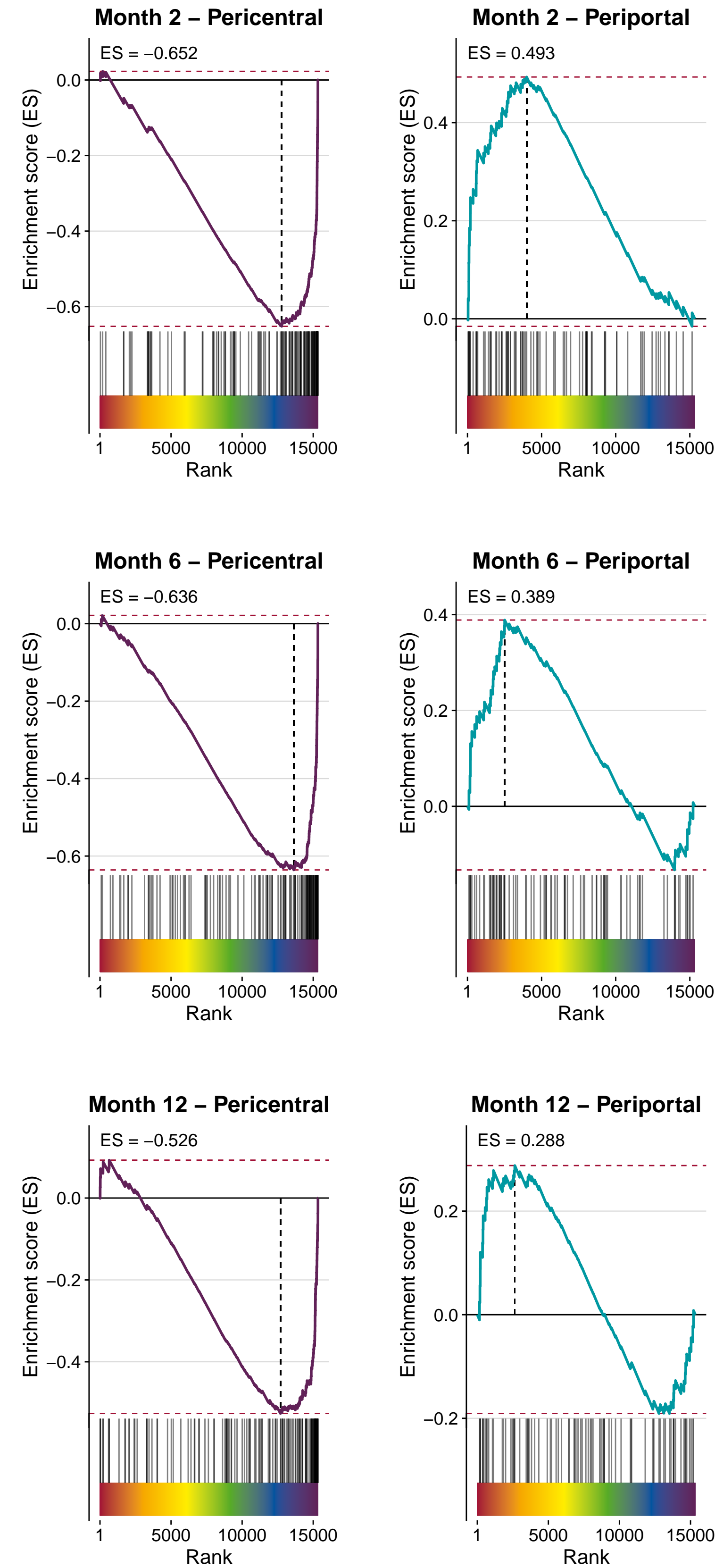
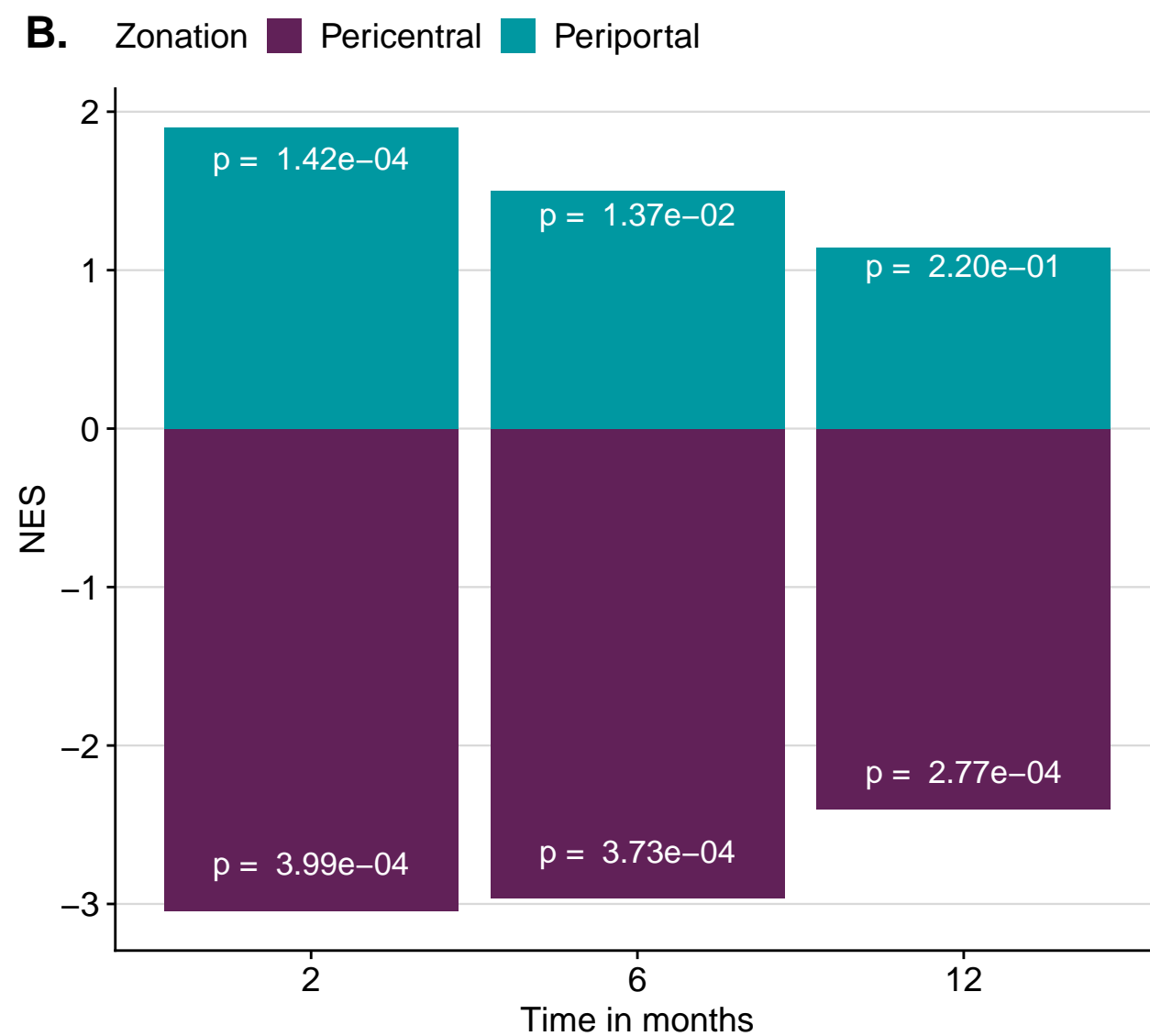


