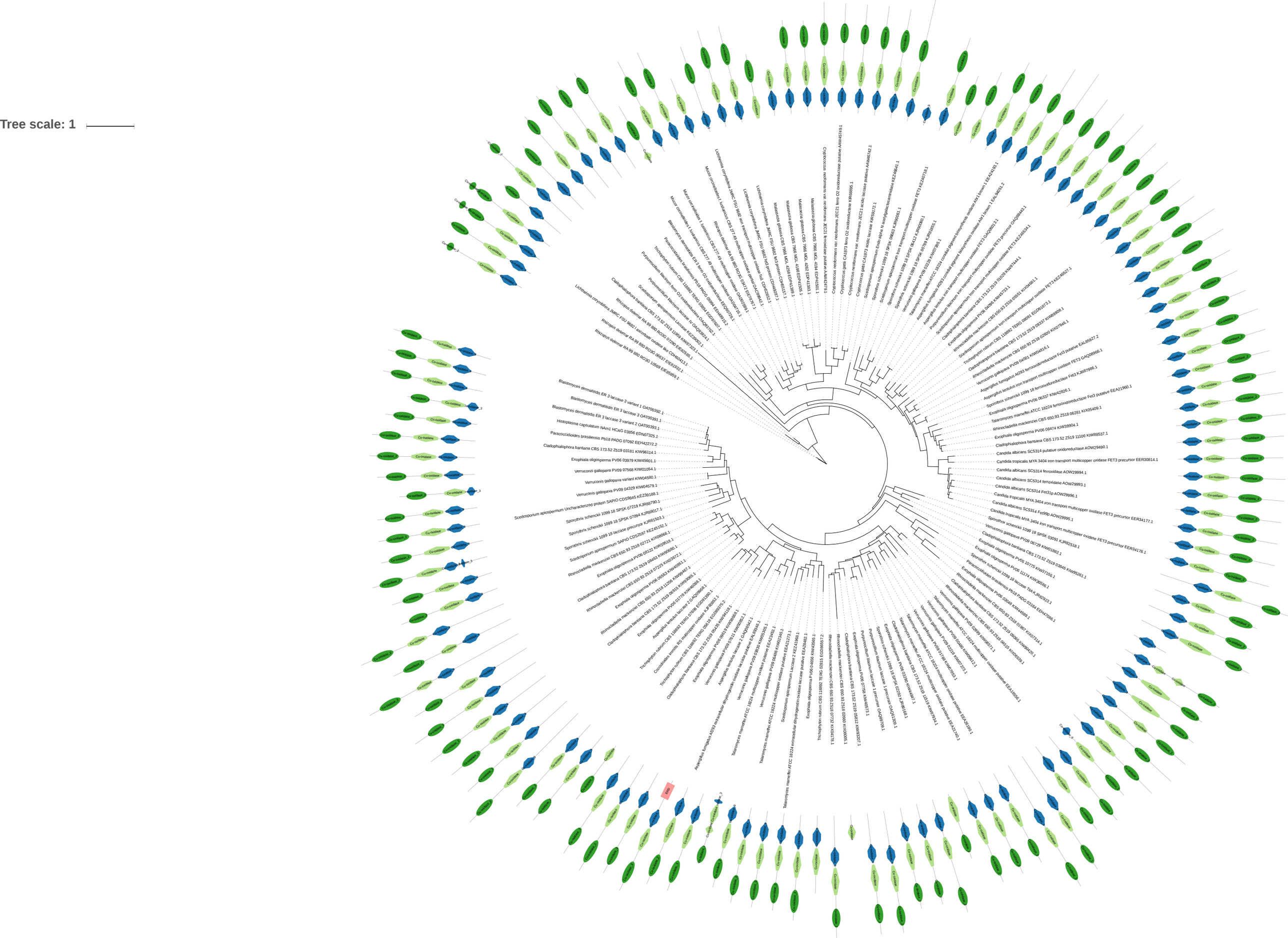
Tree scale: 1


Cu-oxidase
Cu-oxidase_2
Cu-oxidase_3
EPO
Blastomyces dermatitidis ER 3 laccase 3 variant 1 OAT00392.1
Blastomyces dermatitidis ER 3 laccase 3 OAT00391.1
Blastomyces dermatitidis ER 3 laccase 3 variant 2 OAT00393.1
Histoplasma capsulatum NAm1 HCAG 03856 EDN07325.1
Paracoccidioides brasiliensis Pb18 PADG 07092 EEH42272.2
Cladophialophora bantiana CBS 173.52 Z519 03181 KIW96114.1
Exophiala oligosperma PV06 03979 KIW45601.1
Verruconis gallopava PV09 07568 KIW01054.1
Verruconis gallopava variant KIW04580.1
Verruconis gallopava PV09 04329 KIW04579.1
Scedosporium apiospermum Uncharacterized protein SAPIO CDS9845 KEZ39188.1
Sporothrix schenckii 1099 18 SPSK 07219 KJR88790.1
Sporothrix schenckii 1099 18 SPSK 07094 KJR88017.1
Sporothrix schenckii 1099 18 laccase precursor KJR81503.1
Sporothrix schenckii 1099 18 laccase precursor KEZ45152.1
Scedosporium apiospermum SAPIO CDS2597 KEZ45152.1
Rhinocladiella mackenziei CBS 650.93 Z518 02721 KIW39566.1
Exophiala oligosperma PV06 68122 KIW90680.1
Cladophialophora bantiana CBS 173.52 Z519 08493 KIW90680.1
Rhinocladiella mackenziei CBS 650.93 Z518 07225 KIW40961.1
Cladophialophora bantiana CBS 173.52 Z519 07225 KIW40961.1
Rhinocladiella mackenziei CBS 650.93 Z518 06553 KIW89407.1
Exophiala oligosperma PV06 06553 KIW89407.1
Rhinocladiella mackenziei CBS 650.93 Z518 11206 KIW93665.1
Cladophialophora bantiana CBS 173.52 Z519 11206 KIW93665.1
Exophiala oligosperma PV06 06915 KIW96868.1
Exophiala oligosperma PV09 01778 KIW96868.1
Cladophialophora bantiana CBS 173.52 Z519 06915 KIW96868.1
Rhinocladiella mackenziei CBS 650.93 Z518 EGD56375.2
Cladophialophora bantiana CBS 118892 TERG 05818 EGD56375.2
Aspergillus lentulus RI multicopper oxidase 2 GAQ09662.1
Aspergillus lentulus laccase 2 KEZ40662.1
Trichophyton rubrum CBS 118892 TERG 07508 EGD93190.1
Trichophyton rubrum CBS 118892 TERG 05818 EGD56375.2
Coccidioides immitis RS multicopper oxidase EAS30681.1
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Cladophialophora bantiana CBS 173.52 Z519 06425 KIW94119.1
Exophiala oligosperma PV06 06915 KIW96868.1
Aspergillus lentulus laccase 2 GAQ09662.1
Verruconis gallopava PV09 07811 KIW98085.1
Aspergillus lentulus laccase 2 KEZ40662.1
Verruconis gallopava PV09 01830 KIW03305.1
Trichophyton rubrum CBS 118892 TERG 05818 EGD56375.2
Verruconis gallopava PV09 06468 KIW03345.1
Aspergillus lentulus laccase 2 GAQ09662.1
Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21273.1
Scedosporium apiospermum Laccase 2 KEZ41082.1
Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21273.1
Exophiala oligosperma PV06 04656 KIW43566.1
Talaromyces marneffei ATCC 18224 extracellular dihydrogoodin oxidase laccase putative EEA20482.1
Trichophyton rubrum CBS 118892 TERG 03815 EGD86557.2
Rhinocladiella mackenziei CBS 650.93 Z518 07227 KIW34718.1
Rhinocladiella mackenziei CBS 650.93 Z518 03660 KIW08095.1
