

Comprehensive Analysis of Ubiquitously Expressed Genes in Human, From a Data-Driven Perspective

Figure

Fig 1. The Flow Diagram for Systematically Characterization of Ubiquitously Expressed Genes in the Human Genome

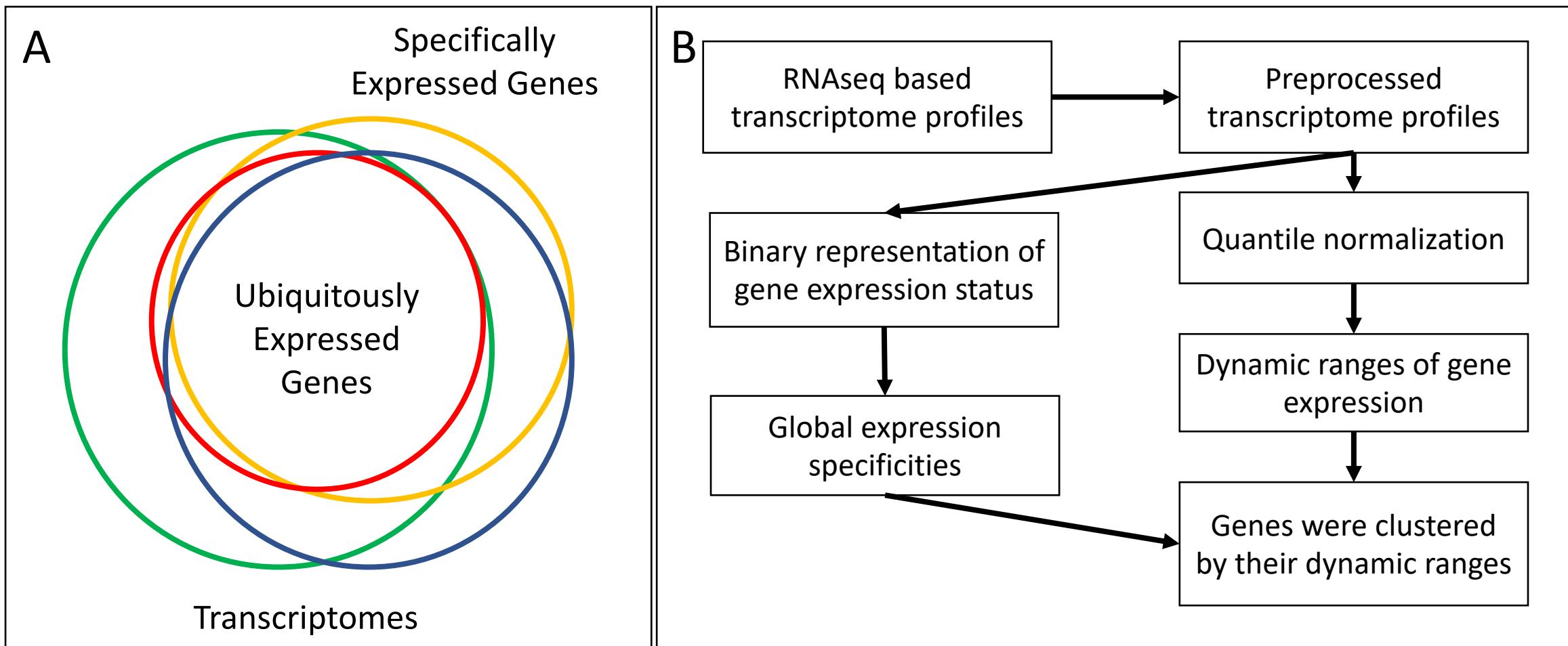


Fig 2. The phenotypic compositions and relatedness among analyzed transcriptomes.

