

Table S7. The SLC52A2⁺NOTCH1⁺ SRCC isolated by different techniques

Related to Fig. S4I

The SLC52A2 ⁺ NOTCH1 ⁺ cells via the WhichCells module		the SLC52A2 ⁺ NOTCH1 ⁺ population via the modified clustering with marker gain		The SLC52A2 ⁺ NOTCH1 ⁺ CD82 ⁺ cells within the SLC52A2 ⁺ NOTCH1 ⁺ compartment determined by the modified clustering with marker	
No.	Cell ID	No.	Cell ID	No.	Cell ID
1	AAATGCCAGAGCCCAA_NS_12	1	AAATGCCAGAGCCCAA_NS_12	1	AAATGCCAGAGCCCAA_NS_12
2	AACCGCGTCTAGAGTC_NS_12	2	AACCGCGTCTAGAGTC_NS_12	2	AACCGCGTCTAGAGTC_NS_12
3	AAGACCTAGCATCATC_NS_12	3	AAGACCTAGCATCATC_NS_12	3	AAGACCTAGCATCATC_NS_12
4	ACACCGGCAGCCAATT_NS_07	4	ACACCGGCAGCCAATT_NS_07	4	ACACCGGCAGCCAATT_NS_07
5	ACAGCTAGTATCTGCA_NS_12	5	ACAGCTAGTATCTGCA_NS_12	5	ACAGCTAGTATCTGCA_NS_12
6	ACAGCTAGTGGTCCGT_NS_12	6	ACAGCTAGTGGTCCGT_NS_12	6	ACAGCTAGTGGTCCGT_NS_12
7	ACATCAGGTGGCGAAT_NS_07	7	ACATCAGGTGGCGAAT_NS_07	7	ACATCAGGTGGCGAAT_NS_07
8	ACATGGTTCGTGTAGT_NS_12	8	ACATGGTTCGTGTAGT_NS_12	8	ACATGGTTCGTGTAGT_NS_12
9	ACCTTTAGTCGCTTTC_NS_12	9	ACCTTTAGTCGCTTTC_NS_12	9	ACCTTTAGTCGCTTTC_NS_12
10	ACGATACCAAAGGCGT_NS_12	10	ACGATACCAAAGGCGT_NS_12	10	ACGATACCAAAGGCGT_NS_12
11	ACGATGTTACCCCTCA_NS_12	11	ACGATGTTACCCCTCA_NS_12	11	ACGATGTTACCCCTCA_NS_12
12	ACGCCGAAGCGCTCCA_NS_12	12	ACGCCGAAGCGCTCCA_NS_12	12	ACGCCGAAGCGCTCCA_NS_12
13	ACTGAACAGTGACATA_NS_12	13	ACTGAACAGTGACATA_NS_12	13	ACTGAACAGTGACATA_NS_12
14	ACTGAACCAGACAAGC_NS_12	14	ACTGAACCAGACAAGC_NS_12	14	ACTGAACCAGACAAGC_NS_12
15	ACTGAACGTGTGACGA_NS_12	15	ACTGAACGTGTGACGA_NS_12	15	ACTGAACGTGTGACGA_NS_12
16	ACTGAACTACCGGGT_NS_07	16	ACTGAACTACCGGGT_NS_07	16	ACTGAACTACCGGGT_NS_07
17	ACTGAGTGTATTAGCC_NS_12	17	ACTGAGTGTATTAGCC_NS_12	17	ACTGAGTGTATTAGCC_NS_12
18	ACTGAGTTCCCGACTT_NS_12	18	ACTGAGTTCCCGACTT_NS_12	18	ACTGAGTTCCCGACTT_NS_12
19	ACTGCTCTCAATACCG_NS_12	19	ACTGCTCTCAATACCG_NS_12	19	ACTGCTCTCAATACCG_NS_12
20	ACTGTCCCATAGGATA_NS_12	20	ACTGTCCCATAGGATA_NS_12	20	ACTGTCCCATAGGATA_NS_12
21	AGACGTTTCTTGCAAT_NS_12	21	AGACGTTTCTTGCAAT_NS_12	21	AGACGTTTCTTGCAAT_NS_12
22	AGATCTGAGACAGACC_NS_07	22	AGATCTGAGACAGACC_NS_07	22	AGATCTGAGACAGACC_NS_07
23	AGATCTGGTGTGTGCC_NS_12	23	AGATCTGGTGTGTGCC_NS_12	23	AGATCTGGTGTGTGCC_NS_12
24	AGCCTAAAGAAGGTGA_NS_12	24	AGCCTAAAGAAGGTGA_NS_12	24	AGCCTAAAGAAGGTGA_NS_12
25	AGCTCCTCAATCACAC_NS_12	25	AGCTCCTCAATCACAC_NS_12	25	AGCTCCTCAATCACAC_NS_12
26	AGCTCTCAGTCGAGTG_NS_07	26	AGCTCTCAGTCGAGTG_NS_07	26	AGCTCTCAGTCGAGTG_NS_07
27	AGGGAGTCACGCCAGT_NS_12	27	AGGGAGTCACGCCAGT_NS_12	27	AGGGAGTCACGCCAGT_NS_12
28	AGGGATGCATCGGGTC_NS_07	28	AGGGATGCATCGGGTC_NS_07	28	AGGGATGCATCGGGTC_NS_07
29	AGGGATGGTAAAGTCA_NS_12	29	AGGGATGGTAAAGTCA_NS_12	29	AGGGATGGTAAAGTCA_NS_12
30	AGGGTGAAGAACAATC_NS_12	30	AGGGTGAAGAACAATC_NS_12	30	AGGGTGAAGAACAATC_NS_12
31	AGGTCATCACACTGCG_NS_12	31	AGGTCATCACACTGCG_NS_12	31	AGGTCATCACACTGCG_NS_12
32	ATAACGCAGCCCAACC_NS_12	32	ATAACGCAGCCCAACC_NS_12	32	ATAACGCAGCCCAACC_NS_12
33	ATAACGCAGTGTACCT_NS_12	33	ATAACGCAGTGTACCT_NS_12	33	ATAACGCAGTGTACCT_NS_12
34	ATCATCTGTAGCCTCG_NS_12	34	ATCATCTGTAGCCTCG_NS_12	34	ATCATCTGTAGCCTCG_NS_12
35	ATCTACTTCGTGGTCG_NS_12	35	ATCTACTTCGTGGTCG_NS_12	35	ATCTACTTCGTGGTCG_NS_12
36	ATCTGCCAGGACAGAA_NS_12	36	ATCTGCCAGGACAGAA_NS_12	36	ATCTGCCAGGACAGAA_NS_12
37	ATCGGATGTCCGAACC_NS_12	37	ATCGGATGTCCGAACC_NS_12	37	ATCGGATGTCCGAACC_NS_12

