channels of 1<sup>st</sup> PDMS chip (Step 3). Barcode B has four functional regions: ligation region, barcode region, UMI region and PCR handle region. Barcode B was also 5' biotin modification. After ligation, tissue was lysed, and cDNAs were purified with streptavidin beads. The cDNAs on the beads were templated switched with template switch oligo (Step 4). PCR was used to amplify the cDNA (Step 5). The products were split into two portions, the mRNA derived cDNAs and the ADT derived cDNAs. The library was then built separately. More details were in the method section.



**Figure S2. Spatial mapping of mouse spleen, colon, intestine and kidney with Spatial-CITE-seq.** A 189 antibodies cocktail was used for all four mouse samples. The bright field image, spatial gene heatmap, spatial gene UMI heatmap, spatial protein heatmap and spatial protein UMI heatmap of spleen (a), colon (b), intestine (c) and kidney (d). (e) gene and gene UMI count per pixel of all four mouse samples. (f) Protein and protein UMI count per pixel of all four mouse samples.



**Figure S3. Immunostaining validation of spatial protein profiles.** Sequential IF staining data using the FFeX technology by Lunaphore Technologies were compared side by side with Spatial-CITE-seq data. (a) CD45RA, (b) PDL1, (c) PD1. Note: the staining is not on exactly the same tonsil block.

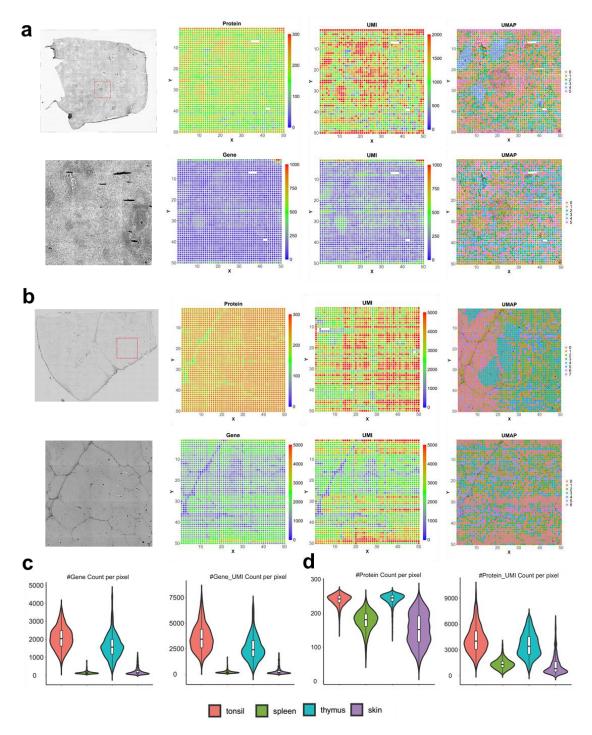


Figure S4. Spatial mapping of human spleen and thymus with Spatial-CITE-seq. A 273 antibodies cocktail was used for all four human samples. The bright field image, spatial gene heatmap, spatial gene UMI heatmap, spatial protein heatmap, spatial protein UMI heatmap, spatial clustering (based protein) and spatial clustering (based on RNA) of spleen (a) and thymus (b). (c) gene and gene UMI count per pixel of all four human samples. (d) Protein and protein UMI count per pixel of all four human samples.

