Summary results DE analysis Miscanthus 2014 AU-IBERS

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Description of the data

- Counts table for the 2014 experiment at AU-IBERS (produced by IGATS)
- RNAseq data from Miscanthus mapped to Sorghum Bicolor
- \bullet Originally 33032 genes in the table \to after data cleaning 1 16343 genes in table
- 96 samples were taken from this green house experiment in two harvests (May 31, 2014 and June 15, 2014). The experiment ran from May 5, 2014 to June 18, 2014 (application of drought treatment as of May 12, 2014).
- Two treatments: control and drought (check %); five genotypes (WAT03 -sacchariflorus; WAT04 -sacchariflorus; WAT09 -giganteus; WAT10 sinensis; WAT11 -sinensis).
- The greenhouse design was not officially randomized, but it seems fairly "random". We have information on the row/column position and a block number.
- Replications: each genotype was harvest replicated 9-10 times per harvest which leads to a replication of about 4-5 samples per harvest/genotype/treatment combination.

¹Genes with no expression in any of the samples were removed. Additionally all genes with a mean expression for the non-zero counts of 10 or less were removed.

DE analysis using DESeq2

Controlling for Harvest

Before starting the statistical analysis using DESeq2 we ran some check on whether the greenhouse design or sampling scheme (through 2 harvests) might have an effect on the expression (analysis). To this end we used glm.nb including the design terms and checked for the significance of these factors. Based on the results we decided to include Harvest into the model in order to control for it. I also for block or for row/column, but due to difficulties with the fitting procedure and less definite results, we decided to keep Harvest. (UPDATE: after discussion on June 15, 2015 we identified that it might be a good idea to include section of the greenhouse into the model. This still needs to be done. See section below.)

- The differential expression analysis was done with the R packages DESeq2² to be downloaded from Bioconductor (on R3.2.0 using DESeq2_1.8.1).
- The (gene-wise) full model is stated as follows: counts \sim Harvest + Treatment + Genotype + Treatment \times Genotype
- Based on this model we proceeded to make a series of LRT (likelihood ratio tests) in order to test for the different terms in the model.
 - 1. LRT test for the interaction: full model vs reduced model (excluding the interaction)
 - 2. LRT test for a general effect of treatment: full model vs reduced model (only with Genotype and Harvest in it)
 - 3. LRT for main effect of the treatment: full model (with only the main effects and design effect) vs reduced model (only with Genotype and Harvest in it)
- Results for (1) LRT for interaction We find 61 (out of 16343) differentially expressed genes (with an adjusted p-value of p=0.0001). The smallest adjusted p-value that we found is in the magnitude of $3*10^{-17}$. Most DE genes are on chromosomes 1 and 2. (Do not pay attention to the first column of the table.)
- Results for (2) LRT for a general treatment effect This analysis was mainly done for a general check. We found 1311 differentially expressed

²Michael I Love, Wolfgang Huber and Simon Anders (2014): Moderated estimation of fold change and dispersion for RNA-Seq data with DESeq2. Genome Biology

genes (where it cannot be determined if the DE comes from the main effect or the interaction). Genes are not reported here.

Results for (3) LRT for treatment main effect We find 1219 differentially expressed genes. The results are exported to a table. 18 of those genes have been found as differentially expressed for interaction, therefore we exclude those from the list.

