

Supplementary information 1 – Almeida et al. Physiological and molecular characterization of yeast cultures pre-adapted for fermentation of lignocellulosic hydrolysate

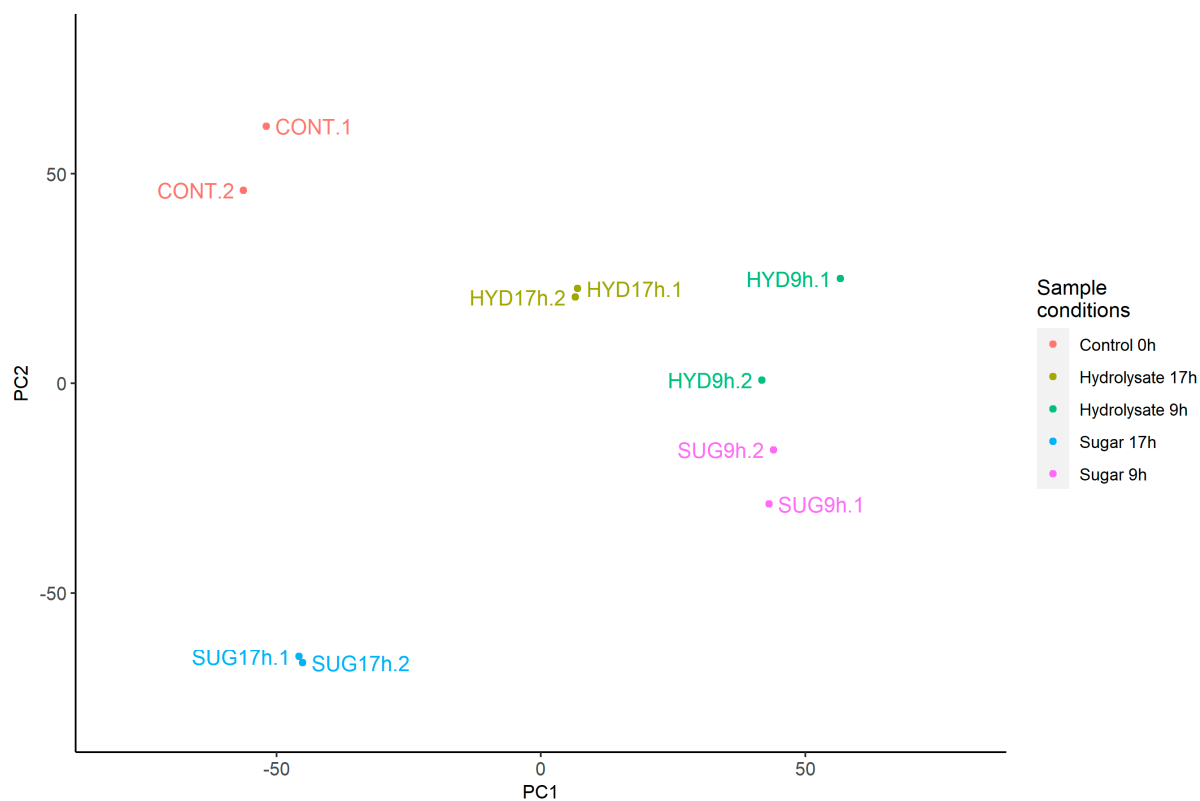


Figure S1. Principal component analysis of the *S. cerevisiae* probes from the 10 microarray samples, after RMA normalization. The biological replicates (.1 and .2) clustered together per sample, which suggest good reproducibility in the experimental workflow. Other trends include how the four hydrolysate samples, and the samples from time point 9h generally clustered together, respectively.

