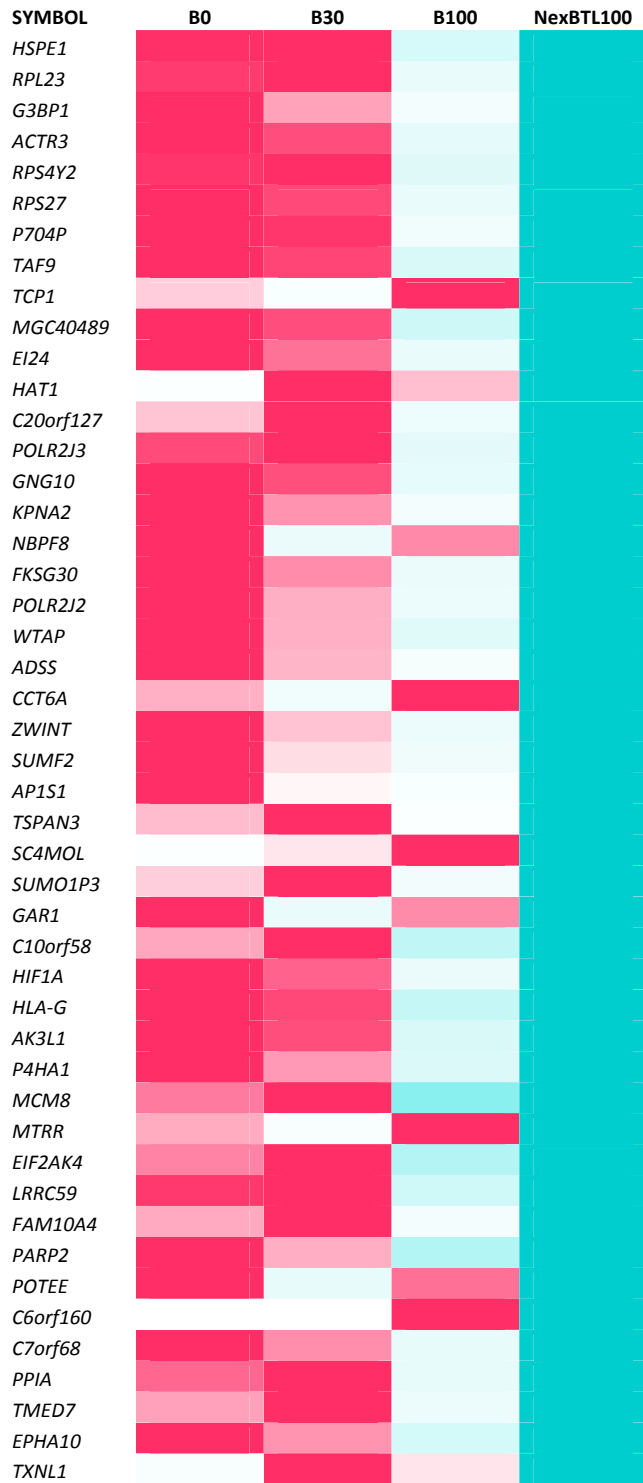
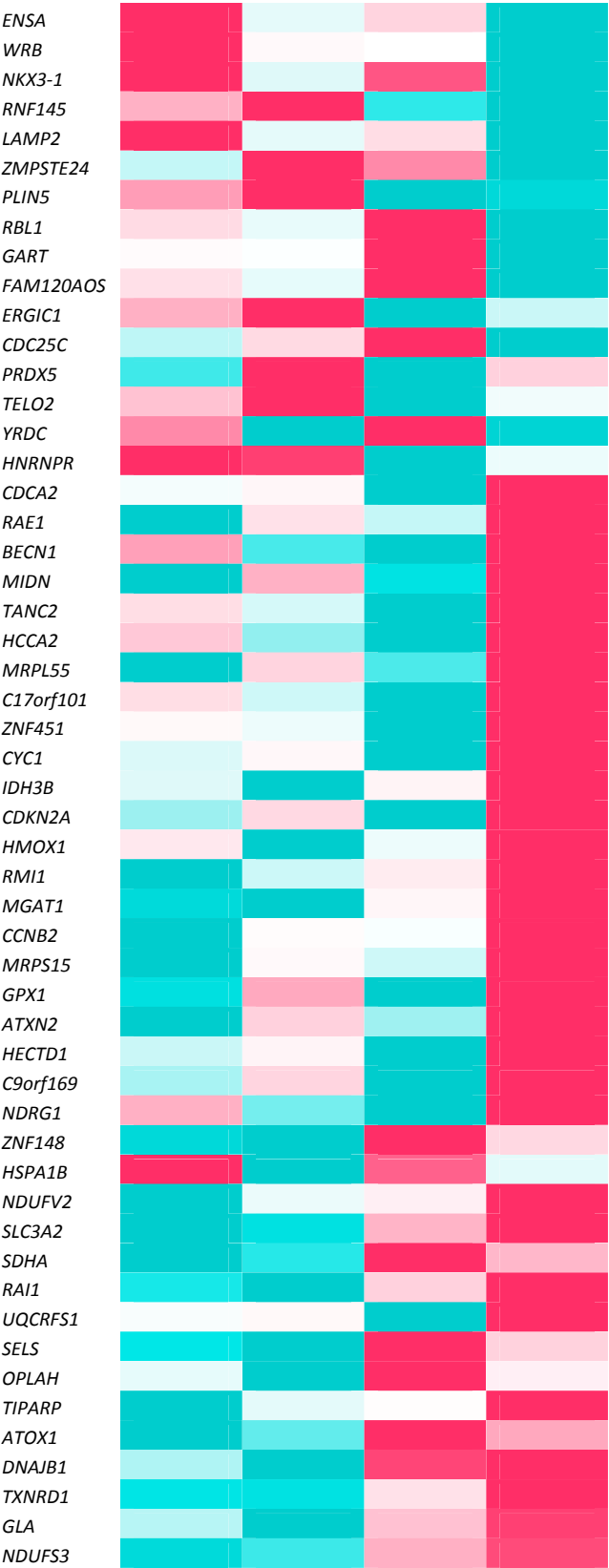


