

Molecular design and Mechanism Analysis of Phthalic Acid Ester Substitutes: Improved Biodegradability in Processes of Sewage Treatment And Soil Remediation

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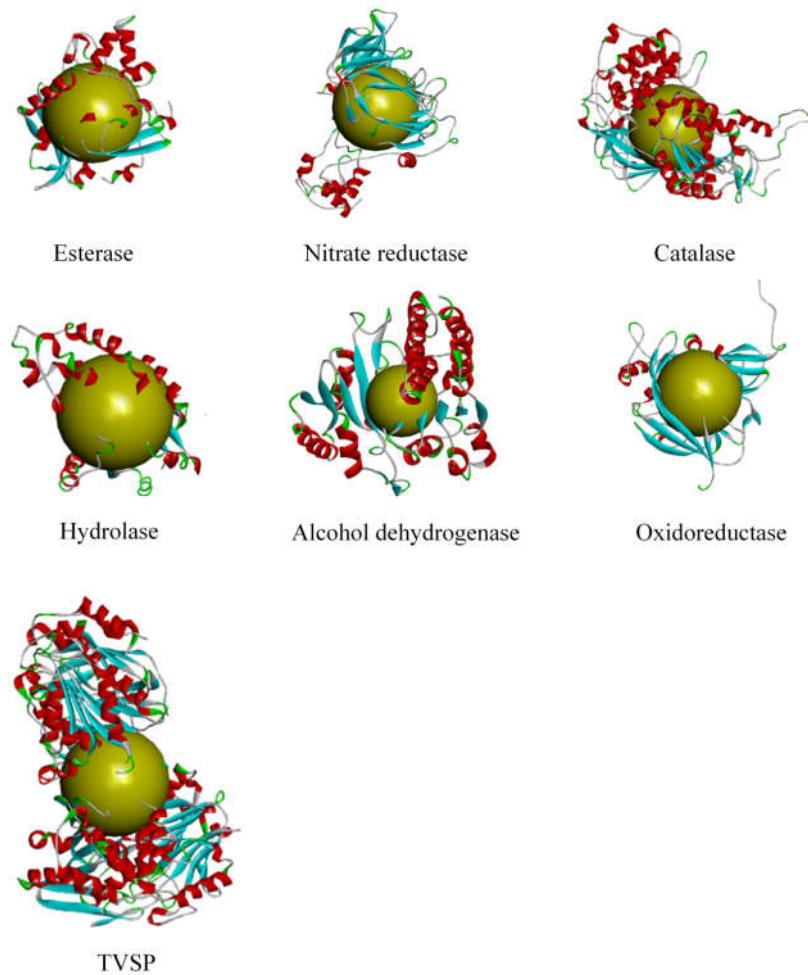


Figure S1. The docking site of proteins.

