

**Table S1. Statistical analyses. Related to Figures 1 and 2.**

<b>Figure 1</b>	Textured NORT	1 Hour Ret.	tPPI	Air Puff Alone	Open Field Time in Center	EPM Time in Open Arms	EPM Total Arm Crosses	Habituation to Startle Noise	Marble Burying	Sociability	Social Novelty Preference	Grooming
Shank3B <sup>+/+</sup> (control) vs ShankB <sup>+/-</sup>	N = 17, 16	N = 17, 17	N = 31, 37	N = 31, 37	N = 14, 10	N = 11, 12	N = 11, 12	N = 31, 37	N = 8, 10	N = 18, 14	N = 18, 14	N = 8, 10
	p = 0.0190	p = 0.0486	p = 0.0029	p = 0.0083	p = 0.0091	p = 0.0125	p = 0.9477	p = 0.0068	p < 0.0001	p = 0.0237	p = 0.0110	p = 0.0016
Shank3 <sup>fl+</sup> (control) vs Cdx2 <sup>Cre</sup> ; Shank3 <sup>fl+</sup>	N = 13, 13	N = 11, 13	N = 11, 13	N = 11, 13	N = 11, 13	N = 11, 13	N = 11, 13	N = 11, 13	N = 13, 12	N = 13, 13	N = 13, 13	N = 11, 11
	p = 0.0437	p = 0.7919	p = 0.0145	p = 0.0172	p = 0.0286	p = 0.0174	p = 0.2810	p = 0.0327	p = 0.0471	p = 0.0447	p = 0.0458	p = 0.1925
Shank3 <sup>fl+</sup> (control) vs Advillin <sup>Cre</sup> ; Shank3 <sup>fl+</sup> vs Advillin <sup>Cre</sup> ; Shank3 <sup>fl+</sup> vs Advillin <sup>Cre</sup> ; Shank3 <sup>fl+</sup>	N = 37, 20, 9	N = 34, 26, 9	N = 30, 25, 9	N = 30, 25, 9	N = 38, 25, 9	N = 34, 20, 9	N = 34, 20, 9	N = 30, 25, 9	N = 34, 21, 9	N = 36, 26, 9	N = 36, 26, 9	N = 8, ---, 9
	F(2,63)=5.826	F(2,66)=0.3597	F(2,61)=6.812	F(2,61)=5.44	F(2,69)=8.502	F(2,60)=8.807	F(2,60)=0.153	F(2,61)=9.049	F(2,61)=4.324	F(2,68)=16.36	F(2,68)=12.11	p = 0.3436
	P = 0.0048	P = 0.6993	P = 0.0021	P = 0.0067	P = 0.0005	P = 0.0004	P = 0.8583	P = 0.0004	P = 0.0165	P < 0.0001	P < 0.0001	
Mecp2 <sup>fl+</sup> or fl/y (control) vs Advillin <sup>Cre</sup> ; Mecp2 <sup>fl+</sup> vs Advillin <sup>Cre</sup> ; Mecp2 <sup>fl/y</sup>	N = 35, 20, 18	N = 33, 17, 18	N = 41, 24, 20	N = 41, 24, 20	N = 38, 20, 20	N = 32,13,11	N = 32,13,11	N = 41, 24, 20	N = 26, 11, 16	N = 33, 24, 20	N = 33, 24, 20	N = 9, ---, 7
	F(2,70)=5.996	F(2,65)=0.4695	F(2,82)=5.104	F(2,82)=6.226	F(2,75)=16.5	F(2,53)=12.02	F(2,53)=0.214	F(2,82)=6.391	F(2,50)=4.219	F(2,62)=11.73	F(2,62)=18.8	p = 0.4667
	P = 0.0039	P = 0.6274	P = 0.0081	P = 0.0030	P < 0.0001	P < 0.0001	P = 0.8083	P = 0.0026	P = 0.0203	P < 0.0001	P < 0.0001	
Shank3 <sup>fl+</sup> (control) vs Shank3 <sup>FX/+</sup> vs Cdx2 <sup>Cre</sup> ; Shank3 <sup>FX/+</sup>	N = 22, 24, 23	N = 24, 24, 29	N = 22, 21, 26	N = 22, 21, 26	N = 21, 24, 22	N = 22, 23, 22	N = 22, 23, 22	N = 22, 21, 26	N = 21, 18, 21	N = 21, 25, 29	N = 21, 25, 29	N = 21, 26, 29
	F(2,66)=4.231	F(2,74)=3.924	F(2,66)=6.496	F(2,66)=4.316	F(2,64)=5.694	F(2,64)=6.679	F(2,64)=0.914	F(2,66)=3.488	F(2,57)=10.4	F(2,72)=10.25	F(2,72)=5.61	F(2,73)=7.824
	P = 0.0187	P = 0.0240	P = 0.0027	P = 0.0173	P = 0.0053	P = 0.0023	P = 0.4061	P = 0.0363	P = 0.0001	P = 0.0001	P = 0.0055	P = 0.0008
Shank3 <sup>fl+</sup> (control) vs Shank3 <sup>FX/+</sup> vs Advillin <sup>Cre</sup> ; Shank3 <sup>FX/+</sup>	N = 25, 26, 22	N = 26, 25, 25	N = 26, 29, 25	N = 26, 29, 25	N = 26, 28, 26	N = 21, 22, 16	N = 21, 22, 16	N = 26, 29, 25	N = 22, 25, 19	N = 26, 29, 24	N = 26, 29, 24	N = 24, 28, 19
	F(2,70)=4.589	F(2,73)=1.631	F(2,77)=7.408	F(2,77)=7.184	F(2,77)=4.49	F(2,56)=5.105	F(2,56)=1.861	F(2,77)=3.19	F(2,63)=11.83	F(2,75)=4.385	F(2,75)=5.208	F(2,68)=4.709
	P = 0.0134	P = 0.2027	P = 0.0011	P = 0.0014	P = 0.0143	P = 0.0092	P = 0.1649	P = 0.0466	P < 0.0001	P = 0.0158	P = 0.0076	P = 0.0122
HCN1 <sup>+/+</sup> (control) vs HCN1 <sup>+/-</sup> vs HCN1 <sup>-/-</sup>	N = 16, 14, 6	N = 16, 14, 6	N = 16, 14, 6	N = 16, 14, 6	not determined (n.d.)	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
	F(2,33)=7.604	F(2,33)=6.308	F(2,33)=7.064	F(2,33)=0.2048	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
	P = 0.0028	P = 0.0048	P = 0.0028	P = 0.8158	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.

