

Figure 2c	IPSC Amplitude				
	dCre	Cre	Cre + Nrnx3a (-SS4)	Cre + Nrnx3a (LNS4-LNS6)	Cre + Nrnx3a (LNS1-LNS4)
	1.878397583	0.372478027	3.935572998	0.084747612	5.267785645
	5.794429199	0.794836426	3.807510986	0.194543289	1.907627197
	0.894480286	1.380685059	1.49530481	0.446087555	0.6237901
	5.245019531	5.650089844	4.662871582	0.920987366	4.714319824
		0.591476196	7.262024902	0.118982117	7.95752832
	4.927504395	0.032868874	4.601295898	0.219041168	1.520331665
	4.504977051	0.122413185	1.351060913	1.0060495	2.651093018
	1.323639771	0.288438599	7.58114502	0.724992615	0.709278564
	4.025512695	0.040847393	5.081791504	1.107179688	2.391909668
	9.749810547	0.198694687	0.528076721	0.321287048	6.24501123
	0.553428223	0.257778564		0.076472359	9.093652344
	1.923399048	0.725104797		0.502667694	3.482282471
	6.204882813	0.943583984		0.090415604	2.928464355
	4.113476074	5.684670898		4.848932129	8.100895996
	5.459125	2.19826123		0.018875826	2.741332275
Mean	4.042720158	1.285481851	4.030665533	0.712084105	4.022353511
Number (# cells)	14	15	10	15	15
Statistical Test	ANOVA with Dunnett's Multiple comparisons test				
Significance		to dCre, ** or 0.0047	ns	to dCre, *** or 0.0005	ns

Figure 2f	IPSC Amplitude				
	dCre	Cre	Cre + Nrnx3a (-SS4)	Cre + Nrnx3a (LNS2/3-only)	Cre + Nrnx3a (LNS2-only)
	0.785692322	0.387728516	1.122364136	5.786537109	4.743918457
	3.587024902	1.143088867	6.007285645	1.671965454	11.80621484
	9.452959961	4.14034668	1.61983728	9.397269531	4.07598877
	4.60894043	0.044391834	5.159187012	6.176229492	4.21179541
	7.380458496	0.020140263	2.636102783	1.918957275	3.855185059
	12.42824121	7.956678711	2.636219971	12.20743262	2.10840332
	1.617584229	0.181040451	10.22994629	1.535411133	6.234746094
	5.631896973	1.97139856	6.554281738	2.549123291	4.081328613
	5.50968457	0.753565491	7.999151855	3.142222656	7.790749512
	7.648881348	2.853336914	2.290608398	5.764934082	12.49784766
	7.243889648	2.134730225	3.952739502	2.064831055	5.605489746
	1.673033813	0.082260223	0.716276001	4.443207031	4.680748047
	3.737975342	0.168303391	6.779841309	5.776748535	3.65961499
	3.903456055	2.590674805	3.476205566	1.791719849	6.68042041
		0.53862561	4.490854004	8.86315918	
		0.445560425	4.730633301		
		5.383587891	12.25826758		
		1.393002686	1.725601196		
		0.598703308	6.535405273		
		0.125680862	1.734318115		
		4.091167725	0.943911194		
		0.189941406	8.049019531		
		1.30984729			
		1.825361938			
		2.967205322			
		0.407705048			
		0.494904663			
		0.279454132			
		0.317594727			
		0.238563858			
Mean	5.372122807	1.501153061	4.620366258	4.872649886	5.858032209
Number (# cells)	14	30	22	15	14
Statistical Test	ANOVA with Dunnett's Multiple comparisons test				
Significance		to dCre, *** or 0.0003	ns	ns	ns

