

SUPPLEMENTARY MATERIAL

Transcriptome sequencing of 3,3',4,4',5-Pentachlorobiphenyl (PCB126)-treated human preadipocytes demonstrates progressive changes in pathways associated with inflammation and diabetes.

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Table S1: Chemical signature pathways (top 10) of PCB126-treated preadipocytes after 9 hours.

| Chemical Id | Chemical Name | pv_p_fdr | zscore |
|--------------------|--|-----------------|---------------|
| D013749 | Tetrachlorodibenzodioxin | 4.96E-10 | 4.15862 |
| C023035 | 3,4,5,3',4'-pentachlorobiphenyl (PCB126) | 4.00E-13 | 3.53009 |
| C086401 | pentabromodiphenyl ether | 1.17E-06 | 4.351941 |
| D008748 | Methylcholanthrene | 2.55E-07 | 2.886751 |
| D000336 | Aerosols | 1.79E-06 | 3.152963 |
| D001564 | Benzo(a)pyrene | 0.000206 | 4.004543 |
| C022838 | nickel chloride | 0.000179 | 3.577709 |
| D017632 | Asbestos, Serpentine | 2.51E-05 | 2.683282 |
| D009853 | Omeprazole | 1.96E-05 | 2.236068 |
| D004137 | Dinitrochlorobenzene | 0.00027 | 3.464102 |

Table S2: Pathways in common between different timepoints (9-hour, 1-day, 3-day) of PCB126 exposure of preadipocytes. $P \leq 0.05$, $Fdr \leq 0.05$.

| Name of Pathway | 9-Hour pv_fdr | 1-Day pv_fdr | 3-Day pv_fdr |
|--|------------------|-----------------|-----------------|
| Cytokine-cytokine receptor interaction | 0.000213 | 1.58E-06 | 3.69E-06 |
| MAPK signaling pathway | 0.000213 | 0.000499 | 1.10E-05 |
| Osteoclast differentiation | 0.002083 | 0.013481 | 7.33E-05 |
| Neuroactive ligand-receptor interaction | 0.04661 | 0.000775 | 7.33E-05 |
| AGE-RAGE signaling pathway in diabetic complications | 0.00959 | 8.53E-05 | 0.000132 |
| Pathways in cancer | 0.011362 | 0.000813 | 0.000177 |
| Ras signaling pathway | 0.037673 | 0.043676 | 0.000197 |
| Complement and coagulation cascades | 0.015962 | 0.049746 | 0.000197 |
| Rheumatoid arthritis | 0.001351 | 0.000393 | 0.000214 |
| NF-kappa B signaling pathway | 0.017713 | 0.004984 | 0.000553 |
| Hematopoietic cell lineage | 0.043771 | 0.012131 | 0.000975 |
| Fluid shear stress and atherosclerosis | 0.04661 | 0.010475 | 0.00362 |