<b>Figures 2D-F and R-V</b>	PV+ Neuron #, S1	PV+ Neuron #, V1	PV+ Neuron #, BLA	E/I Ratio (S1 and V1)	Event Frequency, S1	Event Frequency, V1	Event Amplitude, S1	Event Amplitude, V1	<b>Figures 2I-K</b>	PV+ Neuron #, S1	PV+ Neuron #, V1	PV+ Neuron #, BLA
Shank3 <sup>fl+</sup> (control) vs Advillin <sup>Cre</sup> ; Shank3 <sup>fl+</sup>	N = 6, 6	N = 6, 6	N = 6, 6	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	Shank3 <sup>fl+</sup> (control) vs Shank3 <sup>FX/+</sup> vs Advillin <sup>Cre</sup> ; Shank3 <sup>FX/+</sup>	N = 4, 4, 4	N = 4, 4, 4	N = 4, 4, 4
	p = 0.0115	p = 0.5561	p = 0.0012	F(1,12)=6.254	F(1,12) = 4.698	F(1,12)=0.0030	F(1,12)=9.572	F(1,12)=0.0739		F(2,9)=6.206	F(2,9)=5.794	F(2,9)=6.945
				P = 0.0254	P = 0.0459	P = 0.9569	P = 0.0093	P = 0.7904		P = 0.0202	P = 0.0241	P = 0.0150
Mecp2 <sup>fl+</sup> or fl/y (control) vs Advillin <sup>Cre</sup> ; Mecp2 <sup>fl/y</sup>	N = 5, 4	N = 5, 4	N = 5, 4	N = 5, 4	N = 5, 4	N = 5, 4	N = 5, 4	N = 5, 4	Mecp2 <sup>fl/y</sup> (control) vs Mecp2 <sup>STOP/y</sup> vs Advillin <sup>Cre</sup> ; Mecp2 <sup>STOP/y</sup>	N = 4, 5, 4	N = 4, 5, 4	N = 4, 5, 4
	p = 0.0380	p = 0.5726	p = 0.0108	N(1,14)=5.776	F(1,14) = 6.822	F(1,14)=0.3446	F(1,14)=1.736	F(1,14)=0.3476		F(2,10)=30.95	F(2,10)=4.833	F(2,10)=7.180
				P = 0.0296	P = 0.0205	P = 0.5654	P = 0.2088	P = 0.5648		P < 0.0001	P = 0.0340	P = 0.0117

