



Figure 1 – figure supplement 5. Persistence of glycolytic and hypoxia genes under disturbed or unidirectional flow. HAECs were subjected to either DF or UF for 24, 48, or 72h prior to cell lysis, RNA purification, and qRT-PCR. Fold change of expression of *HK2*, *SLC2A1*, *EGLN3*, and *VEGFA*, relative to cells not treated with flow at each time point, are given above according to the color-code legend (n = 3 for each time point and each condition).