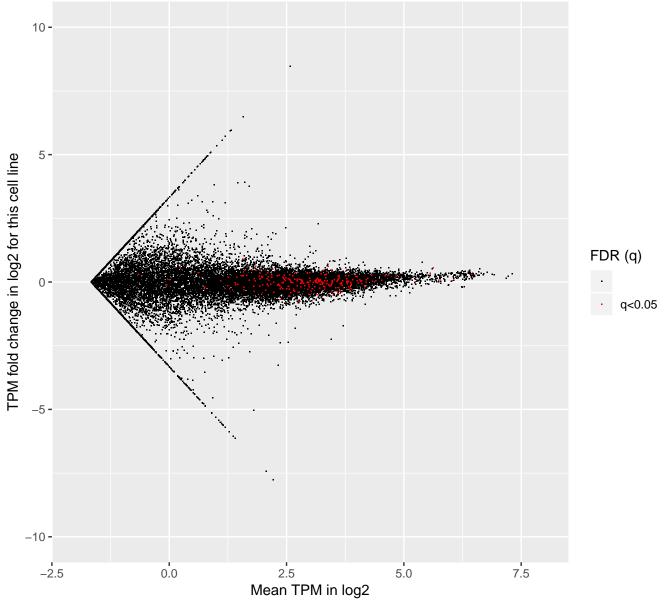
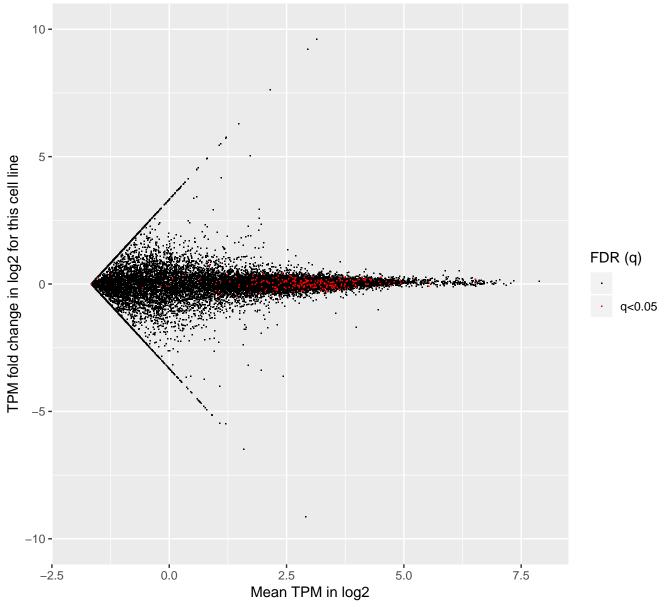
A375
Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line)

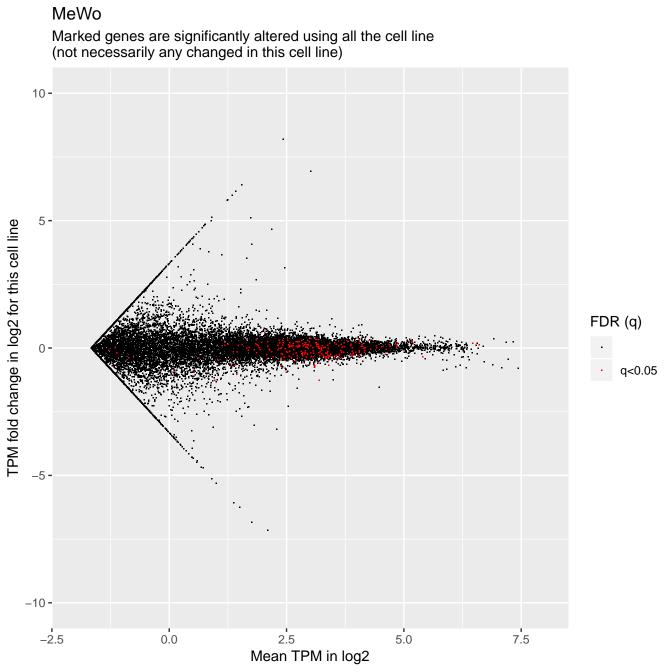


Femxl Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10-5 -TPM fold change in log2 for this cell line FDR (q) 0 q<0.05 -5 **-**. -10 **-**0.0 2.5 5.0 -2.5 7.5 Mean TPM in log2

