

Quantitative comparison of the biomass-degrading enzyme repertoires of five filamentous fungi

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Fig S1 - Aspergillus terreus

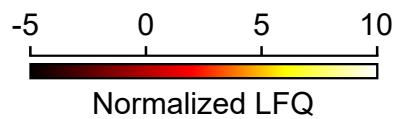


Fig S2 - *Trichoderma reesei*

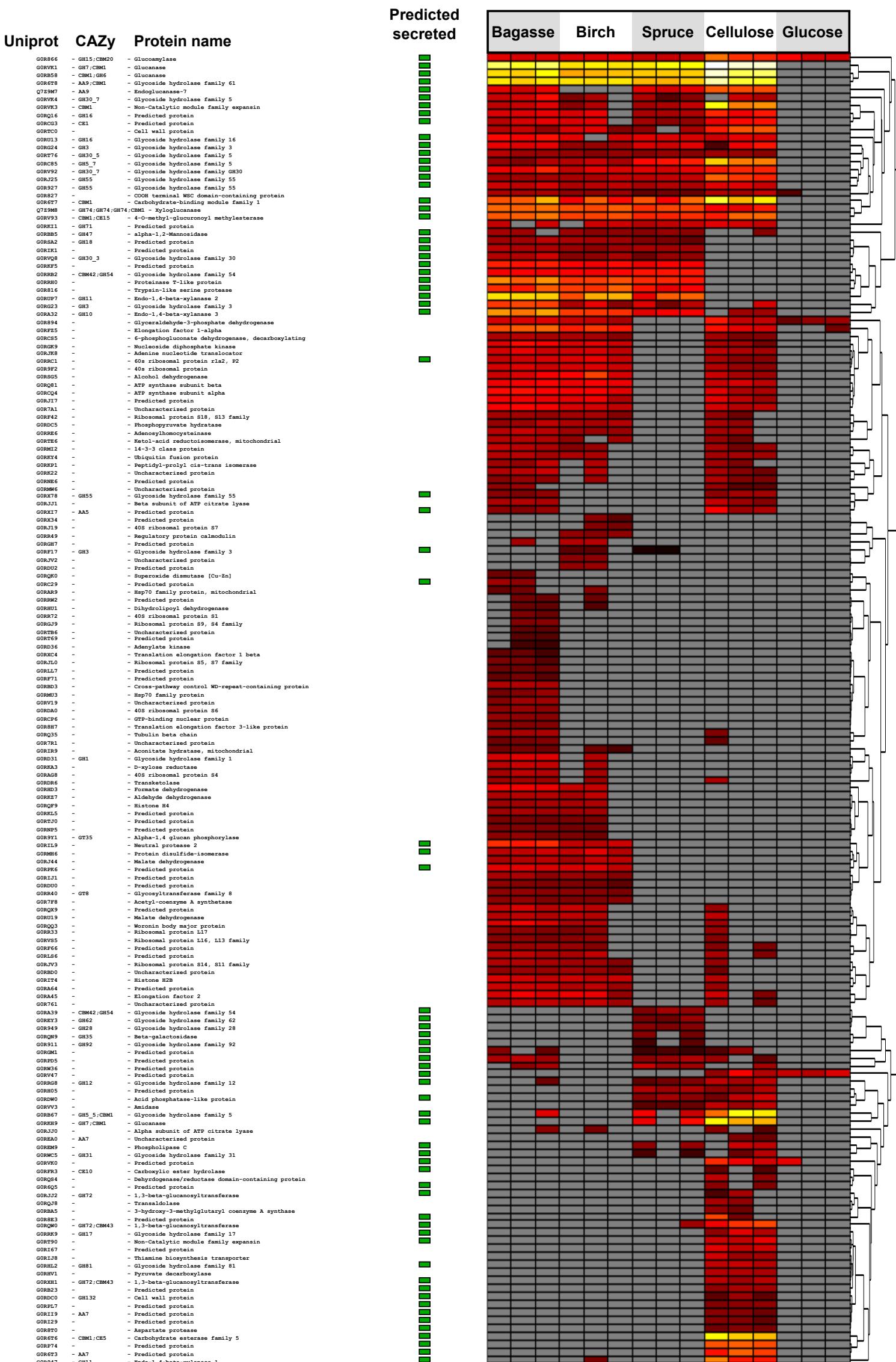
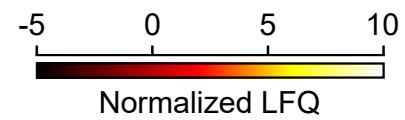


Fig S3 - Myceliophthora thermophila



Normalized LFC

The figure displays a heatmap of protein expression patterns across five samples: Bagasse, Birch, Spruce, Cellulose, and Glucose. The rows represent individual proteins, and the columns represent the samples. A color scale on the left indicates expression levels, with dark red representing high expression and light yellow representing low expression.

