

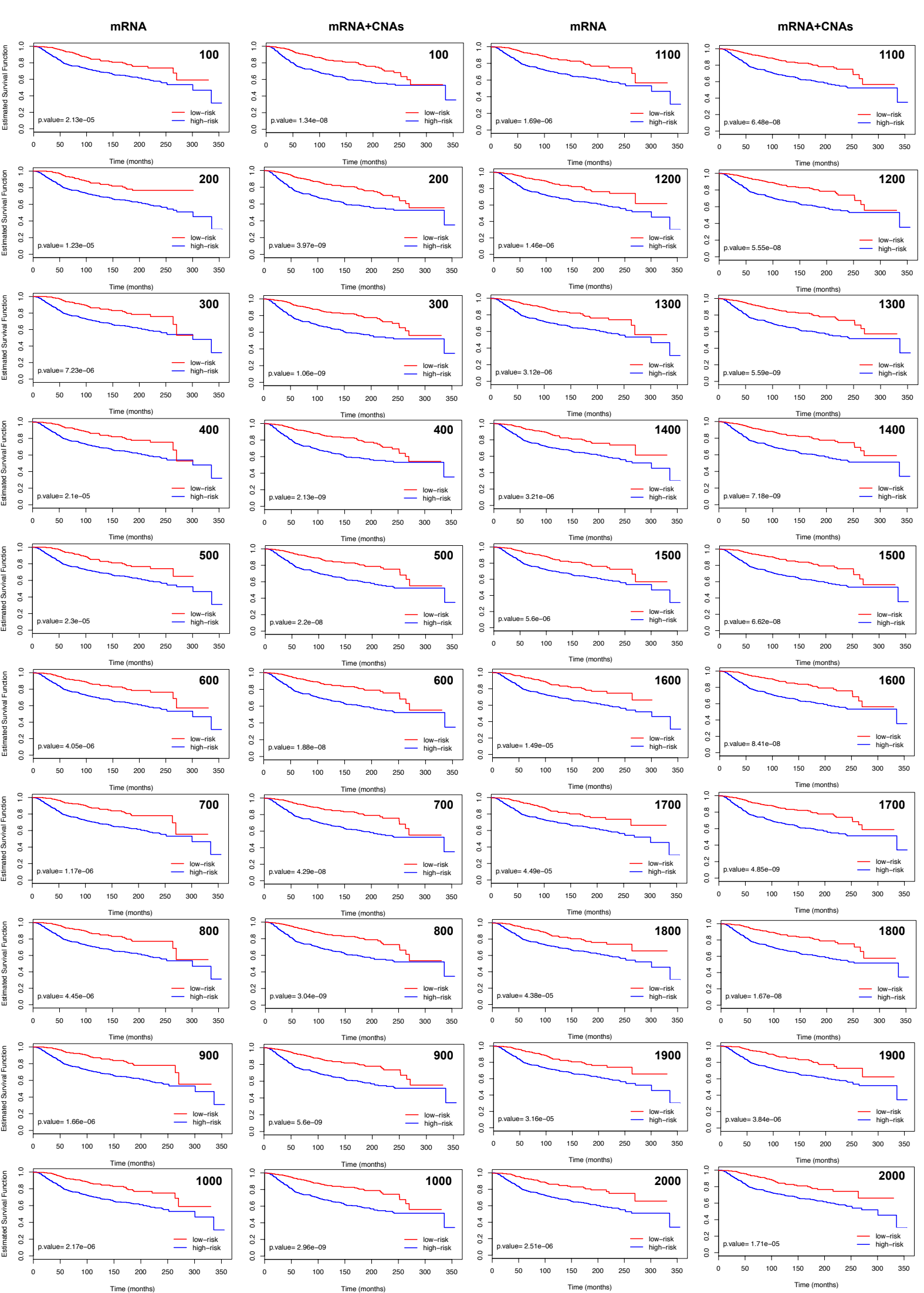


Figure S3. Kaplan-Meier plots of mRNA data and the adjusted mRNA data based on CNAs profiles. For each case, patient samples were separated into two groups according to the predicted risk scores from the selected genes by combining BMD+DAD-screening and network Cox regression method (Coxnet). High-risk group is labelled in blue d and low-risk group is labelled in red. The X-axis represents time and the Y-axis represents survival rate.