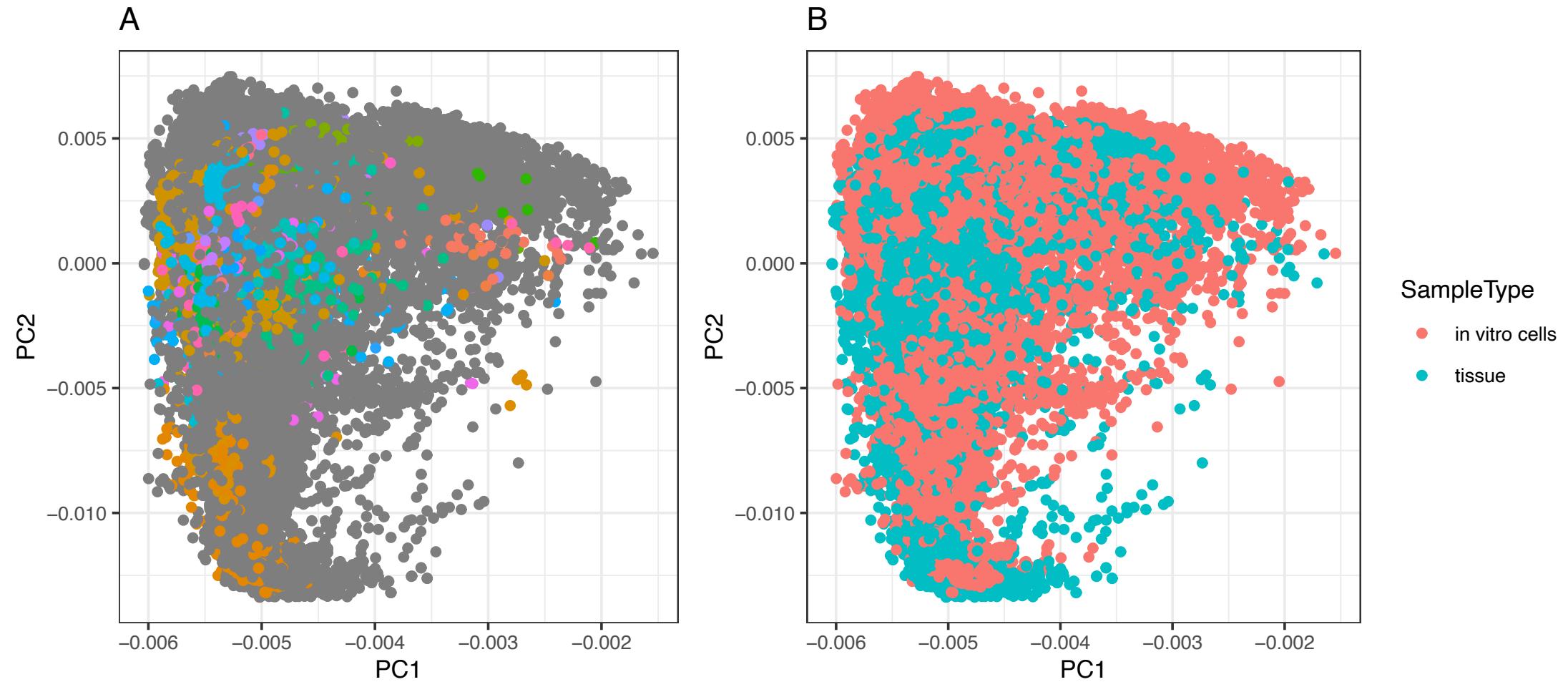


Fig 3. The Global Expression Specificity and Global Dynamic Ranges of Expression Values

