

Elevated plus maze

Percent time in open arms		Means \pm SEM			
		WT	HET	t ratio	p value
Control males	25.91 \pm 4.78	26.31 \pm 3.12	0.06472	17	0.964152
ELSD males	24.98 \pm 4.98	23.19 \pm 5.37	0.2429	19	0.964152
PASD males	33.40 \pm 6.84	38.49 \pm 2.86	0.6866	14	0.877635
Control females	17.69 \pm 1.94	22.49 \pm 6.21	0.8093	16	0.43021
** ELSD females	23.83 \pm 2.75	36.56 \pm 2.13	3.51	16	0.008685
# PASD females	32.75 \pm 4.40	46.10 \pm 4.40	2.148	22	0.084182

Percent open arm entrances		Means \pm SEM			
		WT	HET	t ratio	p value
Control males	36.04 \pm 3.47	34.51 \pm 2.28	0.3363	17	0.766385
ELSD males	32.95 \pm 2.40	30.21 \pm 2.94	0.7133	19	0.766385
PASD males	37.13 \pm 5.02	41.75 \pm 1.13	0.8985	14	0.766385
Control females	31.08 \pm 3.31	31.61 \pm 1.77	0.1317	16	0.896834
# ELSD females	34.55 \pm 2.41	43.38 \pm 2.41	2.557	16	0.061947
PASD females	37.20 \pm 2.93	42.44 \pm 2.87	1.28	22	0.38217

Open field test

Total distance traveled (cm)		Means \pm SEM			
		WT	HET	t ratio	p value
Control males	6117.64 \pm 519.83	6812.57 \pm 407.81	0.9489	16	0.356784
** ELSD males	7675.30 \pm 687.29	5211.27 \pm 478.36	2.988	19	0.022514
PASD males	7402.25 \pm 584.49	6329.25 \pm 167.60	1.765	14	0.188948
Control females	5804.90 \pm 505.58	6679.00 \pm 458.05	1.25	16	0.405805
ELSD females	6831.00 \pm 846.82	6798.75 \pm 530.14	0.03132	15	0.975429
PASD females	8946.58 \pm 845.93	6863.42 \pm 480.23	2.142	22	0.125066

Prepulse inhibition

Percent startle inhibition		Means \pm SEM				Sidak's multiple comparisons test	
		WT	HET	2-way ANOVA			
				Factor	% variation	F(DFn, DFd)	p value
Control males				dB x genotype	0.2831	F (4, 68) = 0.2549	0.9057
Startle tone (dB)	74	12.27 \pm 11.57	13.05 \pm 6.73	dB trial	65.31	F (1,59, 26.67) = 58.81	<0.0001
	78	55.21 \pm 3.30	48.98 \pm 4.62	genotype	0.2335	F (1, 17) = 0.3079	0.5862
	82	65.48 \pm 3.92	66.49 \pm 3.60	subject	12.9	F (17, 68) = 2.733	0.0018
	86	76.72 \pm 3.58	72.65 \pm 3.66				
	90	81.45 \pm 2.93	75.70 \pm 3.05				
ELSD males				Factor	% variation	F(DFn, DFd)	p value
	74	28.69 \pm 5.09	18.16 \pm 9.87	dB x genotype	0.2965	F (4, 76) = 0.4587	0.7658
	78	65.11 \pm 3.87	46.78 \pm 8.62	dB trial	51.84	F (2,234, 42.44) = 80.18	<0.0001
	82	78.92 \pm 2.75	59.67 \pm 6.59	genotype	8.412	F (1, 19) = 5.847	0.0258
	86	84.00 \pm 2.70	67.71 \pm 3.89	subject	27.34	F (19, 76) = 8.902	<0.0001
	*90	86.19 \pm 1.26	69.75 \pm 3.98				
PASD males				Factor	% variation	F(DFn, DFd)	p value
	74	22.40 \pm 7.47	25.41 \pm 10.32	dB x genotype	2.183	F (4, 56) = 2.374	0.063
	78	42.78 \pm 8.39	66.73 \pm 6.99	dB trial	47.33	F (1,909, 26.72) = 51.48	<0.0001
	82	63.65 \pm 8.94	73.04 \pm 5.32	genotype	2.788	F (1, 14) = 1.121	0.3077
	86	67.36 \pm 7.45	77.08 \pm 5.23	subject	34.83	F (14, 56) = 10.82	<0.0001
	90	76.48 \pm 5.87	76.74 \pm 4.15				
Control females				Factor	% variation	F(DFn, DFd)	p value
	74	25.86 \pm 7.56	28.38 \pm 5.30	dB x genotype	1.26	F (4, 64) = 1.107	0.3609
	78	62.79 \pm 3.91	52.80 \pm 6.68	dB trial	53.81	F (2,093, 33.48) = 47.29	<0.0001
	82	68.70 \pm 3.54	57.49 \pm 5.99	genotype	1.74	F (1, 16) = 1.209	0.2879
	86	71.20 \pm 3.64	67.46 \pm 3.28	subject	23.03	F (16, 64) = 5.060	<0.0001
	90	74.65 \pm 3.01	68.21 \pm 4.64				
ELSD females				Factor	% variation	F(DFn, DFd)	p value
	74	30.28 \pm 10.84	11.98 \pm 9.72	dB x genotype	1.577	F (4, 64) = 1.896	0.122
	78	59.81 \pm 6.21	60.11 \pm 5.13	dB trial	51.54	F (1,53, 24.85) = 61.94	<0.0001
	82	66.80 \pm 6.62	67.88 \pm 5.80	genotype	0.7215	F (1, 16) = 0.3407	0.5676
	86	78.05 \pm 3.90	72.46 \pm 6.42	subject	33.88	F (16, 64) = 10.18	<0.0001
	90	77.42 \pm 4.73	75.84 \pm 5.05				
PASD females				Factor	% variation	F(DFn, DFd)	p value
	*74	41.56 \pm 8.21	11.48 \pm 6.32	dB x genotype	3.45	F (4, 88) = 4.895	0.0013
	78	64.63 \pm 3.36	59.28 \pm 3.20	dB trial	59.59	F (1,363, 29.98) = 84.56	<0.0001
	82	73.93 \pm 3.15	65.39 \pm 3.47	genotype	5.679	F (1, 22) = 7.916	0.0101
	86	80.34 \pm 1.70	72.93 \pm 3.14	subject	15.78	F (22, 88) = 4.072	<0.0001
	90	83.07 \pm 1.88	75.45 \pm 2.69				

