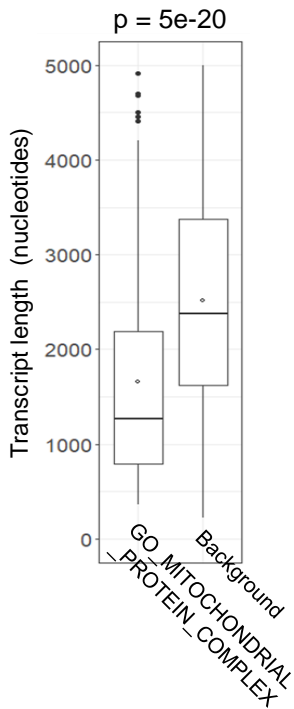
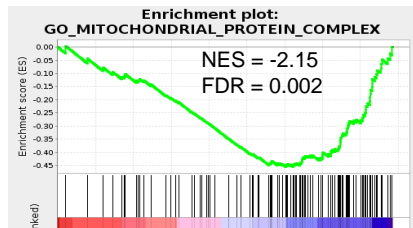


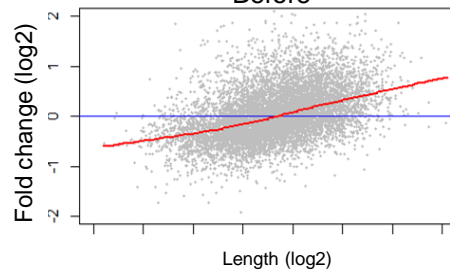
a Dataset #1: TNF (GSE64233): Replicate comparison

Original data

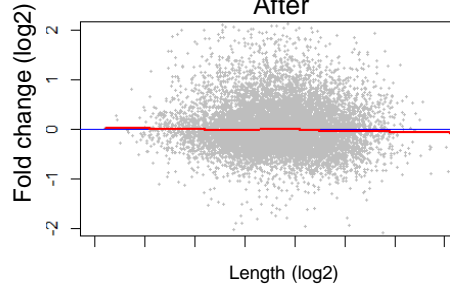


cqn normalization

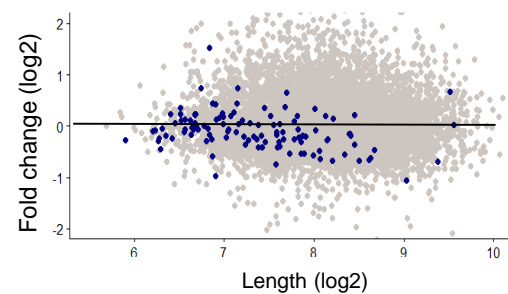
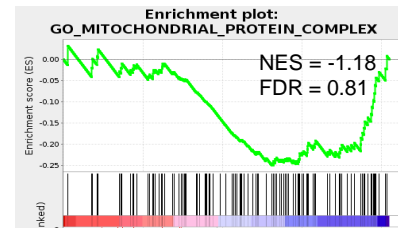
Before



After

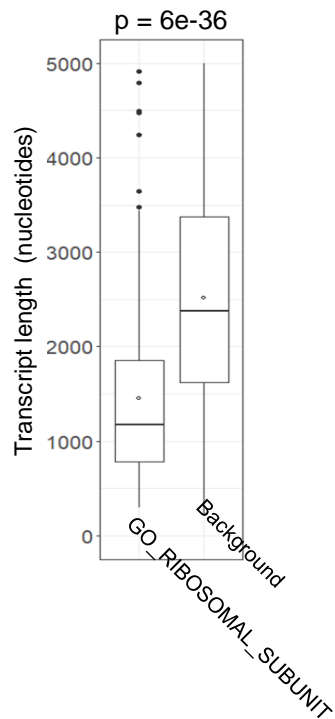
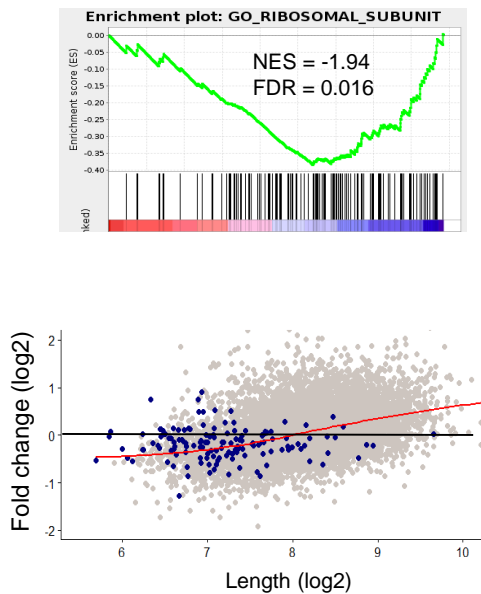