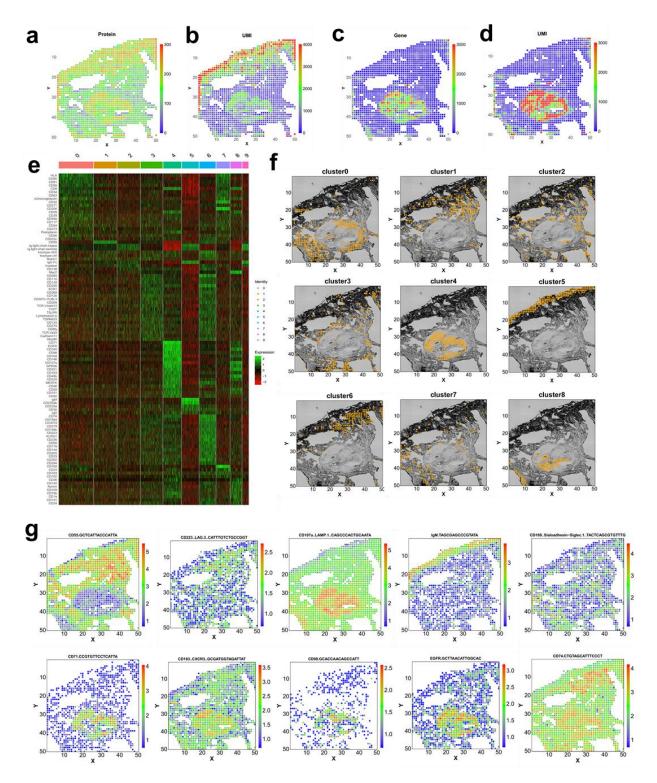
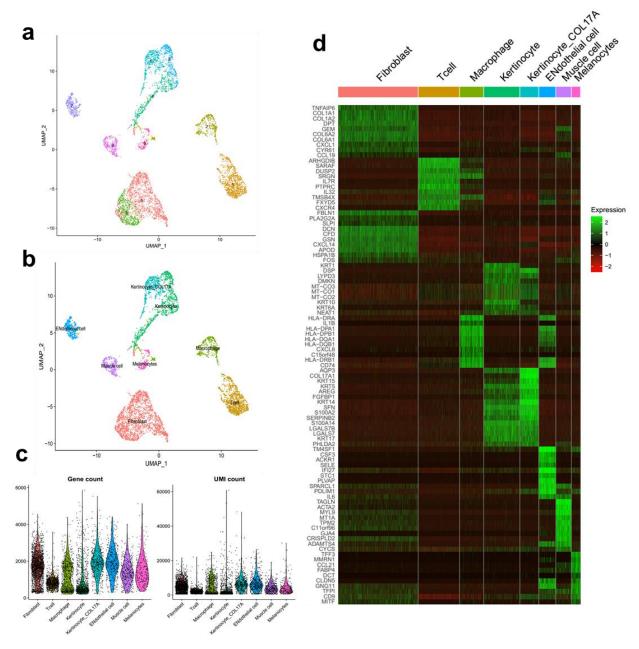


Figure S5. Spatial profiling of human skin biopsy tissue collected from the COVID-19 mRNA vaccine injection site. Spatial heatmap of gene (a), gene UMI (b), protein (c) and protein UMI (d). (e) Expression heatmap of the 10 clusters identified in skin biopsy sample. (f) the individual clusters plotted. (g) spatial distribution of some representative proteins.



**Figure S6. scRNA-seq sequencing data of skin biopsy sample.** (a) spatial clusters of scRNA-seq data. (b) annotated cell types using canonical marker genes. (c) violin plot of genes and UMIs for each cell type. (d) Expression heatmap of different cell types.

Table S1. Summary of gene and protein counts for all the samples sequenced.

Sample	# Useful Pixels	# Unique genes present	# Unique proteins present	Average # genes per pixel	Average # gene UMIs per pixel	Average # proteins per pixel	Average # protein  UMIs per pixel
Spleen (Mouse)	1303	19923	189	1166	1972	118	885
Colon (Mouse)	2037	19468	189	258	462	142	1213
Intestine (Mouse)	902	20444	189	172	447	129	1796
Kidney (Mouse)	2419	23750	189	1367	2675	137	1324
Tonsil (Human)	2492	28417	273	2079	3639	239	4309
Spleen (Human)	2494	20236	273	132	212	177	1403
Thymus (Human)	2500	28278	273	1647	2633	241	3700
Skin (Human)	1691	15486	273	411	815	153	1340

 Table S2. DNA oligos for PCR, ligation and library preparation.
 All Oligos were HPLC purified.