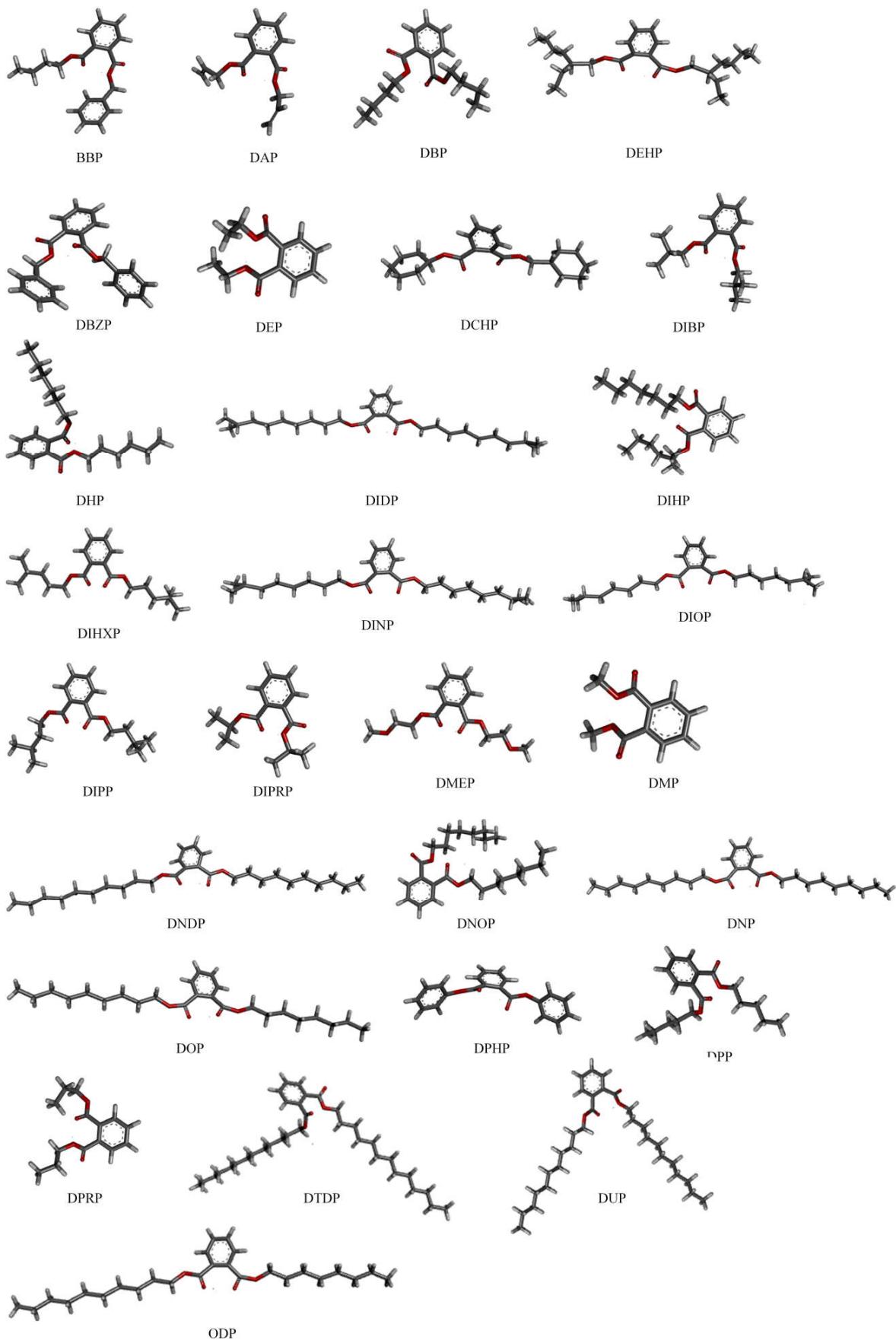
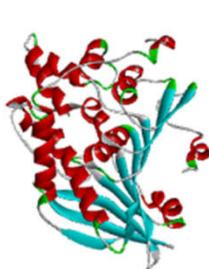
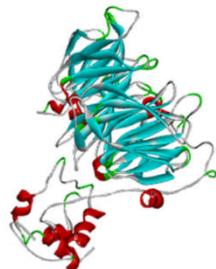


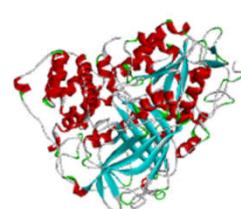
Figure S2. Molecular structure diagram of PAEs.

Aerobic Bacteria:

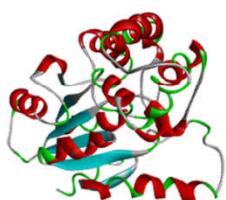
Esterase
PDB ID: 7CUZ



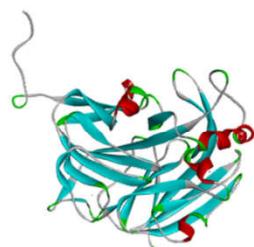
Nitrate reductase
PDB ID: 1HZV



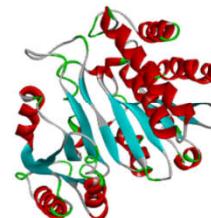
Catalase
PDB ID: 1P80

Anaerobic Bacteria:

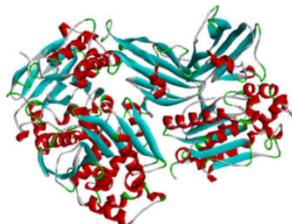
Hydrolase
PDB ID: 2IOF



Oxidoreductase
PDB ID: 1NIC



Alcohol dehydrogenase
PDB ID: 3FMX

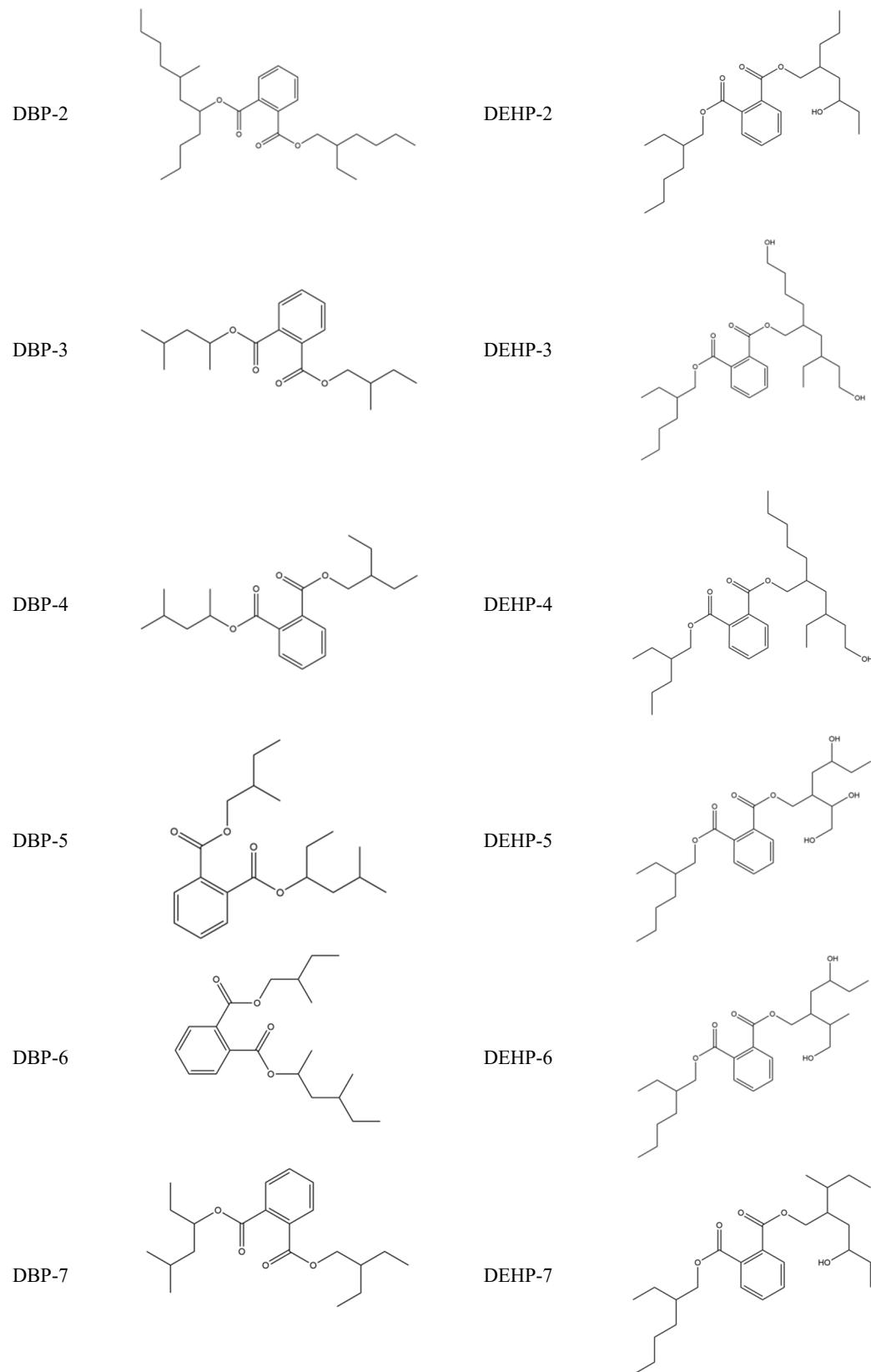
Biodegradable Complex Proteins:

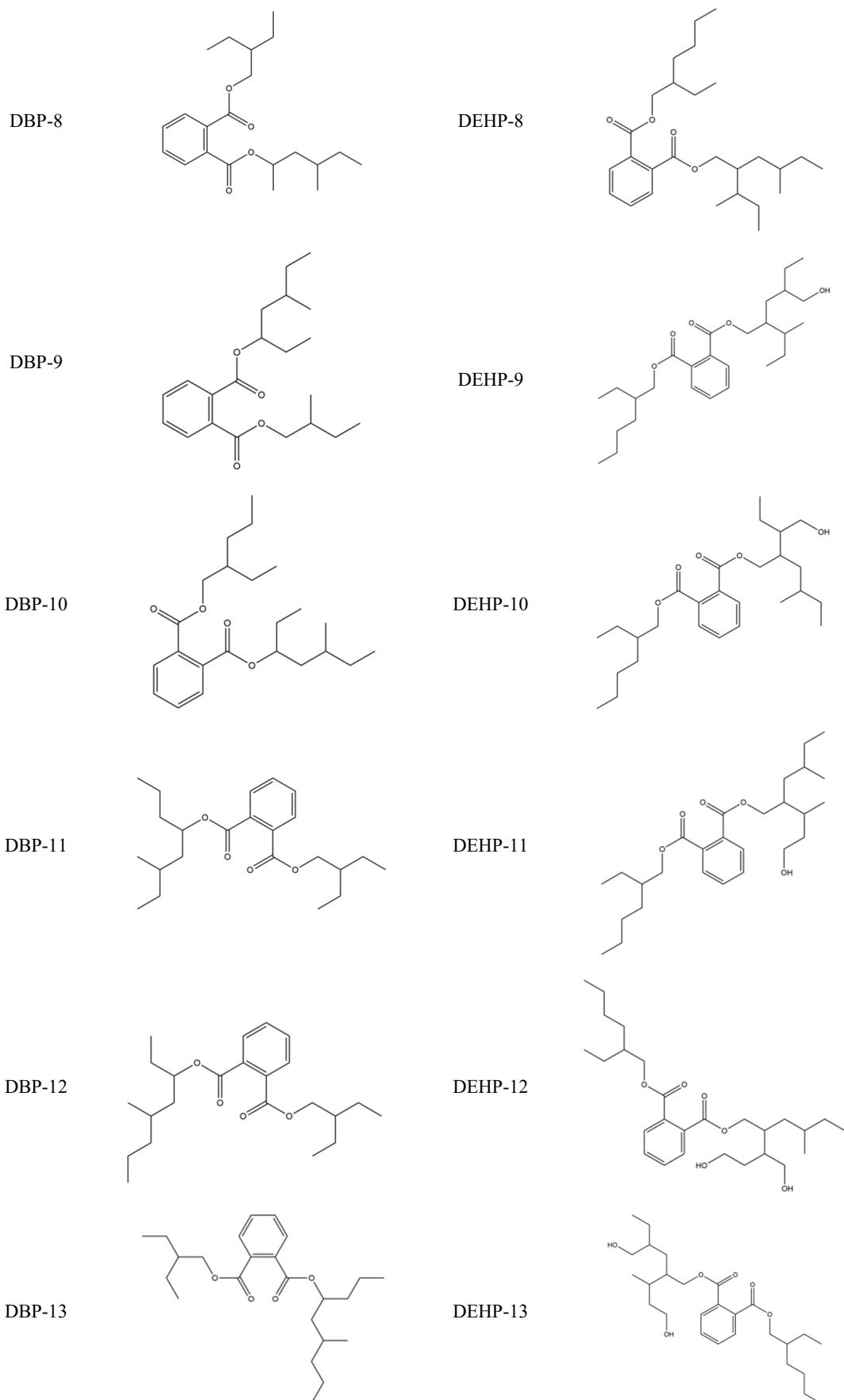
Soil composite degradation enzyme

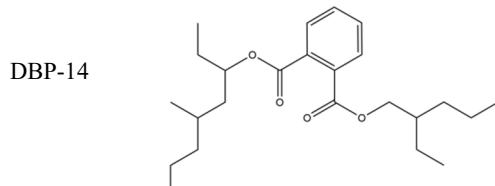
Figure S3. The structure of proteins.