Sociability

Seconds in proximity		Means ± SEM		Paired t-test with Holm-Šídák correction			
		non-social	stranger 1	t	df	p value	
Control males	WT	75.17 ± 8.25	145.26 ± 18.84	WT	3.161	10	0.020179
	HET	81.18 ± 8.68	161.48 ± 24.76	HET	2.544	7	0.038447
ELSD males	WT	78.54 ± 7.35	157.93 ± 12.21	WT	5.723	9	0.000571
	HET	92.20 ± 5.54	117.01 ± 11.81	HET	1.674	10	0.125092
PASD males	WT	71.30 ± 5.61	158.71 ± 11.65	WT	7.071	7	0.000201
	HET	68.65 ± 6.41	144.16 ± 7.28	HET	7.881	7	0.000201
Control females	WT	72.93 ± 7.83	152.36 ± 15.69	WT	4.812	8	0.001334
	HET	75.69 ± 7.29	147.53 ± 6.10	HET	9.265	7	0.000071
ELSD females	WT	80.74 ± 6.40	168.75 ± 19.48	WT	3.823	9	0.00812
	HET	99.78 ± 15.06	182.89 ± 17.83	HET	2.791	7	0.026867
PASD females	WT	73.63 ± 7.00	111.64 ± 10.06	WT	2.807	11	0.017072
	HET	76.98 ± 6.84	132.80 ± 9.93	HET	4.263	11	0.002672

Social novelty preference

Seconds in proximity		Means ± SEM		Paired t-test with Holm-Šídák correction			
		stranger 1	stranger 2	t	df	p value	
Control males	WT	68.88 ± 10.20	125.45 ± 9.88	WT	4.588	10	0.001995
	HET	64.55 ± 10.04	133.73 ± 17.43	HET	2.663	7	0.032307
ELSD males	WT	71.78 ± 10.06	137.16 ± 14.21	WT	3.581	9	0.005919
	HET	56.25 ± 8.98	167.87 ± 16.85	HET	5.955	10	0.000281
PASD males	WT	62.48 ± 7.41	120.98 ± 14.94	WT	3.955	7	0.010959
	HET	68.65 ± 10.37	95.31 ± 5.22	HET	2.003	7	0.085182
Control females	WT	64.03 ± 10.73	123.38 ± 10.09	WT	3.93	8	0.008699
	HET	98.38 ± 16.17	121.80 ± 13.61	HET	1.057	7	0.325736
ELSD females	WT	64.92 ± 7.41	118.66 ± 8.99	WT	5.147	9	0.001211
	HET	75.29 ± 13.98	111.31 ± 9.76	HET	1.677	7	0.137533
PASD females	WT	69.54 ± 10.67	90.07 ± 12.13	WT	1.092	11	0.298247
	HET	82.92 ± 13.20	117.41 ± 18.03	HET	1.781	11	0.194496