Oligo Name	Sequence
Primer 1	CAAGCGTTGGCTTCTCGCATCT
Primer 2	AAGCAGTGGTATCAACGCAGAGT
Primer 2-citeseq	CCTTGGCACCCGAGAATT*C*C
Ligation Linker	CGAATGCTCTGGCCTCTCAAGCACGTGGAT
Template Switch Oligo	AAGCAGTGGTATCAACGCAGAGTGAATrGrG+G
N501	AATGATACGGCGACCACCGAGATCTACACTAGATCGCTCGTCGGCAG CGTCAGATGTGTATAAGAGACAG
N501-citeseq	AATGATACGGCGACCACCGAGATCTACACTAGATCGCTCGTCGGCAG CGTCAGATGTGTATAAGAGACAGCCTTGGCACCCGAGAATTCCA
N701	CAAGCAGAAGACGGCATACGAGATTCGCCTTAGTCTCGTGGGCTCGG AGATGTGTATAAGAGACAGCAAGCGTTGGCTTCTCGCATCT
N702	CAAGCAGAAGACGGCATACGAGATCTAGTACGGTCTCGTGGGCTCGG AGATGTGTATAAGAGACAGCAAGCGTTGGCTTCTCGCATCT
N703	CAAGCAGAAGACGGCATACGAGATTTCTGCCTGTCTCGTGGGCTCGG AGATGTGTATAAGAGACAGCAAGCGTTGGCTTCTCGCATCT
N704	CAAGCAGAAGACGGCATACGAGATGCTCAGGAGTCTCGTGGGCTCGG AGATGTGTATAAGAGACAGCAAGCGTTGGCTTCTCGCATCT
N705	CAAGCAGAAGACGGCATACGAGATAGGAGTCCGTCTCGTGGGCTCGG AGATGTGTATAAGAGACAGCAAGCGTTGGCTTCTCGCATCT

Table S3. The list of 50 DNA Barcode As and 50 Barcode Bs.

