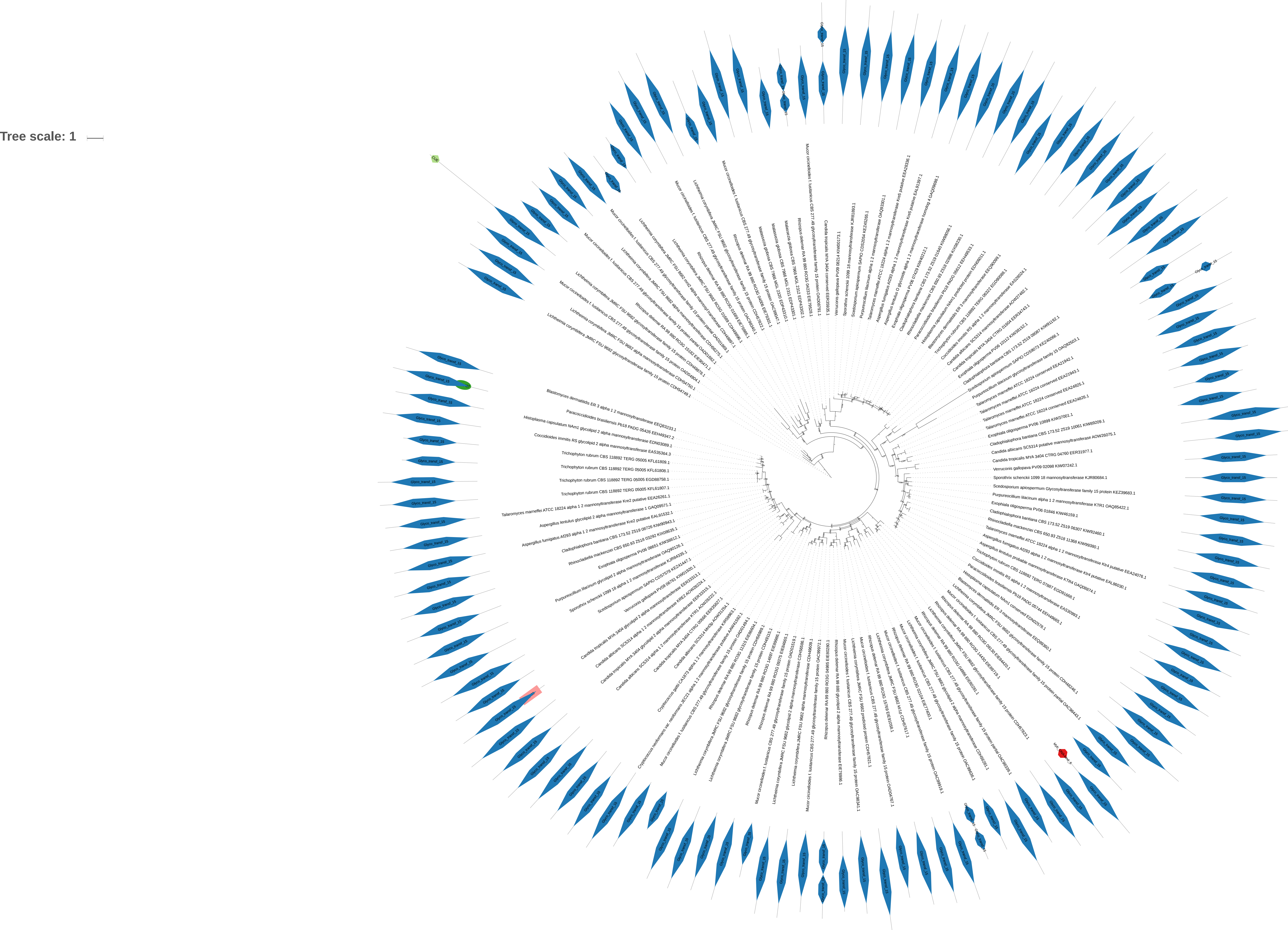
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CUE
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Lichtheimia corymbifera JMRC FSU 9682 alpha mannosyltransferase CDH54750.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAD04904.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH49878.1
Rhizopus delemar RA 99 880 RO3G 15182 EIE90471.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein partial OAD01882.1
Lichtheimia corymbifera JMRC FSU 9682 alpha mannosyltransferase CDH56275.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein partial OAD01869.1
Lichtheimia corymbifera JMRC FSU 9682 mnt2 alpha mannosyl transferase CDH49897.1
Lichtheimia corymbifera JMRC FSU 9682 RO3G 01689 CDH49896.1
Rhizopus delemar RA 99 880 RO3G 01689 EIE76985.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC98340.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH57822.1
Rhizopus delemar RA 99 880 RO3G 04006 EIE79301.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC99047.1
Malassezia globosa CBS 7966 MGL 2320 EDP43310.1
Malassezia globosa CBS 7966 MGL 2311 EDP43301.1
Malassezia globosa CBS 7966 MGL 2312 EDP43302.1
Rhizopus delemar RA 99 880 RO3G 04233 EIE79528.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAD08791.1
Candida tropicalis MYA 3404 conserved EER35935.1
Verruconis gallopava PV09 08214 KIW00173.1
Sporothrix schenckii 1099 18 mannosyltransferase KJR81893.1
Scedosporium apiospermum SAPIO CDS2054 KEZ45265.1
Purpureocillium lilacinum alpha 1 2 mannosyltransferase OAQ93301.1
Talaromyces marneffei ATCC 18224 alpha 1 2 mannosyltransferase Kre5 putative EEA28336.1
Aspergillus fumigatus Af293 alpha 1 2 mannosyltransferase Kre5 putative EAL91397.1
Aspergillus lentulus O glycoside alpha 1 2 mannosyltransferase homolog 4 GAQ09698.1
Exophiala oligosperma PV06 07429 KIW40212.1
Cladophialophora bantiana CBS 173.52 Z519 01640 KIW98056.1
Rhinocladiella mackenziei CBS 650.93 Z518 02886 KIX08230.1
