

Figure S1. Correlation between symptom severity and lncRNA expression; r=Spearman correlations; *p<0.05; **p<0.02; ***p<0.001. †Chronic conditions and first episode <math>‡Inpatients and outpatients.

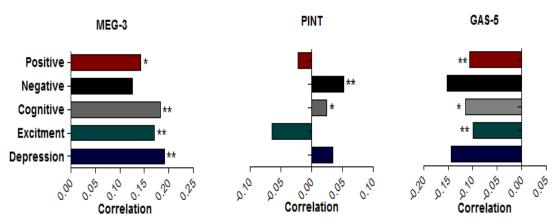


Figure S2. Correlational strength between lncRNA expression and disease severity (PANSS five factor sub-scale) *Spearman correlations* **p*<0.05; ***p*<0.02.

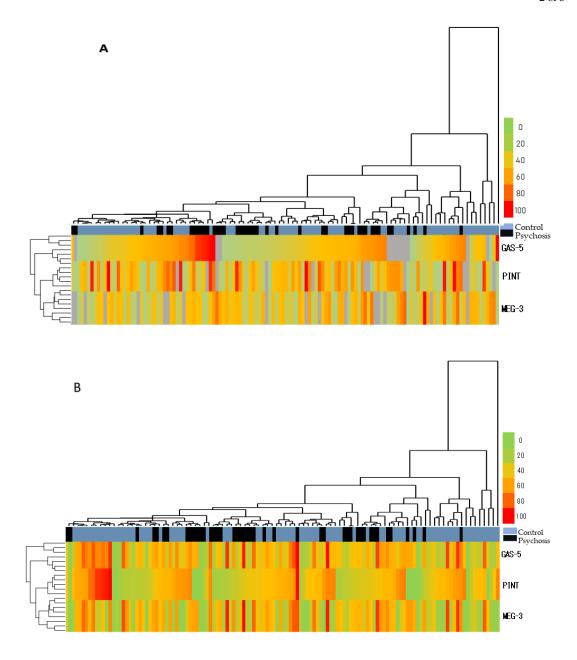


Figure S3: Dendrograms and heat map of hierarchical clustering analysis (complete-linkage) by the expression of lncRNAs MEG-3, PINT and GAS-5, (**A**) predicting clinical diagnosis and, (**B**) predicting symptom acuity

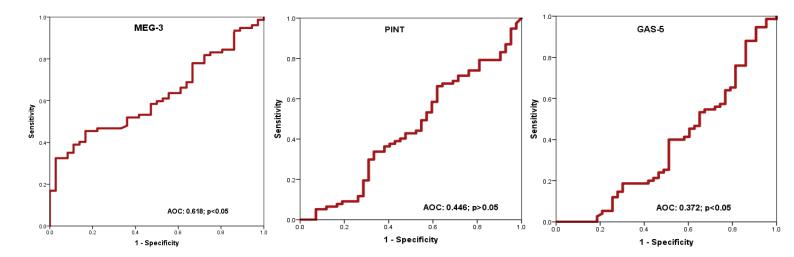


Figure S4: Receiver Operating Characteristic Curve (ROC) for predicting clinical diagnosis with lncRNAs (MEG3, PINT and GAS5) expression.