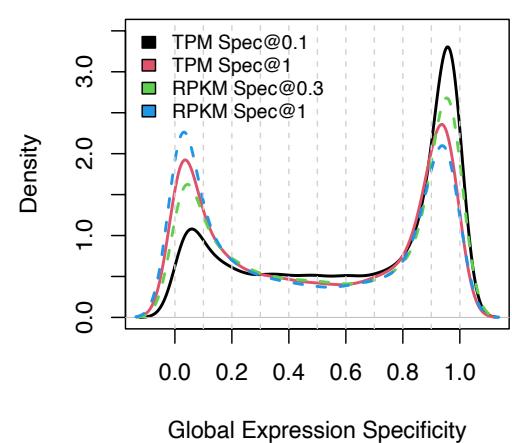
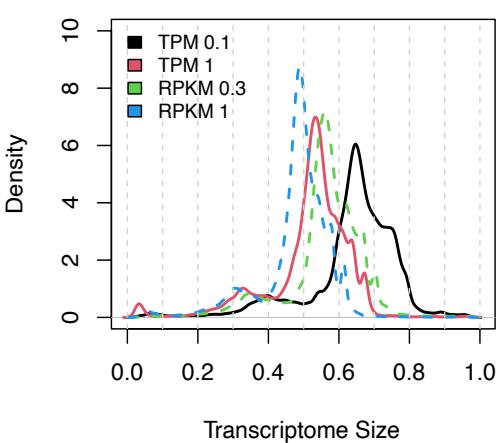
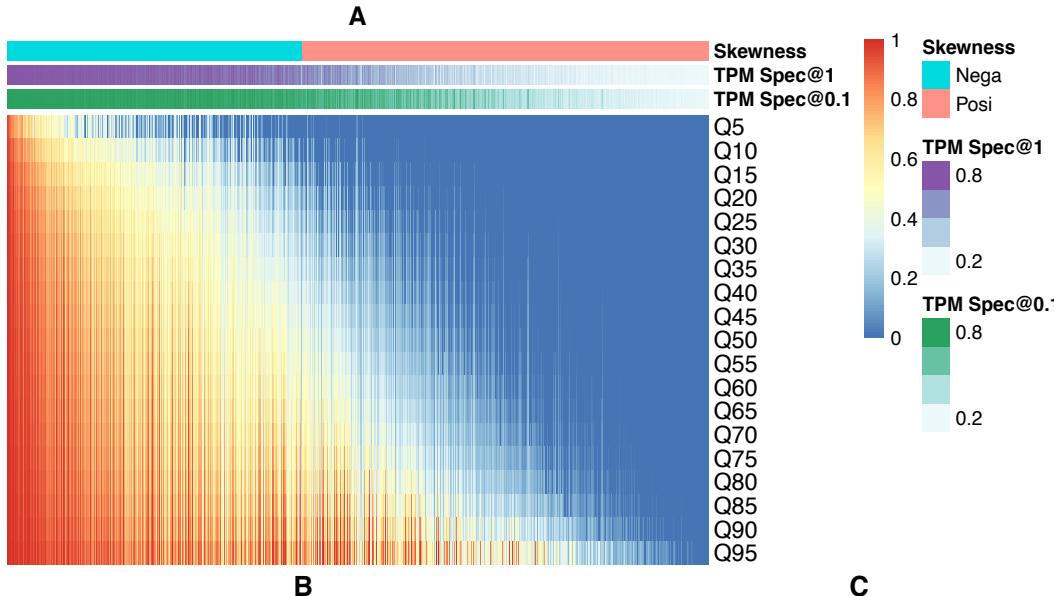


Fig 4. The Relationships between Expression Specificity, Expression Level and Expression Variability

