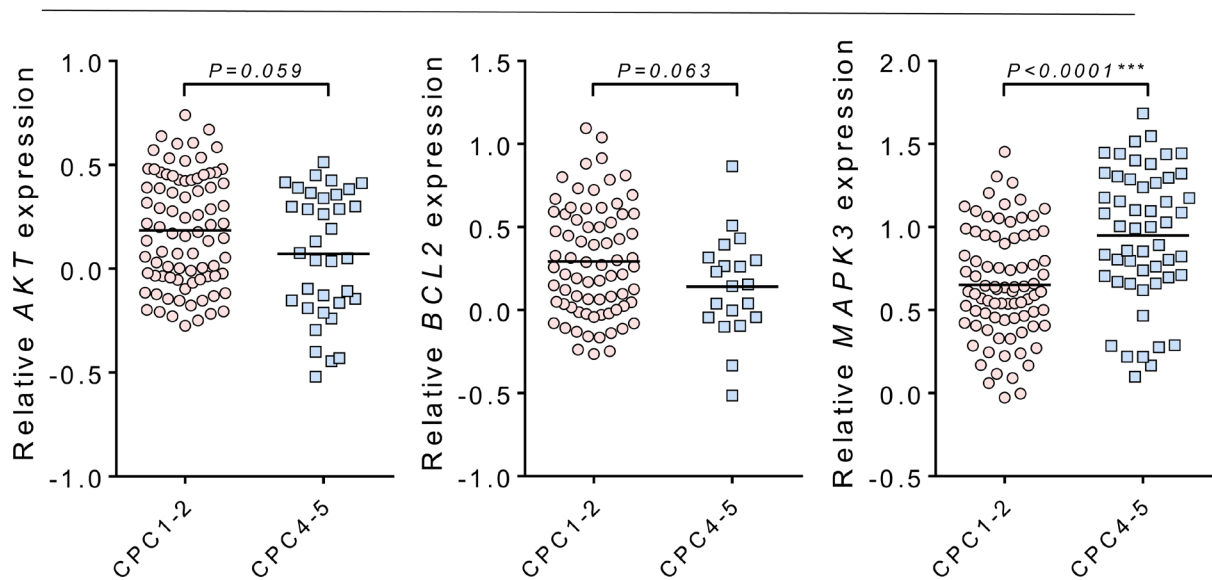# Identification of novel biomarkers for prediction of neurological prognosis following cardiac arrest

Supplementary Figure 1

To validate the expression of *AKT1*, *BCL2*, and *MAPK3* in the large cohort of cardiac arrest patients, we recapitulated gene expression levels of *AKT1*, *BCL2*, and *MAPK3* in the datasets available from the NCBI, GEO database (GSE29540)