Corrected data

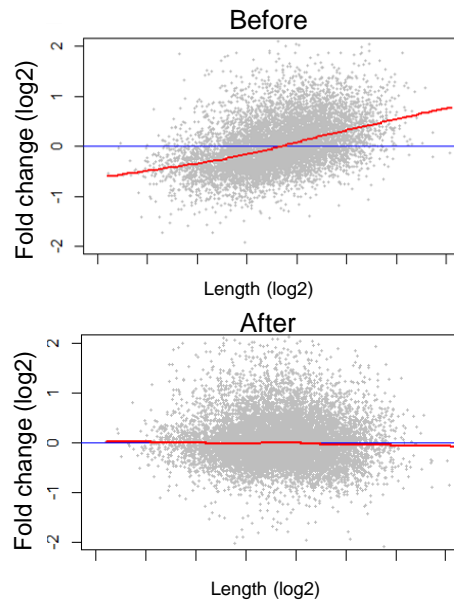


b Dataset #1: TNF (GSE64233): Replicate comparison

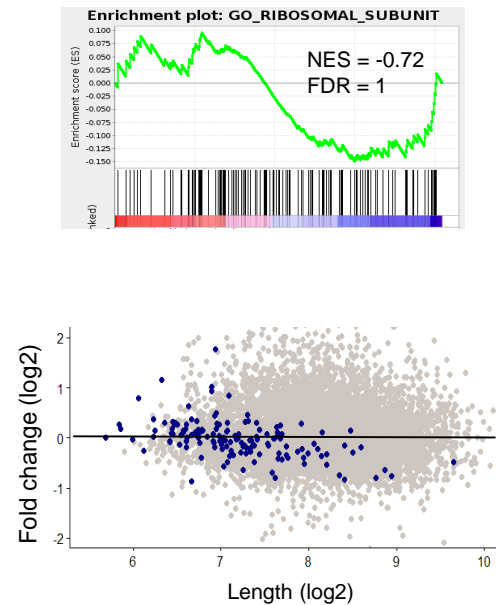
Original data



cqn normalization

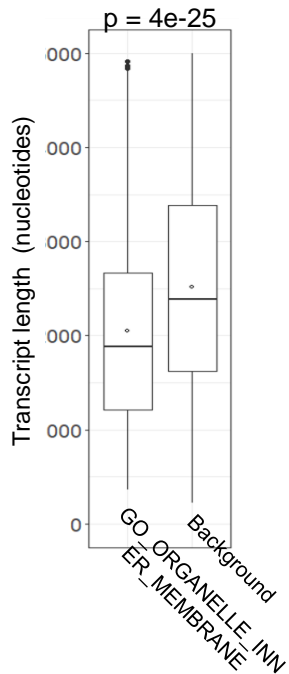
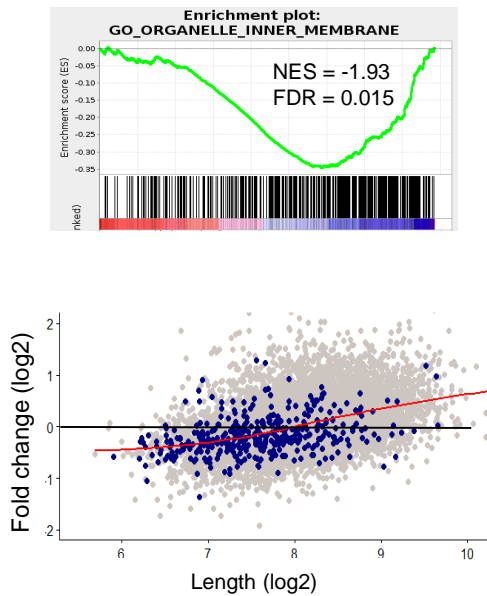


Corrected data

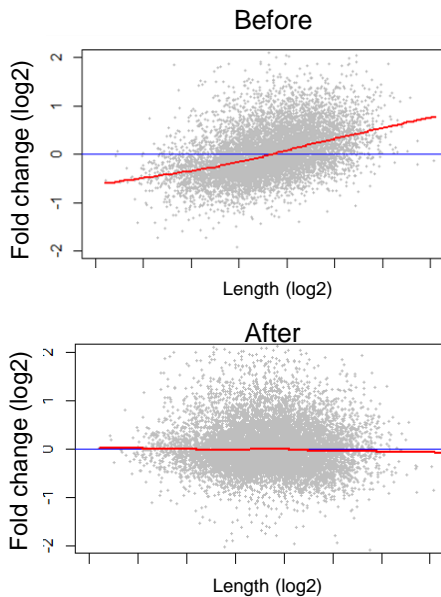


C Dataset #1: TNF (GSE64233): Replicate comparison

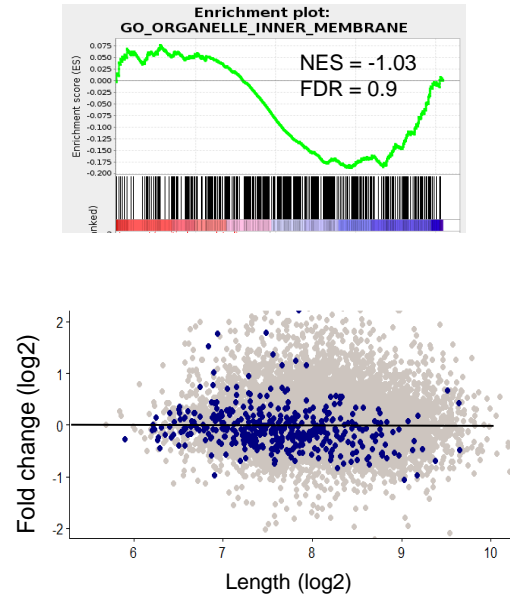
Original data



cqn normalization

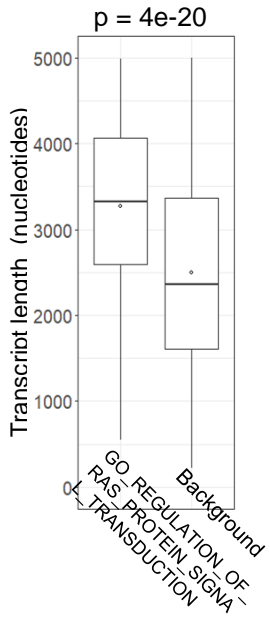
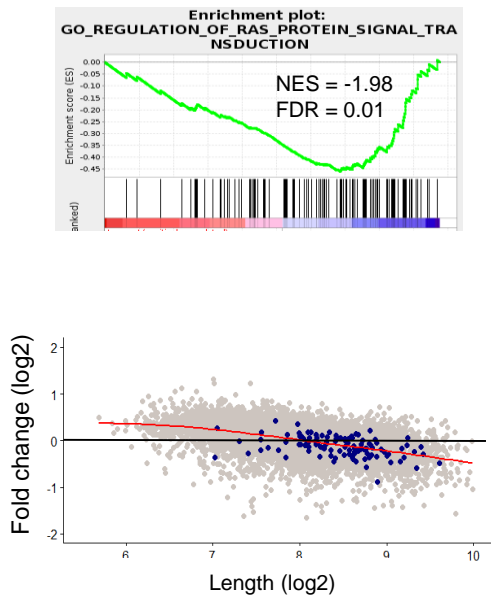


