

**CDH1 Overexpression Predicts Bladder Cancer from Early Stage and Inversely Correlates
with Immune Infiltration**

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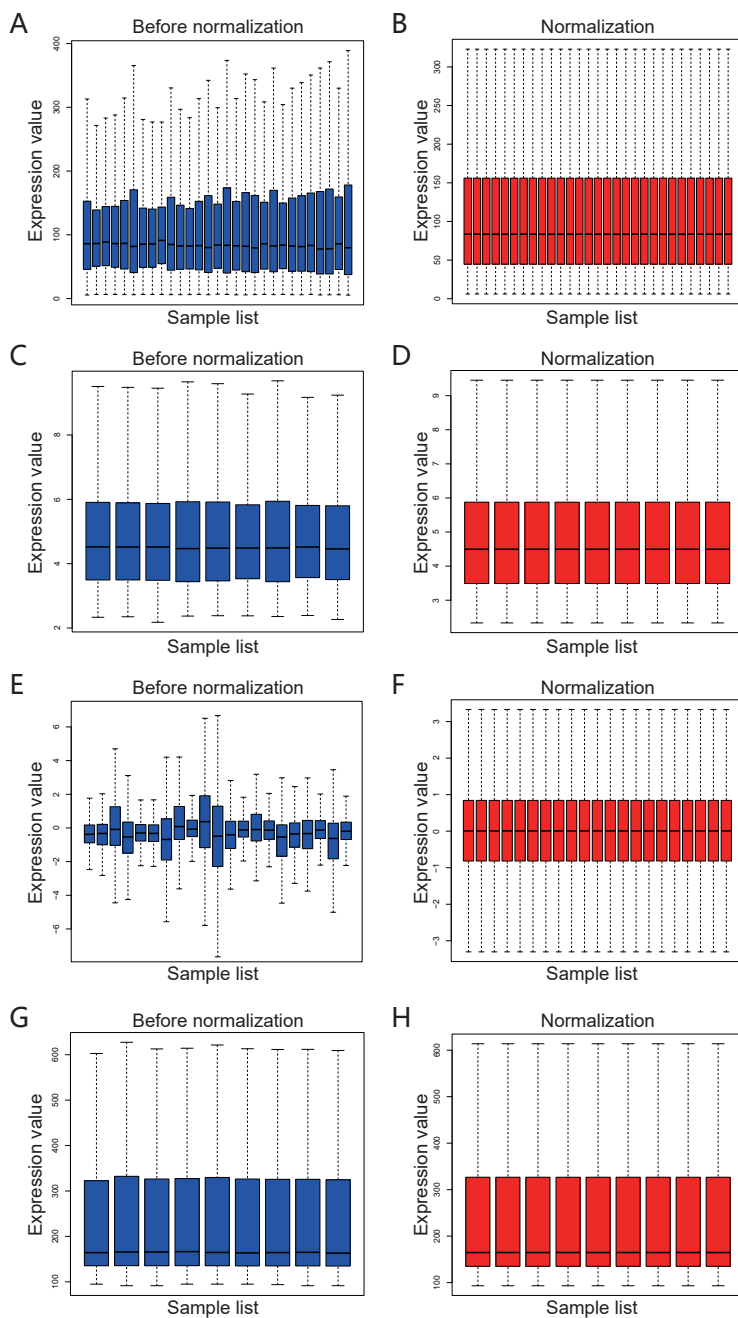
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Additional file