UniProt	CAZy	Protein name	secreted	Bagasse	Birch	Spruce	Cellulose	Glucose
G2QJ91	-	- CBM1;GH10	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJC3	-	- AA9;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q913	-	- CBM1;GH11	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QNNH	-	- GH7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAP3	-	- CBM1;GH6	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q665	-	- CBM1;GH7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G74240	-	- AA3_1;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q5D8	-	- GH5_5;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QLE4	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QKQ1	-	- GH12	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QE4	-	- Catalase	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QNTU	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QHRT	-	- GH74;GH74:GH74	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q998	-	- GH6	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q184	-	- CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q9F7	-	- AA9;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCS4	-	- CBM1;GH7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QNK3	-	- GH131;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QQ09	-	- GH43_36	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QFL3	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAH7	-	- GH26;CBM35	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ27	-	- CE16	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QH79	-	- PL1_7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q3G9	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD29	-	- CE1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ35	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAM0	-	- CBM1;AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q9T3	-	- CBM1;AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QIR2	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD41	-	- GH18	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAZ9	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QIV4	-	- GH62;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q4S5	-	- GH11	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ94	-	- CE5;CBM1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGJ5	-	- GH125	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGP4	-	- AA3	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCM6	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ7D	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QDD9	-	- GH43_24;CBM35	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QFG7	-	- PL4_1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QZJ2	-	- AA3_2	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q3A2	-	- GH53	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGN6	-	- GH10	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QLD1	-	- GH16	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QDN2	-	- GH3	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAI1	-	- GH30_5	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q7A5	-	- AA9	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCR1	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QM7	-	- GH3	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QH4	-	- GH47	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q494	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPS0	-	- CBM20;GH15	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q3S5	-	- AA5_1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q925	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q714	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAD1	-	- GH72;CBM43	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QKY8	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGL4	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD13	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q2P9	-	- GH5_7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2C0X0	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QFL3	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
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G2QPA19	-	- AA3_1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
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G2QIM4	-	- GH55	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPH1	-	- GH43_26	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCT4	-	- PL3_2	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPV9	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ72	-	- GH79	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPF5	-	- CB10	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QH09	-	- GH43_21	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD15	-	- GH95	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q506	-	- GH93	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCC8	-	- GH43_36	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QLDN	-	- CE8	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QIK5	-	- PL1_7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJQ6	-	- GH62	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QBV8	-	- CE1	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJ92	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD7D	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q9T6	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q221	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QQA3	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJU3	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QRR8	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q8B8	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q8R8	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q8B18	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAY2	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QBN2	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QCN5	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QG550	-	- PL26	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPK7	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QB0U9	-	- CE10	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q2N20	-	- CE3	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
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G2QN15	-	- GH109	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QZU8	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2PZQ3	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q9D2	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QBV2	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
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G2Q3L4	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q624	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q974	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QMB3	-	- CE12	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJRS	-	- CE15	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGS8	-	- GH2	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QG16	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QAP6	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QHT5	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QJN4	-	- GH30_7	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QPT1	-	- AA2;AA2	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QZP7	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QGF9	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q610	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q7Q6	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2QD76	-	-	[green]	[yellow]	[yellow]	[red]	[yellow]	[yellow]
G2Q9T6	-	-	[green]	[yellow]	[yellow]			

