#### Supplemental Material for:

Heterotypic stressors unmask behavioral influences of PMAT deficiency in mice

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#### Phase 1 Cued Females

#### Training

	F statistic	р	partial η²
Genotype	F(1,35)=0.531	0.471	0.015
Swim	F(1,35)=0.003	0.957	0.000
Genotype × Swim	F(1,35)=0.881	0.354	0.025
Time	F(3.122,109.266)=183.4	<0.001	0.840
Time × Genotype	F(3.122,109.266)=0.676	0.574	0.019
Time × Swim	F(3.122,109.266)=0.585	0.633	0.016
Time × Genotype × Swim	F(3.122,109.266)=0.964	0.415	0.027

#### Expression Testing & Extinction Training

	F statistic	р	partial η²
Genotype	F(1,35)=0.471	0.497	0.013
Swim	F(1,35)=0.455	0.504	0.013
Genotype × Swim	F(1,35)=0.751	0.392	0.021
Time	F(9.728,340.487)=8.714	<0.001	0.199
Time × Genotype	F(9.728,340.487)=3.865	<0.001	0.099
Time × Swim	F(9.728,340.487)=0.802	0.624	0.022
Time × Genotype × Swim	F(9.728,340.487)=1.138	0.334	0.031

#### Extinction Retention Testing & More Extinction Training

	F statistic	р	partial η²
Genotype	F(1,35)=1.000	0.324	0.028
Swim	F(1,35)=0.477	0.494	0.013
Genotype × Swim	F(1,35)=0.508	0.481	0.014
Time	F(9.774,342.097)=8.024	<0.001	0.186
Time × Genotype	F(9.774,342.097)=1.034	0.414	0.029
Time × Swim	F(9.774,342.097)=1.523	0.131	0.042
Time × Genotype × Swim	F(9.774,342.097)=1.191	0.296	0.033

#### Context Fear Expression - Timecourse F statistic partial $\eta^2$ р Genotype F(1,35)=0.001 0.976 0.000 F(1,35)=0.110 Swim 0.742 0.003 Genotype × Swim F(1,35)=0.027 0.871 0.001 Time F(6.926,242.403)=5.297 <0.001 0.131 Time × Genotype F(6.926,242.403)=0.641 0.720 0.018 F(6.926,242.403)=0.586 Time x Swim 0.765 0.016

Time × Genotype × Swim	F(6.926,242.403)=0.531	0.809	0.015

### Cued Fear Renewal

	F statistic	р	partial η²
Genotype	F(1,35)=0.009	0.926	0.000
Swim	F(1,35)=0.075	0.786	0.002
Genotype × Swim	F(1,35)=0.421	0.521	0.012
Time	F(3.712,129.923)=12.40	<0.001	0.262
Time × Genotype	F(3.712,129.923)=0.574	0.670	0.016
Time × Swim	F(3.712,129.923)=0.745	0.554	0.021
Time × Genotype × Swim	F(3.712,129.923)=0.184	0.937	0.005

Supplemental Table 2. Repeated measures ANOVAs of Phase 1 Cued Male data.

### Phase 1 Cued Males

### Training

	F statistic	р	partial η²
Genotype	F(1,54)=3.148	0.082	0.055
Swim	F(1,54)=1.825	0.182	0.033
Genotype × Swim	F(1,54)=0.166	0.685	0.003
Time	F(3.156,170.433)=205.7	<0.001	0.792
Time × Genotype	F(3.156,170.433)=1.554	0.200	0.028
Time × Swim	F(3.156,170.433)=0.475	0.710	0.009
Time × Genotype × Swim	F(3.156,170.433)=0.477	0.708	0.009

### Expression Testing & Extinction Training

	F statistic	р	partial η²
Genotype	F(1,54)=1.051	0.310	0.019
Swim	F(1,54)=0.074	0.787	0.001
Genotype × Swim	F(1,54)=2.257	0.139	0.040
Time	F(10.594,572.057)=9.786	<0.001	0.153
Time × Genotype	F(10.594,572.057)=0.538	0.872	0.010
Time × Swim	F(10.594,572.057)=0.554	0.860	0.010
Time × Genotype × Swim	F(10.594,572.057)=0.718	0.716	0.013

### Extinction Retention Testing & More Extinction Training

	F statistic	р	partial η²
Genotype	F(1,53)=0.173	0.679	0.003
Swim	F(1,53)=0.035	0.852	0.001
Genotype × Swim	F(1,53)=1.543	0.220	0.028
Time	F(10.125,536.64)=11.09	<0.001	0.173
Time × Genotype	F(10.125,536.64)=1.004	0.439	0.019
Time × Swim	F(10.125,536.64)=1.339	0.205	0.025
Time × Genotype × Swim	F(10.125,536.64)=0.633	0.788	0.012

