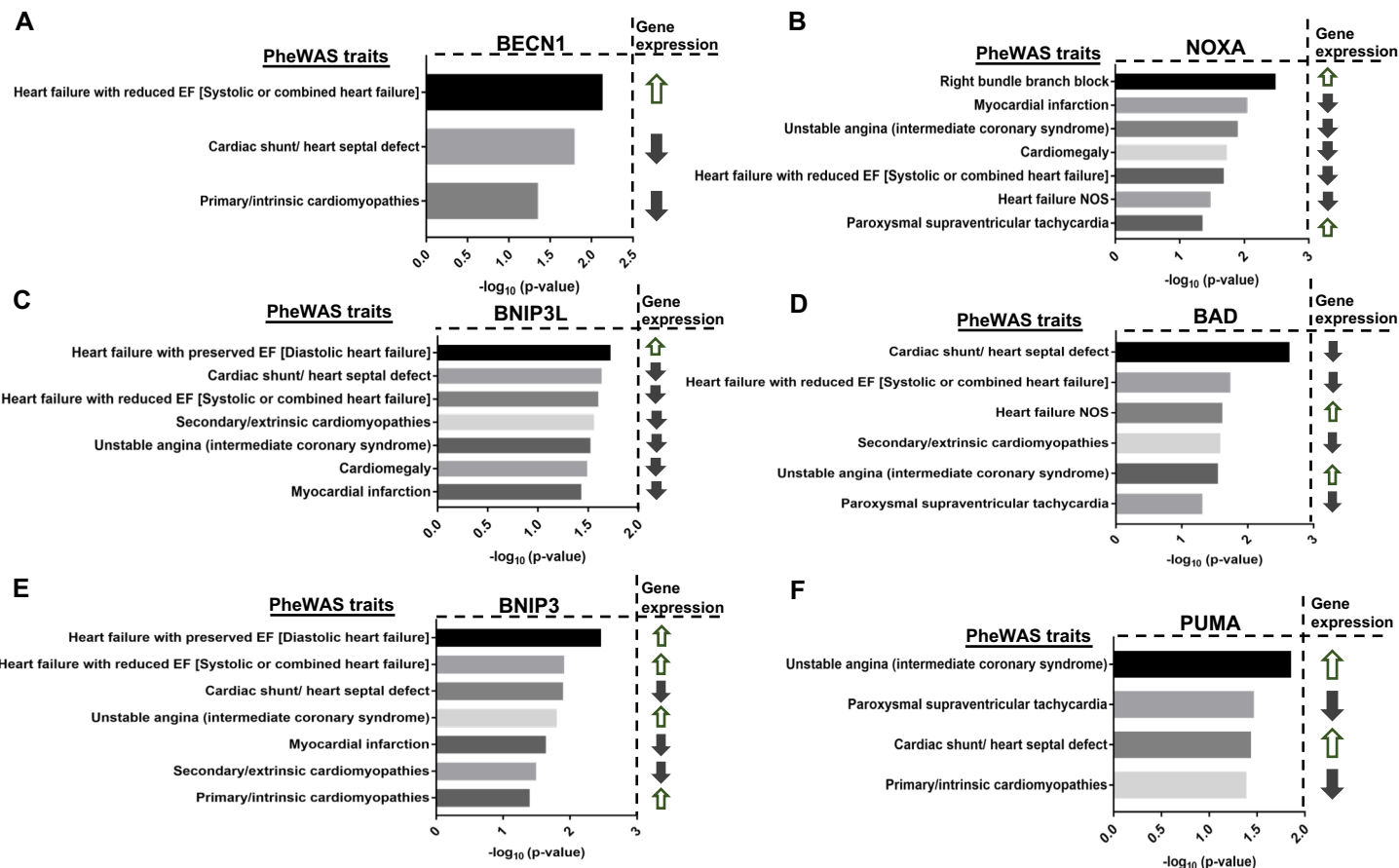


**Figure 7-figure supplement 7**



**Figure 7-figure supplement 7. Secondary PrediXcan analysis of BH3-only Bcl-2 family genes shows discordant direction of effect with heart conditions.**

(A-F) Graphs of significant cardiac traits in the BioVU validation cohort ranked by p-values ( $-\log_{10}$ ) for BECN1, NOXA, BNIP3L, BAD, BNIP3, and PUMA and the corresponding direction of gene expression. For all genes, cardiac traits that appeared more than once were represented in the graph by the most significant p-value. P-values were determined by logistic regression with disease status as response variable and imputed gene expression as predictor.