38 ATTACTCGTGTGAGG_NS_12
39 CAAGTTGGTCACCCAG_NS_12
40 CACAAACCAATGCCAT_NS_12
41 CACACCTCAGCTGTGA_NS_12
42 CACACCTCATGTTGAC_NS_12
43 CACACCTTCGGCCGAT_NS_12
44 CAGCATAGTCATATCG_NS_12
45 CAGCCGAGTCTAAAGA_NS_12
46 CAGCGACTCCTTAATC_NS_12
47 CAGTAACTCCGCGGTA_NS_07
48 CATCAAGCATTACCTT_NS_12
49 CATCAAGTCCTTGACC_NS_12
50 CATCGGGGTAACGCGA_NS_12
51 CATTGCGCAACTGGCC_NS_07
52 CCACTACCATGGAATA_NS_12
53 CCATGTCCATTGACA_NS_12
54 CCATTGCGACAGCCCA_NS_12
55 CCCAGTTCAGTGAGTG_NS_07
56 CCCAGTTTCATGCTCC_NS_12
57 CCGGTAGGTTGGACCC_NS_12
58 CCGTTCACACATGGGA_NS_12
59 CCGTTCAGTACAAGTA_NS_12
60 CCTAAAGGTACATCCA_NS_12
61 CCTAAAGGTGAACCTT_NS_12
62 CCTATTAAGCTAGCCC_NS_07
63 CCTCAGTCAATCACAC_NS_12
64 CCTCTGACAGGCTCAC_NS_12
65 CCTTACGTCGTTTGCC_NS_12
66 CCTTCGAAGACAGACC_NS_12
67 CCTTTCTGTGATAAGT_NS_12
68 CGAACATGTAAGCACG_NS_12
69 CGATCGGAGACGCACA_NS_12
70 CGCCAAGCAACGATGG_NS_12
71 CGCGTTTGTAAGTCAAC_NS_12
72 CGGACACGTAGAAGGA_NS_07
73 CGTAGGCCACAGACTT_NS_07
74 CTAATGGAGGAATGGA_NS_12
75 CTAATGGGTAGAGGAA_NS_07
76 CTACACCGTAAACACA_NS_12
77 CTACATTTCCACTCCA_NS_12
78 CTACATTTTCGTGGGAA_NS_12
79 CTACGTCGTTTGACTG_NS_12
80 CTCATTAGTGCAGTAG_NS_12
81 CTCGAAACATACTACG_NS_07
82 CTCGAGGTCCACTCCA_NS_12
83 CTCGTACCAAGCGAGT_NS_07
84 CTCGTCAGTAAAGTCA_NS_07
85 CTGAAACGTAGGCATG_NS_07
86 CTGCCTATCCTAGGGC_NS_12
87 CTGCTGTAGGGTCTCC_NS_07