	F statistic	р	partial η²
Genotype	F(1,49)=0.250	0.619	0.005
Swim	F(1,49)=0.684	0.412	0.014
Genotype × Swim	F(1,49)=1.147	0.289	0.023
Time	F(10.889,533.558)=11.58	<0.001	0.191
Time × Genotype	F(10.889,533.558)=0.631	0.801	0.013
Time × Swim	F(10.889,533.558)=1.197	0.286	0.024

Time × Genotype × Swim	F(10.889,533.558)=1.319	0.210	0.026
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### Cued Fear Renewal

	F statistic	р	partial η²
Genotype	F(1,49)=0.033	0.856	0.001
Swim	F(1,49)=0.874	0.354	0.018
Genotype × Swim	F(1,49)=0.081	0.777	0.002
Time	F(3.82,187.198)=11.17	<0.001	0.186
Time × Genotype	F(3.82,187.198)=0.190	0.938	0.004
Time × Swim	F(3.82,187.198)=0.182	0.942	0.004
Time × Genotype × Swim	F(3.82,187.198)=0.999	0.407	0.020

Supplemental Table 3. Repeated measures ANOVAs of Phase 1 Context Female data.

## Phase 1 Context Females

### Training

-	F statistic	р	partial η²
Genotype	F(1,34)=0.904	0.348	0.026
Swim	F(1,34)=0.014	0.907	0.000
Genotype × Swim	F(1,34)=0.061	0.806	0.002
Time	F(3.626,123.285)=125.3	<0.001	0.787
Time × Genotype	F(3.626,123.285)=0.252	0.893	0.007
Time × Swim	F(3.626,123.285)=0.818	0.506	0.023
Time x Genotype x Swim	F(3.626,123.285)=0.412	0.782	0.012

	F statistic	р	partial η²
Genotype	F(1,34)=2.936	0.096	0.079
Swim	F(1,34)=0.540	0.468	0.016
Genotype × Swim	F(1,34)=0.555	0.462	0.016
Time	F(9.138,310.686)=8.874	<0.001	0.207
Time × Genotype	F(9.138,310.686)=1.369	0.200	0.039
Time × Swim	F(9.138,310.686)=0.494	0.881	0.014
Time × Genotype × Swim	F(9.138,310.686)=0.984	0.454	0.028

Supplemental Table 4. Repeated measures ANOVAs of Phase 1 Context Male data.

## Phase 1 Context Males

### Training

	F statistic	р	partial η²
Genotype	F(1,39)=0.553	0.462	0.014
Swim	F(1,39)=2.221	0.144	0.054
Genotype × Swim	F(1,39)=0.081	0.777	0.002
Time	F(3.226,125.803)=12.92	<0.001	0.249
Time × Genotype	F(3.226,125.803)=1.424	0.237	0.035
Time × Swim	F(3.226,125.803)=0.441	0.738	0.011
Time × Genotype × Swim	F(3.226,125.803)=0.158	0.934	0.004

	F statistic	р	partial η²
Genotype	F(1,39)=4.555	0.039	0.105
Swim	F(1,39)=0.027	0.871	0.001
Genotype × Swim	F(1,39)=1.280	0.265	0.032
Time	F(9.546,372.282)=13.94	<0.001	0.263
Time × Genotype	F(9.546,372.282)=1.142	0.331	0.028
Time × Swim	F(9.546,372.282)=1.319	0.221	0.033
Time × Genotype × Swim	F(9.546,372.282)=0.840	0.586	0.021

Supplemental Table 5. Repeated measures ANOVAs of Phase 2 Cued Female data.

### Phase 2 Cued Females

### Training

	F statistic	р	partial η²
Genotype	F(1,41)=0.676	0.416	0.016
Swim	F(1,41)=1.273	0.266	0.030
Genotype × Swim	F(1,41)=0.922	0.343	0.022
Time	F(3.605,147.811)=159.1	<0.001	0.795
Time × Genotype	F(3.605,147.811)=0.679	0.593	0.016
Time × Swim	F(3.605,147.811)=0.970	0.420	0.023
Time × Genotype × Swim	F(3.605,147.811)=0.169	0.942	0.004

### Expression Testing & Extinction Training

	F statistic	р	partial η²
Genotype	F(1,41)=0.299	0.587	0.007
Swim	F(1,41)=0.036	0.850	0.001
Genotype × Swim	F(1,41)=0.951	0.335	0.023
Time	F(8.741,358.393)=6.623	<0.001	0.139
Time × Genotype	F(8.741,358.393)=0.852	0.566	0.020
Time × Swim	F(8.741,358.393)=0.726	0.681	0.017
Time × Genotype × Swim	F(8.741,358.393)=0.437	0.911	0.011