Figure 2h	IPSC Amplitude				
	dCre	Cre	Cre + Nrnx3a (LNS2-only, -SS2)	Cre + Nrnx3a (LNS2-only, +SS2a)	Cre + Nrnx3a (LNS2-only, +SS2ab)
	5964.409668	4606.71	7361.821289	3639.122803	3097.981934
	4745.520508	5399.395	11019.78516	3819.175049	7974.344238
	11629.30176	6389.247	7548.390625	1165.312012	8083.233398
	5912.60791	4517.534	5775.598145	4328.794922	5331.562988
	6221.105957		6929.060059	6349.186523	6319.923828
	13302.37695	536.1362	11310.85352	499.245544	266.343872
	4464.449707	1430.944	4805.038574	8412.263672	174.491028
	4620.115723	4690.004	8723.996094	298.91629	59.955173
				1489.80542	7368.539063
				2932.223877	3609.56665
	2691.438721	174.148	2283.496094	506.367737	2053.677246
	5187.203613	308.1678	1365.148071	1636.432861	629.925537
				411.405151	1003.198242
				570.505127	2115.99707
					699.965393
Mean	6473.853052	3116.920667	6712.318763	2575.625499	3252.580377
Number (# cells)	10	9	10	14	15
Statistical Test	ANOVA with Dunnett's Multiple comparisons test				
Significance		ns, p = 0.0511	ns	** , p = 0.0078	* , p = 0.0314

Figure 6a	Ctrl	Dag1 gRNA
	100	39.78243606
	100	41.18625945
	100	31.15536592
Average Count (cultures)	100	37.37568714
Statistical Test	two-tailed, one sample t test	
Tvalue	19.97	
df	2	
Significance	**, p = 0.0025	

Figure 6c	Ctrl	Dag1 gRNA
	1.393716675	4.538832031
	2.448588967	0.374032013
	1.557436279	0.319219849
	6.824246094	1.741697021
	3.690708008	0.605207947
	1.576948242	0.570067383
	5.698196777	2.974362334
	2.13142041	1.538900391
	4.416081543	3.182121338
	4.263709961	0.683313416
	0.848329895	2.760336914
	1.689602783	1.110234009
	6.952677734	0.441584839
Average Count (cells)	3.498358713	1.603070183
Statistical Test	unpaired two-tailed T-test with Welch's correction	
Tvalue	2.439	
df	18.69	
Significance	*, p = 0.0249	

Figure 6e	dCre	dCre + Dag1 gRNA	Cre	Cre + Dag1 gRNA
	1.51827417	2.10462793	2.110289795	3.119438414
	3.787624268	0.245552733	0.368918573	4.967742188
		0.655594175	2.347226271	0.920279834
		0.498259551	1.312881528	1.540763447
		0.285121552	1.465384644	0.973251953
	5.331669922	0.73066095	8.572758859	3.184894531
	4.191088867	0.601045288	1.769262939	5.042607422
	5.6967334	0.25992604	2.769459473	1.875074219
	0.689091675	4.245503906	2.910873047	
	9.86595703		0.152849976	
	4.40634668	3.223001221	2.037180176	5.77991748
	9.017279297	2.340318848	1.687684937	4.436753906
	6.38684226	0.480786571	2.844393311	5.753077637
	5.571657715	4.732151367	6.844233368	0.659419676
	6.601068848	2.016201172	0.380742035	0.409260254
Average Count (cells)	5.256267894	1.603272836	2.505934517	2.977253643
Statistical Test	2-way ANOVA with Tukey's multiple comparisons test			
df	12	14	15	13
Significance	***, p = 0.0004 (to dCre) **, p = 0.0096 (to dCre) ns, p = 0.0521 (to dCre)			

Figure 6f	dCre	dCre + Dag1 gRNA	Cre	Cre + Dag1 gRNA
	0.07399437	0.152970875	0.105920813	0.189186156
	0.240058	0.078600055	0.023582816	0.343229219
		0.276714656	0.235149234	0.049740852
		0.084741336	0.083423867	0.083755261
		0.024394848	0.062349836	0.097094117
	1.113047	0.07453825	0.505035844	0.236606031
	0.2177181	0.060650957	0.077243258	0.44477025
	0.430611	0.008706488	0.203099038	0.255611781
	0.03278019	0.341865844	0.05429418	
	1.353603		0.013444664	
	0.3251208	0.328706344	0.149212828	0.57552875
	0.7693861	0.150327672	0.073906609	0.320148438
	0.3028154	0.027874043	0.138047297	0.482904125
	0.3496759	0.408728156	0.406146375	0.046886703
	0.274544	0.211028719	0.013951248	
Average Count (cells)	0.45668888	0.159989232	0.14300792	0.280872142
Statistical Test	2-way ANOVA with Tukey's multiple comparisons test			
df	12	14	15	12
Significance	*, p = 0.0123 (to dCre) **, p = 0.0063 (to dCre) ns, p = 4023 (to dCre)			