**Table S2. Time spent per compartment in the three chamber social interaction test (seconds). Related to Figure 1.**

	Novel Mouse	Novel Object	Center	Pref. for novel mouse	Novel Mouse	Familiar Mouse	Center	Pref. for novel mouse
Control (n = 18)	298.27 +/-15.70	232.95 +/- 13.01	68.76 +/- 8.26	Yes p = 0.0044	301.41 +/- 16.63	220.12 +/- 17.69	78.43 +/- 9.72	Yes p = 0.0010
Shank3B <sup>+/+</sup> (n=14)	259.26 +/- 18.38	272.32 +/- 18.32	68.38 +/- 17.16	No p = 0.8332	262.77 +/- 24.67	258.38 +/- 23.28	78.81 +/- 17.39	No p = 0.9839
Control (n=13)	320.14 +/- 20.70	229.85 +/- 16.07	50.01 +/- 5.52	Yes p = 0.0002	302.08 +/- 22.70	225.07 +/- 20.20	72.85 +/- 8.79	Yes p = 0.0093
Cdx2 <sup>Cre</sup> ; Shank3 <sup>f/+</sup> (n = 13)	296.29 +/- 19.2	249.12 +/- 16.43	54.56 +/- 7.80	No p = 0.0826	283.41 +/- 15.03	250.77 +/- 16.05	65.82 +/- 6.72	No p = 0.4217
Control (n = 36)	318.59 +/- 7.61	229.55 +/- 7.95	51.83 +/- 3.51	Yes p < 0.0001	302.31 +/- 10.59	233.82 +/- 10.46	63.87 +/- 4.38	Yes p < 0.0001
Advillin <sup>Cre</sup> ; Shank3 <sup>f/+</sup> (n = 26)	286.31 +/- 13.00	258.17 +/- 12.72	55.51 +/- 3.75	No p = 0.0957	275.71 +/- 17.50	258.88 +/- 18.17	65.37 +/- 7.11	No p = 0.6429
Advillin <sup>Cre</sup> ; Shank3 <sup>f/f</sup> (n = 9)	271.12 +/- 21.95	248.43 +/- 23.47	80.40 +/- 12.17	No p = 0.5463	250.53 +/- 24.67	277.25 +/- 29.16	72.22 +/- 16.79	No p = 0.6647
Control (n = 33)	292.60 +/- 9.63	219.51 +/- 8.59	87.79 +/- 6.22	Yes p < 0.0001	300.86 +/- 13.57	220.08 +/- 10.79	79.05 +/- 4.45	Yes p < 0.0001
Advillin <sup>Cre</sup> ; Mecp2 <sup>f/+</sup> (n = 24)	264.45 +/- 12.41	243.93 +/- 12.20	90.95 +/- 9.22	No p = 0.5892	260.96 +/- 9.83	244.33 +/- 10.15	94.69 +/- 7.15	No p = 0.5861
Advillin <sup>Cre</sup> ; Mecp2 <sup>f/y</sup> (n = 20)	265.88 +/- 9.71	270.10 +/- 8.46	64.02 +/- 10.94	No p = 0.9517	236.62 +/- 16.53	264.85 +/- 21.73	98.53 +/- 14.45	No p = 0.3159
Control (n = 21)	315.54 +/- 10.12	233.93 +/- 10.08	50.53 +/- 5.27	Yes p < 0.0001	299.85 +/- 16.41	233.89 +/- 16.86	66.26 +/- 8.22	Yes p = 0.0102
Shank3 <sup>FX/+</sup> (n = 25)	267.10 +/- 15.00	275.67 +/- 15.30	57.22 +/- 4.28	No p = 0.8334	264.64 +/- 15.83	253.43 +/- 12.56	81.89 +/- 7.41	No p = 0.8429
Cdx2 <sup>Cre</sup> ; Shank3 <sup>FX/+</sup> (n = 29)	304.00 +/- 10.74	243.52 +/- 11.61	52.48 +/- 3.85	Yes p < 0.0001	278.90 +/- 17.34	249.11 +/- 19.26	71.98 +/- 7.43	No p = 0.2631
Control (n = 26)	308.01 +/- 11.38	253.89 +/- 11.24	38.10 +/- 4.87	Yes p = 0.0043	307.56 +/- 15.49	234.56 +/- 15.06	57.87 +/- 5.50	Yes p = 0.0007
Shank3 <sup>FX/+</sup> (n = 29)	279.66 +/- 17.66	276.31 +/- 18.40	44.03 +/- 3.72	No p = 0.9761	275.36 +/- 17.66	260.55 +/- 17.30	64.08 +/- 7.82	No p = 0.7030
Advillin <sup>Cre</sup> ; Shank3 <sup>FX/+</sup> (n = 24)	304.88 +/- 10.25	250.61 +/- 10.03	44.51 +/- 4.01	Yes p = 0.0064	294.60 +/- 14.00	244.51 +/- 14.63	60.88 +/- 7.86	Yes p = 0.0380