### Extinction Retention Testing & More Extinction Training

	F statistic	р	partial η²
Genotype	F(1,35)=0.400	0.531	0.011
Swim	F(1,35)=0.093	0.763	0.003
Genotype × Swim	F(1,35)=0.009	0.924	0.000
Time	F(8.864,310.252)=11.15	<0.001	0.242
Time × Genotype	F(8.864,310.252)=0.590	0.803	0.017
Time × Swim	F(8.864,310.252)=0.822	0.594	0.023
Time × Genotype × Swim	F(8.864,310.252)=1.133	0.339	0.031

	F statistic	р	partial η²
Genotype	F(1,41)=0.256	0.615	0.006
Swim	F(1,41)=0.498	0.484	0.012
Genotype × Swim	F(1,41)=0.060	0.807	0.001
Time	F(10.762,441.257)=5.923	<0.001	0.126
Time × Genotype	F(10.762,441.257)=1.001	0.444	0.024
Time × Swim	F(10.762,441.257)=1.15	0.321	0.027

Time × Genotype × Swim	F(10.762,441.257)=1.064	0.389	0.025
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## Cued Fear Renewal

	F statistic	р	partial η²
Genotype	F(1,39)=0.637	0.430	0.016
Swim	F(1,39)=0.050	0.825	0.001
Genotype × Swim	F(1,39)=0.066	0.799	0.002
Time	F(3.57,139.219)=11.74	<0.001	0.231
Time × Genotype	F(3.57,139.219)=0.690	0.584	0.017
Time × Swim	F(3.57,139.219)=3.592	0.011	0.084
Time × Genotype × Swim	F(3.57,139.219)=0.282	0.870	0.007

Supplemental Table 6. Repeated measures ANOVAs of Phase 2 Cued Male data.

#### Phase 2 Cued Males

#### Training

	F statistic	р	partial η²
Genotype	F(1,36)=0.033	0.858	0.001
Swim	F(1,36)=0.140	0.710	0.004
Genotype × Swim	F(1,36)=1.226	0.276	0.033
Time	F(2.700,97.197)=123.1	<0.001	0.774
Time × Genotype	F(2.700,97.197)=1.581	0.203	0.042
Time × Swim	F(2.700,97.197)=0.872	0.449	0.024
Time × Genotype × Swim	F(2.700,97.197)=0.978	0.400	0.026

### Expression Testing & Extinction Training

	F statistic	р	partial η²
Genotype	F(1,30)=0.196	0.661	0.006
Swim	F(1,30)=1.649	0.209	0.052
Genotype × Swim	F(1,30)=0.725	0.401	0.024
Time	F(8.933,267.996)=10.55	<0.001	0.260
Time × Genotype	F(8.933,267.996)=1.029	0.417	0.033
Time × Swim	F(8.933,267.996)=0.987	0.451	0.032
Time × Genotype × Swim	F(8.933,267.996)=1.631	0.107	0.052

## Extinction Retention Testing & More Extinction Training

	F statistic	р	partial η²
Genotype	F(1,38)=6.914	0.012	0.154
Swim	F(1,38)=0.052	0.820	0.001
Genotype × Swim	F(1,38)=1.033	0.316	0.026
Time	F(7.79,296.005)=8.583	<0.001	0.184
Time × Genotype	F(7.79,296.005)=0.955	0.470	0.025
Time × Swim	F(7.79,296.005)=1.211	0.293	0.031
Time × Genotype × Swim	F(7.79,296.005)=1.249	0.272	0.032

	F statistic	р	partial η²
Genotype	F(1,37)=4.175	0.048	0.101
Swim	F(1,37)=0.273	0.604	0.007
Genotype × Swim	F(1,37)=2.728	0.107	0.069
Time	F(9.633,356.405)=5.218	<0.001	0.124
Time × Genotype	F(9.633,356.405)=0.390	0.947	0.010
Time × Swim	F(9.633,356.405)=1.470	0.152	0.038

Time × Genotype × Swim	F(9.633,356.405)=1.183	0.302	0.031
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## Cued Fear Renewal

	F statistic	р	partial η²
Genotype	F(1,34)=0.164	0.688	0.005
Swim	F(1,34)=1.950	0.172	0.054
Genotype × Swim	F(1,34)=0.625	0.435	0.018
Time	F(3.835,130.383)=24.30	<0.001	0.417
Time × Genotype	F(3.835,130.383)=0.851	0.492	0.024
Time × Swim	F(3.835,130.383)=0.898	0.464	0.026
Time × Genotype × Swim	F(3.835,130.383)=1.335	0.262	0.038

Supplemental Table 7. Repeated measures ANOVAs of Phase 2 Context Female data.