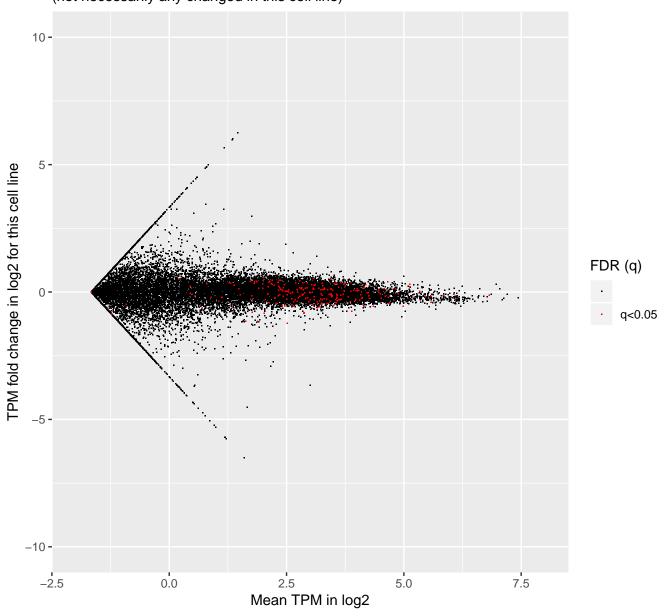
FemxV Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10-5 -TPM fold change in log2 for this cell line FDR (q) 0 q<0.05 -5 **-**-10 **-**0.0 2.5 5.0 -2.5 7.5 Mean TPM in log2

