Supplementary Figure S1. GO Annotation (according to DAVID v2.1, level 3) by Metabolic Function for the 330 transcripts that were present at significantly higher levels in unstimulated retinal endothelial cells (left) and the 449 transcripts that were present at significantly higher levels in unstimulated choroidal endothelial cells (right). Each horizontal bar indicates the percentage of transcripts falling within the stated functional category with respect to the total number of significantly up-regulated genes for that tissue. (A) Results for the 10 categories that contained the highest percentages of the 330 transcripts that were present at higher levels in retinal endothelial cells. (B) Results for the 10 categories that contained the highest percentages of the 449 transcripts that were present at higher levels in choroidal endothelial cells.