Supplementary Materials: Comparative Analysis of Toxic Responses of Organic Extracts from Diesel and Selected Alternative Fuels Engine Emissions in Human Lung BEAS-2B Cells

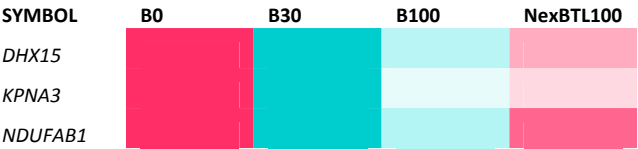
Helena Libalova, Pavel Rossner, Jr., Kristyna Vrbova, Tana Brzicova, Jitka Sikorova, Michal Vojtisek-Lom, Vit Beranek, Jiri Klema, Miroslav Ciganek, Jiri Neca, Katerina Pencikova, Miroslav Machala and Jan Topinka

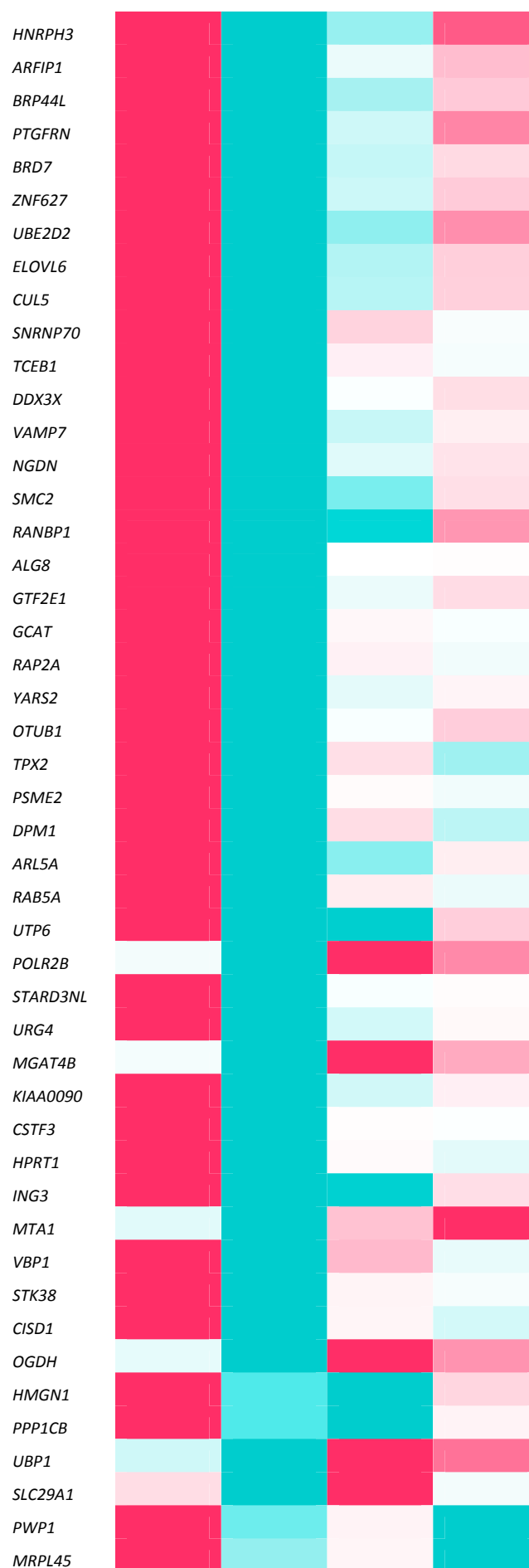
(A)