Corrected data

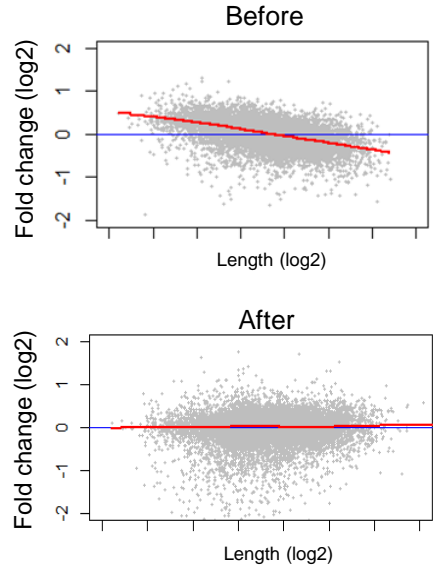


d Dataset #10: shSRC-1 in BC cells (GSE99647): Replicate comparison

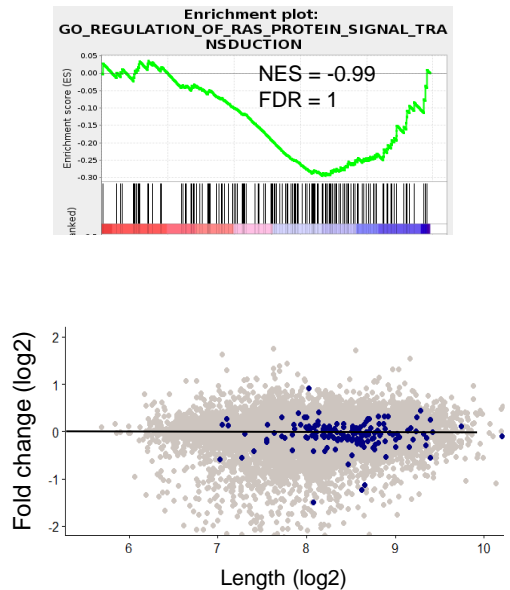
Original data



cqn normalization

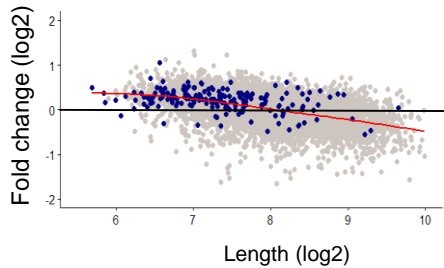
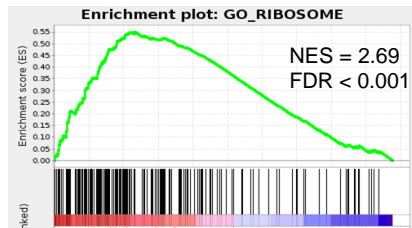


Corrected data



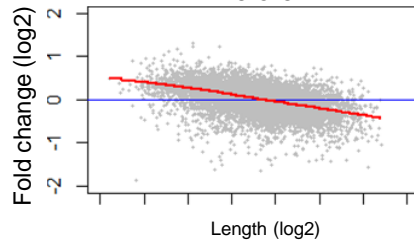
e Dataset #10: shSRC-1 in BC cells (GSE99647): Replicate comparison

Original data

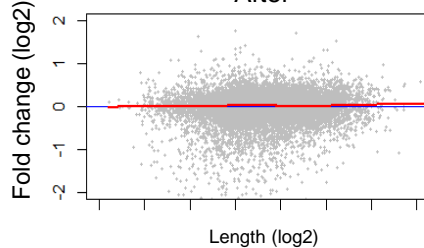


cqn normalization

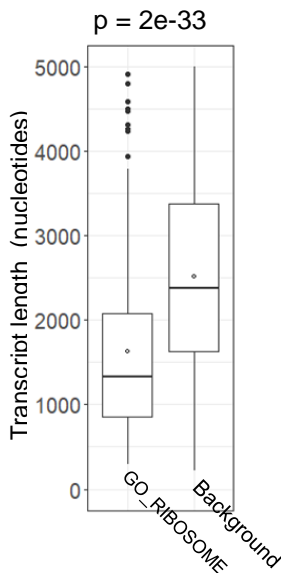
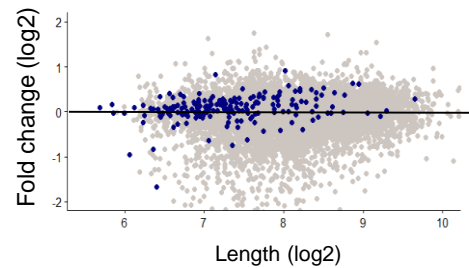
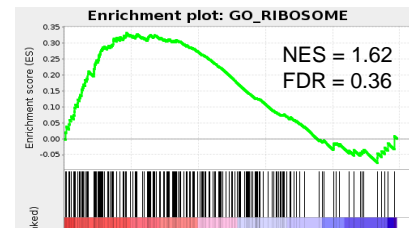
Before