Figure 6h	NMDAR-EPSCs			
	dCre	dCre + Dag1 gRNA	Cre	Cre + Dag1 gRNA
	0.37112735	1.83625	0.813398804	0.076017822
	0.694535767	0.148152313	2.629472412	0.581522644
	0.816024912	0.905601609	0.617316675	1.580326359
	1.315511841	1.648802612	0.250404358	0.532439087
	0.057595222	0.062072693	0.078903465	0.045541519
	0.128634232	0.048939213	0.35528775	0.071145027
	0.096125938	0.093399841		0.185211075
	0.023359358	0.008191372	0.050013329	0.325438599
	0.167145096	0.008552102	0.015279064	0.11013488
	0.096614426	0.093797729	0.306264832	0.059980179
	0.075727943			
Average Count (cells)	0.345242008	0.487980698	0.568137854	0.455675592
Statistical Test	2-way ANOVA with Tukey's multiple comparisons test			
df	11	10	9	10
Significance	ns ns ns ns			

Figure 6i	AMPA-EPSCs			
	dCre	dCre + Dag1 gRNA	Cre	Cre + Dag1 gRNA
	2.358732666	2.915829346	0.583679503	0.979468201
	0.656214722	2.160535156	1.657297363	2.89213501
	1.105677812	1.045551147	5.009924805	1.786452637
	0.75789989	2.304394287	0.848229553	1.108194214
	0.741980042	0.805942688	0.534341003	1.164858887
	0.816252441	0.244198593	0.37730896	1.044040039
	0.30530304	0.283946869	0.644206543	0.468593994
	0.294131012	0.545852234	0.24519519	0.104488419
		0.32795343	1.767139771	1.128227783
	0.883407043	0.689507324	0.629792236	0.425990571
	0.634854187		0.313576599	
	0.922574707			
Average Count (cells)	0.84352976	1.132371107	1.164607993	1.108041976
Statistical Test	2-way ANOVA with Tukey's multiple comparisons test			
df	11	10	11	10
Significance	ns ns ns ns			

Newt 254 Inclusion			Newt 254 Inclusion			Newt 254 Inclusion 2		
Transect	Number 1	Number 2	Transect	Number 1	Number 2	Transect	Number 1	Number 2
OB	71.720875	73.444181	OB	88.203743	84.521521	OB	98.427009	98.250735
Ca	40.260268	41.138425	Ca	38.344878	38.800844	Ca	78.177099	80.814478
Car	84.370478	84.108957	Car	81.149241	82.130521	Car	89.051287	84.983787
BS	61.503871	60.940111	BS	61.143123	51.449078	BS	98.115148	98.205296
Bys	60.380007	61.203195	Bys	51.847466	53.670587	Bys	52.822365	36.47627
Hpp	40.82897	42.899145	Hpp	40.263763	41.150302	Hpp	55.622652	56.3281549

OB			Hpp			OB						
Newt 254	47.509941	47.730576	47.509941	Newt 254	33.792243	37.88884	36.777151	Newt 254	67.6337451	70.5328449	69.1133255	67.8828137
Newt 254	40.682049	43.159953	40.026177	Newt 254	26.2884525	36.4577755	37.5427854	Newt 254	71.324378	75.628915	69.8024798	71.1623393
Newt 254	78.8518467	78.410613	79.150171	Newt 254	48.990823	58.02328	84.860195	Newt 254	91.205928	94.807604	93.818761	93.548424

