

S5 Table. Principal Component Analysis of the brain gene expression profile in TBT-exposed zebrafish.

Gene correlations	PC1 (28.1%)	PC2 (15.2%)	PC3 (10.4%)	PC4 (9.5%)	PC5 (7.3%)
RXR α/a	0.692*	0.0197	0.448*	-0.2111	0.1965
PPAR γ	0.0557	0.5354*	0.1227	-0.5517*	0.2979
DGAT2	0.4546*	0.278	0.6361*	0.2442	-0.1452
FASn	0.8679*	-0.246	-0.0803	0.0073	0.1142
C/EBP α	0.7077*	0.3657	-0.1426	0.1436	0.1898
C/EBP β	-0.1968	0.5737*	0.1577	-0.2567	0.3772*
ACOX1	0.6966*	-0.1349	0.0291	-0.2656	-0.0978
ACC α	0.8134*	0.0577	-0.1067	0.0905	-0.3071*
SREBP1	0.4417*	-0.3527	-0.4722*	0.2334	0.5201*
ChREBP	0.7317*	0.028	0.0749	0.0716	-0.131
IGF-I	-0.1795	-0.2139	0.4257*	0.6146*	0.4865*
IGF-II α	-0.0861	0.5393*	0.0679	0.2879	-0.0206
11 β -HSD2	-0.1781	-0.6243*	0.5882*	-0.0919	-0.0886
11 β -HSD3 α	-0.038	0.6041	-0.0768	0.4738*	-0.2139
TBT (ng/L)		Average scores on PCs			
<i>Males</i>					
Control	1.2452	1.0309	0.5326	0.2819	-0.3656
10	-1.3753 $^{\#}$	-0.2094	-0.5757	-0.5171	-0.4775
50	0.764	-0.9313 $^{\#}$	-1.4772 $^{\#}$	0.8872	0.4224
<i>Females</i>					
Control	1.159	0.7599	0.6676	0.1025	-0.3231
10	-1.9967 $^{\#}$	0.1849	0.3335	-0.5143	0.3634
50	0.2037	-0.835 $^{\#}$	0.5191	-0.2402	0.3804

*indicates significant correlations and $^{\#}$ significant differences compared to the control (n=8, $p<0.05$; one-way ANOVA followed by Dunnett's test)