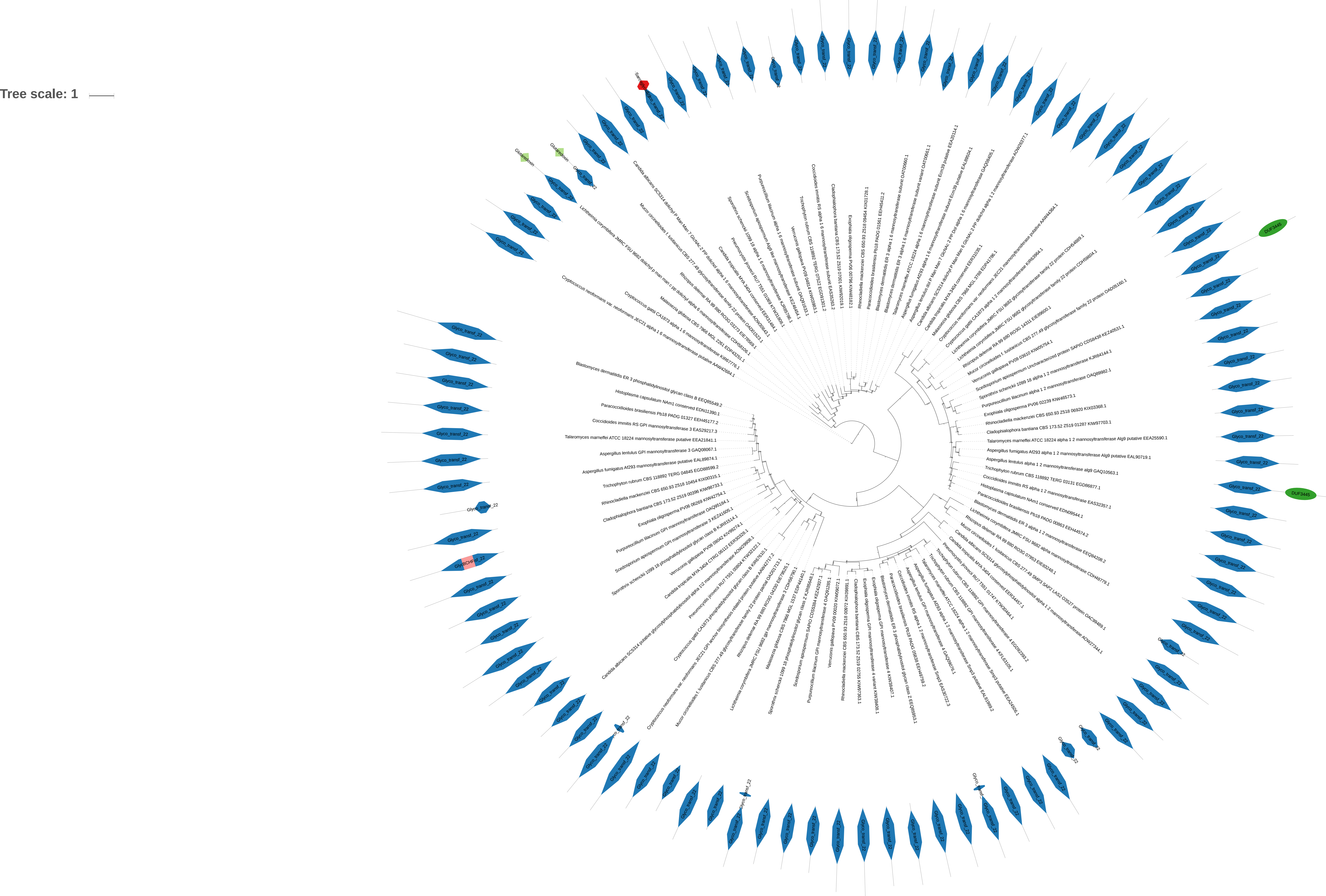
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Blastomyces dermatitidis ER 3 phosphatidylinositol glycan class B EEQ85549.2
Histoplasma capsulatum NAm1 conserved EDN11390.1
Paracoccidioides brasiliensis Pb18 PADG 01327 EEH45177.2
Coccidioides immitis RS GPI mannosyltransferase 3 EAS29217.3
Talaromyces marneffei ATCC 18224 mannosyltransferase putative EEA21841.1
Aspergillus lentulus GPI mannosyltransferase 3 GAQ08067.1
Aspergillus fumigatus Af293 mannosyltransferase putative EAL89874.1
Trichophyton rubrum CBS 118892 TERG 04845 EGD88599.2
Rhinocladiella mackenziei CBS 650.93 Z518 10454 KIX00315.1
Cladophialophora bantiana CBS 173.52 Z519 00396 KIW98733.1
Exophiala oligosperma PV06 06269 KIW42754.1
Purpureocillium lilacinum GPI mannosyltransferase OAQ95184.1
Scedosporium apiospermum GPI mannosyltransferase 3 KEZ41605.1
Sporothrix schenckii 1099 18 phosphatidylinositol glycan class B KJR81514.1
Verruconis gallopava PV09 09042 KIV99274.1
Candida tropicalis MYA 3404 CTRG 06112 EER30328.1
Candida albicans SC5314 putative glycosylphosphatidylinositol alpha 1l2 mannosyltransferase AOW29908.1
Pneumocystis jirovecii RU7 T551 00804 KTW32122.1
Cryptococcus gattii CA1873 phosphatidylinositol glycan class B KIR67610.1
Cryptococcus neoformans var. neoformans JEC21 GPI anchor biosynthesis related protein putative AAW42717.2
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 22 protein partial OAD01713.1
Rhizopus delemar RA 99 880 RO3G 04230 EIE79525.1
Lichtheimia corymbifera JMRC FSU 9682 gpi mannosyltransferase 3 CDH56790.1
Malassezia globosa CBS 7966 MGL 1537 EDP44140.1
Sporothrix schenckii 1099 18 phosphatidylinositol glycan class Z KJR85648.1
Scedosporium apiospermum SAPIO CDS5384 KEZ42937.1
Purpureocillium lilacinum GPI mannosyltransferase 4 OAQ91285.1
Verruconis gallopava PV09 00020 KIW09072.1
Rhinocladiella mackenziei CBS 650.93 Z518 00972 KIX09891.1
Cladophialophora bantiana CBS 173.52 Z519 02755 KIW97363.1
Exophiala oligosperma GPI mannosyltransferase 4 variant KIW38408.1
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Blastomyces dermatitidis ER 3 phosphatidylinositol glycan class Z EEQ85953.1