## Phase 2 Context Females

### Training

	F statistic	р	partial η²
Genotype	F(1,34)=0.713	0.404	0.021
Swim	F(1,34)=0.346	0.560	0.010
Genotype × Swim	F(1,34)=0.065	0.800	0.002
Time	F(3.029,102.984)=76.52	<0.001	0.692
Time × Genotype	F(3.029,102.984)=1.116	0.346	0.032
Time × Swim	F(3.029,102.984)=1.963	0.124	0.055
Time × Genotype × Swim	F(3.029,102.984)=0.978	0.407	0.028

	F statistic	р	partial η²
Genotype	F(1,30)=0.362	0.552	0.012
Swim	F(1,30)=2.066	0.161	0.064
Genotype × Swim	F(1,30)=1.549	0.223	0.049
Time	F(9.209,276.265)=8.773	<0.001	0.226
Time × Genotype	F(9.209,276.265)=1.094	0.367	0.035
Time × Swim	F(9.209,276.265)=0.985	0.454	0.032
Time × Genotype × Swim	F(9.209,276.265)=1.251	0.263	0.040

Supplemental Table 8. Repeated measures ANOVAs of Phase 2 Context Male data.

## Phase 2 Context Males

### Training

	F statistic	р	partial η²
Genotype	F(1,33)=0.042	0.839	0.001
Swim	F(1,33)=0.075	0.786	0.002
Genotype × Swim	F(1,33)=0.180	0.674	0.005
Time	F(3.284,108.38)=66.40	<0.001	0.668
Time × Genotype	F(3.284,108.38)=0.190	0.917	0.006
Time × Swim	F(3.284,108.38)=0.979	0.411	0.029
Time × Genotype × Swim	F(3.284,108.38)=1.062	0.372	0.031

	F statistic	р	partial η²
Genotype	F(1,33)=0.524	0.474	0.016
Swim	F(1,33)=0.024	0.877	0.001
Genotype × Swim	F(1,33)=3.400	0.074	0.093
Time	F(8.625,284.634)=14.87	<0.001	0.311
Time × Genotype	F(8.625,284.634)=1.447	0.171	0.042
Time × Swim	F(8.625,284.634)=1.160	0.322	0.034
Time × Genotype × Swim	F(8.625,284.634)=0.763	0.646	0.023

Supplemental Table 9. Three-way ANOVAs of log-transformed corticosterone levels taken 2 h following final testing (Phase 1 – swim stress; Phase 2 – context testing & cued fear renewal).

Phase 1 Cued			
	F statistic	р	partial η²
Genotype	F(1,82)=0.962	0.330	0.012
Sex	F(1,82)=17.77	<0.001	0.178
Swim	F(1,82)=0.301	0.584	0.004
Genotype × Sex	F(1,82)=0.119	0.731	0.001
Genotype × Swim	F(1,82)=0.170	0.681	0.002
Sex × Swim	F(1,82)=0.000	0.992	0.000
Genotype × Sex × Swim	F(1,82)=0.639	0.426	0.008

### Phase 1 Context

	F statistic	р	partial $\eta^2$
Genotype	F(1,72)=2.007	0.161	0.027
Sex	F(1,72)=45.11	<0.001	0.385
Swim	F(1,72)=8.382	0.005	0.104
Genotype × Sex	F(1,72)=4.839	0.031	0.063
Genotype × Swim	F(1,72)=0.239	0.626	0.003
Sex × Swim	F(1,72)=5.938	0.017	0.076
Genotype × Sex × Swim	F(1,72)=6.029	0.016	0.077

## Phase 2 Cued

	F statistic	р	partial η <sup>2</sup>
Genotype	F(1,79)=2.285	0.135	0.028
Sex	F(1,79)=17.40	<0.001	0.180
Swim	F(1,79)=0.967	0.328	0.012
Genotype × Sex	F(1,79)=1.157	0.285	0.014
Genotype × Swim	F(1,79)=1.748	0.190	0.022
Sex × Swim	F(1,79)=0.390	0.534	0.005
Genotype × Sex × Swim	F(1,79)=1.235	0.270	0.015

### Phase 2 Context

	F statistic	р	partial η²
Genotype	F(1,63)=0.001	0.979	0.000
Sex	F(1,63)=1.437	0.235	0.022
Swim	F(1,63)=0.670	0.416	0.011
Genotype × Sex	F(1,63)=0.186	0.668	0.003
Genotype × Swim	F(1,63)=4.377	0.040	0.065
Sex × Swim	F(1,63)=0.100	0.753	0.002

Genotype × Sex × Swim	F(1,63)=0.282	0.597	0.004

Supplemental Table 10. Two-way ANOVAs for behaviors during swim stress in Phase 1 Cued mice.

## Phase 1 Cued

### <u>Swimming</u>

-	F statistic	p	partial η²
Genotype	F(1,39)=0.010	0.922	0.000
Sex	F(1,39)=1.825	0.185	0.045
Genotype × Sex	F(1,39)=0.419	0.521	0.011

### Immobility

	F statistic	р	partial η²
Genotype	F(1,39)=0.801	0.376	0.020
Sex	F(1,39)=0.065	0.801	0.002
Genotype × Sex	F(1,39)=0.570	0.455	0.014

## <u>Climbing</u>

	F statistic	р	partial η²
Genotype	F(1,39)=0.403	0.529	0.010
Sex	F(1,39)=0.626	0.434	0.016
Genotype × Sex	F(1,39)=0.134	0.716	0.003

	F statistic	р	partial η²
Genotype	F(1,39)=0.356	0.554	0.009
Sex	F(1,39)=1.051	0.312	0.026
Genotype × Sex	F(1,39)=0.003	0.957	0.000

Supplemental Table 11. Two-way ANOVAs for behaviors during swim stress in Phase 1 Context mice.