After

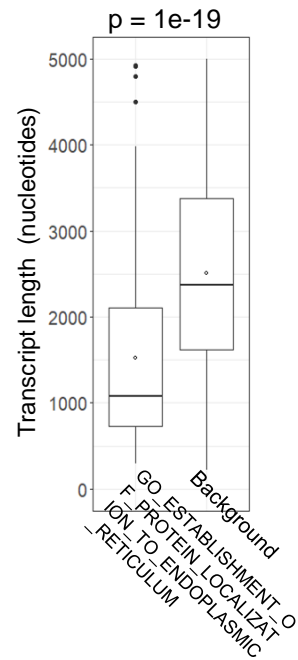
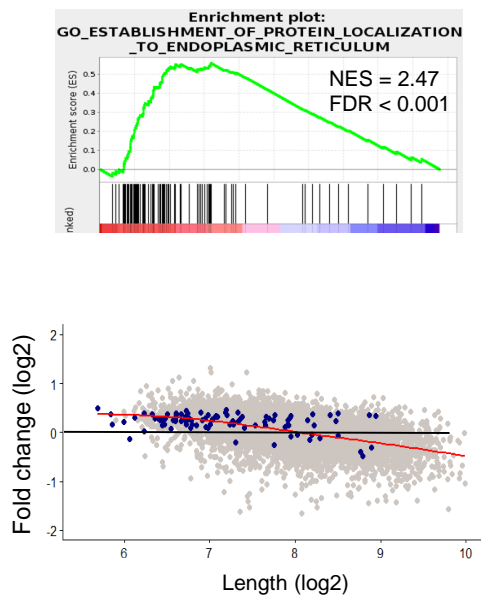


Corrected data

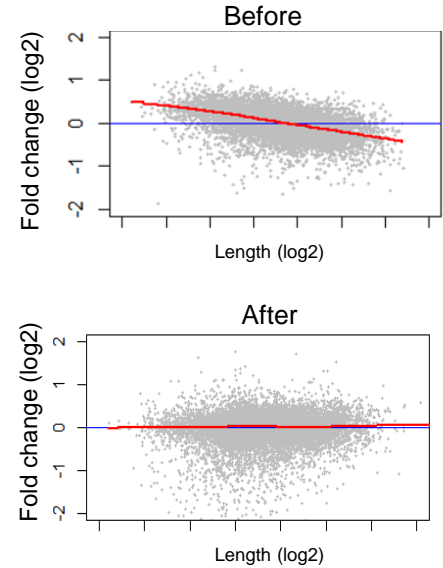


f Dataset #10: shSRC-1 in BC cells (GSE99647): Replicate comparison

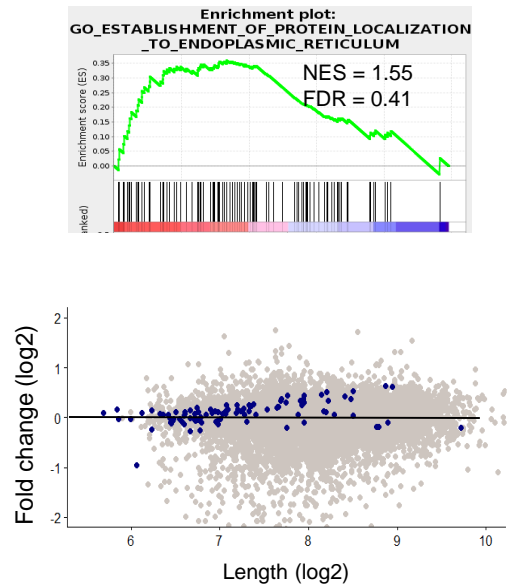
Original data



cqn normalization

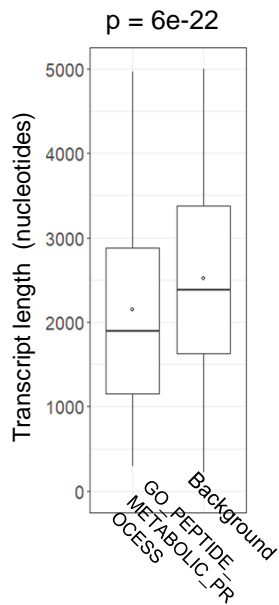
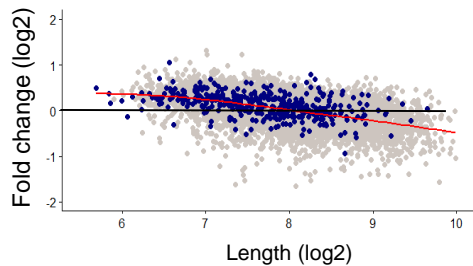
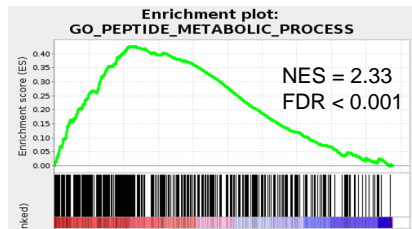


Corrected data



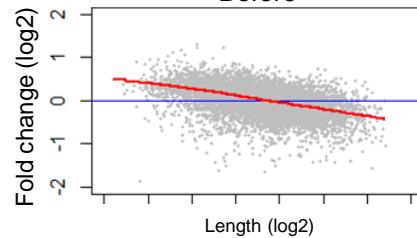
9 Dataset #10: shSRC-1 in BC cells (GSE99647): Replicate comparison