Paracoccidioides brasiliensis Pb18 PADG 05838 EEH49759.2
Coccidioides immitis RS alpha 1 2 mannosyltransferase Smp3 EAS30722.3
Aspergillus lentulus GPI mannosyltransferase 4 GAQ09076.1
Aspergillus fumigatus Af293 alpha 1 2 mannosyltransferase Smp3 putative EAL91989.2
Talaromyces marneffei ATCC 18224 alpha 1 2 mannosyltransferase Smp3 putative EEA24506.1
Trichophyton rubrum CBS 118892 GPI mannosyltransferase 4 KFL63105.1
Trichophyton rubrum CBS 118892 GPI mannosyltransferase 4 EGD92393.2
Pneumocystis jirovecii RU7 T551 01747 KTW30464.1
Candida tropicalis MYA 3404 conserved EER34457.1
Candida albicans SC5314 glycosylphosphatidylinositol alpha 1 2 mannosyltransferase AOW27344.1
Mucor circinelloides f. lusitanicus CBS 277.49 SMP3 SAP2 LAS2 O3527 protein OAC99489.1
Rhizopus delemar RA 99 880 RO3G 07953 EIE83248.1
Lichtheimia corymbifera JMRC FSU 9682 alpha mannosyltransferase CDH48779.1
Blastomyces dermatitidis ER 3 alpha 1 2 mannosyltransferase EEQ84208.2
Paracoccidioides brasiliensis Pb18 PADG 00863 EEH44574.2
Histoplasma capsulatum NAm1 conserved EDN09544.1
Coccidioides immitis RS alpha 1 2 mannosyltransferase EAS32357.1
Trichophyton rubrum CBS 118892 TERG 03131 EGD86877.1
Aspergillus lentulus alpha 1 2 mannosyltransferase alg9 GAQ10563.1
Aspergillus fumigatus Af293 alpha 1 2 mannosyltransferase Alg9 putative EAL90719.1
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Cladophialophora bantiana CBS 173.52 Z519 01287 KIW97703.1
Rhinocladiella mackenziei CBS 650.93 Z518 06920 KIX03368.1
Exophiala oligosperma PV06 02239 KIW46573.1
Purpureocillium lilacinum alpha 1 2 mannosyltransferase OAQ89982.1
Sporothrix schenckii 1099 18 alpha 1 2 mannosyltransferase KJR84144.1
Scedosporium apiospermum Uncharacterized protein SAPIO CDS8438 KEZ40531.1
Verruconis gallopava PV09 03610 KIW05754.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 22 protein OAD05160.1
Rhizopus delemar RA 99 880 RO3G 14311 EIE89600.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 22 protein CDH59604.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 22 protein CDH54889.1