Cell #	Cell Type	Ni Transmitter	Cell #	% Total	Cell Type	Ni Transmitter	Cell #	% Total
1	GC-1 (a)	VGAT	1814	8.85182888	GC-1 (a)	VGAT	1814	18.28424993
2	GC-2 (b)	VGAT	3608	18.93252071	GC-2 (b)	VGAT	3608	34.49849483
3	GC-3 (c)	VGAT	273	1.25483249	GC-3 (c)	VGAT	273	2.83794802
4	GC-4 (d)	VGAT	24	0.11689999	GC-4 (d)	VGAT	24	2.25234428
5	GC-5 (e)	VGAT	1589	7.81452476	GC-5 (e)	VGAT	1589	15.94382881
6	GC-6 (f)	VGAT	237	1.09989505	GC-6 (f)	VGAT	237	2.28424727
7	GC-7 (g)	VGAT	979	4.72413917	GC-7 (g)	VGAT	979	9.48648463
8	PGC-1 (h)	VGAT	1037	4.94889302	PGC-1 (h)	VGAT	1037	9.96381351
9	PGC-2 (i)	VGAT	337	1.61788718	PGC-2 (i)	VGAT	337	3.18140244
10	PGC-3 (j)	VGAT	279	1.33928474	PGC-3 (j)	VGAT	279	2.68526571
11	EPN-2b (k)	VGAT	161	0.76739844	EPN-2b (k)	VGAT	161	1.53820263
12	MTC-1 (l)	vGLUT1	69	0.31788977	None			19.668
13	MTC-2 (m)	vGLUT2	116	0.53421438				
14	MTC-3 (n)	vGLUT3	67	0.31828228				
15	OSN (o)	nr	1200	5.81826631	% Total			82.288804
16	Trachea (p)	nr	2085	9.87922021	PGC			16.1289897
17	Immature (q)	nr	3810	17.8951922	EPN-1 (r)			1.53820263
18	Anterior-Like (s)	nr	2650	12.58571323				

Figure 53a	Neuro	Neuro2	Neuro3	Neuro4	Neuro5	Neuro6
71.12621854	99.1814017	124.191117	173.743021	98.876267	182.511533	13.3611074
64.1262373	86.0275944	101.488104	163.842245	99.734982	188.932559	18.1544874

Newt 254		Newt 254		Newt 254	
VGAT-Cre x Rhodopsin-DTR	87.780782	84.5185716	80.2605538	81.5178844	89.9598789
VGAT-Cre x Rhodopsin-DTR	78.882742	77.7919786	68.0426768	79.83392	89.9598782

Figure 53b	sga4	hsp	vglut1	vglut2	Dgat1-cre6	GRD1	GRD2	Ngn2	Ngn3	LMTM1	LMTM2	LMTM3	LMTM4	Chol1	Chol2	Chol4	Ngh1	Ngh2	Ngh3	Ngh4	Cart0	Cart1	Fam19a1	Fam19a2	Fam19a3
1.3248151	2.358993	216.06481	1.44791383	74.05410775	79.1848688	13.262568	77.4622652	79.4991862	105.028284	10.6024128	113.412944	142.828714	110.915268	28.7720436	130.869255	2.9720944	151.823795	110.262878	89.4568832	43.5784137	81.731355	46.4374028	2.96889737	81.8818885	31.934031
7.34348152	7.0280725	195.94888	8.31610714	71.71963751	74.2983743	15.6517838	77.4680276	80.7705708	108.462088	18.6879289	95.7435501	132.564143	97.8978882	28.3073886	118.048155	14.4385551	155.28374	116.088718	64.2376581	56.348118	91.0590841	54.8881818	5.63181432	70.4320831	61.6188838

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New4-182a			
OB	22.297238	22.297232	0.000006
CV	34.02442	34.793863	0.247989
COV	14.320026	15.122889	0.000006
ES	31.202026	30.823275	0.000006
EW	28.148812	28.148812	0.000000
MSD	22.187672	22.894489	0.103801

