Database	Q. lobata	Q. alba EST	Q. robur EST
Q. lobata (this work)	—	79%	63%
Q. alba EST	19%	—	32%
Q. robur EST	55%	87%	—

Additional file 4: Overlap in gene content among oak RNA data sets.
Pairwise comparison of percentage of transcripts from each transcriptome appearing in the other.