

ES = 0.426 NES = 1.08 Nom. p-val= 0.333 FWER= 0.996 FDR= 1

Heatmap visualization showing gene expression data across two clusters: **clus1** and **NON.clus1**. The y-axis lists 40 genes, and the x-axis lists 40 samples. The color scale ranges from -2 (blue) to 2 (red).

Gene List (Y-axis):

- Class
- CYP26C1
- CYP26A1
- ADH1C
- LRAT
- CYP1A2
- ALDH1A2
- ADH7
- CYP1A1
- ADH6
- RDH11
- CYP4A11
- RDH10
- CYP2C9
- ADH4
- UGT2B7
- ALDH1A1
- ADH1A
- CYP2C8
- CYP2C18
- DHRS3
- UGT1A6
- CYP3A43
- RDH16
- CYP3A7
- UGT1A9
- DHRS4
- ADH5
- DHRS4L2
- CYP2A6
- DHRS9
- RETSA1
- UGT2A3
- PNPLA4
- UGT2B10
- DGAT1
- DGAT2
- RPE65
- RDH5
- ADH1B
- CYP26B1
- RDH12
- UGT2B11
- CYP3A5
- UGT2B28
- UGT2A1
- BCMO1
- UGT2B4

Sample List (X-axis):

- TCGA_KL_1322_01
- TCGA_KL_1327_01
- TCGA_KL_1328_01
- TCGA_KL_1334_01
- TCGA_KL_1338_01
- TCGA_AL_1341_01
- TCGA_KL_1345_01
- TCGA_KM_1345_01
- TCGA_KM_1346_01
- TCGA_KN_1348_01
- TCGA_KN_1354_01
- TCGA_KM_1355_01
- TCGA_KM_1356_01
- TCGA_KL_1357_01
- TCGA_KL_1358_01
- TCGA_KL_1359_01
- TCGA_KL_1360_01
- TCGA_KL_1361_01
- TCGA_KL_1362_01
- TCGA_KL_1363_01
- TCGA_KL_1364_01
- TCGA_KL_1365_01
- TCGA_KM_1366_01
- TCGA_KM_1367_01
- TCGA_KM_1368_01
- TCGA_KL_1369_01
- TCGA_KL_1370_01
- TCGA_KL_1371_01
- TCGA_KL_1372_01
- TCGA_KL_1373_01
- TCGA_KL_1374_01
- TCGA_KL_1375_01
- TCGA_KL_1376_01
- TCGA_KL_1377_01
- TCGA_KM_1378_01
- TCGA_KM_1379_01
- TCGA_KN_1380_01
- TCGA_KN_1381_01
- TCGA_KN_1382_01
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- TCGA_KN_1408_01
- TCGA_KN_1409_01
- TCGA_KN_1410_01
- TCGA_KN_1411_01
- TCGA_KN_1412_01
- TCGA_KN_1413_01
- TCGA_KN_1414_01
- TCGA_KN_1415_01