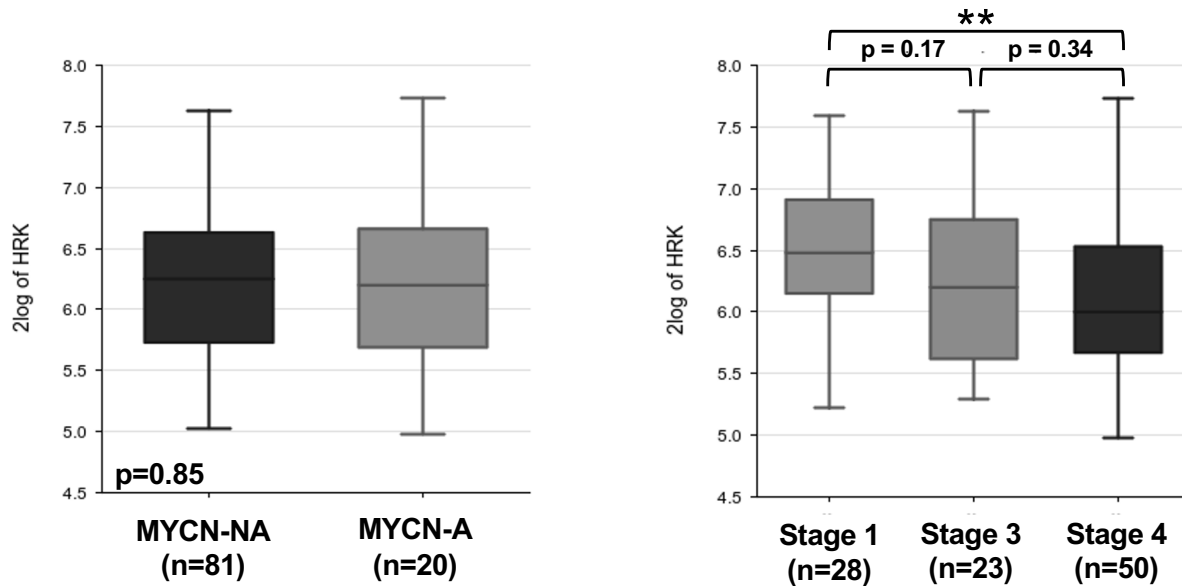
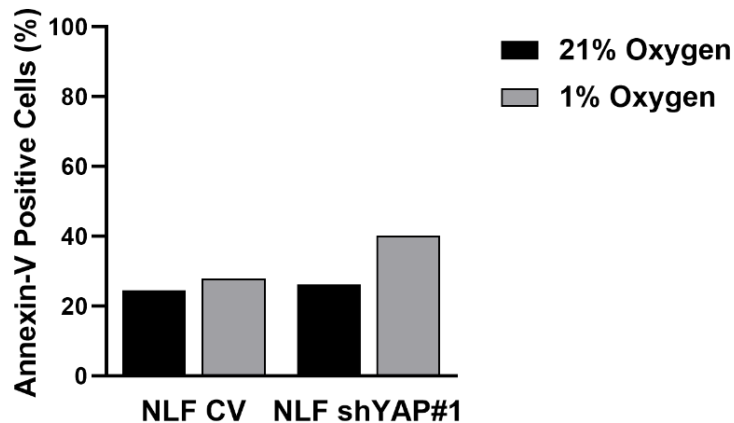


Supplementary Data



Supplementary Figure S7. *HRK* gene expression was evaluated in tumors based on *MYCN* status (left) and INSS staging (right) in the *Maris* cohort from the R2 database (<http://r2.amc.nl/>) as shown in box plots. **, $p < 0.01$.



Supplementary Figure S8. Apoptosis quantified via Annexin-V flow in NLF CV and shYAP#1 cells in normoxic (21% oxygen) or hypoxic (1% oxygen) conditions for 48 hours. Data represent $n=1$.