Original data

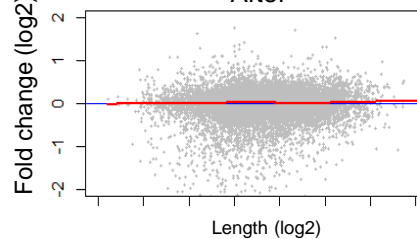


cqn normalization

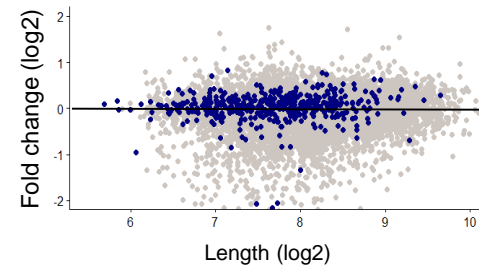
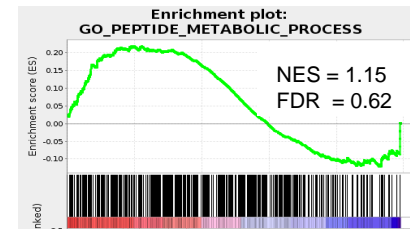
Before



After

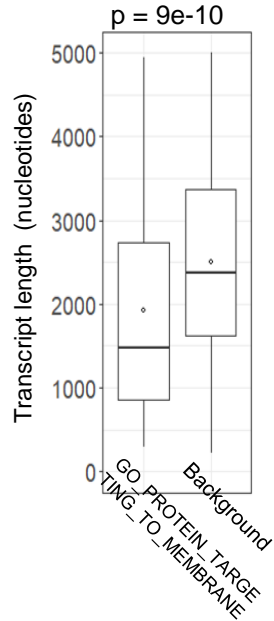
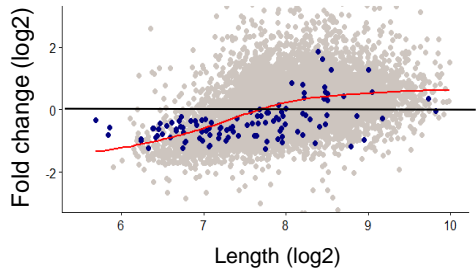
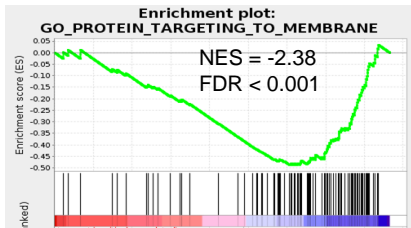


Corrected data

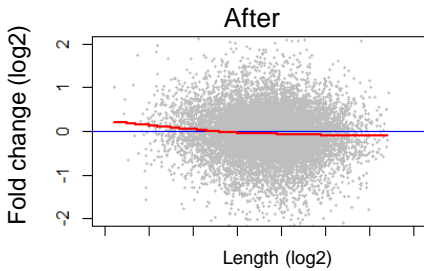
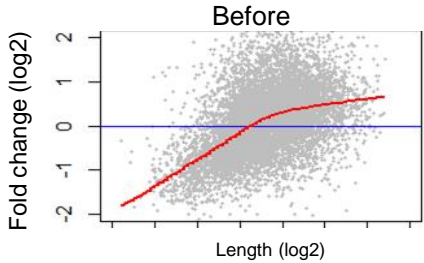


h Dataset #8: NCS treated vs. untreated MCF-7 cells (GSE101738): Replicate comparison

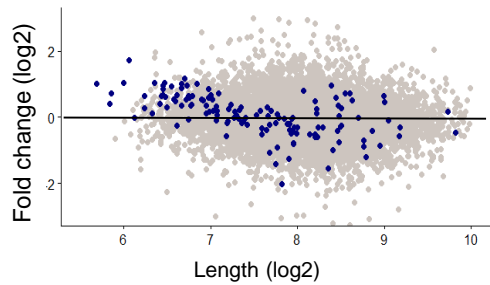
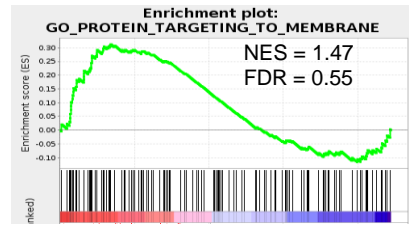
Original data



cqn normalization



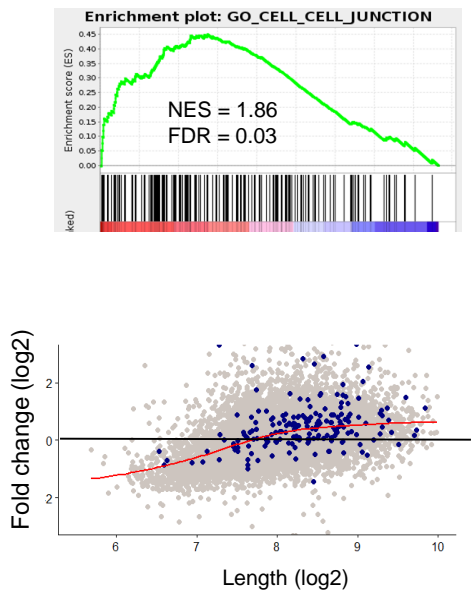
Corrected data



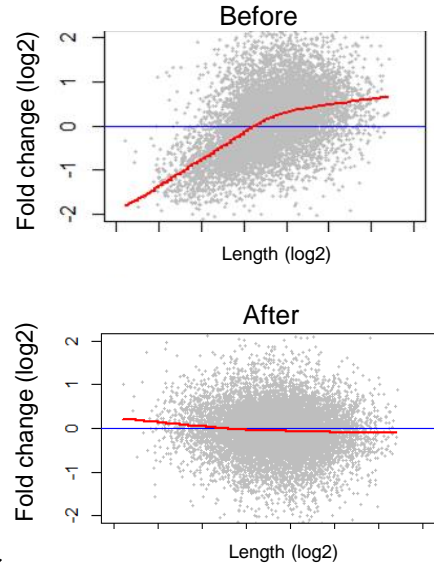
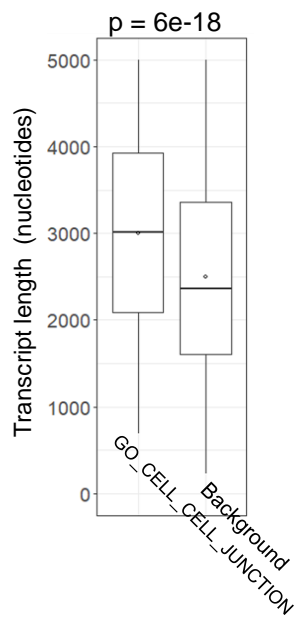


i Dataset #8: NCS treated vs. untreated MCF-7 cells (GSE101738): Replicate comparison