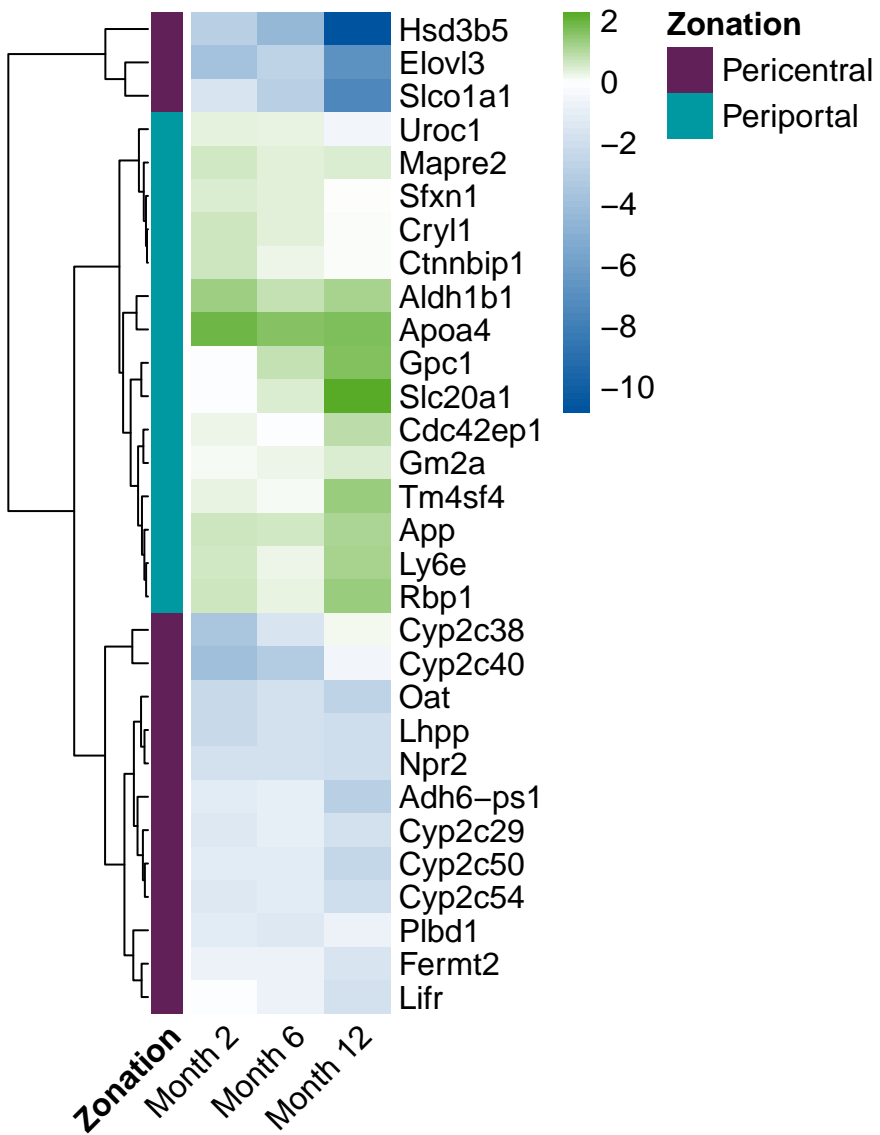
A.