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43 CACACCTTCGGCCGAT_NS_12
44 CAGCATAGTCATATCG_NS_12
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46 CAGCGACTCCTTAATC_NS_12
47 CAGTAACTCCGCGGTA_NS_07
48 CATCAAGCATTACCTT_NS_12
49 CATCAAGTCCTTGACC_NS_12
50 CATCGGGGTAACGCGA_NS_12
51 CATTGCGCAACTGGCC_NS_07
52 CCACTACCATGGAATA_NS_12
53 CCATGTCCATTGACA_NS_12
54 CCATTGCGACAGCCCA_NS_12
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58 CCGTTCACACATGGGA_NS_12
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60 CCTAAAGGTACATCCA_NS_12
61 CCTAAAGGTGAACCTT_NS_12
62 CCTATTAAGCTAGCCC_NS_07
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64 CCTCTGACAGGCTCAC_NS_12
65 CCTTACGTCGTTTGCC_NS_12
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67 CCTTTCTGTGATAAGT_NS_12
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69 CGATCGGAGACGCACA_NS_12
70 CGCCAAGCAACGATGG_NS_12
71 CGCGTTTGTAAGTCAAC_NS_12
72 CGGACACGTAGAAGGA_NS_07
73 CGTAGGCCACAGACTT_NS_07
74 CTAATGGAGGAATGGA_NS_12
75 CTAATGGGTAGAGGAA_NS_07
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85 CTGAAACGTAGGCATG_NS_07
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87 CTGCTGTAGGGTCTCC_NS_07

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46 CAGCGACTCCTTAATC_NS_12
47 CAGTAACTCCGCGGTA_NS_07
48 CATCAAGCATTACCTT_NS_12
49 CATCAAGTCCTTGACC_NS_12
50 CATCGGGGTAACGCGA_NS_12
51 CATTGCGCAACTGGCC_NS_07
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72 CGGACACGTAGAAGGA_NS_07
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 128 TATCAGGCAGTAACGG_NS_12
 129 TATTACCTCGGCGCAT_NS_07
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 131 TCACGAAAGTCAGGACA_NS_12
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182 CATCAAGAGATCGATA_NS_07
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186 CATTGCGCTAGGTT_NS_07
187 CCGTACTGTCTAACGT_NS_07

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139 TGCCCATAGGCGTACA_NS_12
140 TGCGCAGAGATGTTAG_NS_12
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146 TGTGTTTCAAGTACCT_NS_07
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149 TTTACTGAGTCAAGCG_NS_12
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Footnote:

The cells highlighted in orange are the same