## Phase 1 Context

Swimming			
_	F statistic	р	partial η²
Genotype	F(1,36)=0.013	0.911	0.000
Sex	F(1,36)=4.250	0.047	0.106
Genotype × Sex	F(1,36)=5.572	0.024	0.134

### Immobility

	F statistic	р	partial η²
Genotype	F(1,36)=0.283	0.598	0.008
Sex	F(1,36)=0.181	0.673	0.005
Genotype × Sex	F(1,36)=1.360	0.251	0.036

## <u>Climbing</u>

	F statistic	р	partial η²
Genotype	F(1,36)=0.765	0.388	0.021
Sex	F(1,36)=1.538	0.223	0.041
Genotype × Sex	F(1,36)=0.189	0.667	0.005

	F statistic	р	partial η²
Genotype	F(1,36)=0.151	0.700	0.004
Sex	F(1,36)=0.259	0.614	0.007
Genotype × Sex	F(1,36)=2.022	0.164	0.053

Supplemental Table 12. Two-way ANOVAs for behaviors during swim stress in Phase 2 Cued mice.

## Phase 2 Cued

### <u>Swimming</u>

_	F statistic	р	partial η²
Genotype	F(1,42)=1.062	0.309	0.025
Sex	F(1,42)=0.048	0.828	0.001
Genotype × Sex	F(1,42)=0.271	0.605	0.006

### Immobility

	F statistic	р	partial η²
Genotype	F(1,42)=3.221	0.080	0.071
Sex	F(1,42)=0.065	0.800	0.002
Genotype × Sex	F(1,42)=1.161	0.287	0.027

## <u>Climbing</u>

	F statistic	р	partial η²
Genotype	F(1,42)=3.024	0.089	0.067
Sex	F(1,42)=0.867	0.357	0.020
Genotype × Sex	F(1,42)=1.483	0.230	0.034

	F statistic	р	partial η²
Genotype	F(1,42)=4.679	0.036	0.100
Sex	F(1,42)=2.722	0.106	0.061
Genotype × Sex	F(1,42)=0.030	0.863	0.001

Supplemental Table 13. Two-way ANOVAs for behaviors during swim stress in Phase 2 Context mice.

## Phase 2 Context

Swimming			
	F statistic	р	partial η²
Genotype	F(1,34)=0.694	0.411	0.020
Sex	F(1,34)=4.996	0.032	0.128
Genotype × Sex	F(1,34)=0.003	0.954	0.000

### Immobility

	F statistic	р	partial η²
Genotype	F(1,34)=1.512	0.227	0.043
Sex	F(1,34)=10.09	0.003	0.229
Genotype × Sex	F(1,34)=0.013	0.909	0.000

## <u>Climbing</u>

	F statistic	р	partial η²
Genotype	F(1,34)=0.930	0.342	0.027
Sex	F(1,34)=5.611	0.024	0.142
Genotype × Sex	F(1,34)=0.020	0.890	0.001

	F statistic	р	partial η²
Genotype	F(1,34)=0.845	0.364	0.024
Sex	F(1,34)=18.76	<0.001	0.356
Genotype × Sex	F(1,34)=0.087	0.770	0.003

Supplemental Table S14. Two-way ANOVA of log-transformed corticosterone levels in mice 30 min following an acute swim stressor. Refer to Supplemental Figure S1.

Log-transformed Corticosterone Levels			
	F statistic	р	partial η²
Genotype	F(2,62)=0.285	0.753	0.009
Sex	F(1,62)=0.368	0.546	0.006
Genotype × Sex	F(2,62)=0.706	0.497	0.022

Supplemental Table S15. Two-way ANOVAs of average of context fear expression during minutes 2-6. Refer to Supplemental Figures S2, S3.

Phase 1 Cued			
	F statistic	р	partial η²
Genotype	F(1,88)=0.006	0.938	0.000
Sex	F(1,88)=1.056	0.307	0.012
Genotype × Sex	F(1,88)=0.196	0.659	0.002

Phase 1 Context			
	F statistic	р	partial η²
Genotype	F(1,77)=5.646	0.020	0.068
Sex	F(1,77)=1.238	0.269	0.016
Genotype × Sex	F(1,77)=0.646	0.424	0.008

Phase 2 Cued			
	F statistic	р	partial η²
Genotype	F(1,88)=0.006	0.938	0.000
Sex	F(1,88)=1.056	0.307	0.012
Genotype × Sex	F(1,88)=0.196	0.659	0.002

Phase 2 Context			
	F statistic	р	partial η²
Genotype	F(1,88)=0.006	0.938	0.000
Sex	F(1,88)=1.056	0.307	0.012
Genotype × Sex	F(1,88)=0.196	0.659	0.002

Supplemental Table S16. Two-way ANOVAs of fecal boli during swim stress. Refer to Supplemental Figure S4.