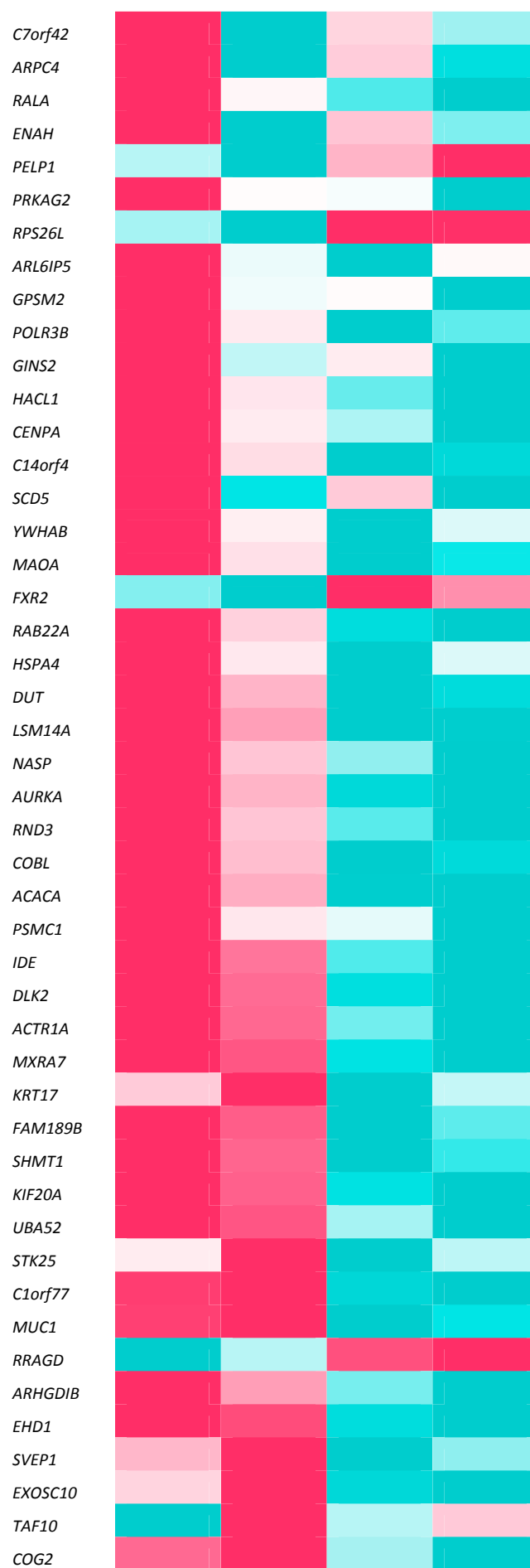


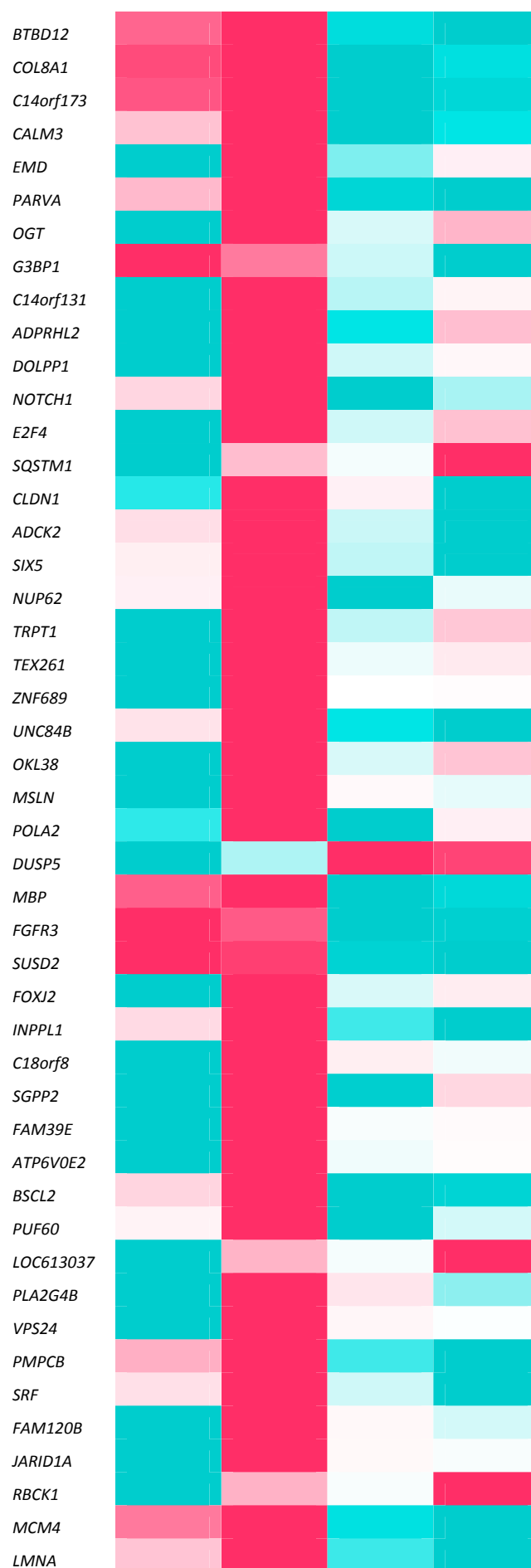


(B)









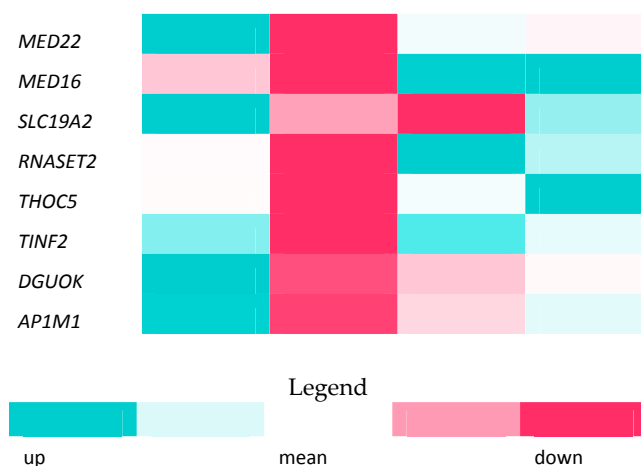


Figure S1. Top differentially expressed genes among diesel exhaust particles (DEP) extract treatments identified by one-way ANOVA (p value < 0.05 , $\log FC > \pm 1.5$). Heat maps show relative expression of significantly modulated genes after (A) 4 h and (B) 24 h exposure to all DEP extracts. Each row corresponds to a gene (gene symbols are listed to the right of each row) and each column to a sample. The colors are scaled by row; red and blue indicate two standard deviations below or above the mean (white), respectively.