New4-182a			
OB	28.285445	28.497395	0.000006
CV	50.271348	48.484811	1.064238
COV	41.40404	48.447174	0.000006
ES	54.251411	51.887782	1.681766
EW	51.41789	48.887178	0.000006
MSD	41.581789	41.428187	0.022285

New4-182a			
OB	1.9727023	1.4819113	0.1770281
CV	15.028129	14.20128	1.000007
COV	8.191719	8.247702	0.000006
ES	17.029129	17.300429	0.000006
EW	17.029129	17.300429	0.000006
MSD	1.5006929	1.7708178	0.113388

New4-182a			
OB	11.426208	10.287121	0.211786
CV	3.1526273	9.2026126	0.0736626
COV	4.820252	6.0281852	1.7719246
ES	5.380215	2.2182864	0.0464728
EW	0.9992512	1.4883984	1.4883984
MSD	4.3812874	6.153854	0.5710284

New4-182a			
OB	2.9925281	2.1225287	0.1077804
CV	4.8742688	3.9713827	1.0817111
COV	8.1626708	7.2787788	1.4883984
ES	8.810148	8.4922285	0.0772814
EW	3.177878	2.6966717	1.6944117
MSD	4.3898134	6.4762882	1.6944117

New4-182a			
OB	2.3049048	2.3776968	1.0887914
CV	3.1534884	3.3847514	1.3428828
COV	4.8428708	4.2327368	4.9440114
ES	3.177778	3.702728	0
EW	2.8171878	2.9747368	1.5442228
MSD	2.3892581	2.8882127	2.8188828

Figure 54D

	OB	CV	COV	ES	EW	MSD
New1	31.802567	42.185483	39.058421	0	0	0
	37.47172	24.424891	24.424891	0	0	0
	57.291204	44.572304	56.818333	0	0	0
New2	81.80789	82.963986	81.80789	0	0	0
	73.78178	71.824839	71.824839	0	0	0
	41.588143	47.320219	46.344534	0	0	0
New3	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54E

	OB	CV	COV	ES	EW	MSD
New4	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54F

	OB	CV	COV	ES	EW	MSD
New5	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54G

	OB	CV	COV	ES	EW	MSD
New6	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54H

	OB	CV	COV	ES	EW	MSD
New7	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54I

	OB	CV	COV	ES	EW	MSD
New8	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786

Figure 54J

	OB	CV	COV	ES	EW	MSD
New9	1.831178	4.897847	3.756458	0.353707	0.363774	0.6712421
	1.831178	1.238484	1.6558487	4.796209	3.678163	5.2911884
	13.84178	16.721864	23.455867	2.828242	5.839121	11.118786



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Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Average Count: 142.389, 142.330
 Statistical Test: Two-tailed unpaired parametric t-test
 t value: 0.000000
 df: 18
 Significance: 0.999999

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Average Count: 142.389, 142.330
 Statistical Test: Two-tailed unpaired parametric t-test
 t value: 0.0775
 df: 18
 Significance: 0.9822

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Group	Count	Average	Count	Average	Count	Significance
1	0.00125	10	0.00125	10		0.0000
2	0.00125	10	0.00125	10		0.0000
3	0.00125	10	0.00125	10		0.0000
4	0.00125	10	0.00125	10		0.0000
5	0.00125	10	0.00125	10		0.0000
6	0.00125	10	0.00125	10		0.0000
7	0.00125	10	0.00125	10		0.0000
8	0.00125	10	0.00125	10		0.0000
9	0.00125	10	0.00125	10		0.0000
10	0.00125	10	0.00125	10		0.0000

Two-way ANOVA with Bonferroni Multiple comparisons test at each stimulus intensity (df=9 vs. Cor p value=0.0000)

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Average Count: 142.389, 142.330
 Statistical Test: Two-tailed unpaired parametric t-test
 t value: 0.2734
 df: 18
 Significance: 0.7845

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Average Count: 142.389, 142.330
 Statistical Test: Two-tailed unpaired parametric t-test
 t value: 0.1170
 df: 18
 Significance: 0.9129

