**S4 Table.** Species-specific CAZymes of *M. coronaria*, *M. brunnea* and *M. rosae*.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| CAZyme familya | Gene locus/ name | Potential substrateb | Enzyme activityb | Taxonomy of the top10 best hitsc |
| *M.coronaria*b |
| GH3 | B2J93\_4547 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 9, D 1 |
| GH16 | B2J93\_7882 | Hemicellulose | Xyloglucanase | L 9, D 1 |
| GH16 | B2J93\_9182 | Hemicellulose | Xyloglucanase | L 2, D 7, S 1 |
| GH18 | B2J93\_4402 | N-linked oligosaccharides | endo-β-N-acetylglucosaminidase | L 4, D 3, X 1, E 1, C 1 |
| GH89 | B2J93\_2177 | N-linked oligosaccharides | α-N-acetylglucosaminidase | L 9, D 1 |
| GH92 | B2J93\_5902 | Oligosaccharides  | α-1,2-mannosidase | L 8, D 1, E 1 |
| PL1 | B2J93\_4840 | pectin | pectate lyaseexo-pectate lyase | L 5, D 2, E 2, S 1 |
| PL3 | B2J93\_5418 | Pectin | Pectate lyase | L 1, D 7, S 2 |
| PL26 | B2J93\_9264 | pectin | rhamnogalacturonan exolyase | L 3, S 3, E 3, D 1 |
| CE2 | B2J93\_2098 | hemicellulose | acetyl xylan esterase | L 3, D 7 |
| CE4 | B2J93\_6888 | HemicelluloseN-linked oligosaccharides | acetyl xylan esterasepeptidoglycan GlcNAc deacetylase | L 7, D 3 |
| AA1 | B2J93\_5665 | lignin | Multicopper oxidases | L 5, S 4, D 1 |
| AA2 | B2J93\_2861 | Lignin | Lignin peroxidase | L 2, D 7, S 1 |
| AA3 | B2J93\_7165 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 6, E 3, D 1 |
| AA3 | B2J93\_1229 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 4, D 6 |
| AA3 | B2J93\_3628 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | D 10 |
| AA7 | B2J93\_6765 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 2, D 3, S 5 |
| AA7 | B2J93\_6390 | See above | See above | L 3, S 6, D 1 |
| AA12 | B2J93\_4172 | cellulose | pyrroloquinoline quinone-dependent oxidoreductase | L 9, D 1 |
| GT1 | B2J93\_5387 | NA | NA | L 10 |
| GT2 | B2J93\_7586 | NA | NA | L 6, D 4 |
| GT31 | B2J93\_7585 | NA | NA | L 6, D 4 |
| GT69 | B2J93\_6811 | NA | NA | L 4, D 6 |
| GT71 | B2J93\_8885 | NA | NA | L 10 |
| *M．brunnea*b |
| GH1 | MBM\_02089 | cellulosehemicellulose pectin | β-glucosidaseexo-β-1,4-glucanaseβ-galactosidase | L 9, E 1 |
| GH3 | MBM\_00209 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 4, D 5, E 1 |
| GH3 | MBM\_09941 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 10 |
| GH3 | MBM\_05263 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 10 |
| GH3 | MBM\_06277 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 3, A 1, S 3, E 2, D 1 |
| GH3 | MBM\_03435 | CelluloseHemicellulosePectin  | β-glucosidase β-xylosidase exo-β-1,4-glucanase  | L 10 |
| GH5 | MBM\_02055 | CelluloseHemicelluloseβ-1,3-glucans | endo-β-1,4-glucanaseendo-β-1,4-xylanaseglucan β-1,3-glucosidase | L 7, S 3 |
| GH6 | MBM\_04333 | cellulose | endo-β-1,4-glucanasecellobiohydrolase | L 8, D 2 |
| GH10 | MBM\_03633 | hemicellulose | endo-β-1,4-β-xylanase | L 4, E 3, S 3 |
| GH15 | MBM\_08967 | Polysaccharides | Glucoamylase | L 10 |
| GH16 | MBM\_08515 | Hemicellulose | Xyloglucanase | L 10 |
| GH16 | MBM\_00319 | Hemicellulose | Xyloglucanase | L 10 |
| GH28 | MBM\_02037 | Pectin | Polygalacturonase | L 1, D 8, E 1 |
| GH28 | MBM\_03901 | Pectin | Polygalacturonase | L 10 |
| GH30 | MBM\_04774 | cellulosehemicellulosepectin | β-glucosidaseendo-β-1,4-xylanaseendo-β-1,6-galactanase | L 6, D 3, S 1 |
| GH30 | MBM\_08263 | cellulosehemicellulosepectin | β-glucosidaseendo-β-1,4-xylanaseendo-β-1,6-galactanase | L 4, S 5, D 1 |
| GH31 | MBM\_03122 | Hemicellulose | α-xylosidase | P 1, D 6, E 2, S 1 |
| GH35 | MBM\_08774 | HemicellulosePectin | β-galactosidaseexo-β-1,4-galactanase | L 10 |
| GH35 | MBM\_05705 | HemicellulosePectin | β-galactosidaseexo-β-1,4-galactanase | L 9, D 1 |
| GH39 | MBM\_08295 | hemicellulose | β-xylosidase | L 9, E 1 |
| GH43 | MBM\_04126 | Hemicellulose Pectin | β-xylosidaseα-L-arabinofuranosidase | L 1, S 9 |
| GH43 | MBM\_01191 | Hemicellulose Pectin | β-xylosidaseα-L-arabinofuranosidase | L 7, E 1, S 1, D 1 |
| GH43 | MBM\_05474 | Hemicellulose Pectin | β-xylosidaseα-L-arabinofuranosidase | L 4, D 2, E 2, S 2 |
