

Supplemental FIGURES

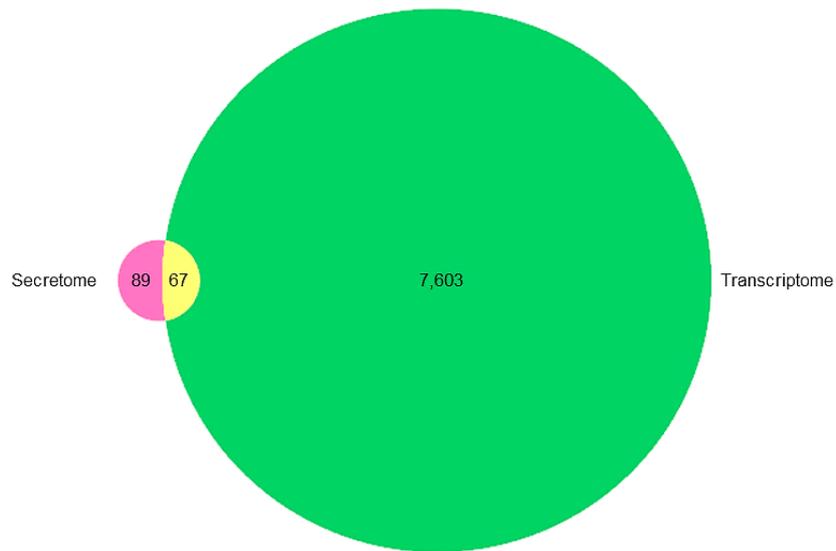


FIGURE S2: Venn diagram showing the number of uniquely regulated genes of the transcriptome-dataset (RPKM 10.0, green) versus the expressed proteins identified by the secretome-dataset (red) of *Scenedesmus quadricauda* (MZCH 10104).