Cladophialophora bantiana CBS 173.52 Z519 05812 KIW92207.1
Exophiala oligosperma PV06 07758 KIW40573.1
Purpureocillium lilacinum laccase 1 precursor OAQ97709.1
Sporothrix schenckii 1099 18 SPSK 02292 EJR66186.1
Exophiala oligosperma PV06 02360 KIW44847.1
Cladophialophora bantiana CBS 173.52 Z519 1519 KIW97924.1
Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21273.1
Verruconis gallopava PV09 01728 KIW07373.1
Verruconis gallopava PV09 01728 KIW07353.1
Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21273.1
Verruconis gallopava PV09 02696 KIW06571.1
Verruconis gallopava PV09 04680 KIW08012.1
Rhinocladiella mackenziei CBS 650.93 Z518 01967 KIW27314.1
Cladophialophora bantiana CBS 173.52 Z519 06365 KIW96430.1
Verruconis gallopava PV09 04515 KIW06339.1
Rhinocladiella mackenziei CBS 650.93 Z518 02360 KIW44568.1
Exophiala oligosperma PV06 02648 KIW44568.1
Cladophialophora bantiana CBS 173.52 Z519 03845 KIW85261.1
Exophiala oligosperma PV06 10775 KIW37156.1
Exophiala oligosperma PV06 11174 KIW38586.1
Paracoccidioides brasiliensis Pb18 PADG 03184 EEH47086.1
Sporothrix schenckii 1099 18 laccase TIAKJR82923.1
Verruconis gallopava PV09 06729 KIW01882.1
Cladophialophora bantiana CBS 173.52 Z519 11105 KIW88537.1
Exophiala oligosperma PV06 08474 KIW39904.1
Rhinocladiella mackenziei CBS 650.93 Z518 06281 KIX05409.1
Talaromyces marneffei ATCC 18224 ferrooxidoreductase Fet3 putative EEA21960.1
Sporothrix schenckii 1099 18 ferrooxidoreductase Fet3 KJR87896.1
Exophiala oligosperma PV06 06337 KIW42826.1
Aspergillus lentulus iron transport multicopper oxidase FET3 GAQ09896.1
Aspergillus fumigatus A093 ferrooxidoreductase Fet3 putative EAL85927.2
Verruconis gallopava PV09 04001 KIW04816.1
Rhinocladiella mackenziei CBS 650.93 Z518 05500 KIW07846.1
Trichophyton rubrum CBS 118892 TERG 08091 EGD92873.1
Cladophialophora bantiana CBS 173.52 Z519 09337 KIW89698.1
Exophiala oligosperma PV06 04086 KIW45733.1
Rhinocladiella mackenziei CBS 650.93 Z518 09531 KIX04681.1
Scedosporium apiospermum iron transport multicopper oxidase FET3 KEZ46527.1
Trichophyton rubrum CBS 118892 TERG 03815 EGD86557.2
Scedosporium apiospermum iron transport multicopper oxidase FET3 KEZ46534.1
Purpureocillium lilacinum iron transport multicopper oxidase FET3 KWZ97444.1
Aspergillus lentulus iron transport multicopper oxidase FET3 GAQ09441.1
Aspergillus fumigatus A0291 conidial pigment biosynthesis oxidase Abr1 isoform 1 EEA24168.1
Aspergillus fumigatus A0291 conidial pigment biosynthesis oxidase Abr1 isoform 1 EAL94052.2
Cryptococcus neoformans var. neoformans JEC21 laccase putative AAW47471.1
Cryptococcus neoformans var. neoformans JEC21 ferro O2 oxidoreductase putative AAW45749.1
Cryptococcus gattii CA1873 ferro O2 oxidoreductase KIR68895.1
Cryptococcus gattii CA1873 laccase putative AAW46742.1
Cryptococcus gattii CA1873 acidic laccase KIR59172.1
Scedosporium apiospermum Endo alpha N acetylgalactosaminidase KEZ46641.1