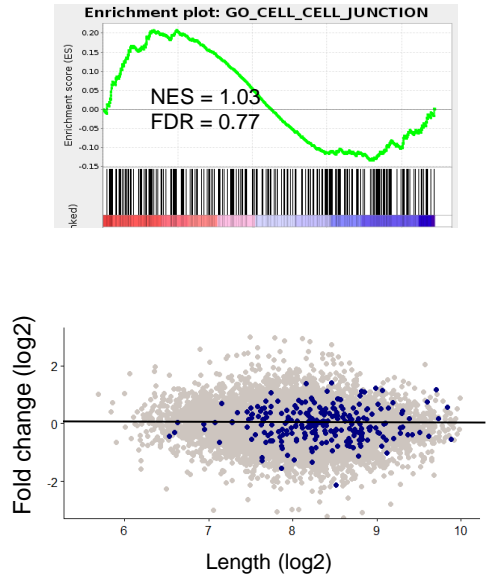
Original data



cqn normalization

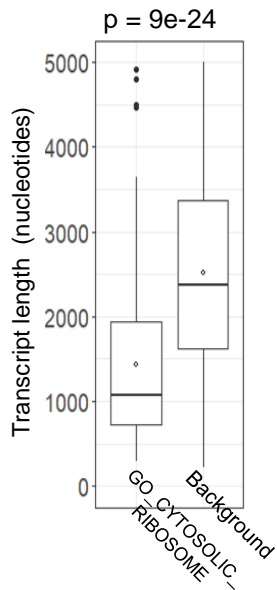
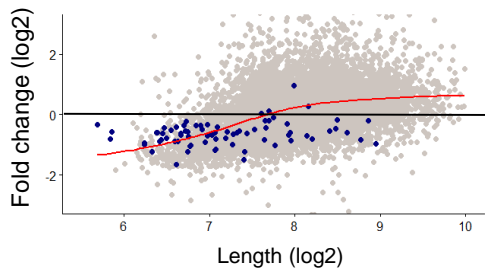
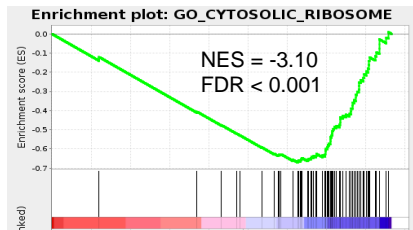


Corrected data

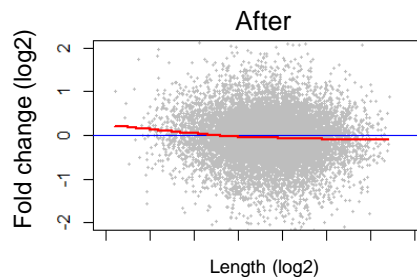
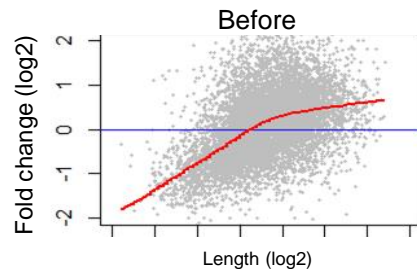


j Dataset #8: NCS treated vs. untreated MCF-7 cells (GSE101738): Replicate comparison

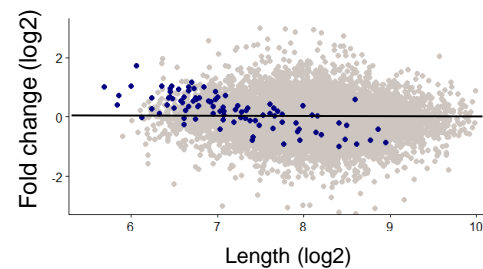
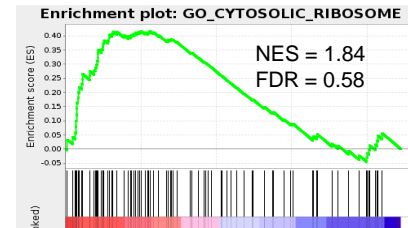
Original data



cqn normalization

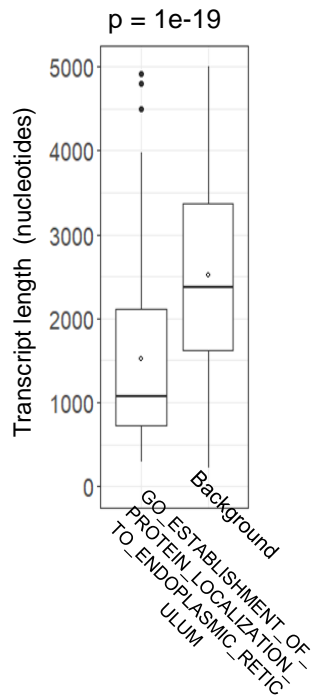
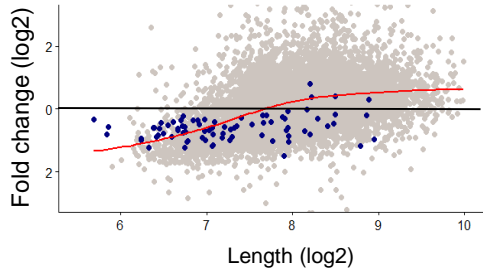
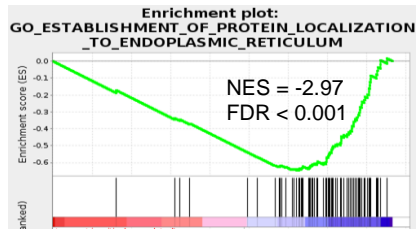