**Table S3. Time spent per compartment in the three chamber social interaction test (seconds). Related to Figure 3.**

	Novel Mouse	Novel Object	Center	Pref. for novel mouse	Novel Mouse	Familiar Mouse	Center	Pref. for novel mouse
Control (TAM P5, n = 13)	349.62 +/- 16.86	206.53 +/- 14.34	43.80 +/- 5.79	Yes p < 0.0001	322.74 +/- 15.59	230.75 +/- 12.99	46.51 +/- 5.33	Yes p < 0.0001
Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P5, n = 15)	286.43 +/- 19.87	249.20 +/- 19.90	64.37 +/- 9.48	No p = 0.1944	275.27 +/- 19.37	257.81 +/- 20.95	66.92 +/- 7.10	No p = 0.6809
Control (TAM P10, n = 21)	303.06 +/- 10.55	226.48 +/- 8.99	70.46 +/- 5.79	Yes p < 0.0001	289.51 +/- 10.38	222.13 +/- 9.50	88.36 +/- 6.40	Yes p < 0.0001
Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P10, n = 15)	271.91 +/- 16.14	239.42 +/- 16.49	88.67 +/- 8.89	No p = 0.1413	250.75 +/- 16.59	247.23 +/- 16.91	102.02 +/- 9.15	No p = 0.9782
Control (TAM P28, n = 19)	344.57 +/- 14.61	200.88 +/- 12.22	54.54 +/- 5.06	Yes p < 0.0001	307.33 +/- 14.13	221.20 +/- 13.32	71.47 +/- 6.20	Yes p < 0.0001
Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P28, n = 11)	282.90 +/- 16.43	228.27 +/- 19.03	88.83 +/- 8.00	Yes p = 0.0321	268.47 +/- 21.98	255.38 +/- 18.22	76.14 +/- 8.23	No p = 0.8342
Control (TAM P5, n = 14)	340.81 +/- 6.10	217.48 +/- 8.16	41.71 +/- 10.47	Yes p < 0.0001	347.97 +/- 24.46	182.43 +/- 20.13	69.59 +/- 6.94	Yes p < 0.0001
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P5, n = 10)	265.07 +/- 10.00	254.04 +/- 15.60	80.89 +/- 8.90	No p = 0.7007	271.75 +/- 14.27	232.13 +/- 15.78	96.12 +/- 2.59	No p = 0.3015
Control (TAM P10, n = 28)	315.17 +/- 12.39	215.41 +/- 11.61	69.43 +/- 8.84	Yes p < 0.0001	295.56 +/- 11.41	233.80 +/- 12.50	70.64 +/- 6.46	Yes p = 0.0004
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P10, n = 19)	293.58 +/- 12.40	239.46 +/- 13.44	66.95 +/- 9.35	Yes p = 0.0108	243.66 +/- 15.43	279.71 +/- 17.94	76.64 +/- 6.13	No p = 0.9980
Control (TAM P28, n = 19)	294.83 +/- 7.72	221.32 +/- 8.27	83.85 +/- 6.45	Yes p < 0.0001	278.09 +/- 9.62	218.26 +/- 9.30	103.65 +/- 7.45	Yes p = 0.0029
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P28, n = 15)	271.80 +/- 13.30	233.79 +/- 11.70	94.41 +/- 13.82	Yes p = 0.0355	258.26 +/- 19.11	250.20 +/- 22.49	91.54 +/- 9.97	No p = 0.9133
Control (TAM P28, n = 27)	311.54 +/- 8.03	228.78 +/- 7.46	59.66 +/- 5.10	Yes p < 0.0001	321.87 +/- 12.15	200.02 +/- 9.20	78.10 +/- 6.71	Yes p < 0.0001
Shank3 <sup>FX/+</sup> (TAM P28, n = 16)	288.40 +/- 23.65	261.07 +/- 21.29	50.52 +/- 5.20	No p = 0.3343	269.44 +/- 25.01	242.85 +/- 21.55	87.70 +/- 22.80	No p = 0.5104
Advillin <sup>CreERT2</sup> ; Shank3 <sup>FX/+</sup> (TAM P28, n = 10)	322.14 +/- 18.54	209.10 +/- 24.33	68.76 +/- 8.48	Yes p < 0.0001	261.30 +/- 24.87	238.59 +/- 22.72	100.11 +/- 12.60	No p = 0.7423
Control (TAM P28, n = 23)	306.43 +/- 12.56	195.60 +/- 12.48	97.97 +/- 8.83	Yes p < 0.0001	287.16 +/- 18.43	220.98 +/- 15.86	91.86 +/- 7.81	Yes p < 0.0014
Mecp2 <sup>STOP/y</sup> (TAM P28, n = 13)	266.50 +/- 20.00	241.09 +/- 17.69	92.41 +/- 11.19	No p = 0.5220	225.78 +/- 25.47	256.42 +/- 32.13	117.78 +/- 30.65	No p = 0.6635
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>STOP/y</sup> (TAM P28, n = 10)	330.98 +/- 20.74	203.65 +/- 21.45	65.38 +/- 21.45	Yes p < 0.0001	267.65 +/- 37.29	239.57 +/- 36.78	92.78 +/- 37.43	No p = 0.7678