Phase 1 Cued			
	F statistic	р	partial η²
Genotype	F(1,45)=0.013	0.910	0.000
Sex	F(1,45)=0.015	0.902	0.000
Genotype × Sex	F(1,45)=0.604	0.441	0.013

Phase 1 Context				
	F statistic	р	partial η²	
Genotype	F(1,36)=0.032	0.858	0.001	
Sex	F(1,36)=2.235	0.144	0.058	
Genotype × Sex	F(1,36)=0.122	0.729	0.003	

Phase 2 Cued			
	F statistic	р	partial η²
Genotype	F(1,43)=1.661	0.204	0.037
Sex	F(1,43)=6.354	0.016	0.129
Genotype × Sex	F(1,43)=0.042	0.839	0.001

Phase 2 Contex	t		
	F statistic	р	partial η²
Genotype	F(1,34)=0.297	0.589	0.009
Sex	F(1,34)=1.126	0.296	0.032
Genotype × Sex	F(1,34)=1.126	0.296	0.032

Supplemental Table 17. Specific numbers of mice graphed in each Figure.

Figure 2: Phase 1 Cued	Females		Males	
	Wildtype	Heterozygous	Wildtype	Heterozygous
Training	20	19	25	33
Cued Expression Testing &	20	19	25	33
Extinction Training				
Extinction Retention Testing	20	19	25	32
Context Fear Expression	20	19	23	30
Cued Fear Renewal	20	19	22	31

Figure 3: Phase 1 Context	Females		Males	
	Wildtype	Heterozygous	Wildtype	Heterozygous
Training	20	18	20	23
Context Fear Expression	20	18	20	23

Figures 4 & 5: Phase 2 Cued	Females		Males	
Training	Wildtype	Heterozygous	Wildtype	Heterozygous
No Swim	11	11	9	11
Swim	12	11	9	11
Cued Expression Testing &				
Extinction Training				
No Swim	11	11	7	10
Swim	12	11	7	10
Extinction Retention Testing				
No Swim	10	9	9	11
Swim	10	10	10	12
Context Fear Expression				
No Swim	11	11	9	11
Swim	12	11	10	12
Cued Fear Renewal				
No Swim	10	11	7	10
Swim	11	11	10	12

Figures 6 & 7: Phase 2 Context	Females		Males	
Training	Wildtype	Heterozygous	Wildtype	Heterozygous
No Swim	9	10	9	9
Swim	10	9	9	10
Context Fear Expression				
No Swim	9	8	9	9
Swim	8	9	9	10

Figure 8: Corticosterone	Females		Males	
Phase 1 Cued	Wildtype	Heterozygous	Wildtype	Heterozygous
No Swim	10	8	12	15
Swim	10	10	15	16
Phase 1 Context				
No Swim	10	8	10	11
Swim	10	10	10	11

Figure 9: Corticosterone		Females		Males	
Phase 1 Cued		Wildtype	Heterozygous	Wildtype	Heterozygous
	No Swim	11	11	10	11
	Swim	12	11	12	12
Phase 1 Context					
	No Swim	9	10	9	9
	Swim	10	9	9	10

Figures 10 & 11: Swim behaviors	Females		Males	
	Wildtype	Heterozygous	Wildtype	Heterozygous
Phase 1 Cued	10	10	9	14
Phase 1 Context	10	10	9	11
Phase 2 Cued	12	11	12	11
Phase 2 Context	10	8	10	10

Supplemental Figure S1.

30 min Post-swim



Supplemental Figure S1. *Log-transformed corticosterone levels 30 min after an acute swim stress*.

Wildtypes are represented by black squares, heterozygotes by grey diamonds, and knockouts by clear circles. All mice underwent a six min swim stress, and blood was collected 30 min later to quantify serum corticosterone levels. Data are log-transformed cort levels [1–4]. Female wildtypes, n=10; female heterozygotes, n=9; female knockouts, n=10; male wildtypes, n=12; male heterozygotes, n=13, male knockouts, n=14. Data are graphed as mean (horizontal line) with 95% confidence interval (vertical line).

Supplemental Figure S2.



Supplemental Figure S2. Average context fear expression during minutes 2 through 6 of testing for Phase 1 mice.