Fig S4 - *Neurospora crassa*

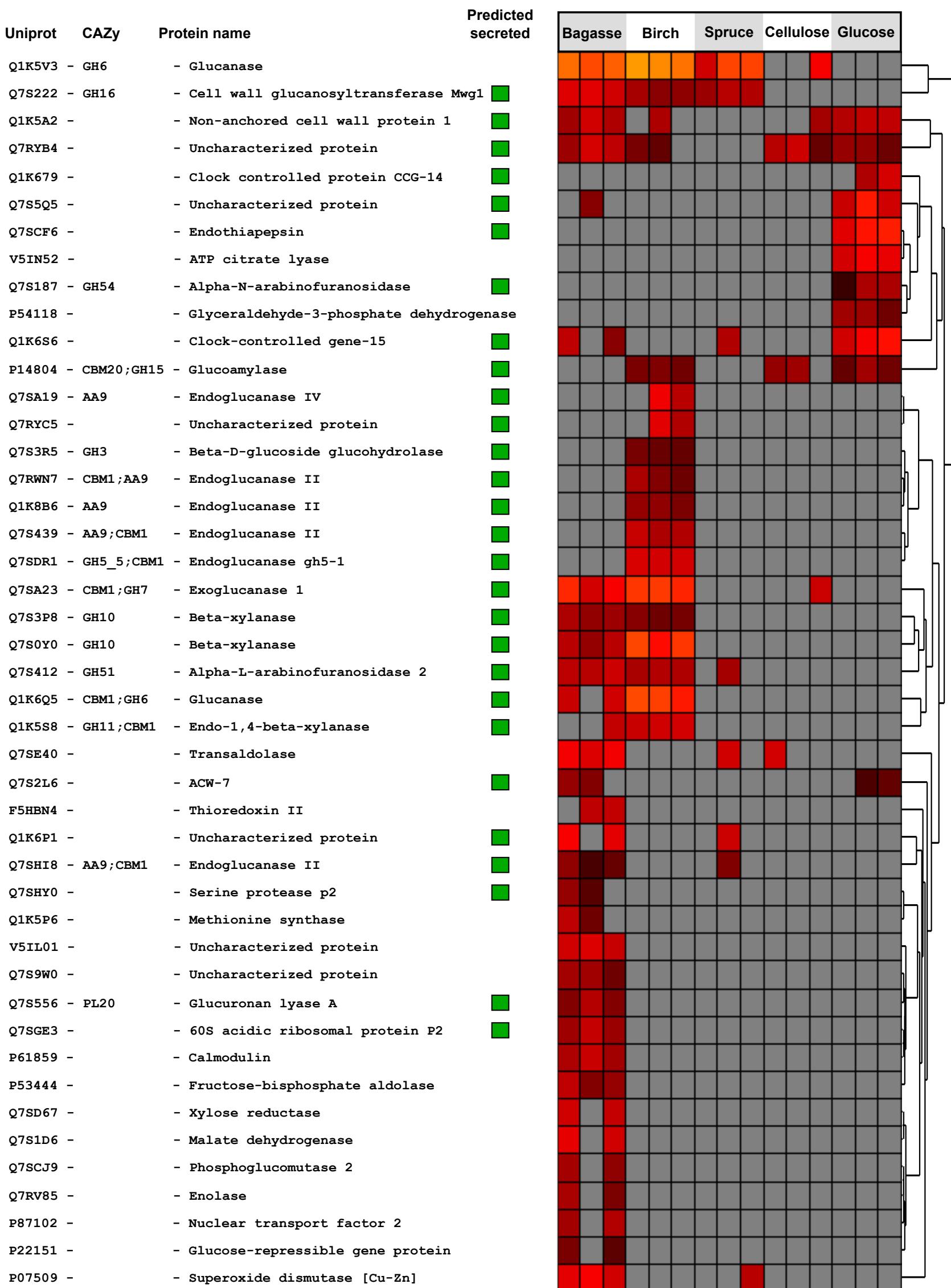
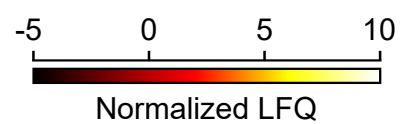