**Table S4. Statistical analyses. Related to Figures 3 and S3**

	Textured NORT	1 Hour Ret.	tPPI	Air Puff Alone	Open Field Time in Center	EPM Time in Open Arms	EPM Total Arm Crosses	Habituation to Startle Noise	Marble Burying	Sociability	Social Novelty Preference
Shank3 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P5)	N = 8, 15	N = 8, 15	N = 10, 15	N = 10, 15	N = 8, 14	N = 13, 15	N = 13, 15	N = 13, 15	N = 12, 15	N = 13, 15	N = 13, 15
	p = 0.0319	p = 0.9657	p = 0.0126	p = 0.0306	p = 0.0024	p = 0.0007	p = 0.6128	p = 0.0057	p = 0.0483	p = 0.0004	p = 0.0007
Shank3 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P10)	N = 20, 14	N = 17, 14	N = 20, 14	N = 20, 14	N = 25, 14	N = 17, 11	N = 17, 11	N = 23, 14	N = 23, 14	N = 21, 15	N = 21, 15
	p = 0.0128	p = 0.9789	p = 0.0294	p = 0.0362	p = 0.0489	p = 0.0189	p = 0.6092	p = 0.5227	p = 0.0070	p = 0.0219	p = 0.0013
Shank3 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Shank3 <sup>fl/y</sup> (TAM P28)	N = 20, 14	N = 17, 14	N = 20, 14	N = 20, 14	N = 21, 14	N = 21, 11	N = 21, 11	N = 19, 11	N = 18, 11	N = 19, 11	N = 19, 11
	p = 0.0292	p = 0.9122	p = 0.0126	p = 0.0115	p = 0.1732	p = 0.1144	p = 0.2031	p = 0.8368	p = 0.6816	p = 0.1592	p < 0.0001
Mecp2 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P5)	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10	N = 14, 10
	p = 0.0074	p = 0.7196	p = 0.0021	p = 0.0010	p = 0.0026	p < 0.0001	p = 0.3922	p = 0.0012	p = 0.0010	p = 0.0004	p < 0.0001
Mecp2 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P10)	N = 27, 29	N = 27, 28	N = 16, 22	N = 16, 22	N = 22, 20	N = 28, 29	N = 28, 29	N = 27, 25	N = 20, 15	N = 28, 19	N = 28, 19
	p = 0.0355	p = 0.6588	p = 0.0120	p = 0.0124	p = 0.0398	p = 0.0026	p = 0.1291	p = 0.1242	p = 0.0269	p = 0.0064	p = 0.0003
Mecp2 <sup>fl/y</sup> (control) vs Advillin <sup>CreERT2</sup> ; Mecp2 <sup>fl/y</sup> (TAM P28)	N = 25, 13	N = 22, 12	N = 22, 12	N = 22, 12	N = 22, 12	N = 22, 12	N = 22, 12	N = 19, 15	N = 17, 15	N = 19, 15	N = 19, 15
	p = 0.0051	p = 0.8460	p = 0.0338	p = 0.0005	p = 0.8042	p = 0.5364	P = 0.9674	p = 0.8970	p = 0.6079	p = 0.0272	p = 0.0024
Shank3 <sup>fl/y</sup> (control) vs Shank3 <sup>FX/+</sup> vs Advillin <sup>CreERT2</sup> ; Shank3 <sup>FX/+</sup> (TAM P28)	N = 27, 16, 10	N = 27, 16, 10	N = 27, 18, 14	N = 27, 18, 14	N = 25, 15, 10	N = 29, 16, 10	N = 29, 16, 10	N = 27, 16, 10	N = 27, 16, 10	N = 27, 16, 10	N = 27, 16, 10
	F(2,50)=3.648	F(2,50)=6.130	F(2,56)=4.193	F(2,56)=4.318	F(2,47)=6.597	F(2,52)=7.242	F(2,52)=1.758	F(2,50)=8.214	F(2,50)=8.548	F(2,50)=8.028	F(2,50)=13.87
	P = 0.0332	P = 0.0042	P = 0.0201	P = 0.0180	P = 0.0030	P = 0.0017	P = 0.1825	P = 0.0008	P = 0.0006	P = 0.0009	p < 0.0001
Mecp2 <sup>fl/y</sup> (control) vs Mecp2 <sup>STOP/y</sup> vs Advillin <sup>CreERT2</sup> ; Mecp2 <sup>STOP/y</sup> (TAM P28)	N = 24, 14, 11	N = 25, 15, 11	N = 25, 18, 12	N = 25, 18, 12	N = 32, 22, 12	N = 25, 14, 14	N = 25, 14, 14	N = 25, 18, 11	N = 19, 15, 10	N = 23, 13, 10	N = 23, 13, 10
	F(2,46)=3.970	F(2,48)=4.783	F(2,52)=5.151	F(2,52)=8.054	F(2,63)=10.84	F(2,50)=7.014	F(2,50)=0.243	F(2,54)=4.480	F(2,41)=17.17	F(2,43)=10.42	F(2,43)=8.417
	P = 0.0257	P = 0.0128	P = 0.0091	P = 0.0009	P < 0.0001	P = 0.0021	P = 0.7856	P = 0.0158	P < 0.0001	P = 0.0002	P = 0.0008

**Table S5. Statistical analyses. Related to Figures 4, 5 and S5**

Figures 4 and S5	GABRB3/vGLUT1 coloc.	Textured NORT	1 Hour Ret.	tPPI	Air Puff Alone	Open Field Time in Center	EPM Time in Open Arms	EPM Total Arm Crosses	Habituation to Startle Noise	Sociability	Social Novelty Preference	PV+ Neuron #, S1	PV+ Neuron #, V1	PV+ Neuron #, BLA
Mecp2 <sup>R/y</sup> (control) vs Mecp2 <sup>C/y</sup> vs Advillin <sup>Cre</sup> ; Mecp2 <sup>C/y</sup>		N = 30, 22, 10	N = 32, 21, 10	N = 30, 16, 8	N = 30, 16, 8	N = 31, 22, 12	N = 31, 16, 8	N = 31, 16, 8	N = 30, 22, 10	N = 26, 15, 10	N = 26, 15, 10	N = 6, 6, 6	N = 6, 6, 6	N = 6, 6, 6
	N = 5, 5, 19	F(2,59)=4.671	F(2,60)=5.974	F(2,51)=21.42	F(2,51)=25.90	F(2,55)=14.53	F(2,52)=6.615	F(2,52)=0.729	F(2,59)=4.689	F(2,48)=9.428	F(2,48)=6.188	F(2,15)=40.32	F(2,15)=8.162	F(2,15)=21.18
	F(2,26)=14.88	P = 0.0131	P = 0.0043	P < 0.0001	P < 0.0001	P < 0.0001	P = 0.0028	P = 0.4872	P = 0.0129	P = 0.0004	P = 0.0041	P < 0.0001	P = 0.0040	P < 0.0001
Mecp2 <sup>R/R</sup> (control) vs Mecp2 <sup>R/C</sup> vs Advillin <sup>Cre</sup> ; Mecp2 <sup>R/C</sup>		N = 25, 14, 12	N = 26, 14, 8	N = 29, 15, 8	N = 29, 15, 8	N = 28, 15, 8	N = 22, 14, 8	N = 22, 14, 8	N = 29, 15, 8	N = 26, 13, 8	N = 26, 13, 8	N = 7, 7, 6	N = 7, 7, 6	N = 6, 6, 6
	not determined (n.d.)	F(2,48)=4.234	F(2,45)=4.803	F(2,49)=13.74	F(2,49)=16.24	F(2,48)=5.746	F(2,41)=4.664	F(2,41)=0.113	F(2,48)=5.544	F(2,44)=9.207	F(2,44)=9.962	F(2,17)=5.977	F(2,17)=7.372	F(2,15)=8.452
		P = 0.0202	P = 0.0129	P < 0.0001	P < 0.0001	P = 0.0058	P = 0.0150	P = 0.8936	P = 0.0068	P = 0.0005	P = 0.0003	P = 0.0108	P = 0.0050	P = 0.0035