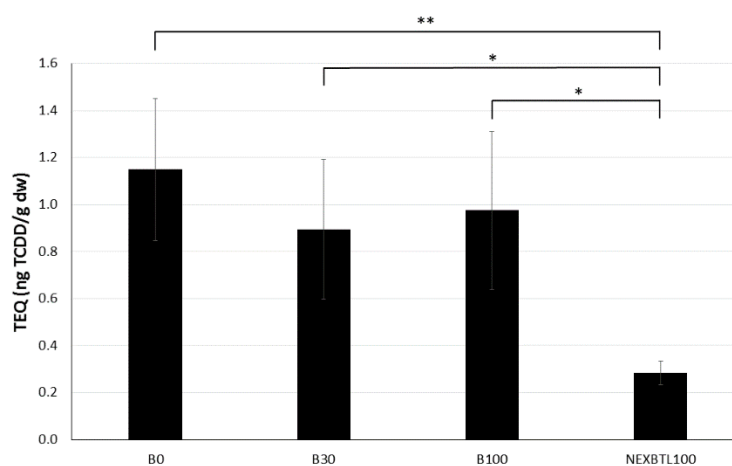


Figure S2. The aryl hydrocarbon receptor (AhR)-mediated activity of B0, B30, B100 and NEXBT100 extracts determined in human hepatoma HepG2 stable transfected by luciferase reporter gene under the control by the AhR (DR-CALUX assay). The activities were expressed as 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) equivalents (TEQs). * p -value < 0.05 , ** p -value < 0.01 .

Table S2. Sequences of primers used in RT-qPCR.

Symbol	RefSeq ID	Oligonucleotide	
<i>AKR1C2</i>	NM_001354.4	sense	GGTATTTATCAGTCAGTGCCTCTC
		antisense	TGCTAGTCAATATCGCTCATCCT
<i>TXNRD1</i>	NM_182743.1	sense	CTTGGATAGGAGTTGGTGAATAGAA
		antisense	GGGCTTGAGACTGGTGACTT
<i>HMOX1</i>	NM_002133.1	sense	GGAAGCCCCCACTCAACA
		antisense	GCATAAAGCCCTACAGCAACT
<i>CYP1A1</i>	NM_000499.2	sense	CAAGGTGTTAAGTGAGAAGGTG
		antisense	AGCAGGATAGCCAGGAAGAG
<i>CYP1B1</i>	NM_000104.2	sense	CACTGGAAACCGCACCTC
		antisense	AGCACCGACAGGAGTAGC
<i>TPX2</i>	NM_012112.4	sense	CAGGAGGAATATAAGGAAGTGAAC
		antisense	TGGGACAGGTTGAAAGGCTTA
<i>CCNB2</i>	NM_004701.2	sense	TGGAGAATATTGACACAGGAGTTAAT
		antisense	TGGGTTGAACTGGAACCTTGG
<i>CDKN2A</i>	NM_058195.2	sense	ATGTCCTGCCTTTTAACGTAGATA
		antisense	CTCACTCCAGAAAACCTCCAACA
<i>HSPE1</i>	NM_002157.1	sense	ATTATGCTTCCAGAAAAATCTCAACG
		antisense	CTTTATCTCCAACCTTTCACGCTAAC
<i>FOSL1</i>	NM_005438.2	sense	AGCCCAGAGACTTTGTAGATCCTT
		antisense	TTCTGTCAGGAGATAGGGTTGGG
<i>BNIP3</i>	NM_004052.2	sense	AGACCCACAGGACACTAAC
		antisense	GACCTCTTTCCTCCTCTCCAT