

Research article

Relative contribution of three transporters to D-xylose uptake in *Aspergillus niger*

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Supplementary

Table S1. *Aspergillus niger* strains used in this study.

Strain	CBS number	Genotype	Reference
N593 Δ ku70	CBS 138852	<i>cspA1</i> , <i>kusA</i> :: <i>amdS</i> , <i>pyrG</i> ⁻	Alazi et al, 2016
Δ xltA	CBS 147732	<i>cspA1</i> , <i>kusA</i> :: <i>amdS</i> , <i>pyrG</i> ⁻ , <i>xltA</i> ⁻	This study
Δ xltB	CBS 147733	<i>cspA1</i> , <i>kusA</i> :: <i>amdS</i> , <i>pyrG</i> ⁻ , <i>xltB</i> ⁻	This study
Δ xltD	CBS 147734	<i>cspA1</i> , <i>kusA</i> :: <i>amdS</i> , <i>pyrG</i> ⁻ , <i>xltD</i> ⁻	This study
Δ xltA Δ xltB Δ xltD	CBS 147741	<i>cspA1</i> , <i>kusA</i> :: <i>amdS</i> , <i>pyrG</i> ⁻ , <i>xltA</i> ⁻ , <i>xltB</i> ⁻ , <i>xltD</i> ⁻	This study

Reference: Alazi, E., et al., 2016. The transcriptional activator GaaR of *Aspergillus niger* is required for release and utilization of D-galacturonic acid from pectin. FEBS Lett. 590, 1804-1815.

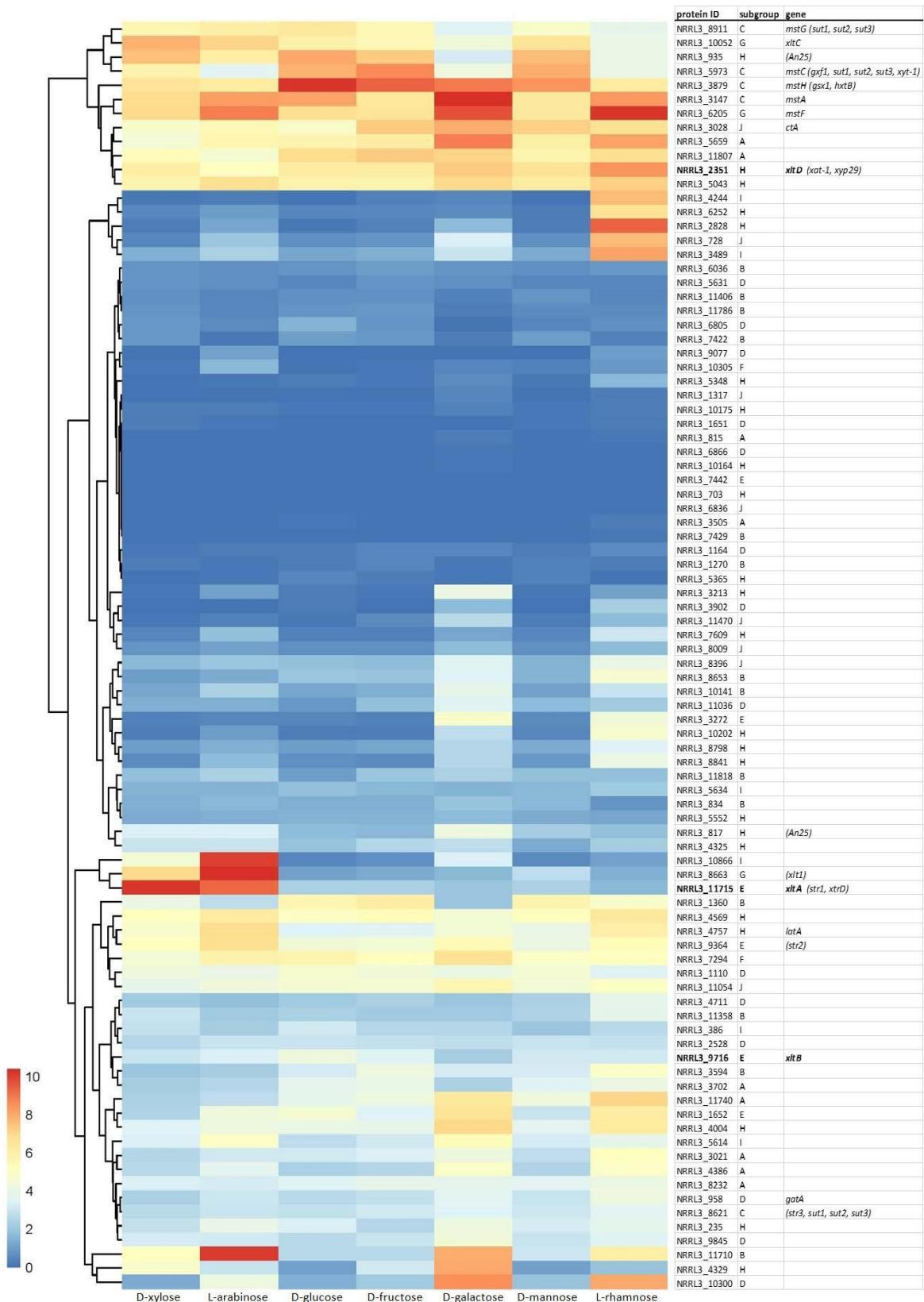


Figure S1. Expression profile of *A. niger* candidate sugar transporters. RNAseq analysis was performed on triplicate cultures of *A. niger* on 25 mM of the indicated sugars. *A. niger* gene names are indicated in the right column, while gene names of orthologs in other fungi are in brackets. Transporters analyzed in this study are in bold. The grouping in the middle column is based on [18] and reflects putative functions. The color bar indicates the average log₂ FPKM values. A = inositol/hexose, B = maltose/sucrose, C = hexose/pentose/polyol, D = uronic/quinic acid, E = xylose, F = unknown, G = pentose/hexose, H = glycerol/arabitol/pentose, I = unknown, J = lactose/cellobextrin/xylobiose.



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