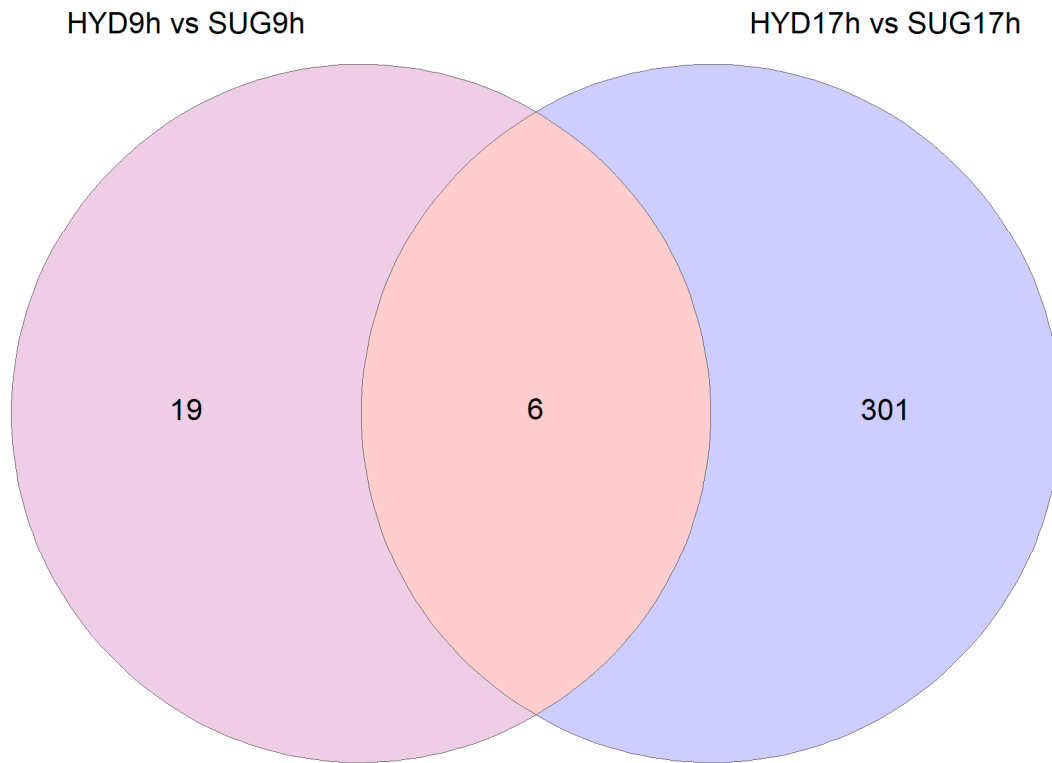


Figure S2. Venn diagram showing the intersection of the significantly differentially expressed genes identified in the 9h and 17h comparisons. Six genes were found to differentially expressed in both comparisons: *YHK8*, *OYE3*, *HBN1* were up-regulated in both conditions; *CTR1* and *RCK1* were down-regulated in both conditions. For more details on the gene names, see Table S1 below, and Supplemental Information 2 for the full processed microarray dataset.

Table S1. List of the genes that were found to be significantly up- or down-regulated in hydrolysate as compared to the sugar medium. Genes that were found at both times are highlighted in bold.

