



Supplementary Figure S2. Venn diagram showing the number of differentially expressed genes (DEGs) in the transcriptome of human coronary artery endothelial cells (HCAEC, 1,014 DEGs) and human internal thoracic artery endothelial cells (HITAEC, 849 DEGs) under laminar flow. DEGs were defined as those with logarithmical fold change ≥ 1 and FDR-corrected p value ≤ 0.05 .