

Heatmap visualization of gene expression data. The color scale ranges from -2 (blue) to 2 (red), with 0 being white. The dendrogram on the left shows hierarchical clustering of genes, while the dendrogram on the top shows hierarchical clustering of samples. The samples are labeled at the top: Ba Ba Ba Ba Ba Ba Ba Ba Ba Ba LB HE HE HE HE HE No No No No HE HE HE LB LB No HE LB HE LA No LB LB LB LB LA LB LA LA LA LA LA LA.

Gene list (from top to bottom):

- MLPH
- FOXA1
- ESR1
- SLC39A6
- BLVRA
- BAG1
- ACTR3B
- BCL2
- GRB7
- ERBB2
- MMP11
- EGFR
- CDH3
- SFRP1
- PHGDH
- MYC
- CDC20
- ANLN
- TYMS
- UBE2C
- RRM2
- CENPF
- CCNB1
- NUF2
- NDC80
- KIF2C
- MKI67
- UBE2T
- CEP55
- MELK
- BIRC5
- ORC6
- KRT5
- KRT17
- KRT14
- PGR
- NAT1

Sample labels (from left to right):

- Ba Ba Ba Ba Ba Ba Ba Ba Ba Ba
- LB HE HE HE HE HE No No No No
- HE HE HE LB LB No HE LB HE LA No
- LB LB LB LB LA LB LA LA LA LA LA

