# Online Supporting Material for Chapter 7

# Simultaneous RNA-seq to investigate microbe mediated protection against an algicidal bacterium in *Nannochloropsis oceanica* CCAP849/10 co-cultures

Frederik De Boever

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# Tables

**Table 7.1 –** Differential gene expression analysis for *Marinobacter* sp. FDB33 using DESEQ2 (Love et al., 2014), indicated are Log2foldchanges for all comparisons. For example, N33 (treatment) versus F33 (control). Time point is indicated by TP1 or TP2.

**Table 7.2 –** Differential gene expression analysis for *Alteromonas* sp. FDB36 using DESEQ2 (Love et al., 2014), indicated are Log2foldchanges for all comparisons. For example, N36 (treatment) versus F36 (control). Time point is indicated by TP1 or TP2.

**Table 7.3 –** Differential gene expression analysis for *Nannochloropsis oceanica* CCAP849/10 using DESEQ2 (Love et al., 2014), indicated are Log2foldchanges for all comparisons. For example, N33 (treatment) versus NAX (control). Time point is indicated by TP1 or TP2.

# References

Love, M. I., Huber, W., and Anders, S. (2014). Moderated estimation of fold change and dispersion for RNA-seq data with DESeq2. *Genome Biol.* doi:10.1186/s13059-014-0550-8.