Cryptococcus gattii CA1873 alpha 1 2 mannosyltransferase KIR63964.1
Cryptococcus neoformans var. neoformans JEC21 mannosyltransferase putative AAW44364.1
Malassezia globosa CBS 7966 MGL 3788 EDP41786.1
Candida tropicalis MYA 3404 conserved EER31035.1
Candida albicans SC5314 dolichyl P Man Man 6 GlcNAc 2 PP dolichol alpha 1 2 mannosyltransferase AOW26377.1
Aspergillus lentulus dol P Man Man 7 GlcNAc 2 PP Dol alpha 1 6 mannosyltransferase GAQ08405.1
Aspergillus fumigatus Af293 alpha 1 6 mannosyltransferase subunit Ecm39 putative EAL89504.1
Talaromyces marneffei ATCC 18224 alpha 1 6 mannosyltransferase subunit Ecm39 putative EEA20114.1
Blastomyces dermatitidis ER 3 alpha 1 6 mannosyltransferase subunit variant OAT00661.1
Blastomyces dermatitidis ER 3 alpha 1 6 mannosyltransferase subunit OAT00660.1
Paracoccidioides brasiliensis Pb18 PADG 01561 EEH45411.2
Rhinocladiella mackenziei CBS 650.93 Z518 09454 KIX01728.1
Exophiala oligosperma PV06 00796 KIW48182.1
Cladophialophora bantiana CBS 173.52 Z519 07001 KIW92019.1
Coccidioides immitis RS alpha 1 6 mannosyltransferase subunit EAS35263.2
Trichophyton rubrum CBS 118892 TERG 07522 EGD91301.2
Verruconis gallopava PV09 04814 KIW03983.1
Purpureocillium lilacinum alpha 1 6 mannosyltransferase subunit OAQ91633.1
Scedosporium apiospermum Alg9 like mannosyltransferase KEZ46454.1
Sporothrix schenckii 1099 18 alpha 1 6 mannosyltransferase KJR85786.1
Pneumocystis jirovecii RU7 T551 01069 KTW31808.1
Candida tropicalis MYA 3404 conserved EER31484.1
Candida albicans SC5314 dolichyl P Man Man 7 GlcNAc 2 PP dolichol alpha 1 6 mannosyltransferase AOW30643.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 22 protein OAD01523.1
Rhizopus delemar RA 99 880 RO3G 03273 EIE78569.1
Lichtheimia corymbifera JMRC FSU 9682 dolichyl p man man c pp dolichyl alpha 6 mannosyltransferase CDH58106.1
Malassezia globosa CBS 7966 MGL 2261 EDP43251.1
Cryptococcus gattii CA1873 alpha 1 6 mannosyltransferase KIR67776.1
Cryptococcus neoformans var. neoformans JEC21 alpha 1 6 mannosyltransferase putative AAW42684.1
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Glutaredoxin
Sarcoglycan
GlycBCHFisf_22
DUF3445