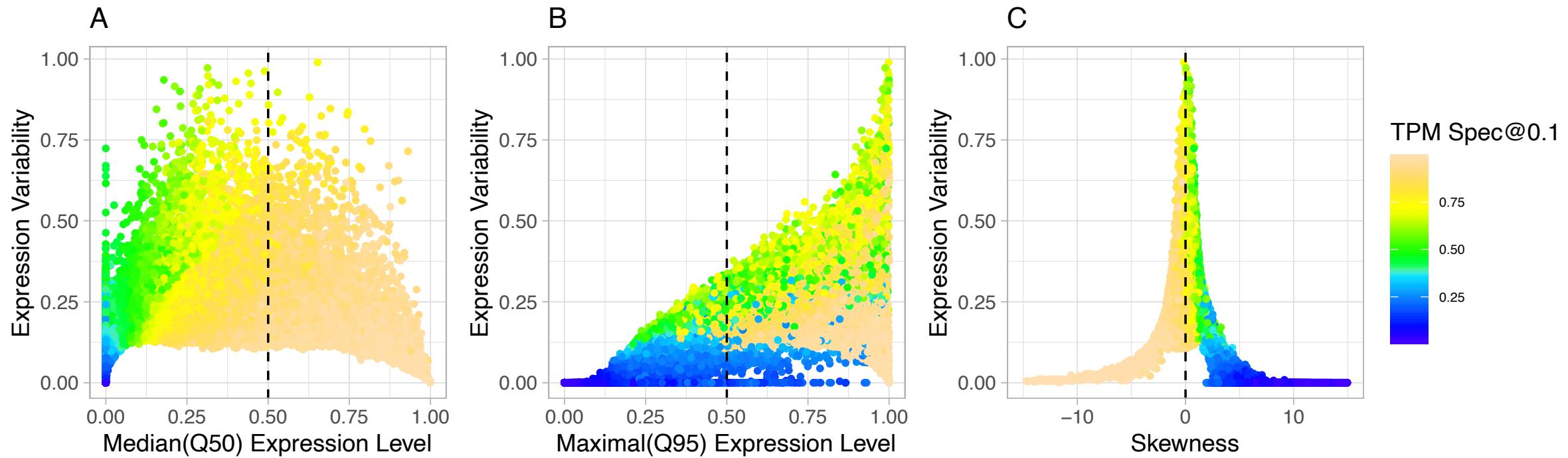


Fig 5. Global Expression Specificity Categories

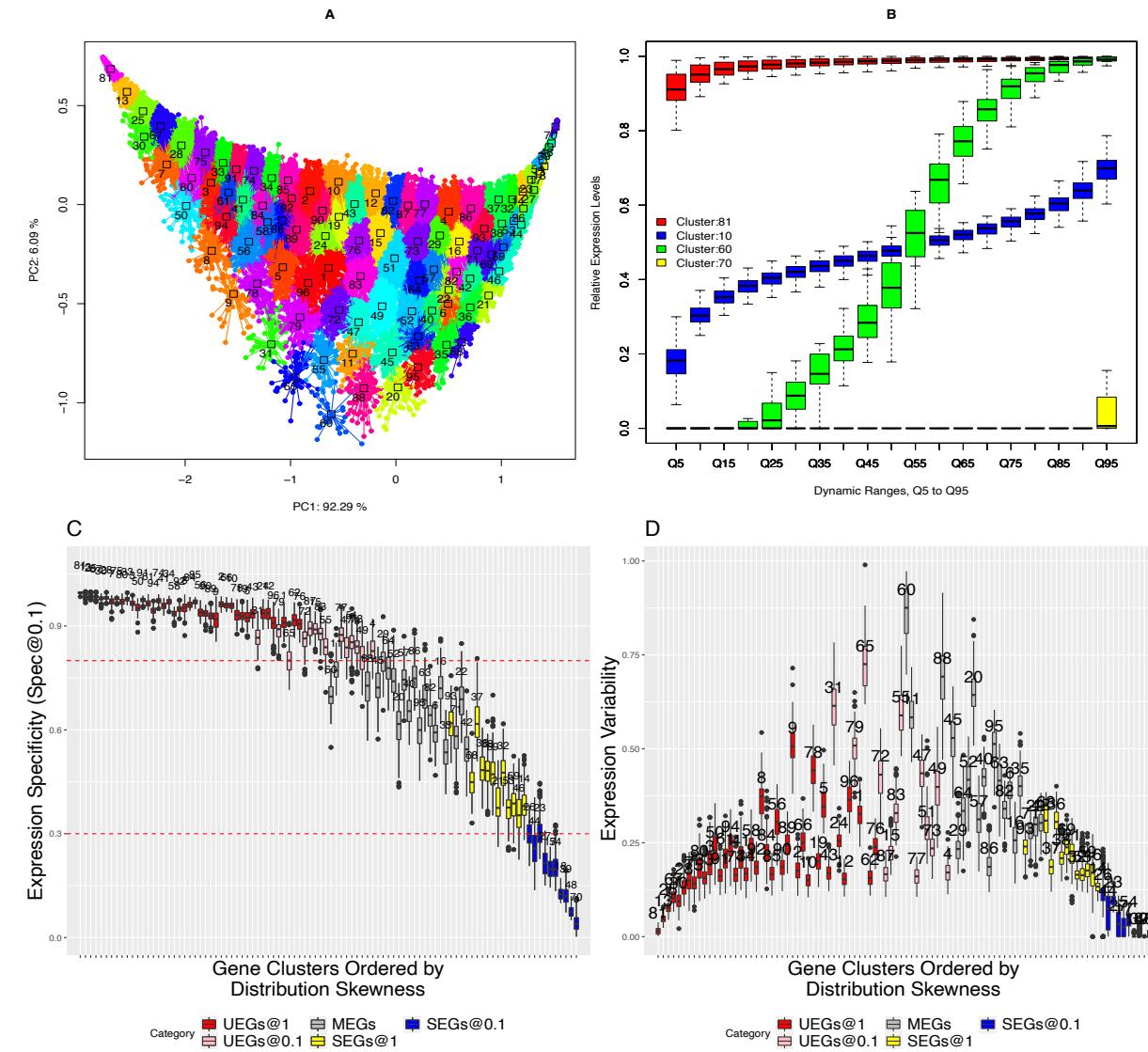


Fig 6. Evaluating the Uniqueness of Repression for Putative Disallowed Genes in Islets Beta Cells