Wildtypes are represented by black squares, and heterozygotes are represented by grey diamonds. Context fear testing occurred on Day 5 for mice that underwent cued fear conditioning (A), whereas context fear testing occurred on Day 2 for mice that underwent context fear conditioning (B). Data are average percent time spent freezing during minutes two through six of the 10 min context fear testing period [2,5]. Phase 1 Cued: Female wildtypes, n=20; female heterozygotes, n=19; male wildtypes, n=23; male heterozygotes, n=30. Phase 1 Context: Female wildtypes, n=20; female heterozygotes, n=20; female heterozygotes, n=20; male heterozygotes, n=20; male heterozygotes, n=20; female heterozygotes, n=20; male heterozygo

Supplemental Figure S3.



# Supplemental Figure S3. Average context fear expression during minutes 2 through 6 of testing for Phase 2 mice.

Female (A-B) wildtypes are represented by teal squares, and female heterozygotes are represented by blue diamonds. Male (C-D) wildtypes are represented by orange squares, and male heterozygotes are represented by yellow diamonds. Context fear testing occurred on Day 5 for mice that underwent cued fear conditioning (A,C), whereas context fear testing occurred on Day 2 for mice that underwent context fear conditioning (B,D). Data are average percent time spent freezing during minutes two through six of the 10 min context fear testing period [2,5]. Phase 2 Cued No Swim: Female wildtypes, n=11; female heterozygotes, n=11; male wildtypes, n=9; male heterozygotes, n=11. Phase 2 Cued Swim: Female wildtypes, n=12; female heterozygotes, n=11; male wildtypes, n=3; female wildtypes, n=9; male heterozygotes, n=3; male heterozygotes, n=9; male heterozygotes, n=10. Data are graphed as mean (horizontal line) with 95% confidence interval (vertical line). \*p=0.019 indicates difference between

heterozygous and wildtype within same sex and swim condition. <sup>o</sup>p=0.001 indicates difference between sexes within same genotype and swim condition.

Supplemental Figure S4.



#### Supplemental Figure S4. Fecal boli from swim stressors across Phases.

Female wildtypes are represented by teal squares, and female heterozygotes are represented by blue diamonds. Male wildtypes are represented by orange squares, and male heterozygotes are represented by yellow diamonds. Mice were exposed to six minutes of swim stress after (Phase 1) or before (Phase 2) cued or context fear conditioning. Data are fecal boli produced during swim stress. Phase 1 Cued: Female wildtypes, n=10; female heterozygotes, n=10; male wildtypes, n=12; male heterozygotes, n=17. Phase 1 Context: Female wildtypes, n=10; female heterozygotes, n=10; male wildtypes, n=12; female heterozygotes, n=9; male heterozygotes, n=11. Phase 2 Cued: Female wildtypes, n=12; female heterozygotes, n=11; male wildtypes, n=13; male heterozygotes, n=11. Phase 2 Context: Female wildtypes, n=10; female heterozygotes, n=10; male heterozygotes, n=10; male heterozygotes, n=10; male heterozygotes, n=10; female heterozygotes, n=10; female heterozygotes, n=10; male wildtypes, n=10; female heterozygotes, n=11. Phase 2 Cued: Female wildtypes, n=12; female heterozygotes, n=10; female heterozygotes, n=13; male heterozygotes, n=11. Phase 2 Context: Female wildtypes, n=10; female heterozygotes, n=8; male wildtypes, n=10; male heterozygotes, n=10. Data are graphed as mean (horizontal line) with 95% confidence interval (vertical line).

### Excluded fear behavior data

Phase 1 Cued	Reason
0608	All fear data excluded because cued fear expression never exceeds 25%
0609	Data are excluded for context fear testing & cued fear renewal because camera was laggy for 609, 610, and 616, thus freezing data are inaccurate
0610	Data are excluded for context fear testing & cued fear renewal because camera was laggy for 609, 610, and 616, thus freezing data are inaccurate
0616	Data are excluded for context fear testing & cued fear renewal because camera was laggy for 609, 610, and 616, thus freezing data are inaccurate
0664	First 5 tones of cued fear expression <25%, so all data excluded
0681	All fear data excluded because cued fear training never exceeds 25%
0768	First 5 tones of cued fear expression <25%, so all data excluded
0770	No freezing during context testing or cued fear renewal>25%
1057	No freezing during context testing >25%, context testing data excluded
1182	All fear data excluded because cued fear training never exceeds 25%
1168	Cued fear renewal data excluded because cued fear renewal never exceeds 25%
0828	First 5 tones of cued fear expression <25%, so all data excluded
1255	First 5 tones of cued fear expression <25%, so all data excluded
1264	First 5 tones of cued fear expression <25%, so all data excluded
1284	First 5 tones of cued fear expression <25%, so all data excluded

Phase 1 Context	Reason
0713	Shock did not work, was accidentally turned off; exclude all data because never trained
0771	Excluding data because mouse exhibited >75% freezing prior to first shock
0775	Excluding data because mouse exhibited >75% freezing prior to first shock
0776	No freezing during testing >25%, all data excluded
0832	No freezing during testing >25%, all data excluded
1081	Exceeded 75% freezing prior to first shock being administered; excluding all data
1070	No freezing during testing >25%, all data excluded
1080	No freezing during testing >25%, all data excluded

