

S2 Fig. Genes specifically expressed between L-DC and CD8⁺ cDC.

Gene Symbol	Fold-Change (L-DC vs. CD8 ⁺ cDC) ⁺	L-DC		CD8 ⁺ cDC	
		Mean*	Standard Error	Mean*	Standard Error
<i>Gm10673</i>	7.387	272.591	56.193	36.702	7.355
<i>Fcgr3</i>	7.119	300.711	40.598	42.165	3.019
<i>Irak3</i>	6.566	264.136	22.644	40.625	5.823
<i>Map3k15</i>	6.537	247.350	9.595	37.823	0.491
<i>Hp</i>	6.532	182.405	36.805	27.146	1.271
<i>Tmem26</i>	6.344	244.988	38.620	38.677	4.480
<i>Mdm1</i>	6.070	260.629	15.696	43.076	3.419
<i>Apoc2</i>	6.045	240.037	10.652	39.798	3.288
<i>Fpr1</i>	5.936	169.278	11.785	28.583	2.002
<i>Thbd</i>	5.904	211.722	22.901	35.616	1.195
<i>Xylt1</i>	5.803	273.414	32.386	46.949	1.091
<i>Cd300lb</i>	5.783	218.947	26.248	38.000	3.319
<i>Svil</i>	5.765	256.716	16.560	44.520	1.425
<i>Trem1</i>	5.757	211.974	20.345	36.820	2.085
<i>Pros1</i>	5.710	179.988	20.266	31.242	0.695
<i>Ccde125</i>	5.227	197.052	15.631	37.781	2.532
<i>Gstm1</i>	5.092	212.010	16.306	41.796	6.312
<i>Lrrc16a</i>	-5.055	35.121	73.789	177.984	16.044
<i>Ptpn3</i>	-5.076	30.921	2.082	157.520	13.583
<i>Ptpn3</i>	-5.165	47.759	7.073	246.474	10.867
<i>Dennd3</i>	-5.320	37.311	4.596	199.139	16.767
<i>Ctnnd2</i>	-5.372	33.902	1.754	182.360	12.929
<i>Flnb</i>	-5.546	48.463	6.019	270.441	29.982
<i>Bub1</i>	-5.601	34.466	5.957	195.833	36.781
<i>Card11</i>	-5.850	36.686	2.685	214.530	18.356
<i>Tns1</i>	-6.049	32.084	2.670	194.485	12.789
<i>Mical3</i>	-6.091	46.421	5.960	282.759	12.329
<i>Clec9a</i>	-6.333	45.057	3.686	285.108	36.356
<i>Gpm6b</i>	-6.405	34.712	5.485	223.304	21.383
<i>Tbc1d4</i>	-6.438	42.156	3.074	272.840	28.020
<i>Siglecg</i>	-6.908	49.420	4.720	338.777	11.596
<i>St8sial1</i>	-7.381	29.334	8.984	217.773	23.448
<i>B4galt6</i>	-7.506	44.314	6.252	333.803	32.710
<i>Cdon</i>	-7.539	41.034	2.337	309.066	12.666
<i>Ciita</i>	-7.679	43.440	2.430	332.968	10.191
<i>Ptpn3</i>	-7.950	33.400	4.627	266.634	30.051
<i>I300002K09Rik</i>	-7.991	41.162	3.429	329.781	24.192

⁺ Genes were selected which showed between 5 to 8 fold change in signal value where one subset has signal value ≤ 50 and the comparison subset has a signal value ≥ 150 , in either L-DC or CD8⁺ cDC assessed in pairwise comparison.

* Data represent mean of duplicate samples (n=2).