| GH43 | MBM\_00969 | Hemicellulose Pectin | β-xylosidaseα-L-arabinofuranosidase | L 8, E 1, D 1 |
| GH43 | MBM\_01508 | Hemicellulose Pectin | β-xylosidaseα-L-arabinofuranosidase | L 6, E 4 |
| GH55 | MBM\_03699 | Polysaccharides | endo-1,3-β-glucosidase | L 5, E 5 |
| GH67 | MBM\_06928 | Hemicelluloses | α-glucuronidase | L 5, D 4, S 1 |
| GH74 | MBM\_05691 | CelluloseHemicellulose | endo-β-1,4-glucanaseXyloglucanase | L 5, T 1, E 2, S 1, Bacteria 1 |
| GH81 | MBM\_01851 | Polysaccharides | endo-1,3-β-glucosidase | L 10 |
| GH105 | MBM\_04106 | Pectin | rhamnogalacturonyl hydrolase | L 1, S 4, D 5 |
| GH135 | MBM\_09138 | exopolysaccharide galactosaminogalactan | α-1,4-galactosaminogalactan hydrolase | L 9, S 1 |
| CE1 | MBM\_05316 | Hemicellulose | Acetyl xylan esteraseFeruloyl esterase | L 10 |
| CE3 | MBM\_01585 | hemicellulose | acetyl xylan esterase | L 5, S 2, D 1, bacteria 2 |
| CE5 | MBM\_09546 | HemicelluloseCutin | Acetyl xylan esteraseCutinase | L 9, E 1 |
| CE5 | MBM\_02401 | HemicelluloseCutin | Acetyl xylan esteraseCutinase | L 8, D 1, B 1 |
| CE5 | MBM\_03163 | HemicelluloseCutin | Acetyl xylan esteraseCutinase | L 4, S 3, O 2, P 1 |
| CE8 | MBM\_01176 | pectin | pectin methylesterase | L 10 |
| CE8 | MBM\_00548 | pectin | pectin methylesterase | L 8, D 2 |
| CE10 | MBM\_03986 | NA | NA | L 9, E 1 |
| CE10 | MBM\_02486 | NA | NA | L 2, D 4, E 4 |
| CE10 | MBM\_06621 | NA | NA | L 10 |
| CE10 | MBM\_08671 | NA | NA | L 2, E 4, S 3, D 1 |
| CE10 | MBM\_03188 | NA | NA | L 10 |
| CE12 | MBM\_05265 | Pectin | Pectin acetylesterase | L 2,D 8 |
| AA1 | MBM\_02253 | lignin | multicopper oxidases | L 9, S 1 |
| AA3 | MBM\_09150 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 7, D 3 |
| AA3 | MBM\_04141 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 7, D 2, E 1 |
| AA3 | MBM\_07353 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 4, E 5, D 1 |
| AA3 | MBM\_08750 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | D 10 |
| AA4 | MBM\_06330 | Lignin | vanillyl-alcohol oxidase | L 5, E 4, D 1 |
| AA7 | MBM\_02730 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 1, S 8, E 1 |
| AA7 | MBM\_04037 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | S 5, Pis 1, B 1, D 1, E 1 |
| AA7 | MBM\_07678 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 1, S 3, D 3, E 2, Pis 1 |
| AA7 | MBM\_04264 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 1, S 8, D 1 |
| AA7 | MBM\_03338 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 2, E 5, D 2, S 1 |
| AA7 | MBM\_04587 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 6, E 4 |
| AA7 | MBM\_00406 | Cellobiosechitin/glycoproteins | glucooligosaccharide oxidasechitooligosaccharide oxidase | L 7, E 2, C 1 |
| AA9 | MBM\_00975 | cellulose | copper-dependent monooxygenase | L 10 |
| GT17 | MBM\_00251 | NA | NA | L 10 |
| *D. rosea*b |
| GH53 | PBP24709PBP28278 | pectin | endo-β-1,4-galactanase | L 10 |
| PL4 | PBP19949PBP28439 | pectin | rhamnogalacturonan lyase | L 7, D 2, E 1 |
| CE16 | PBP27316PBP26860 | Polysaccharides | Acetylesterase | L 10 |
| AA2 | PBP21410 | See above | See above | L 10 |
| AA3 | PBP21841 | Celluloselignin | glucose 1-oxidasearyl alcohol oxidase | L 1, S 2, D 5, E 2 |
| CBM48 | PBP22087 | NA | NA | L 10 |
| CBM48 | PBP22865 | NA | NA | L 1/10, D 9/10 |

a, The CAZyme family was annotated by dbcan2 server

b, The potential substrate and enzyme activity were annotated based on two references[1, 2]

c, E, Eurotiomycetes, S, Sordariomycetes, L, Leotiomycetes, D, Dothideomycetes, X, Xylonomycetes, Pis, Pezizomycotina incertae sedis, C, Lecanoromycetes, B, Basidiomycota; P, Pezizomycetes, O, Orbiliomycetes, A, Saccharomycetes, T, Tremellomycetes

Gray filling indicate atypically distributed genes.

1. Blackman LM, Cullerne DP, Hardham AR: **Bioinformatic characterisation of genes encoding cell wall degrading enzymes in the Phytophthora parasitica genome**. *Bmc Genomics* 2014, **15**.

2. Chang HX, Yendrek CR, Caetano-Anolles G, Hartman GL: **Genomic characterization of plant cell wall degrading enzymes and in silico analysis of xylanses and polygalacturonases of Fusarium virguliforme**. *Bmc Microbiol* 2016, **16**.