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Average Count: 142.389, 142.330
 Statistical Test: Two-tailed unpaired parametric t-test
 t value: 0.0522
 df: 18
 Significance: 0.9583

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Group	Mean	SD
1	142.389	142.330
2	142.389	142.330
3	142.389	142.330
4	142.389	142.330
5	142.389	142.330
6	142.389	142.330
7	142.389	142.330
8	142.389	142.330
9	142.389	142.330
10	142.389	142.330

Group	Count	Average	Count	Average	Count	Significance
1	0.00125	10	0.00125	10		0.0000
2	0.00125	10	0.00125	10		0.0000
3	0.00125	10	0.00125	10		0.0000
4	0.00125	10	0.00125	10		0.0000
5	0.00125	10	0.00125	10		0.0000
6	0.00125	10	0.00125	10		0.0000
7	0.00125	10	0.00125	10		0.0000
8	0.00125	10	0.00125	10		0.0000
9	0.00125	10	0.00125	10		0.0000
10	0.00125	10	0.00125	10		0.0000

Two-way ANOVA with Bonferroni Multiple comparisons test at each PPR interval (df=9 vs. Cor p value=0.0000)

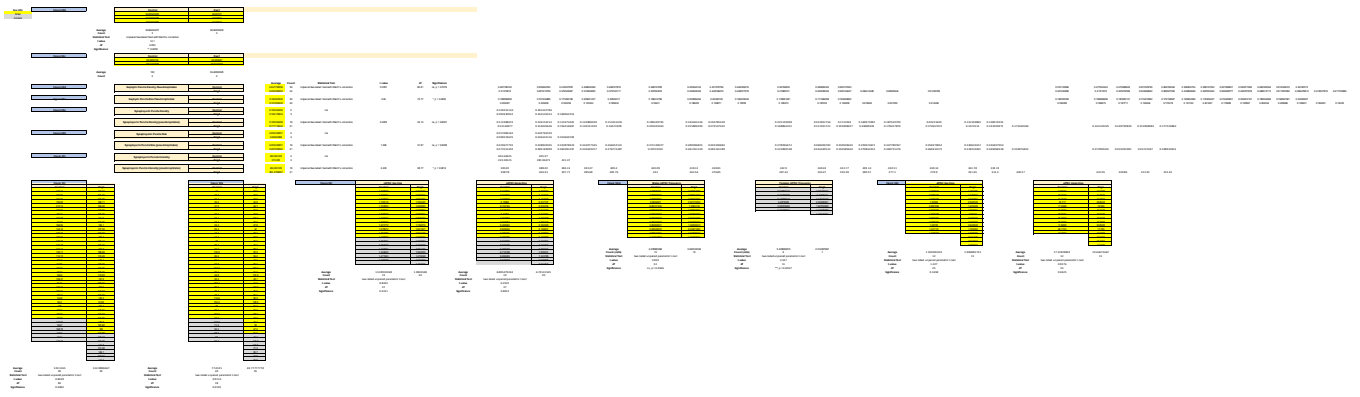
Sex Info

		2 males, 1 female	female	female
Figure S8e	% Gephyrin with Dystroglycan (avg. per animal)	10"	20"	O/N
		36.4929513	26.5676373	5.69961144
		20.72548408		
		25.93236631		
Average	27.7169339	26.5676373	5.69961144	
Count	3	1	1	

		10"	20"	O/N
Figure S8f	Gephyrin Size with Dystroglycan (avg. per animal)	0.048988203	0.047410739	0.038281746
		0.043506626		
		0.049433426		
		Average	0.047309418	0.047410739
Count	3	1	1	

		10"	20"	O/N
	Gephyrin Size without Dystroglycan (avg. per animal)	0.027810297	0.017452976	0.017541977
		0.021896803		
		0.019119668		
		Average	0.022942256	0.017452976
Count	3	1	1	

		10"	20"	O/N
Figure S8g	Dystroglycan Puncta Size (avg. per animal)	0.023284233	0.015653814	0.009445584
		0.018735656		
		0.016995115		
		Average	0.019671668	0.015653814
Count	3	1	1	



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