Fig S5 - Phanerochaete chrysosporium

-5 0 5 10

Normalized LFQ

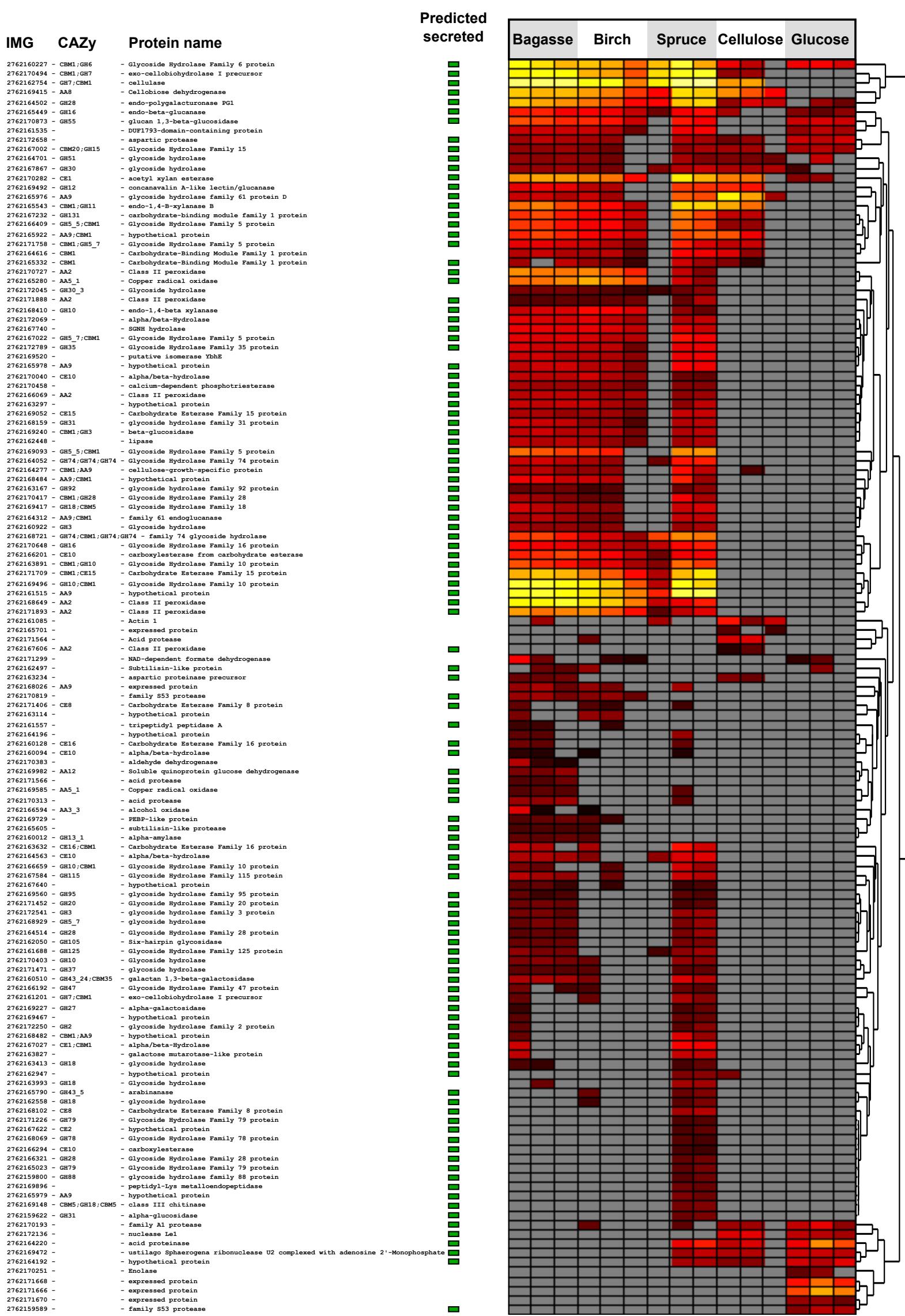
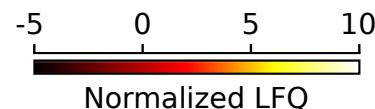