Figure 5A-D	tPPI	Air Puff Alone	Startle Amplitude	Startle Amplitude (compare to iso)
Mecp2 <sup>F/y</sup> (control) vs. Advillin <sup>Cre</sup> ; Mecp2 <sup>F/y</sup>	N = 18, 9	N = 18, 9	N = 18, 9	N = 31
	F(1,50)=38.66	F(1,50)=8.531	F(1,50)=14.03	F(2,90)=3.908
	P < 0.0001	P = 0.0047	P = 0.0004	P = 0.0236

**Table S6. Statistical analyses. Related to Figures 7 and S7**

	Textured NORT	1 Hour Ret.	tPPI	Air Puff Alone	Open Field Time in Center	EPM Time in Open Arms	EPM Total Arm Crosses	Habituation to Startle Noise	Sociability	Social Novelty Preference	Marble Burying	Grooming
Saline-treated: Shank3B <sup>+/+</sup> (control) vs Shank3B <sup>+/-</sup>	N = 12, 16	N = 12, 17	N = 12, 17	N = 12, 17	N = 14, 15	N = 12, 17	N = 12, 17	N = 12, 17	N = 12, 17	N = 12, 17	N = 12, 17	N = 7, 7
	p = 0.0040	p = 0.0349	p = 0.0214	p = 0.0332	p = 0.0038	p = 0.0025	p = 0.8292	p = 0.3355	p = 0.0026	p < 0.0001	p = 0.0004	p = 0.0320
Isoguvacine-treated: Shank3B <sup>+/+</sup> (control) vs Shank3B <sup>+/-</sup>	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 19, 21	N = 10, 11
	p = 0.0447	p = 0.0458	p = 0.8862	p = 0.1553	p = 0.0750	p = 0.9697	p = 0.8744	p = 0.3633	p = 0.3855	p = 0.0220	p = 0.7603	p = 0.0192
Saline-treated: Mecp2 <sup>R/y or R/R</sup> (control) vs Mecp2 <sup>R/C</sup> vs Mecp2 <sup>C/y</sup>	N = 34, 11, 10	N = 34, 11, 10	N = 35, 14, 11	N = 35, 14, 11	N = 34, 16, 11	N = 35, 11, 10	N = 35, 11, 10	N = 35, 11, 10	N = 29, 10, 8	N = 29, 10, 8	N = 29, 9, 9	N = 8, ---, 7
	F(2,52)=8.116	F(2,52)=4.440	F(2,57)=9.182	F(2,57)=18.61	F(2,58)=8.146	F(2,53)=14.55	F(2,53)=0.255	F(2,46)=0.462	F(2,44)=6.203	F(2,45)=14.03	F(2,44)=16.14	p = 0.4319
	P = 0.0009	P = 0.0166	P = 0.0003	P < 0.0001	P = 0.0008	P < 0.0001	P = 0.7756	P = 0.6328	P = 0.0042	p < 0.0001	P < 0.0001	
Isoguvacine-treated: Mecp2 <sup>R/y or R/R</sup> (control) vs Mecp2 <sup>R/C</sup> vs Mecp2 <sup>C/y</sup>	N=30, 14, 8	N = 30, 14, 8	N = 31, 15, 8	N = 31, 15, 8	N = 30, 15, 8	N = 28, 15, 8	N = 28, 15, 8	N = 28, 15, 8	N = 28, 8, 8	N = 28, 8, 8	N = 25, 14, 8	N = 7, ---, 7
	F(2,49)=3.708	F(2,49)=3.251	F(2,51)=0.719	F(2,51)=2.671	F(2,50)=0.376	F(2,48)=0.765	F(2,48)=0.175	F(2,39)=0.206	F(2,41)=0.975	F(2,41)=0.337	F(2,44)=14.53	p = 0.4002
	P = 0.0335	P=0.0472	P = 0.4920	P = 0.0789	P = 0.6886	P = 0.4708	P = 0.8484	P = 0.8144	P = 0.3858	P = 0.7162	P < 0.0001	