LOX

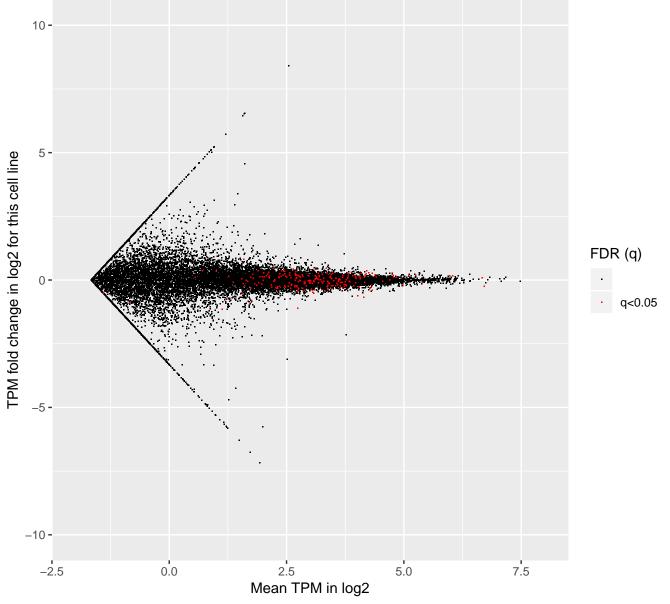




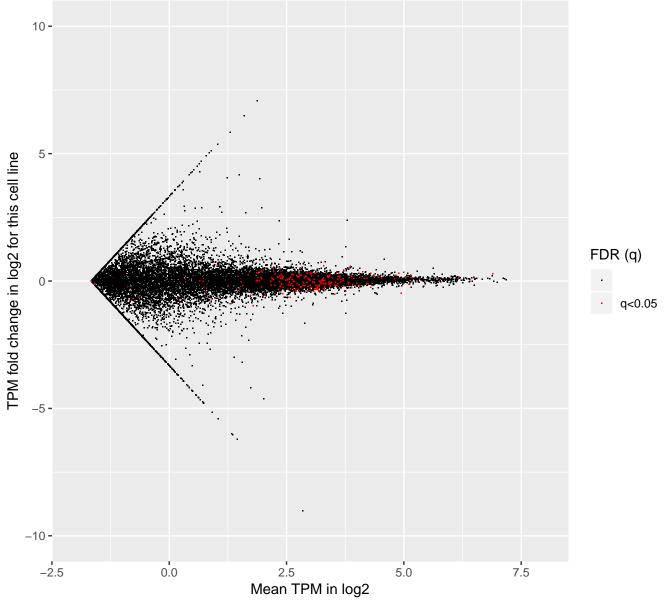
Skmel28



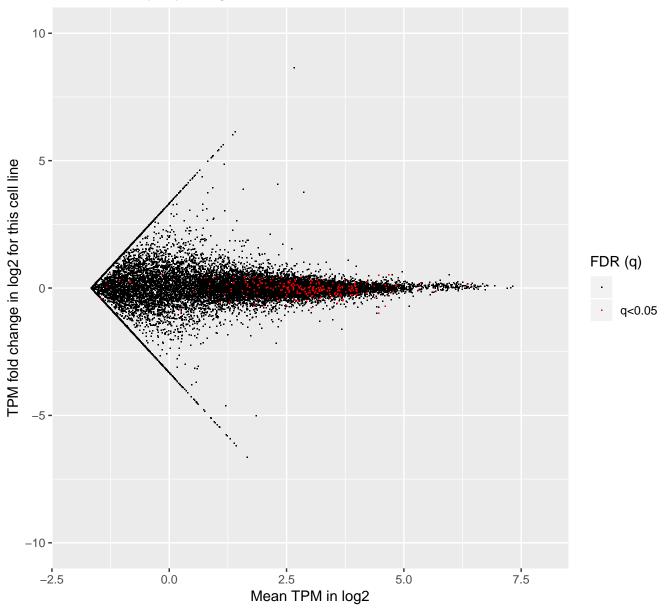
WM115



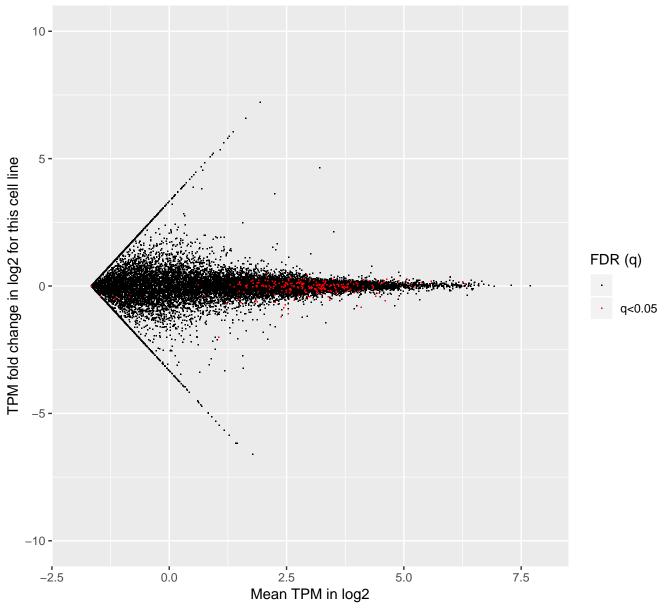
WM1341



WM1366



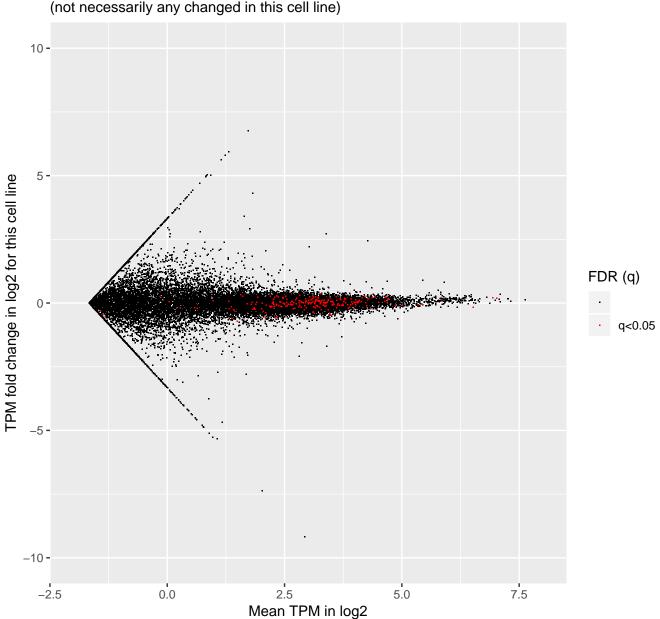
WM1382



WM239 Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10 -5 -TPM fold change in log2 for this cell line FDR (q) 0 q<0.05 -5 **-**-10 **-**0.0 2.5 5.0 -2.5 7.5

Mean TPM in log2

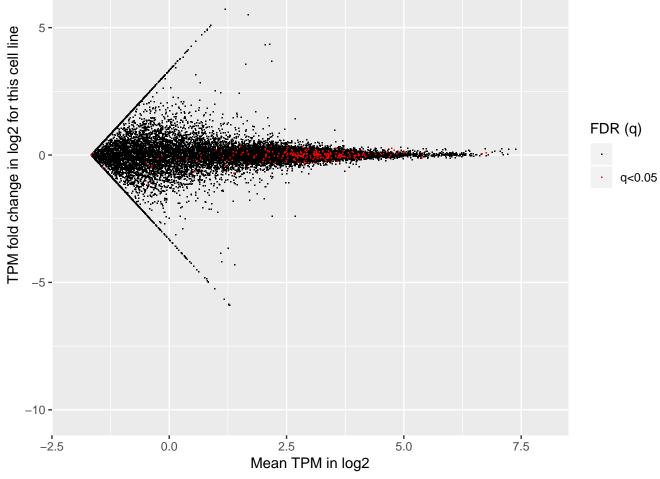
WM266 Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10 -5 -



