



Supplementary Figure S3. Venn diagram showing the number of differentially expressed genes in the transcriptome of human coronary artery endothelial cells (HCAEC, 729 DEGs) and human internal thoracic artery endothelial cells (HITAEC, 926 DEGs) at static cell culture conditions. DEGs were defined as those with logarithmical fold change ≥ 1 and FDR-corrected p value ≤ 0.05 .