Barcode A (RT primer in 1st PDMS chip)	Sequence
Barcode A-1	/5Phos/AGGCCAGAGCATTCGAACGTGATTTTTTTTTTTTTVN
Barcode A-2	/5Phos/AGGCCAGAGCATTCGAAACATCGTTTTTTTTTTTTTVN
Barcode A-3	/5Phos/AGGCCAGAGCATTCGATGCCTAATTTTTTTTTTTTTVN
Barcode A-4	/5Phos/AGGCCAGAGCATTCGAGTGGTCATTTTTTTTTTTTVN
Barcode A-5	/5Phos/AGGCCAGAGCATTCGACCACTGTTTTTTTTTTTTTVN
Barcode A-6	/5Phos/AGGCCAGAGCATTCGACATTGGCTTTTTTTTTTTTVN
Barcode A-7	/5Phos/AGGCCAGAGCATTCGCAGATCTGTTTTTTTTTTTTVN
Barcode A-8	/5Phos/AGGCCAGAGCATTCGCATCAAGTTTTTTTTTTTTTVN
Barcode A-9	/5Phos/AGGCCAGAGCATTCGCGCTGATCTTTTTTTTTTTTVN
Barcode A-10	/5Phos/AGGCCAGAGCATTCGACAAGCTATTTTTTTTTTTTVN
Barcode A-11	/5Phos/AGGCCAGAGCATTCGCTGTAGCCTTTTTTTTTTTTTVN
Barcode A-12	/5Phos/AGGCCAGAGCATTCGAGTACAAGTTTTTTTTTTTTVN
Barcode A-13	/5Phos/AGGCCAGAGCATTCGAACAACCATTTTTTTTTTTTVN
Barcode A-14	/5Phos/AGGCCAGAGCATTCGAACCGAGATTTTTTTTTTTTVN
Barcode A-15	/5Phos/AGGCCAGAGCATTCGAACGCTTATTTTTTTTTTTTTVN
Barcode A-16	/5Phos/AGGCCAGAGCATTCGAAGACGGATTTTTTTTTTTTVN
Barcode A-17	/5Phos/AGGCCAGAGCATTCGAAGGTACATTTTTTTTTTTTVN
Barcode A-18	/5Phos/AGGCCAGAGCATTCGACACAGAATTTTTTTTTTTTVN
Barcode A-19	/5Phos/AGGCCAGAGCATTCGACAGCAGATTTTTTTTTTTTVN
Barcode A-20	/5Phos/AGGCCAGAGCATTCGACCTCCAATTTTTTTTTTTTVN
Barcode A-21	/5Phos/AGGCCAGAGCATTCGACGCTCGATTTTTTTTTTTTVN
Barcode A-22	/5Phos/AGGCCAGAGCATTCGACGTATCATTTTTTTTTTTTVN
Barcode A-23	/5Phos/AGGCCAGAGCATTCGACTATGCATTTTTTTTTTTTVN
Barcode A-24	/5Phos/AGGCCAGAGCATTCGAGAGTCAATTTTTTTTTTTTVN
Barcode A-25	/5Phos/AGGCCAGAGCATTCGAGATCGCATTTTTTTTTTTTVN
Barcode A-26	/5Phos/AGGCCAGAGCATTCGAGCAGGAATTTTTTTTTTTTVN
Barcode A-27	/5Phos/AGGCCAGAGCATTCGAGTCACTATTTTTTTTTTTTVN
Barcode A-28	/5Phos/AGGCCAGAGCATTCGATCCTGTATTTTTTTTTTTTVN
Barcode A-29	/5Phos/AGGCCAGAGCATTCGATTGAGGATTTTTTTTTTTTVN
Barcode A-30	/5Phos/AGGCCAGAGCATTCGCAACCACATTTTTTTTTTTTVN
Barcode A-31	/5Phos/AGGCCAGAGCATTCGGACTAGTATTTTTTTTTTTTVN
Barcode A-32	/5Phos/AGGCCAGAGCATTCGCAATGGAATTTTTTTTTTTTVN
Barcode A-33	/5Phos/AGGCCAGAGCATTCGCACTTCGATTTTTTTTTTTTVN
Barcode A-34	/5Phos/AGGCCAGAGCATTCGCAGCGTTATTTTTTTTTTTTVN
Barcode A-35	/5Phos/AGGCCAGAGCATTCGCATACCAATTTTTTTTTTTTVN
Barcode A-36	/5Phos/AGGCCAGAGCATTCGCCAGTTCATTTTTTTTTTTTVN
Barcode A-37	/5Phos/AGGCCAGAGCATTCGCCGAAGTATTTTTTTTTTTTVN
Barcode A-38	/5Phos/AGGCCAGAGCATTCGCCGTGAGATTTTTTTTTTTTVN

Barcode A-39	/5Phos/AGGCCAGAGCATTCGCCTCCTGATTTTTTTTTTTTVN
Barcode A-40	/5Phos/AGGCCAGAGCATTCGCGAACTTATTTTTTTTTTTTVN
Barcode A-41	/5Phos/AGGCCAGAGCATTCGCGACTGGATTTTTTTTTTTTTVN
Barcode A-42	/5Phos/AGGCCAGAGCATTCGCGCATACATTTTTTTTTTTTVN
Barcode A-43	/5Phos/AGGCCAGAGCATTCGCTCAATGATTTTTTTTTTTTTVN
Barcode A-44	/5Phos/AGGCCAGAGCATTCGCTGAGCCATTTTTTTTTTTTTVN
Barcode A-45	/5Phos/AGGCCAGAGCATTCGCTGGCATATTTTTTTTTTTTVN
Barcode A-46	/5Phos/AGGCCAGAGCATTCGGAATCTGATTTTTTTTTTTTVN
Barcode A-47	/5Phos/AGGCCAGAGCATTCGCAAGACTATTTTTTTTTTTTVN
Barcode A-48	/5Phos/AGGCCAGAGCATTCGGAGCTGAATTTTTTTTTTTTTVN
Barcode A-49	/5Phos/AGGCCAGAGCATTCGGATAGACATTTTTTTTTTTTTVN
Barcode A-50	/5Phos/AGGCCAGAGCATTCGGCCACATATTTTTTTTTTTTVN
Barcode B (Ligation to Barcode A in 2 <sup>nd</sup> PDMS chip)	Sequence
Barcode B-1	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAACGTGATATC CACGTGCTTGAG
Barcode B-2	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAAACATCGATC CACGTGCTTGAG
Barcode B-3	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNATGCCTAAATC CACGTGCTTGAG
Barcode B-4	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAGTGGTCAATC CACGTGCTTGAG
Barcode B-5	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNACCACTGTATC CACGTGCTTGAG
Barcode B-6	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNACATTGGCATC CACGTGCTTGAG
Barcode B-7	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNCAGATCTGATC CACGTGCTTGAG
Barcode B-8	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNCATCAAGTATC CACGTGCTTGAG
Barcode B-9	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNCGCTGATCATC CACGTGCTTGAG
Barcode B-10	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNACAAGCTAATC CACGTGCTTGAG
Barcode B-11	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNCTGTAGCCATC CACGTGCTTGAG
Barcode B-12	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAGTACAAGATC CACGTGCTTGAG
Barcode B-13	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAACAACCAATC CACGTGCTTGAG
Barcode B-14	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAACCGAGAATC CACGTGCTTGAG
Barcode B-15	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAACGCTTAATC CACGTGCTTGAG
Barcode B-16	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAAGACGGAATC CACGTGCTTGAG
Barcode B-17	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAAGGTACAATC CACGTGCTTGAG
Barcode B-18	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNACACAGAAATC CACGTGCTTGAG

