Supplementary Materials: Utilizing a Population-Genetic Framework to Test for Gene-Environment Interactions between Zebrafish Behavior and Chemical Exposure

Preethi Thunga, Lisa Truong, Yvonne Rericha, Jane La Du, Mackenzie Morshead, Robyn L. Tanguay and David M. Reif

Table S1. Results from fitting LMM to Control data

Random effect variances	Control group		
Group	*Variance*	*Standard Deviation*
Genes	1.506e+08 (σ^2_{Gene})	12270.70
Family ID	1.661e+02 (σ^2_{Family})	12.89
Residual	1.348e+07 (σ^2_E)	3671.80
**Random effect parameter estimates	Control group**		
Group	*Estimate*	
Genes (β_1)	3.34	
Family ID (β_2)	0.0035	
Error (ϵ)	3671.80	
**Fixed effect parameter estimates	Control group**		
Group	*Estimate*	*Standard Error*
Intercept	553.92	84.65

Table S2. Results from fitting LMM to Medium exposure group (16.4 uM PFHxA).

Random effect variances	Medium group		
Group	*Variance*	*Standard Deviation*
Genes	1.743e+08 (σ^2_{Gene})	13202.70
Family ID	3.063e+02 (σ^2_{Family})	17.50
Residual	1.809e+07 (σ^2_E)	4252.80
**Random effect parameter estimates	Medium group**		
Group	*Estimate*	
Genes (β_1)	3.10	
Family ID (β_2)	0.0041	
Error (ϵ)	4252.8	
**Fixed effect parameter estimates	Medium group**		
Group	*Estimate*	*Standard Error*
Intercept	564.44	91.24

Table S3. Results from fitting LMM to High exposure group (74.8 uM PFHxA).

Random effect variances	High group		
Group	*Variance*	*Standard Deviation*
Genes	1.849e+08 (σ^2_{Gene})	13596.95
Family ID	2.479e+00 (σ^2_{Family})	1.57
Residual	1.159e+07 (σ^2_E)	3404.12
**Random effect parameter estimates	High group**		
Group	*Estimate*	

Genes (β_1)	3.99	
Family ID (β_2)	0.00046	
Error (ϵ)	3404.12	
Fixed effect parameter estimates | High group		
Group	*Estimate*	*Standard Error*
Intercept	565.09	93.47