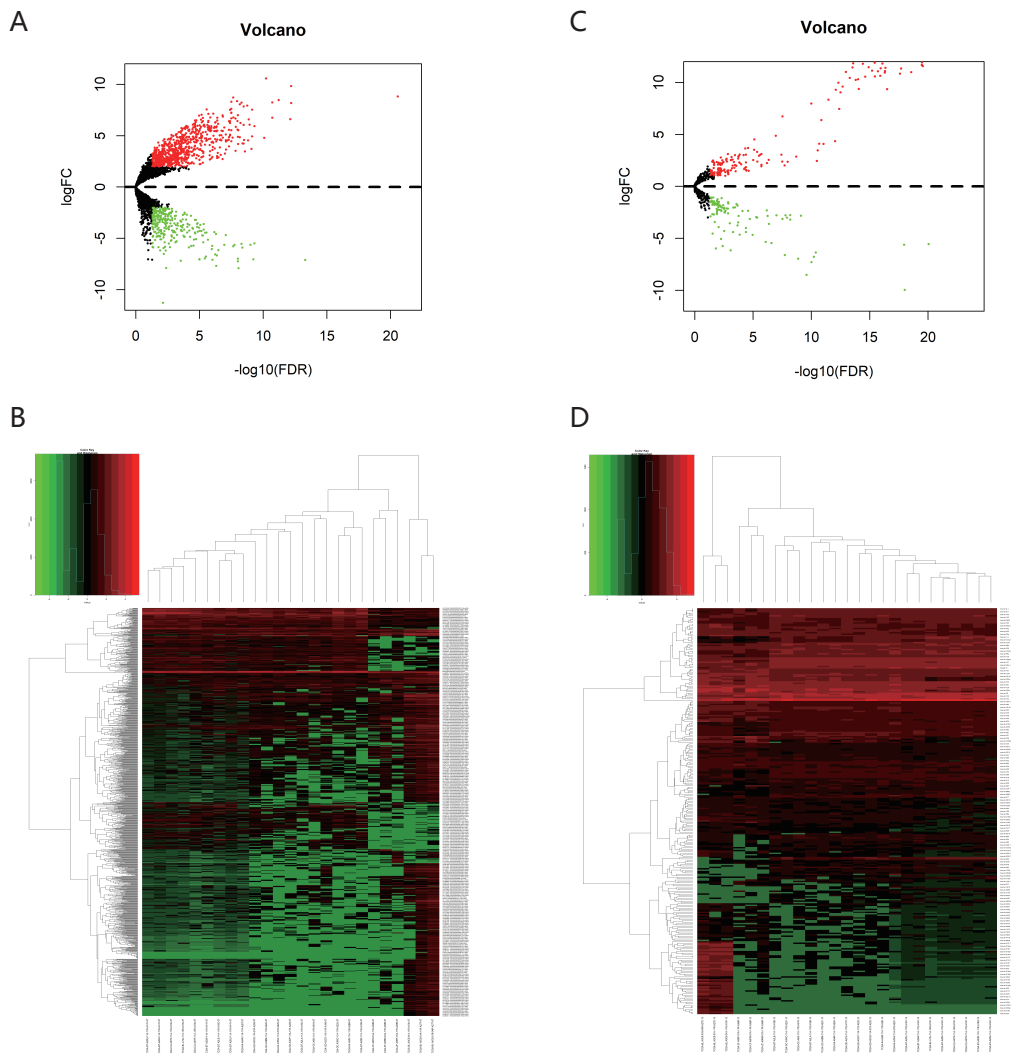
Additional file 1:

Supplementary Figure S1. Normalisation of gene expression.

Supplementary Figure S2. DElncRNAs and DEmiRNAs in patients with BC.



Supplementary Figure S1. Normalization of gene expression. (A–B) Normalization of the GSE3167 data set. (C–D) Normalization of the GSE7476 data set. (E–F) Normalization of the GSE40355 data set. (G–H) Normalization of the GSE65635 data set. Blue represents data before normalization, and red represents data after normalization.



Supplementary Figure S2. DElncRNAs and DEMiRNAs in patients with NMIBC (A) Volcano plot and (B) heatmap of DElncRNAs in NMIBC and adjacent non-carcinoma bladder tissues. The DElncRNAs were identified with the thresholds of $|\log_{2} \text{FC}| > 2.0$ and $\text{adj.P-value} < 0.05$. (C) Volcano plot and (D) heatmap of DEMiRNAs NMIBC and adjacent non-carcinoma bladder tissues. The DEMiRNAs were identified with the thresholds of $|\log_{2} \text{FC}| > 1.0$ and $\text{adj.P-value} < 0.05$.