Barcode B-19	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNACAGCAGAATC CACGTGCTTGAG
Barcode B-20	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNACCTCCAAATC CACGTGCTTGAG
Barcode B-21	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNACGCTCGAATC CACGTGCTTGAG
Barcode B-22	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNACGTATCAATC CACGTGCTTGAG
Barcode B-23	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNACTATGCAATC CACGTGCTTGAG
Barcode B-24	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAGAGTCAAATC CACGTGCTTGAG
Barcode B-25	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNAGATCGCAATC CACGTGCTTGAG
Barcode B-26	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAGCAGGAAATC CACGTGCTTGAG
Barcode B-27	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNAGTCACTAATC CACGTGCTTGAG
Barcode B-28	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNNATCCTGTAATCC ACGTGCTTGAG
Barcode B-29	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNNATTGAGGAATC CACGTGCTTGAG
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Barcode B-35	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNCATACCAAATC CACGTGCTTGAG
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Barcode B-50	/5Biosg/CAAGCGTTGGCTTCTCGCATCTNNNNNNNNNNNNGCCACATAATC CACGTGCTTGAG

Table S4. Chemicals and reagents used.

Name	Catlog Number	Vender	
Maxima H Minus	EP7051	Thermo Fisher	
dNTP mix	R0192	Thermo Fisher	
RNase Inhibitor	Y9240L	Enzymatics	
SUPERase• In™ RNase Inhibitor	AM2694	Thermo Fisher	
T4 DNA Ligase	M0202L	New England Biolabs	
Ampure XP beads	A63880	Beckman Coulter	
Dynabeads MyOne C1	65001	Thermo Fisher	
	65002		
Nextera XT DNA Preparation Kit	FC-131-1024	Illumina	
Kapa Hotstart HiFi ReadyMix	KK2601	Kapa Biosystems	
Proteinase K, recombinant, PCR grade	EO0491	Thermo Fisher	
RNase free water	10977015	Invitrogen	
Ethanol	187380-4L	Sigma	
Formaldehyde solution	F8775-25ML	Sigma	
Triton X-100	T8787-100ML	Sigma	
NEBuffer 3.1	B7203S	New England Biolabs	
T4 DNA Ligase Reaction Buffer	B0202S	New England Biolabs	
Binding and washing	15568025	Thermo Fisher	
(B&W) Buffer (2x)	AM9261	Thermo Fisher	
	AM9760G	Thermo Fisher	
Tween 20	3005	Thermo Fisher	

Table S5. ADT lists for human and mouse. See attached Excel Spreadsheet - Table S5.