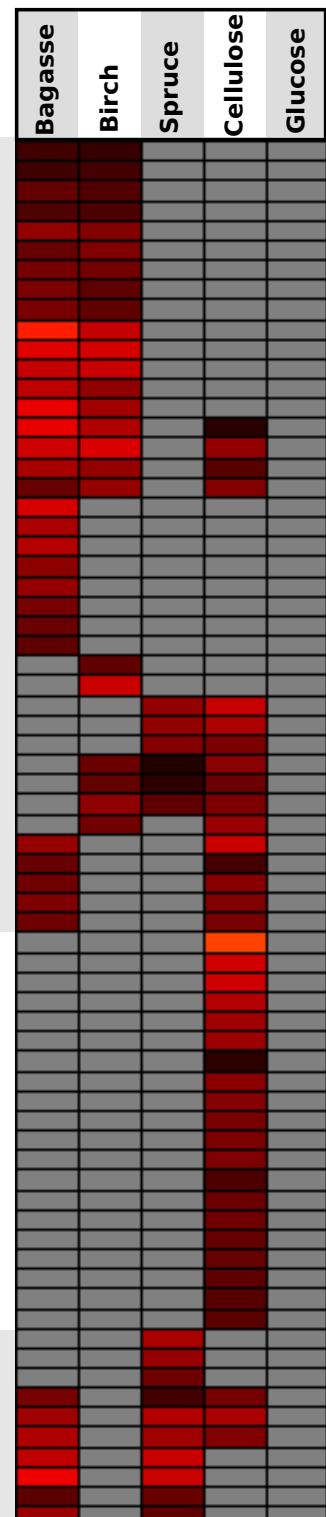
Fig S6



Acc. number **Protein name**

Organism

2762161557	tripeptidyl peptidase A
G2Q8R8	Pectin lyase-like protein
2762169729	PEBP-like protein
2762165605	subtilisin-like protease
2762170819	family S53 protease
G2Q8I8	Uncharacterized protein
G2Q372	Uncharacterized protein
G2QLD3	Tyrosinase-like protein
G2PZJ6	FAD dependent oxidoreductase-like protein
G0RLI9	Neutral protease 2
G0RPK6	Predicted protein
G0RMH6	Protein disulfide-isomerase
G2QMS4	Secretory phospholipase A2
G2QGW5	Uncharacterized protein
G2Q057	Uncharacterized protein
G0RRC1	60s ribosomal protein rla2, P2
G2Q792	Peptide hydrolase
Q0CAZ7	Peptide hydrolase
G2QJM5	Uncharacterized protein
Q7SGE3	60S acidic ribosomal protein P2
G2QBY3	Intradiol ring-cleavage dioxygenase-like protein
G2QNU8	Alpha-galactosidase
G0RC29	Predicted protein
Q7SHY0	Serine protease p2
G2Q650	Uncharacterized protein
2762171566	acid protease
G2Q9T6	Fumarate reductase/succinate dehydrogenase flavo protein-like protein
Q7RYC5	Uncharacterized protein
G0REM9	Phospholipase C
G0RDW0	Acid phosphatase-like protein
2762162947	hypothetical protein
G2Q493	Uncharacterized protein
G2QMP1	Uncharacterized protein
Q0CLA1	Uncharacterized protein
G2QMW2	Uncharacterized protein
Q0CXA9	Uncharacterized protein
G2QP4	Uncharacterized protein
G2QLJ3	Carbohydrate-binding WSC domain protein
G2QCK0	Phospholipase C-like protein
2762163234	aspartic proteinase precursor
G0RP74	Predicted protein
G0RT90	Non-Catalytic module family expansin
G0R8E3	Predicted protein
Q0CVY4	Uncharacterized protein
G0RB23	Predicted protein
Q0CH63	Uncharacterized protein
G2QGG5	Uncharacterized protein
G0RPL7	Predicted protein
G0R6Q5	Predicted protein
G0R129	Predicted protein
G0R8T0	Aspartate protease
Q0CE26	Uncharacterized protein
G2QBZ2	Uncharacterized protein
G2QPH7	Uncharacterized protein
G2Q9P8	Peptidase-like protein
G2QA29	Uncharacterized protein
Q0CIE1	Superoxide dismutase [Cu-Zn]
Q0CR38	Uncharacterized protein
Q0CAE1	Carboxypeptidase
Q0CBH2	Uncharacterized protein
G2QNT3	Peptide hydrolase
G2QPG6	Peptide hydrolase
2762169896	peptidyl-Lys metalloendopeptidase
G0RGM1	Predicted protein
G0RW36	Predicted protein
G0RD5	Predicted protein
2762163827	galactose mutarotase-like protein
Q1K6P1	Uncharacterized protein
2762169467	hypothetical protein
2762170313	acid protease



█ ***A. terreus***

█ ***M. thermophila***

█ ***P. chrysosporium***

█ ***T. reesei***

█ ***N. crassa***