

Figure S6

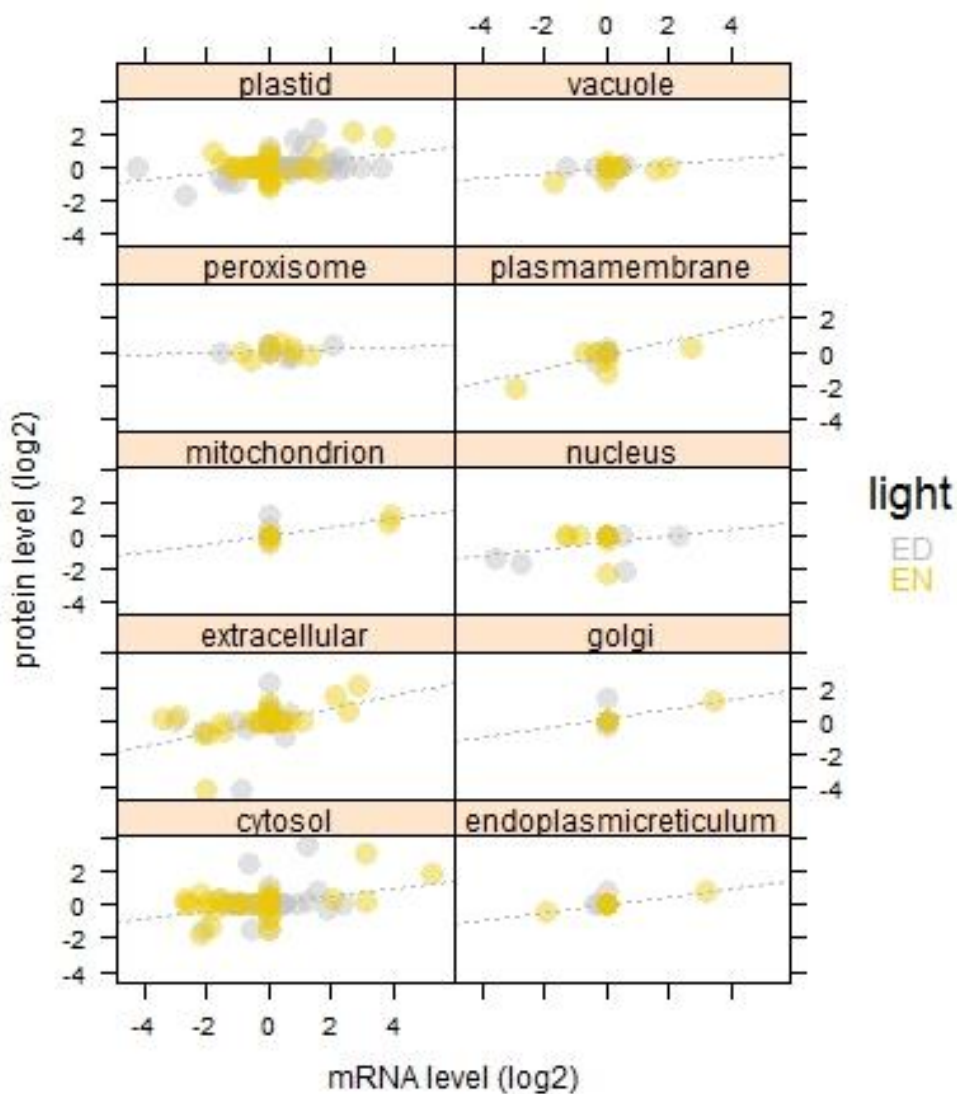


Figure S6. Linear regression analysis of concurrently changing transcripts and proteins across different subcellular locations ED and EN. Depicted are the transcripts (X-axis) and proteins (Y-axis) of the 51 genes in each circadian clock mutant ED and EN across several subcellular locations. Grey lines represent linear regressions. ED samples are shown in grey and EN in yellow.