Sample	Up-regulated genes			Down-regulated genes		
9h	<i>GAL10</i> <i>GAL7</i> <i>GAL2</i> <i>GAL1</i> <i>AAD4</i> <i>YLR108C</i>	<i>FLR1</i> <i>CUP1-1</i> <i>PTR2</i> <i>YOR389W</i> <i>SNG1</i> <i>ARN1</i>	<i>YPL278c</i> <i>ATR</i> <i>YHK8</i> <i>OYE3</i> <i>HBN1</i> <i>YPR159C-A</i>	<i>CTR1</i> <i>RCK1</i> <i>AQY1</i>	<i>FET4</i> <i>HXT2</i>	<i>ZRT1</i> <i>SUL1</i>
17h	<i>YHK8</i> <i>YBR219C</i> <i>YPR159C-A</i> <i>MCH2</i> <i>YBL008W-A</i> <i>AIM33</i> <i>OYE3</i> <i>OYE2</i> <i>FSH1</i> <i>UPS2</i> <i>YBR298C-A</i> <i>HBN1</i> <i>AMF1</i> <i>YGR035C</i> <i>MAL12</i> <i>FET3</i> <i>CUZ1</i> <i>MIG2</i> <i>IMA1</i> <i>HAP4</i> <i>HAP3</i> <i>YNL217W</i> <i>PDR16</i> <i>YPL034W</i> <i>YGL230C</i> <i>POF1</i> <i>MRI1</i> <i>SDD2</i> <i>NUC1</i> <i>VPH2</i> <i>TDA2</i> <i>ASN1</i> <i>POP5</i> <i>ADH6</i> <i>YOL029C</i> <i>LSM5</i> <i>HR11</i> <i>YLR012C</i> <i>ATX2</i> <i>HMLALPHA1</i> <i>TTI2</i> <i>PRM10</i> <i>TIM9</i> <i>MIX14</i> <i>YPL088W</i> <i>NEL1</i>	<i>STL1</i> <i>KRR1</i> <i>KAR1</i> <i>KTR5</i> <i>CWC21</i> <i>APC11</i> <i>ERG4</i> <i>YBR096W</i> <i>YJR011C</i> <i>NRG2</i> <i>YCR087C-A</i> <i>FBP1</i> <i>NRK1</i> <i>YGK3</i> <i>CYC1</i> <i>HIS6</i> <i>TRR1</i> <i>ADO1</i> <i>EPT1</i> <i>DPB3</i> <i>ETT1</i> <i>YDR210W</i> <i>TRS33</i> <i>YKL033W-A</i> <i>ATP20</i> <i>CDA2</i> <i>MHF2</i> <i>DIB1</i> <i>PRO1</i> <i>CIR1</i> <i>CAF16</i> <i>YCR043C</i> <i>SMI1</i> <i>YDL085C-A</i> <i>CBS1</i> <i>AAT1</i> <i>TAD2</i> <i>RCE1</i> <i>MED4</i> <i>URM1</i> <i>MET31</i> <i>YDR222W</i> <i>TRM13</i> <i>SKG6</i> <i>PAN5</i> <i>YLR050C</i>	<i>RPC25</i> <i>ATF2</i> <i>POP8</i> <i>ERG3</i> <i>GCG1</i> <i>YJL107C</i> <i>SMC2</i> <i>CPT1</i> <i>MKC7</i> <i>YPL199C</i> <i>DTD1</i> <i>YER158C</i> <i>ERG26</i> <i>HIS2</i> <i>DUT1</i> <i>ADE17</i> <i>MIS1</i> <i>EMC6</i> <i>MPC2</i> <i>YDL157C</i> <i>SAM4</i> <i>LDB18</i> <i>TPM1</i> <i>YIL027C</i> <i>RPS18B</i> <i>APT1</i> <i>ERG27</i> <i>GUA1</i> <i>JLP2</i> <i>NDE1</i> <i>DAL3</i> <i>MCM5</i> <i>ALD6</i> <i>HPT1</i> <i>ACS2</i> <i>YEF3</i> <i>YSY6</i> <i>HXK2</i> <i>LEU9</i> <i>LCL2</i> <i>CLA4</i> <i>JEM1</i> <i>TKL1</i> <i>INO1</i> <i>YBL081W</i> <i>RPS17B</i>	<i>YNR071C</i> <i>MAN2</i> <i>PDR18</i> <i>MUP1</i> <i>TRA1</i> <i>CIN5</i> <i>YOR019W</i> <i>NUM1</i> <i>YLR352W</i> <i>CMR3</i> <i>YOR032W-A</i> <i>PSA1</i> <i>NCW2</i> <i>YCR101C</i> <i>ASK10</i> <i>SCW11</i> <i>DIA3</i> <i>SWH1</i> <i>OPT1</i> <i>FRK1</i> <i>APC2</i> <i>YNR064C</i> <i>YOR248W</i> <i>SPO20</i> <i>YMR122W-A</i> <i>ECM18</i> <i>ACE2</i> <i>CTS1</i> <i>HOS3</i> <i>DSE3</i> <i>SSD1</i>	<i>TMA108</i> <i>HXK1</i> <i>YOR316C-A</i> <i>IST2</i> <i>YCR100C</i> <i>YNL058C</i> <i>SPS22</i> <i>YBR225W</i> <i>PKH3</i> <i>PHO89</i> <i>ERV2</i> <i>PAU7</i> <i>RAS1</i> <i>CIS3</i> <i>MET30</i> <i>MET1</i> <i>BNA3</i> <i>FLO11</i> <i>NBL1</i> <i>REC107</i> <i>RGS2</i> <i>GEM1</i> <i>MDS3</i> <i>USV1</i> <i>YGR130C</i> <i>GTT3</i> <i>PCA1</i> <i>SIM1</i> <i>PIR1</i> <i>DFG16</i>	<i>FRE4</i> <i>AMA1</i> <i>SIP5</i> <i>BAG7</i> <i>TOS1</i> <i>PLB2</i> <i>YMR114C</i> <i>YGR066C</i> <i>NGR1</i> <i>ASF2</i> <i>TDA1</i> <i>PEX9</i> <i>HHO1</i> <i>GSY1</i> <i>GNP1</i> <i>ECM10</i> <i>JLP1</i> <i>AGP3</i> <i>BOP2</i> <i>ALT2</i> <i>FMP48</i> <i>DBP1</i> <i>CRR1</i> <i>RNR3</i> <i>CWP1</i> <i>SHC1</i> <i>RCK1</i> <i>CTR1</i>

