

Udder from SRA archive and mEV gene set comparison.

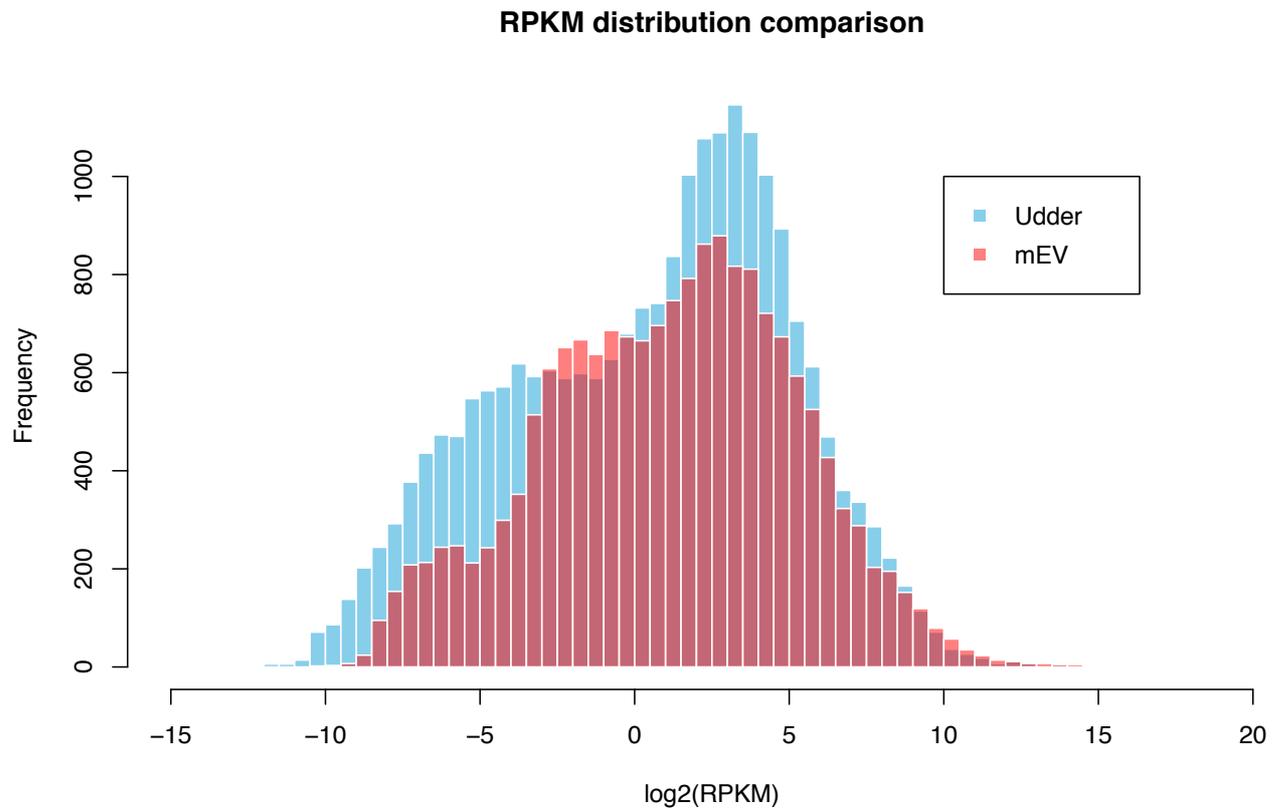


Figure S1. Histogram of RPKM mean for udder data (blue) downloaded from SRA database (BioSamples: SAMN12831050, SAMN12831049, SAMN14600526 and SAMN14600525) and mEV gene set (red).