	PV+ Neuron #, S1	PV+ Neuron #, V1	PV+ Neuron #, BLA	E/I Balance, S1	E/I Balance, V1	Event Amplitude, S1	Event Frequency, S1	Event Amplitude, V1	Event Frequency, V1
Saline-treated: Shank3B <sup>+/+</sup> (control) vs Shank3B <sup>+/-</sup>	N = 4, 5	N = 4, 5	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4
	p = 0.0100	p = 0.0012	p = 0.0154	p < 0.0001	p = 0.0011	F(1,12)=17.52 ; P=0.0013	F(1,12)=4.949 ; P=0.0461	F(1,12)=21.75 ; P=0.0005	F(1,12)=10.07 ; P=0.0080
Isoguvacine-treated: Shank3B <sup>+/+</sup> (control) vs Shank3B <sup>+/-</sup>	N = 6, 4	N = 6, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4
	p = 0.0582	p = 0.0023	p = 0.0066	p = 0.1086	p = 0.0003	F(1,12)=0.2058 ; P=0.6582	F(1,12)=6.922 ; P=0.0219	F(1,12)=17.78 ; P=0.0012	F(1,12)=19.09 ; P=0.0009
Saline-treated: Mecp2 <sup>R/y</sup> (control) vs Mecp2 <sup>C/y</sup>	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4
	p = 0.0384	p = 0.0146	p = 0.0012	p = 0.0006	p = 0.0016	F(1,12)=0.2084 ; P=0.6562	F(1,12)=15.94 ; P=0.0018	F(1,12)=2.551 ; P=0.1362	F(1,12)=9.131 ; P=0.0106
Isoguvacine-treated: Mecp2 <sup>R/y</sup> (control) vs Mecp2 <sup>C/y</sup>	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4	N = 4, 4
	p = 0.1504	p = 0.0089	p = 0.0194	p = 0.3652	p = 0.0117	F(1,12)=1.740 ; P=0.2118	F(1,12)=0.2264 ; P=0.6428	F(1,12)=1.155 ; P=0.3036	F(1,12)=4.863 ; P=0.0477

**Table S7. Summary of findings. Related to all figures.**

	Hairy Skin Sensitivity	Texture Discrimination	Anxiety-like Behaviors	Sociability	Social Novelty	Marble Burying	Grooming	PV+ Neuron #, S1	PV+ Neuron #, BLA	PV+ Neuron #, V1	E/I Balance S1	E/I Balance V1
Germline Mutants												
Shank3 <sup>B<sup>-/-</sup></sup>	↑	↓	↑	↓	↓	↓	↑	↓	↓	↓	↓	↓
Shank3 <sup>FX/+</sup>	↑	↓	↑	↓	↓	↓	↑	↓	↓	↓	n.d.	n.d.
Mecp2 <sup>STOP/y</sup>	↑	↓	↑	↓	↓	↓	---	↑	↓	↑	n.d.	n.d.
Mecp2 <sup>R/C or C/y</sup>	↑	↓	↑	↓	↓	↓	---	↑	↓	↑	↓	↓
Conditional Deletions												
Cdx2 <sup>Cre</sup> ; Shank3 <sup>B<sup>+/+</sup></sup>	↑	↓	↑	↓	↓	↓	---	n.d.	n.d.	n.d.	n.d.	n.d.
Advillin <sup>Cre</sup> ; Shank3 <sup>B<sup>+/+</sup></sup>	↑	↓	↑	↓	↓	↓	n.d.	↓	↓	---	↓	---
Advillin <sup>Cre</sup> ; Shank3 <sup>B<sup>fl</sup></sup>	↑	↓	↑	↓	↓	↓	---	n.d.	n.d.	n.d.	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Shank3 <sup>B<sup>+/+</sup></sup> (TAM P5)	↑	↓	↑	↓	↓	↓	n.d.	↓	↓	---	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Shank3 <sup>B<sup>+/+</sup></sup> (TAM P10)	↑	↓	↓	↓	↓	↑	n.d.	↓	↑	---	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Shank3 <sup>B<sup>+/+</sup></sup> (TAM P28)	↑	↓	---	---	↓	---	n.d.	↓	---	---	n.d.	n.d.
Advillin <sup>Cre</sup> ; Mecp2 <sup>B<sup>+/+</sup></sup>	↑	↓	↑	↓	↓	---	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
Advillin <sup>Cre</sup> ; Mecp2 <sup>B<sup>y</sup></sup>	↑	↓	↑	↓	↓	↓	---	↑	↓	---	↓	---
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>B<sup>y</sup></sup> (TAM P5)	↑	↓	↑	↓	↓	↓	n.d.	↑	↓	---	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>B<sup>y</sup></sup> (TAM P10)	↑	↓	↓	+/-	↓	↑	n.d.	↓	↑	---	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>B<sup>y</sup></sup> (TAM P28)	↑	↓	---	---	↓	---	n.d.	↓	---	---	n.d.	n.d.
Genetic Rescues												
Shank3 <sup>FX/+</sup>	↑	↓	↑	↓	↓	↓	↑	↓	↓	↓	↓	↓
Cdx2 <sup>Cre</sup> ; Shank3 <sup>FX/+</sup>	---	---	---	---	+/-	+/-	↑	n.d.	n.d.	n.d.	n.d.	n.d.
Advillin <sup>Cre</sup> ; Shank3 <sup>FX/+</sup>	---	---	---	---	+/-	+/-	↑	---	---	↓	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Shank3 <sup>FX/+</sup> (TAM P28)	---	↓	↑	---	↓	↓	n.d.	---	↓	↓	n.d.	n.d.
Mecp2 <sup>STOP/y</sup>	↑	↓	↑	↓	↓	↓	---	↑	↓	↑	↓	↓
Advillin <sup>Cre</sup> Mecp2 <sup>STOP/y</sup>	---	---	---	---	---	n.d.	n.d.	+/-	---	↑	n.d.	n.d.
Advillin <sup>CreERT2</sup> ; Mecp2 <sup>STOP/y</sup> (TAM P28)	---	↓	↑	---	↓	↓	n.d.	+/-	↓	↑	n.d.	n.d.
Viral Rescues												
Mecp2 <sup>R/C or C/y</sup>	↑	↓	↑	↓	↓	↓	---	↑	↑	↑	n.d.	n.d.
Advillin <sup>Cre</sup> ; Mecp2 <sup>R/C</sup>	---	---	---	---	---	n.d.	n.d.	---	---	↑	n.d.	n.d.
Advillin <sup>Cre</sup> ; Mecp2 <sup>C/y</sup>	---	---	---	---	+/-	n.d.	n.d.	+/-	+/-	↑	n.d.	n.d.
Pharmacological Rescues												
Shank3B <sup>+/-</sup> + saline	↑	↓	↑	↓	↓	↓	↑	↓	↓	↓	↓	↓
Shank3B <sup>+/-</sup> + iso	---	↓	---	---	---	---	---	---	---	↓	---	↓
Mecp2 <sup>R/C or C/y</sup> + saline	↑	↓	↑	↓	↓	↓	---	↑	↑	↑	↓	↓
Mecp2 <sup>R/C or C/y</sup> + iso	---	↓	---	---	---	↓	---	---	---	↑	---	↓

