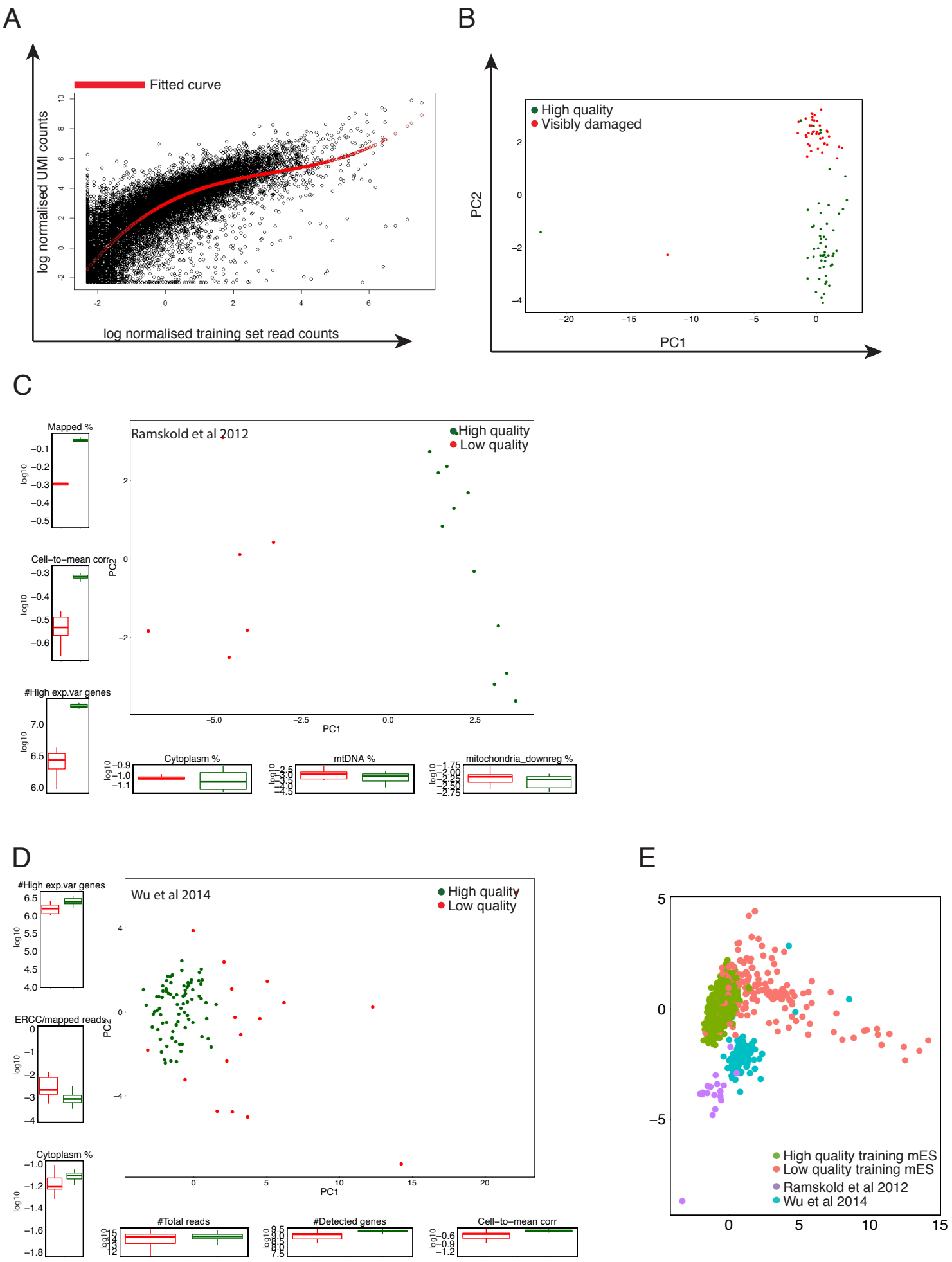
Figure S4


A
Fitted curve
log normalised UMI counts
log normalised training set read counts
B
High quality
Visibly damaged
PC2
PC1
C
Mapped %
Cell−to−mean corr
#High exp.var genes
Ramskold et al 2012
High quality
Low quality
PC2
PC1
Cytoplasm %
mtDNA %
mitochondria_downreg %
D
#High exp.var genes
ERCC/mapped reads
Cytoplasm %
Wu et al 2014
High quality
Low quality
PC2
PC1
#Total reads
#Detected genes
Cell−to−mean corr
E
High quality training mES
Low quality training mES
Ramskold et al 2012
Wu et al 2014