B.



C.



D.

Regulation	Zonation	Gene set	p
Down	Pericentral	Monocarboxylic Acid Metabolic Process	4.30e-06
Down	Pericentral	Nr0b2	1.08e-05
Down	Pericentral	Carboxylic Ester Hydrolase Activity	2.25e-05
Down	Pericentral	Glutamine Family Amino Acid Catabolic Process	3.61e-05
Down	Pericentral	Long Chain Fatty Acid Metabolic Process	4.83e-05
Down	Pericentral	Glutamate Metabolic Process	6.78e-05
Down	Pericentral	Epoxygenase P450 Pathway	8.51e-05
Down	Pericentral	Arginine And Proline Metabolism	3.61e-04
Down	Pericentral	Lipid Metabolic Process	7.16e-04
Down	Pericentral	Primary Bile Acid Biosynthesis	8.76e-04
Down	Pericentral	Nr1h4	3.30e-03
Up	Periportal	Phospholipid Transport	2.61e-05
Up	Periportal	Positive Regulation Of Triglyceride Lipase Activity	6.31e-05
Up	Periportal	Lipid Metabolic Process	7.49e-05
Up	Periportal	Tbx3	2.42e-03
Up	Periportal	Atf1	1.85e-02

■ Biological process (GO) ■ Pathway (KEGG)
■ Molecular function (GO) ■ TF (DoRoThEA)