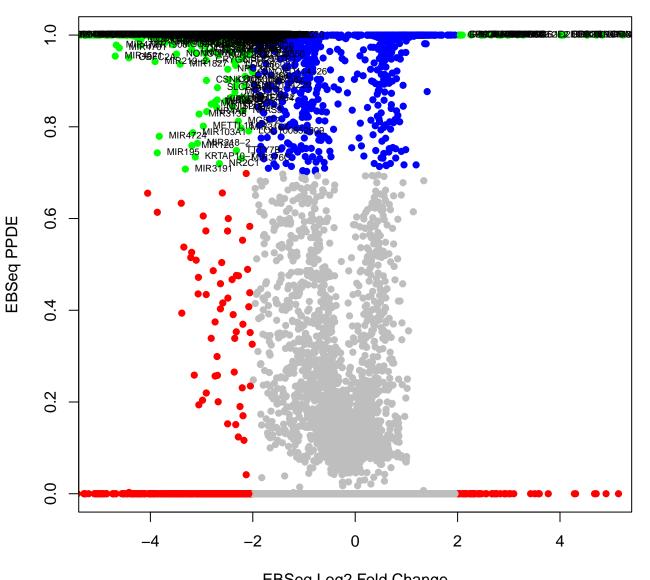
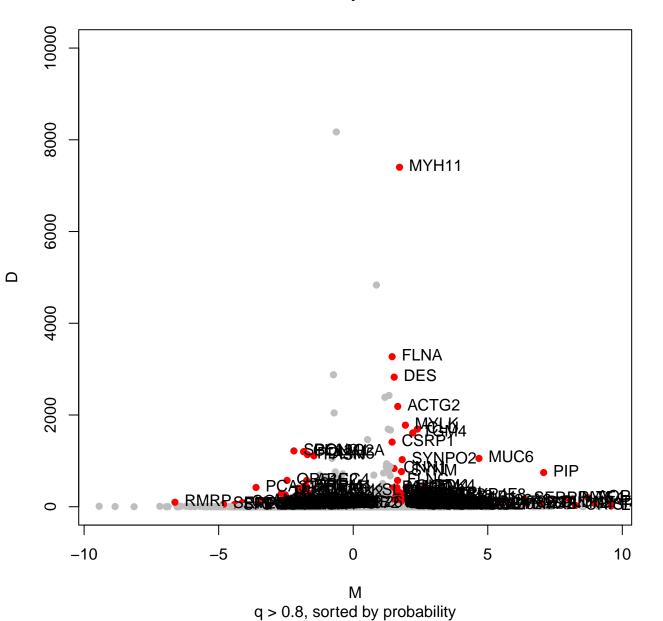
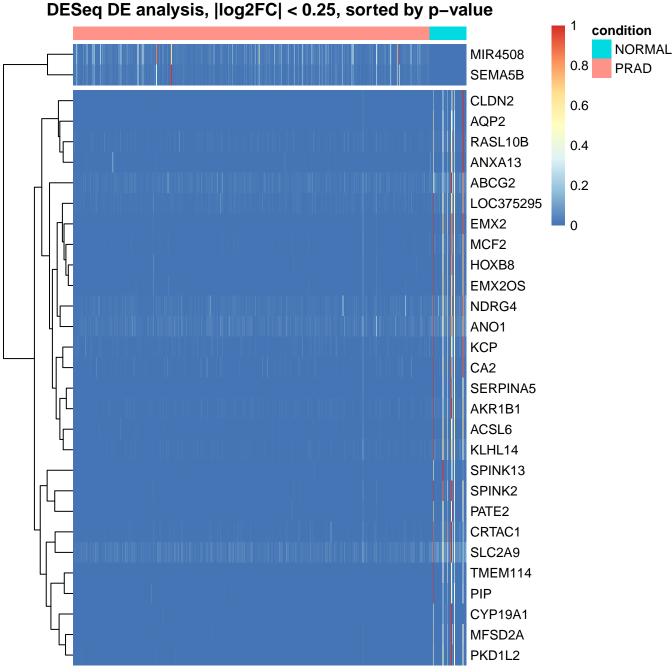
## **EBSeq results**