Phase 2 Cued	Reason
0782	All data excluded because first 5 tones cued fear expression on first testing day never exceed 25%
0792	Cued fear renewal data excluded because cued fear renewal never exceeds 25%
0796	Cued fear renewal data excluded because cued fear renewal never exceeds 25%
0827	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0869	Context testing & cued fear renewal testing, door was open and mouse left the chamber at 532.67, data from 510 s onwards on this day excluded
0870	Cued fear renewal data excluded because cued fear renewal never exceeds 25%
0873	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0879	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0881	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0885	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0886	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0887	Mice were not run under correct protocol for testing day 2, so excluding data from this testing day (were run under context testing/cued fear renewal protocol, though in context B)
0908	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0909	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0910	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0916	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0917	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses; Cued fear renewal data excluded because cued fear renewal never exceeds 25%
0918	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses
0919	Videos did not record for cued expression testing & extinction training (7 May 2022) for these animals, so must exclude from analyses

0923	All data excluded because cued fear expression on first testing day never exceeds 25%
0924	Training video did not save/record; All data excluded because cued fear expression on first testing day never exceeds 25%
0925	Training video did not save/record
0929	Training video did not save/record; All data excluded because cued fear expression on first testing day never exceeds 25%
0944	Training video did not save/record
0945	All data excluded because cued fear expression on first testing day never exceeds 25%
0946	All data excluded because cued fear expression on first testing day never exceeds 25%
0953	Exclude all fear data. These animals were swam in both cohorts 19 and 20. Only Swim scoring of cohort 19 will be used.
0967	Exclude all fear data. These animals were swam in both cohorts 19 and 20. Only Swim scoring of cohort 19 will be used.
0981	All data excluded because cued fear expression to first 5 tones on first testing day never exceeds 25%
1192	All data excluded because first 5 tones cued fear expression on first testing day never exceed 25%
1210	All fear data excluded because cued fear training never exceeds 25%
1218	Cued fear renewal data excluded because cued fear renewal never exceeds 25%
1212	All data excluded because cued fear expression to first 5 tones on first testing day never exceeds 25%
1223	All fear data excluded because cued fear training never exceeds 25%

Phase 2 Context	Reason
1083	All data excluded because context fear expression never exceeds 25%
1092	Testing data didn't record because hard drive was not cleared on schedule
1098	Testing data didn't record because hard drive was not cleared on schedule
1079	Testing data didn't record because hard drive was not cleared on schedule
1106	Testing data didn't record because hard drive was not cleared on schedule
1141	Exceeded 75% freezing prior to first shock being administered; excluding all data

#### Excluded swim data

Male	Male · Phase 1 Context · Wildtype		
Climbing Behavior • (All Swim Data Excluded for 0836)			luded for 0836)
Animal ID	Data	Outlier Det	ermination
0700	44.1		_
0723	0.0	22.0	Avg
0726	16.6	16.7909946	SD
0783	28.5	105.92164	Avg + 5 SD
0819	4.6		
0820	0.9		
0836	121.5	>5 SD beyond m	ean
0857	36.4		
1120	34.0		
1146	32.6		

Ma	le • Pha	se 2 Cued • Wile	dtype
Swimming	Behavio	r • (All Swim Data	Excluded for
		0944)	
Animal ID	Data	Outlier Det	ermination
0881	119.9		
0909	177.1		
0919	138.1		
0944	356.9	>8 SD beyond m	ean
0946	172.1		
0952	118.3		
0967	159.4		
1009	151.4		
1034	132.4	135.1	Avg
1036	104.2	25.7674959	SD
1038	130.0	341.269134	Avg + 8 SD
1045	Record	ling failed	
1212	124.8		
1223	93.9		

Male	Phase	2 Cued · Heterozygous
Climbing Beh	avior • (	All Swim Data Excluded for 0869)
Animal ID	Data	Outlier Determination
0869	153.3	>5 SD beyond mean
0908	4.0	
0910	11.1	
0899	22.8	
0904	63.2	

0925	34.1		
0935	32.5		
0994	11.5		
1007	22.3	33.6	Avg
1016	31.6	23.2230254	SD
1086	67.2	149.706036	Avg + 5 SD
1125	69.2		

1125	69.2			
Female · Phase 2 Context · Heterozygous				
Climbing Behavior • (All Swim Data Excluded for 0955)				
Animal ID	Data	Outlier Det	ermination	
0921	25.9			
0922	4.4	14.4	Avg	
0937	0.5	15.7784437	SD	
0943	44.5	140.65255	Avg + 8 SD	
0949	1.4			
0955	144.0	>8 SD beyond m	ean	
1023	25.4			
1092	7.4			
1158	5.9			

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