| 5 Sobic.001G313100.v2.1 0 39 Sobic.005G165800.v2.1 0 2 Sobic.001G142800.v2.1 0 | sted p-value .0000000000 .00000000000 .00000000000 |
|--------------------------------------------------------------------------------------|-------------------------------------------------------------|
| 39 Sobic.005G165800.v2.1 0 2 Sobic.001G142800.v2.1 0 | .0000000000 |
| 2 Sobic.001G142800.v2.1 0 | |
| | |
| 26 Sobic.003G203600.v2.1 0 | .0000000000 |
| | .0000000000 |
| | .0000000002 |
| | .0000000017 |
| | .0000000199 |
| | .0000000293 |
| | .0000000330 |
| | .0000002337 |
| | .0000003188 |
| | .0000000427 |
| | .0000015078 |
| | .0000015076 |
| | .0000028754 |
| | .0000028060 |
| | .0000032211 |
| | .0000037527 |
| | .0000045595 |
| | .000001000 |
| | .0000075508 |
| | .0000156941 |
| | .0000156941 |
| | .0000168083 |
| | .0000171320 |
| | .0000176362 |
| | .0000176362 |
| 1 Sobic.001G142200.v2.1 0 | .0000178267 |
| 54 Sobic.009G122000.v2.1 0 | .0000178267 |
| 32 Sobic.004G189900.v2.1 0 | .0000201983 |
| 56 Sobic.009G190500.v2.1 0 | .0000201983 |
| 40 Sobic.005G169200.v2.1 0 | .0000238157 |
| 41 Sobic.006G021900.v2.1 0 | .0000255084 |
| 37 Sobic.004G347500.v2.1 0 | .0000255119 |
| | .0000255119 |
| | .0000280657 |
| | .0000280657 |
| | .0000280657 |
| | .0000360883 |
| | .0000360883 |
| | .0000400168 |
| | .0000432178 |
| | .0000447936 |
| | .0000468262 |
| | .0000483192 |
| | .0000485755 |
| | .0000636227 |
| | .0000648755 |
| | .00000659518 |
| | .0000039318 .0000723126 |
| | .0000723126 |
| | .0000760341 |
| | .0000824074 |
| | .0000882305 |
| | .0000882305 |
| | .0000882305 |
| | .0000882305 |
| | .0000929545 |
| | .0000955330 |

Controlling for section*Harvest

In addition to including Harvest as a design factor into the mode we ran some additional checks whether the section (side) of the greenhouse may have an effect on differential expression. On the one side the compartment was bordered by a white painted glass wall and on the other side it had glass towards the neighbouring compartment. Therefore we included as well harvest and section as a main effect as well as the interaction between these two design factors.

- The differential expression analysis was done with the R packages DESeq2 to be downloaded from Bioconductor (on R3.2.0 using DESeq2_1.8.1).
- The (gene-wise) full model is stated as follows: counts ~ section*Harvest + Treatment + Genotype + Treatment × Genotype
- Based on this model we proceeded to make a series of LRT (likelihood ratio tests) in order to test for the different terms in the model.
 - 1. LRT test for the interaction: full model vs reduced model (excluding the interaction)
 - 3. LRT for main effect of the treatment: full model (with only the main effects and design effect) vs reduced model (only with Genotype and section*Harvest in it)
- Results for (1) LRT for interaction We find 64 (out of 16343) differentially expressed genes (with an adjusted p-value of p=0.0001). This list of 64 genes overlaps in 51 genes with the list of genes found when including only Harvest. There are 13 additional genes on the list displayed below (possible explanation: controlling for more design effects improves the analysis). 10 genes dropped off the list (possible explanation: sort of "false positive", artefacts of failing to include structural effects into the model).
- Results for (3) LRT for treatment main effect We find 1171 differentially expressed genes. The results are exported to a table. 20 of those genes have been found as differentially expressed for interaction, therefore we exclude those from the list.