Corrected data

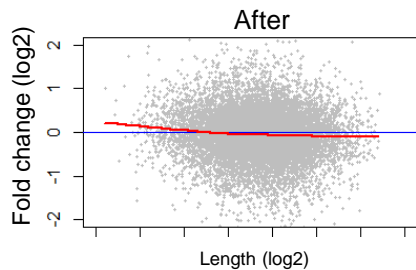
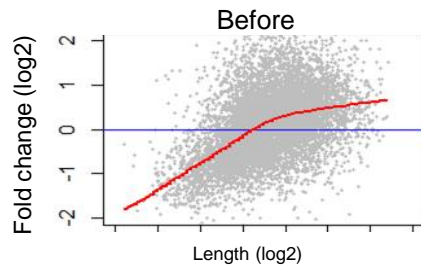


k Dataset #8: NCS treated vs. untreated MCF-7 cells (GSE101738): Replicate comparison

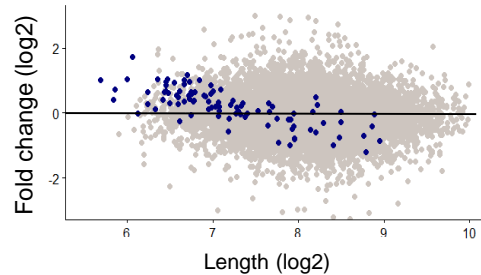
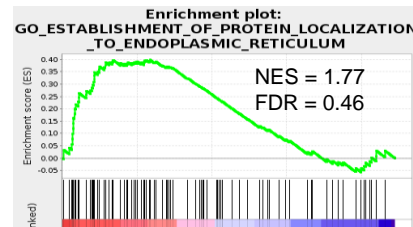
Original data



cqn normalization

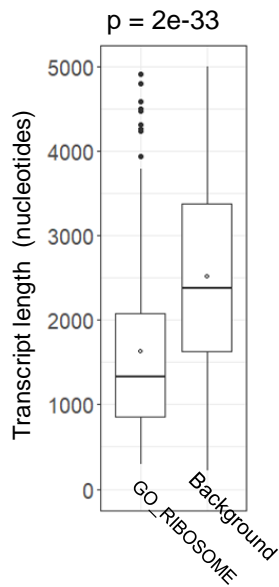
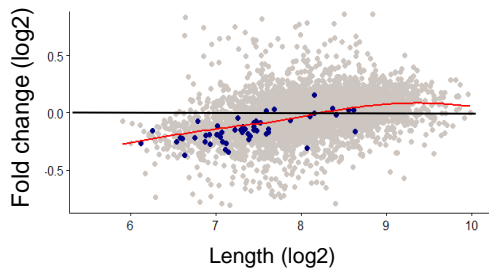
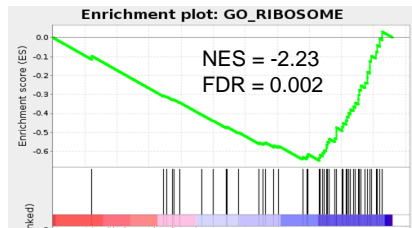


Corrected data

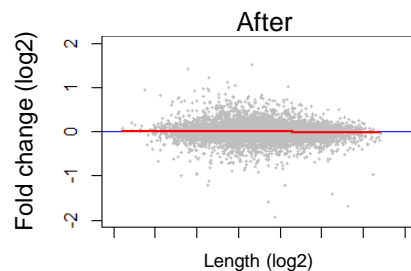
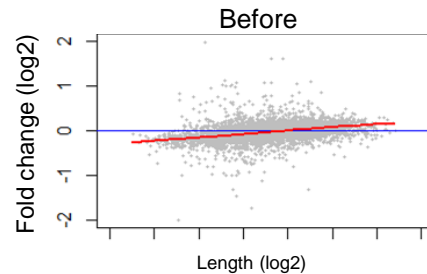


I Dataset #18: AI-10-49 treatment on ME-1 cells (GSE101788): Replicate comparison

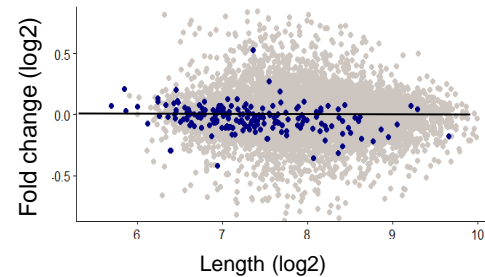
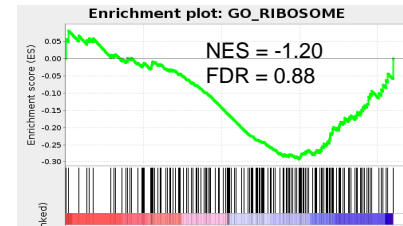
Original data