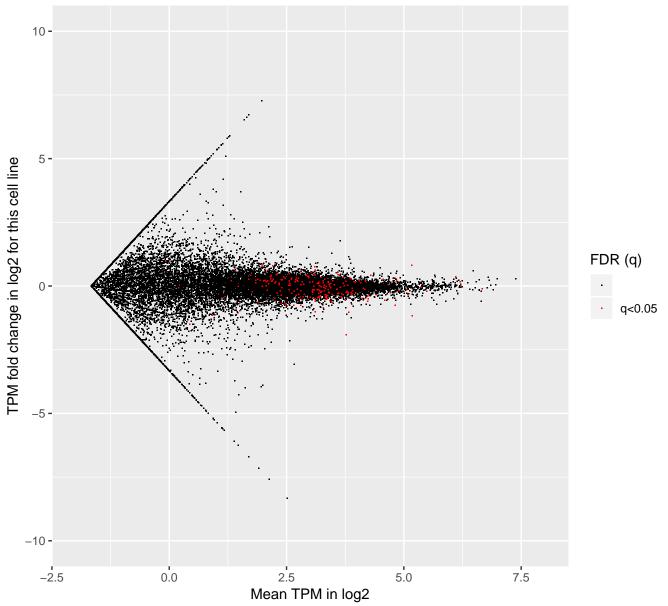
WM35 Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10-5 -TPM fold change in log2 for this cell line FDR (q) 0 q<0.05 -5 **-**-10 **-**0.0 2.5 5.0 -2.5 7.5

Mean TPM in log2

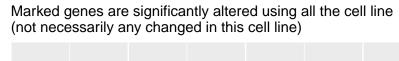
WM45-1 Marked genes are significantly altered using all the cell line (not necessarily any changed in this cell line) 10 -5 -FDR (q) 0 q<0.05 -5 **-**

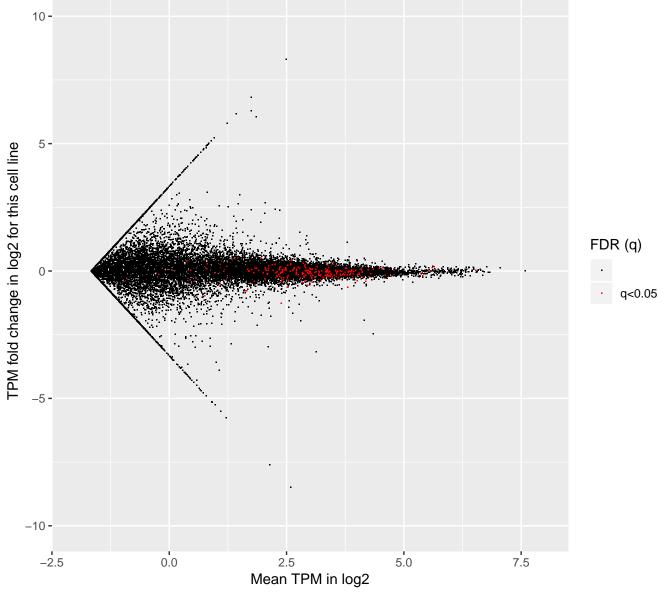


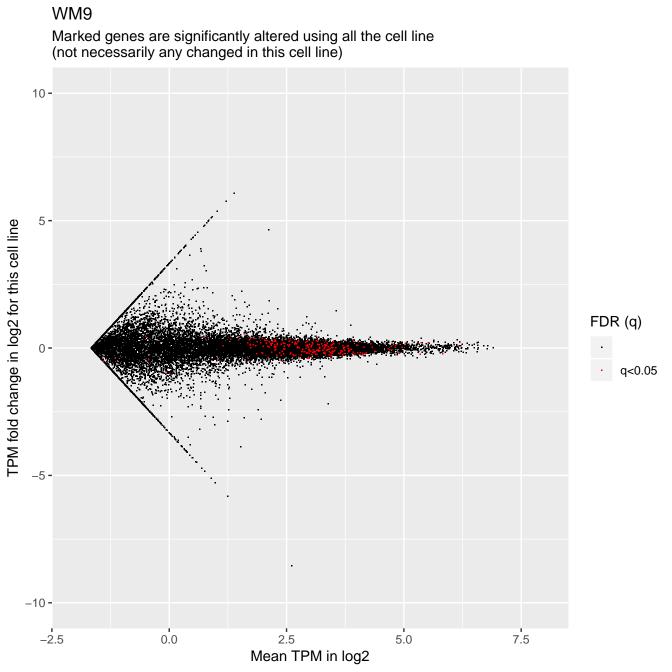
WM793



WM852







WM983