<i>SAP4</i>	<i>RIE1</i>	<i>RPL15A</i>			
<i>PCK1</i>	<i>SMC3</i>	<i>MCM16</i>			
<i>EFM6</i>	<i>COS9</i>	<i>ARG1</i>			
<i>HTL1</i>	<i>HAT1</i>	<i>HIS1</i>			
<i>NMT1</i>	<i>YBL028C</i>	<i>DAS2</i>			
<i>PHM6</i>	<i>APS2</i>	<i>FCY2</i>			
<i>RDH54</i>	<i>YRB1</i>	<i>RK11</i>			
<i>MEP3</i>	<i>UTP11</i>	<i>SPH1</i>			
<i>IST3</i>	<i>MRS3</i>	<i>RPS26B</i>			
<i>MMT1</i>	<i>BUD7</i>	<i>MCD4</i>			
<i>RPC17</i>	<i>CNM67</i>	<i>CDC45</i>			
<i>FCF2</i>	<i>YMR321C</i>	<i>MNN1</i>			
<i>SDO1</i>	<i>IMP2</i>	<i>FRS1</i>			
<i>REX4</i>	<i>NVJ3</i>	<i>PPT2</i>			
<i>BIT61</i>	<i>PDR12</i>	<i>YMC2</i>			
<i>ACH1</i>	<i>ATP17</i>	<i>RPS22A</i>			
<i>YHI9</i>	<i>MIF2</i>	<i>SEN34</i>			
<i>COS3</i>	<i>LTP1</i>	<i>YHM2</i>			
<i>YCL021W-A</i>	<i>DCD1</i>	<i>PAL1</i>			
<i>GIM3</i>	<i>YCR016W</i>	<i>RPL39</i>			
<i>YIL024C</i>	<i>DUO1</i>	<i>RPS29A</i>			
<i>LSM12</i>	<i>TOH1</i>	<i>RPL41A</i>			
<i>TYR1</i>	<i>PFD1</i>	<i>ECO1</i>			
<i>FLC1</i>	<i>RAD33</i>	<i>HAM1</i>			
<i>YML131W</i>	<i>TRM11</i>	<i>YLR053C</i>			
<i>FTR1</i>	<i>RPC11</i>	<i>BSC2</i>			
<i>AGX1</i>	<i>KTR6</i>				