cqn normalization

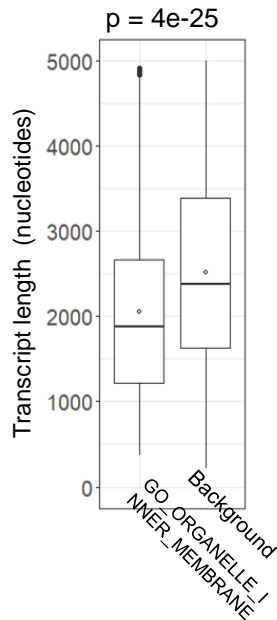
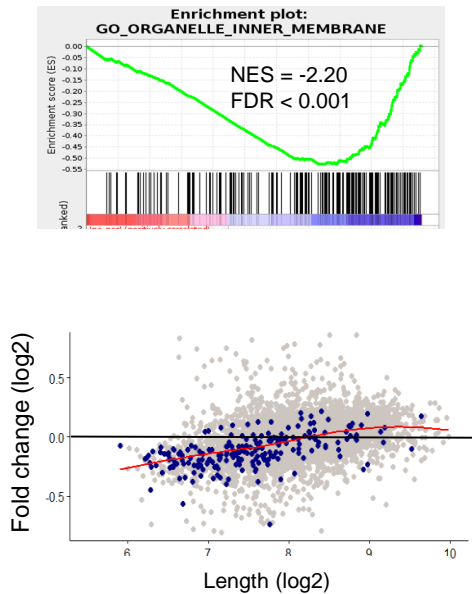


Corrected data

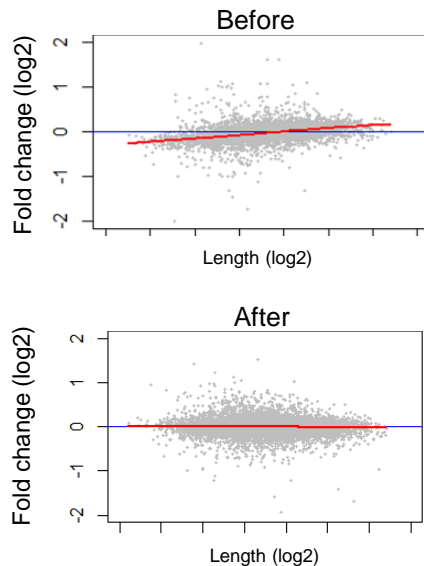


m Dataset #18: AI-10-49 treatment on ME-1 cells (GSE101788): Replicate comparison

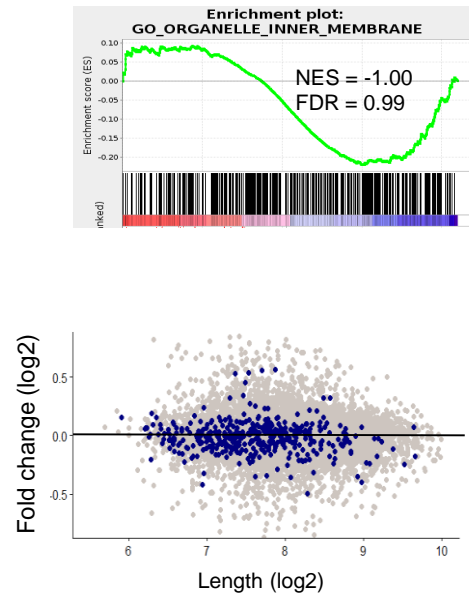
Original data



cqn normalization

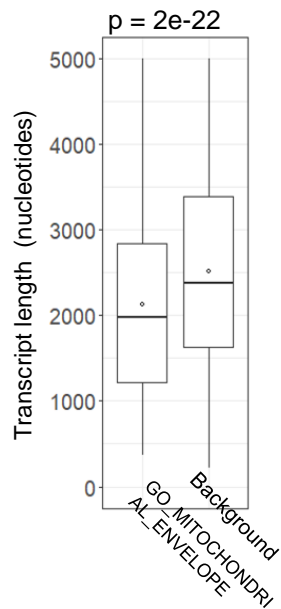
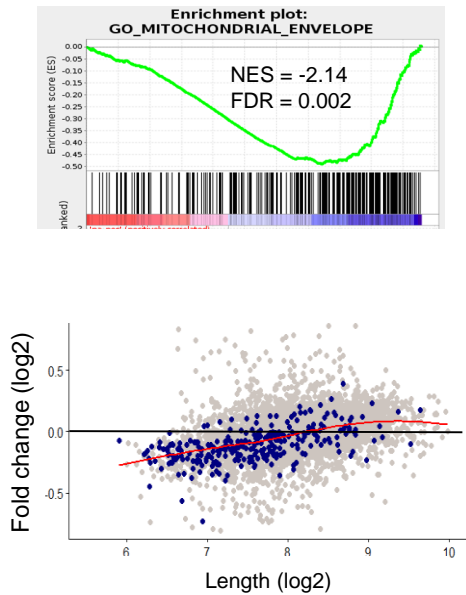


Corrected data

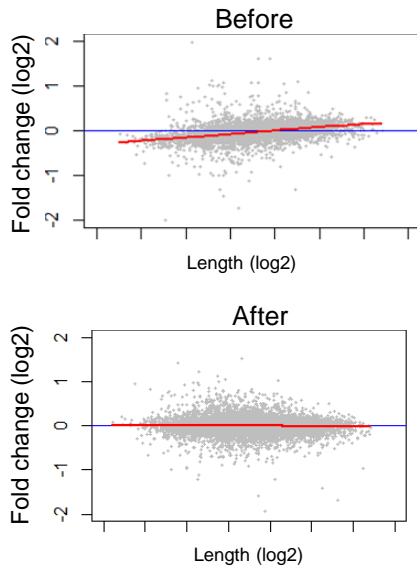


n Dataset #18: AI-10-49 treatment on ME-1 cells (GSE101788): Replicate comparison

Original data



cqn normalization



Corrected data