(↑, increased; ↓, decreased; +/-, intermediate phenotype; ---, not different than control littermates; n.d., not determined; \*, findings observed in Orefice et al., 2016)

**Table S8. Oligonucleotide List. Related to Key Resources Table.**

Mouse Line	Oligo	Sequence (5' to 3')
<i>Advillin<sup>Cre</sup></i>	transgene forward	CCC TGT TCA CTG TGA GTA GG
	transgene reverse	AGT ATC TGG TAG GTG CTT CCA G
	internal control	GCG ATC CCT GAA CAT GTC CAT C
<i>Advillin<sup>CreERT2</sup></i>	AdvilCreERT2_1	CCC TGT TCA CTG TGA GTA GG
	AdvilCreERT2_2	AGT ATC TGG TAG GTG CTT CCA G
	AdvilCreERT2_3	GCG ATC CCT GAA CAT GTC CAT C
<i>Cdx2<sup>Cre</sup></i>	forward	CTC GAC GTC TCC AAC CAT TG
	reverse	ATC TTC AGG TTC TGC GGG AA
<i>Cntnap2 null</i>	wild type forward	TGC CCT CCT AGA AAG TAA ATG C T
	mutant forward	GCC AGA GGC CAC TTG TGT AG
	common reverse	TCC TCT CTT CAT GCA CAC TAT GA
<i>Fmr1 null</i>	mutant forward	CAC GAG ACT AGT GAG ACG TG
	wild type forward	TGT GAT AGA ATA TGC AGC ATG TGA
	common reverse	CTT CTG GCA CCT CCA GCT T
<i>Gabrb3<sup>floxed</sup></i>	forward	ATT CGC CTG AGA CCC GAC T
	reverse	GTT CAT CCC CAC GCA GAC
<i>Hcn1 null</i>	mutant forward	AGA GAA ATC ATT CCC CGT GA
	wild type forward	CAC CTG CTA CGC AAT GTT TG
	common reverse	ATT GGG CAC TAC ACG CTA GG
<i>Mecp2<sup>floxed</sup></i>	forward	TGG TAA AGA CCC ATG TGA CCC AAG
	reverse	GGC TTG CCA CAT GAC AAG AC
	post-Cre excision	TCC ACC TAG CCT GCC TGT ACT TTG
<i>Mecp2<sup>R306C</sup></i>	forward	GGA TTG TGG AAA AGC CAG
	reverse	ATG ACC TGG GCA GAT GTG GTA G
<i>Mecp2<sup>STOP</sup></i>	forward common	AAC AGT GCC AGC TGC TCT TC
	wild type reverse	CTG TAT CCT TGG GTC AAG CTG
	mutant reverse	GCC AGA GGC CAC TTG TGT AG
<i>Shank3<sup>floxed</sup></i>	pFW (Sh3cKO) Gen 1a	CAGCATTTATACCTGACTGTGAAGC
	pRV (Sh3cKO) Gen 3b	GGGAGTAGAGCTCAGATAACC
<i>Shank3B null</i>	common forward	GAG ACT GAT CAG CGC AGT TG
	wild type reverse	TGA CAT AAT CGC TGG CAA AG
	mutant reverse	GCT ATA CGA AGT TAT GTC GAC TAG G
<i>Shank3<sup>Δex4-9</sup></i>	common forward	TGG GAT GTG AGA GTG ACC AG
	wild type reverse	AGG AGG TCA GTG GCG TTG T
	mutant reverse	AAT TCT GTT CAG TCC ACA CAG G
<i>Shank3<sup>FX</sup></i>	common forward	CGT TTG ACA CAC ATA AGC ACC
	wild type reverse	CTC CAC CTA GCT GAA TTT CCC



**Table S8. Oligonucleotide List. Related to Key Resources Table.**

	FX mutant reverse	GCT GAC ATC ACA TTG CTG CC
	FX rescue forward	CGT TTG ACA CAC ATA AGC ACC
	FX rescue reverse (wildtype rvs)	CTC CAC CTA GCT GAA TTT CCC
<i>16<sup>p11.2df</sup> null</i>	mutant forward	ACC TGG TAT TGA ATG CTT GC
	wild type forward	CCT GAG CCT CTT GAG TGT CC
	mutant reverse	TGG TAT CTC CAT AAG ACA GAA TGC
	wild type reverse	GTC GGT TCA GGT GGT AGA CG