| | genes(interaction) | adjusted p-value |
|----------|------------------------------------------------|-----------------------------|
| 5 | Sobic.001G313100.v2.1 | 0.0000000000 |
| 38 | Sobic.005G165800.v2.1 | 0.0000000000 |
| 2 | Sobic.001G142800.v2.1 | 0.0000000000 |
| 25 | Sobic.003G203600.v2.1 | 0.0000000000 |
| 34 | Sobic.004G223500.v2.1 | 0.0000000000 |
| 60 | Sobic.009G199600.v2.1 | 0.0000000008 |
| 33 | Sobic.004G223400.v2.1 | 0.0000000123 |
| 45 | Sobic.006G201700.v2.1 | 0.0000000123 |
| 4 | Sobic.001G218500.v2.1 | 0.0000000232 |
| 48 | Sobic.007G131000.v2.1 | 0.0000000202 |
| 43 | Sobic.006G146000.v2.1 | 0.0000001002 |
| 32 | Sobic.004G189900.v2.1 | 0.0000004124 |
| 10 | Sobic.004G183300.v2.1 Sobic.002G107800.v2.1 | 0.0000004304 |
| 27 | Sobic.002G107600.v2.1 Sobic.003G251700.v2.1 | 0.0000010540 |
| 51 | Sobic.003G231700.v2.1 Sobic.007G215600.v2.1 | 0.0000017639 |
| 13 | Sobic.007G215000.v2.1 Sobic.002G176500.v2.1 | 0.0000017039 |
| 26 | Sobic.002G170300.v2.1 Sobic.003G237600.v2.1 | 0.0000018215 |
| 20 44 | Sobic.005G257600.v2.1 Sobic.006G195800.v2.1 | |
| | | 0.0000018215 |
| 58 | Sobic.009G129300.v2.1 | 0.0000018215 0.0000024368 |
| 46 | Sobic.006G215400.v2.1 | |
| 47 | Sobic.006G260300.v2.1 | 0.0000041982 |
| 62 | Sobic.010G045400.v2.1 | 0.0000050603 |
| 41 | Sobic.006G055400.v2.1 | 0.0000069177 |
| 35 | Sobic.004G269100.v2.1 | 0.0000070020 |
| 54 | Sobic.009G062900.v2.1 | 0.0000075290 |
| 50 | Sobic.007G208700.v2.1 | 0.0000082966 |
| 14 | Sobic.002G176700.v2.1 | 0.0000085121 |
| 22 | Sobic.003G003800.v2.1 | 0.0000098331 |
| 1 | Sobic.001G142200.v2.1 | 0.0000129220 |
| 61 | Sobic.010G015800.v2.1 | 0.0000134534 |
| 18 | Sobic.002G387000.v2.1 | 0.0000165850 |
| 24 | Sobic.003G082000.v2.1 | 0.0000182743 |
| 16 | Sobic.002G237900.v2.1 | 0.0000184003 |
| 15 | Sobic.002G218100.v2.1 | 0.0000207249 |
| 29 | Sobic.003G302100.v2.1 | 0.0000257685 |
| 39 | Sobic.005G169200.v2.1 | 0.0000257685 |
| 55 | Sobic.009G081500.v2.1 | 0.0000268328 |
| 63 | Sobic.010G134000.v2.1 | 0.0000274942 |
| 19 | Sobic.002G393100.v2.1 | 0.0000278982 |
| 52 | Sobic.008G147900.v2.1 | 0.0000284957 |
| 40 | Sobic.006G021900.v2.1 | 0.0000363424 |
| 56 | Sobic.009G082200.v2.1 | 0.0000435715 |
| 42 | Sobic.006G132300.v2.1 | 0.0000470303 |
| 9 | Sobic.001G543500.v2.1 | 0.0000494851 |
| 57 | Sobic.009G122000.v2.1 | 0.0000496454 |
| 3 | Sobic.001G184300.v2.1 | 0.0000496883 |
| 37 | Sobic.005G152200.v2.1 | 0.0000498833 |
| 17 | Sobic.002G275700.v2.1 | 0.0000527150 |
| 59 | Sobic.009G190500.v2.1 | 0.0000538729 |
| 20 | Sobic.002G405400.v2.1 | 0.0000548754 |
| 31 | Sobic.003G371100.v2.1 | 0.0000548754 |
| 7 | Sobic.001G368600.v2.1 | 0.0000631353 |
| 8 | Sobic.001G391400.v2.1 | 0.0000631353 |
| 23 | Sobic.003G073700.v2.1 | 0.0000631353 |
| 30 | Sobic.003G319300.v2.1 | 0.0000631353 |
| 49 | Sobic.007G160600.v2.1 | 0.0000690122 |
| 64 | Sobic.010G178800.v2.1 | 0.0000707546 |
| 36 | Sobic.004G347500.v2.1 | 0.0000739505 |
| 21 | Sobic.002G416700.v2.1 | 0.0000749294 |
| 6 | Sobic.001G329200.v2.1 | 0.0000827363 |
| 12 | Sobic. $002G165400.v2.1$ | 0.0000848367 |
| 11 | Sobic.002G142300.v2.1 | 0.0000867574 |
| 53 | C 1: 000C000C00 0.1 | 0.00000000797 |
| 00 | Sobic.009G009600.v2.1 | 0.0000968735 |
| 28 | Sobic.009G009600.v2.1 Sobic.003G255800ev2.1 | 0.0000995604 |