

SUPPLEMENTARY FIG. S3. Log2FoldChange for microRNA (miRNA) biomarkers that were used for machine learning based analysis. Error bars represent standard error. (A) Fold change (injured vs. sham) of the eight miRNAs identified for injured vs. sham classification. (B) Fold change (Blast vs. CCI) of the four miRNAs identified for Blast vs. CCI classification. (C) Fold change (Blast vs. Blast sham) of the five miRNAs were identified for Blast vs. Blast sham classification. (D) Fold change (CCI vs. CCI sham) of the three miRNAs identified for CCI vs. CCI sham classification.