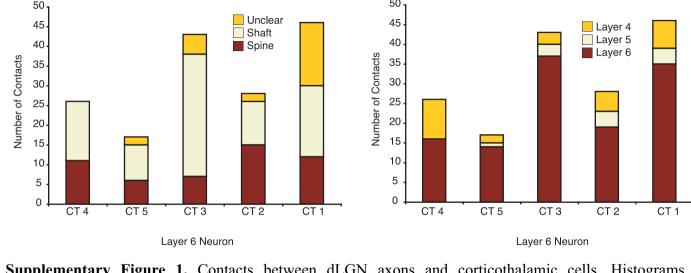
CT neuron	Cat2003 S23C1	Cat2304 - S30C1	Cat2403 S39C1	Cat1804 S50C1	Cat0105 S45C1
Apical Dendrite Length (μm)	3449	7213	4689	4747	4477
Basal Dendrites Length (μm)	3095	4131	3132	3364	2942
Mean Path Length to Termination (±std)[range](μm) (apical)	340 ± 206 (n = 18) [107 to 633]	614 ± 295 (n = 25) [127 to 1080]	386 ± 189 (n = 25) [73 to 885]	465 ± 247 (n = 22) [146 to 931]	402 ± 229 (n = 26) [100 to 843]
Mean Path Length to Termination (±std)[range](μm) (basals)	131 ± 39 (n = 25) [35 to 186]	185 ± 36 (n = 28) [96 to 258]	170 ± 36 (n = 20) [77 to 212]	160 ± 25 (n = 24) [97 to 199]	121 ± 24 (n = 27) [45 to 149]
Number of apical terminations	18	25	25	22	26
Number of Basal Terminations	25	28	20	24	27
Percentage of Dendrite formed by Terminal Segment	94.2%	81.4%	92.1%	87%	91.7%
Length of Dendrite In Layer 2/3 (μm)	0	347	128	134	62
Length of Dendrite In Layer 4 (μm)	692.3	2175	592	2360	95 ^(b) 1391 ^(c)
Length of Dendrite In Layer 5 (μm)	1059.9	2404	3190	2322	1373
Length of Dendrite In Layer 6 (μm)	4223.8	5765	3899	3077	3911

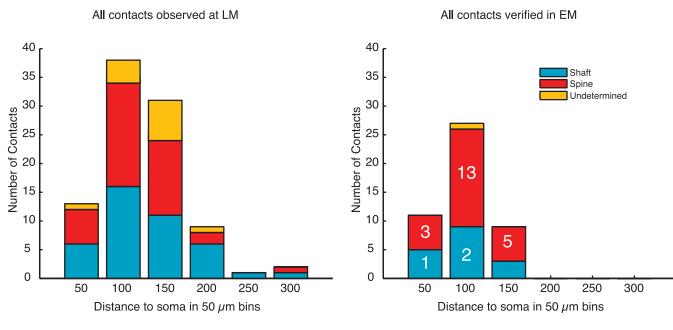
Supplementary Table 1. Topological and metric measure of the dendritic arborizations of the reconstructed layer 6 CT cells. Values were not corrected for shrinkage. (a) Neuron CT5 had a portion of its dendrite in a dense region of dLGN boutons and another portion in a less dense region, and so the dLGN bouton density was calculated separately. (b) Dense region of boutons. (c) Less dense region of bouton.



В

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Supplementary Figure 1. Contacts between dLGN axons and corticothalamic cells. Histograms summarizing the number of contacts made with each reconstructed cell. A. Number of contacts per target location on the dendrite. **B.** Number of contacts per cortical layer.



Supplementary Figure 2. Distribution off contacts with distance to the soma. Histograms showing the distribution of the number of contacts formed with basal dendrites with distance from the soma measured in bins of 50μm. The bars are color coded for the type of target. **A.** Histogram shows the distribution of all the contacts. **B.** Histogram shows the distribution of the contacts investigated in EM. White numbers inside the bar indicate the number of contacts that formed synapses.