Sporothrix schenckii 1099 18 SPSK 06663 KJR90091.1
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Verruconis gallopava PV09 02228 KIW07865.1
Talaromyces marneffei ATCC 18224 conidial pigment biosynthesis oxidase putative EEA24168.1
Aspergillus lentulus iron transport multicopper oxidase FET3 GAQ09441.1
Purpureocillium lilacinum iron transport multicopper oxidase FET3 KWZ97444.1
Scedosporium apiospermum iron transport multicopper oxidase FET3 KEZ46534.1
Cladophialophora bantiana CBS 173.52 Z519 01028 KIW97444.1
Exophiala oligosperma PV06 04086 KIW45733.1
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Scedosporium apiospermum iron transport multicopper oxidase FET3 KEZ46527.1
Trichophyton rubrum CBS 118892 TERG 08091 EGD92873.1
Cladophialophora bantiana CBS 173.52 Z519 09337 KIW89698.1
Rhinocladiella mackenziei CBS 650.93 Z518 05500 KIW07846.1
Aspergillus lentulus iron transport multicopper oxidase FET3 GAQ09896.1
Aspergillus fumigatus A093 ferrooxidoreductase Fet3 putative EAL85927.2
Verruconis gallopava PV09 04001 KIW04816.1
Exophiala oligosperma PV06 06337 KIW42826.1
Sporothrix schenckii 1099 18 ferrooxidoreductase Fet3 putative EEA21960.1
Talaromyces marneffei ATCC 18224 ferrooxidoreductase Fet3 KJR87896.1
Rhinocladiella mackenziei CBS 650.93 Z518 06281 KIX05409.1
Exophiala oligosperma PV06 08474 KIW39904.1
Cladophialophora bantiana CBS 173.52 Z519 11105 KIW88537.1
Candida albicans SC5314 putative oxidoreductase AOW29490.1
Candida tropicalis MYA 3404 iron transport multicopper oxidase FET3 precursor EER30814.1
Candida albicans SC5314 ferroxidase AOW29994.1
Candida albicans SC5314 ferroxidase AOW29993.1
Candida albicans SC5314 Fet31p AOW29996.1
Candida tropicalis MYA 3404 iron transport multicopper oxidase FET3 precursor EER34177.1
Candida albicans SC5314 Fet99p AOW29995.1
Candida tropicalis MYA 3404 iron transport multicopper oxidase FET3 precursor EER34178.1
Sporothrix schenckii 1099 18 SPSK 03091 KJR82518.1
Verruconis gallopava PV09 06729 KIW01882.1
Cladophialophora bantiana CBS 173.52 Z519 03845 KIW85261.1
Exophiala oligosperma PV06 10775 KIW37156.1
Exophiala oligosperma PV06 11174 KIW38586.1
Paracoccidioides brasiliensis Pb18 PADG 03184 EEH47086.1
Sporothrix schenckii 1099 18 laccase TIAKJR82923.1
Rhinocladiella mackenziei CBS 650.93 Z518 02360 KIW44568.1
Exophiala oligosperma PV06 02648 KIW44568.1
Cladophialophora bantiana CBS 173.52 Z519 03845 KIW85261.1
Rhinocladiella mackenziei CBS 650.93 Z518 01967 KIW27314.1
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Verruconis gallopava PV09 04515 KIW06339.1
Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21960.1
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Verruconis gallopava PV09 04680 KIW08012.1
Verruconis gallopava PV09 01728 KIW07373.1
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Talaromyces marneffei ATCC 18224 multicopper oxidase putative EEA21273.1
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