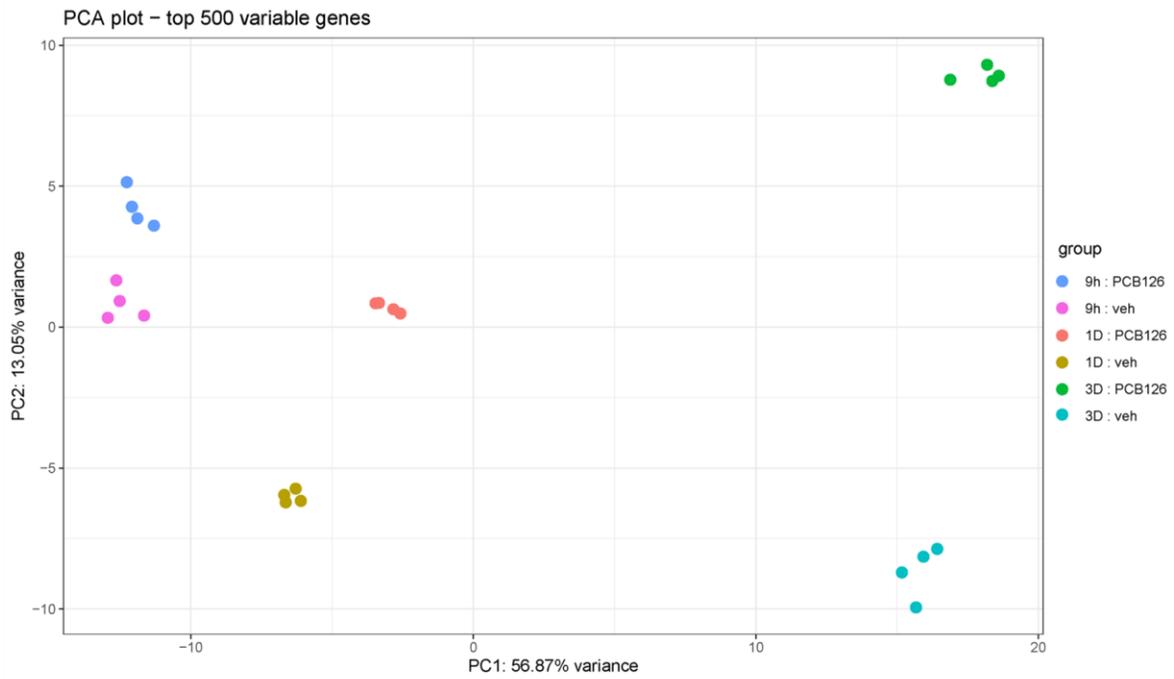
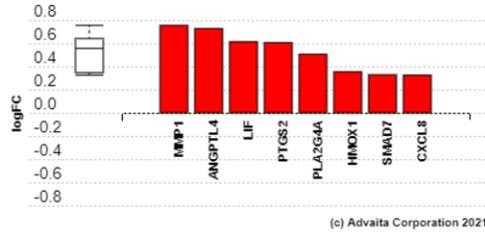
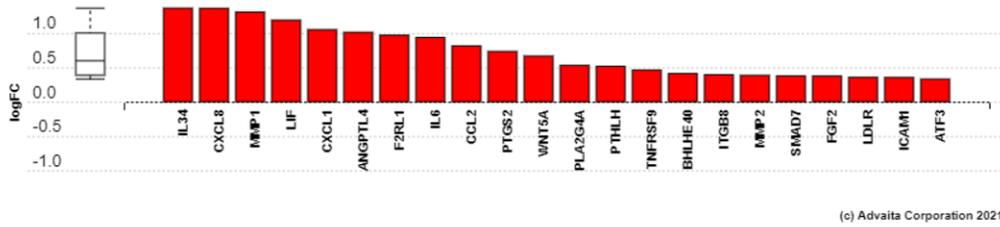


Figure S1: Principal Component Analysis of top 500 variable gene transcripts from replicates of vehicle and PCB126 treated preadipocytes over a time course of 9 hours (9h), 1 day (1D), and 3 days (3D).

9-Hour



1-Day



3-Day

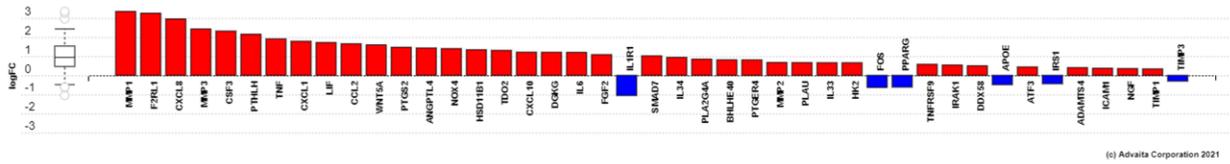


Figure S2: Transcript level changes (log2fold) of genes in the IL-1 β pathway at different timepoints (9-hour, 1-day, 3-day) of PCB126-exposed preadipocytes compared to vehicle controls at the same timepoints. Fdr \leq 0.05.