Table S1. Molecular structure of DBP and DEHP substitutes.

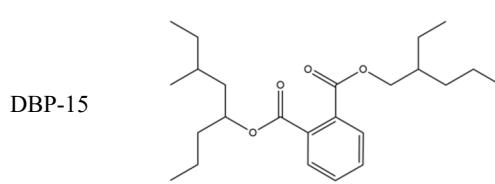
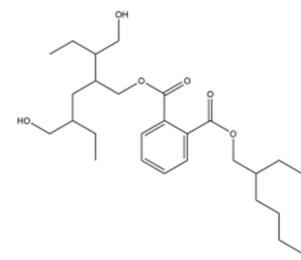
Molecular	Chemical structure diagram	Molecular	Chemical structure diagram
DBP-1		DEHP-1	



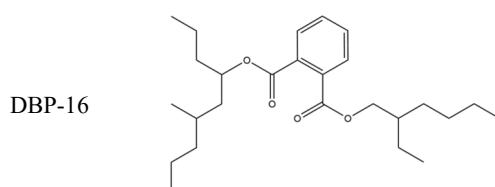
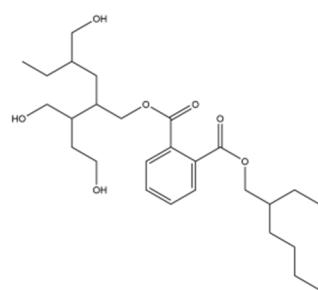




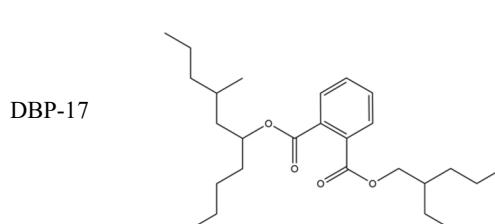
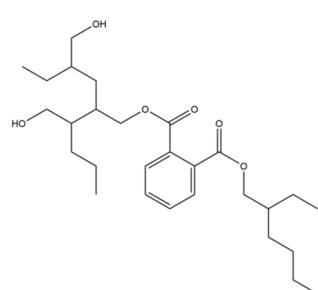
DEHP-14



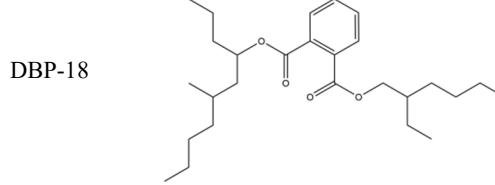
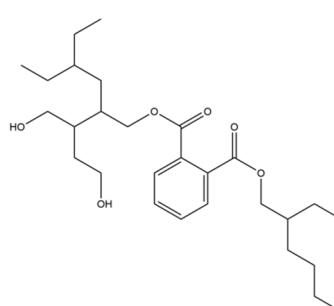
DEHP-15



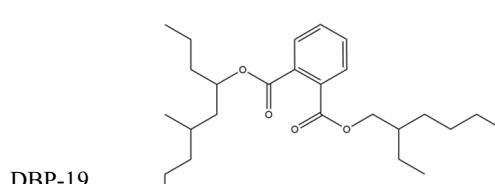
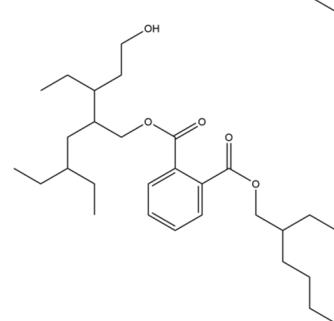
DEHP-16



DEHP-17



DEHP-18



DEHP-19