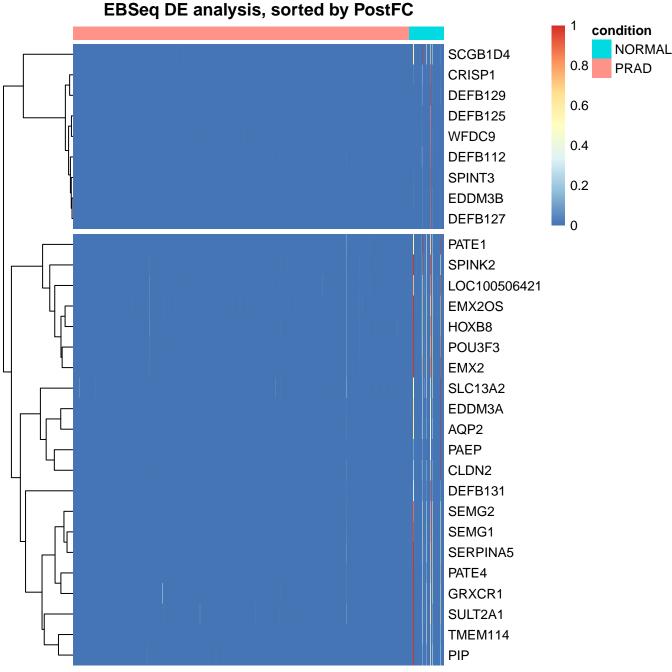


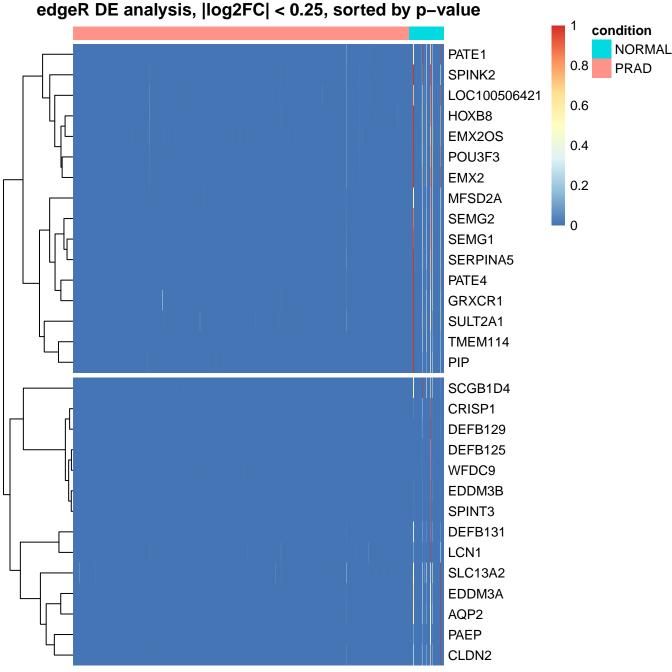
EBSeq Log2 Fold Change sorted by PostFC

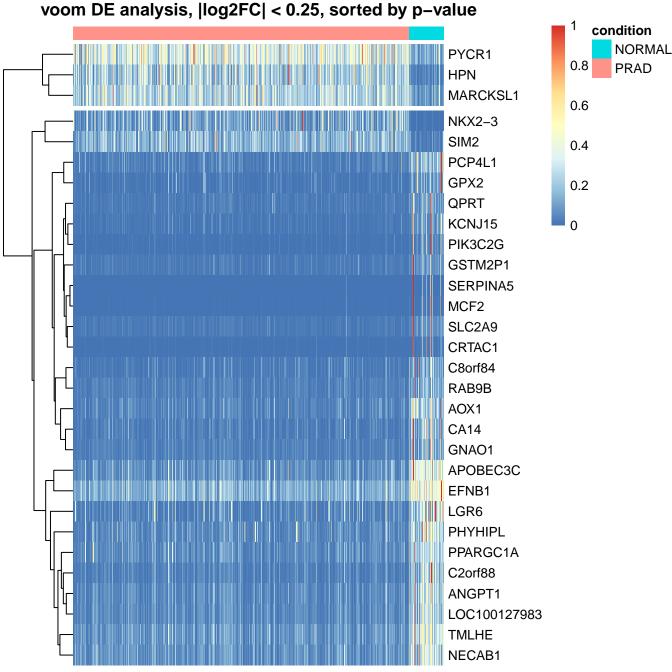
## **NOISeq** results











**NOISeq DE analysis, q > 0.8, sorted by probability** condition **NORMAL** AMACR 8.0 PRAD COL2A1 SLPI 0.6 MUC6 CLDN2 0.4 AQP2 PAEP 0.2 PDK4 PTGS2 0 STAC2 PATE1 SPINK2 CYP4F8 MFSD2A PIP PATE4 SLC26A3 AKR1B1 SEMG2 SEMG1 SERPINA5 SEMA5B KRT13 **RMRP** RPPH1 TFF3 ACSM1 PCA3 OR51E2 HPN