Paracoccidioides brasiliensis Pb18 PADG 05612 EEH49533.1
Histoplasma capsulatum NAm1 predicted protein EDN06011.1
Blastomyces dermatitidis ER 3 mannosyltransferase EEQ90099.1
Trichophyton rubrum CBS 118892 TERG 06322 EGD90088.1
Coccidioides immitis RS alpha 1 2 mannosyltransferase EAS28024.1
Candida albicans SC5314 mannosyltransferase AOW27482.1
Candida tropicalis MYA 3404 CTRG 01604 EER34743.1
Exophiala oligosperma PV06 10112 KIW38152.1
Cladophialophora bantiana CBS 173.52 Z519 08087 KIW91192.1
Scedosporium apiospermum SAPIO CDS9073 KEZ40066.1
Purpureocillium lilacinum glycosyltransferase family 15 OAQ92503.1
Talaromyces marneffei ATCC 18224 conserved EEA21942.1
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Talaromyces marneffei ATCC 18224 conserved EEA24825.1
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Exophiala oligosperma PV06 10899 KIW37001.1
Cladophialophora bantiana CBS 173.52 Z519 10061 KIW89209.1
Candida albicans SC5314 putative mannosyltransferase AOW26075.1
Candida tropicalis MYA 3404 CTRG 04760 EER31977.1
Verruconis gallopava PV09 02098 KIW07242.1
Sporothrix schenckii 1099 18 mannosyltransferase KJR80684.1
Scedosporium apiospermum Glycosyltransferase family 15 protein KEZ39683.1
Purpureocillium lilacinum alpha 1 2 mannosyltransferase KTR1 OAQ85422.1
Exophiala oligosperma PV06 01846 KIW46159.1
Cladophialophora bantiana CBS 173.52 Z519 06307 KIW92460.1
Rhinocladiella mackenziei CBS 650.93 Z518 11368 KIW99380.1
Talaromyces marneffei ATCC 18224 alpha 1 2 mannosyltransferase Ktr4 putative EEA24076.1
Aspergillus fumigatus Af293 alpha 1 2 mannosyltransferase Ktr4 putative EAL86030.1
Aspergillus lentulus probable mannosyltransferase KTR4 GAQ08874.1
Trichophyton rubrum CBS 118892 TERG 07887 EGD91668.1
Coccidioides immitis RS alpha 1 2 mannosyltransferase EAS30993.1
Paracoccidioides brasiliensis Pb18 PADG 05744 EEH49665.1
Histoplasma capsulatum NAm1 conserved EDN02578.1
Blastomyces dermatitidis ER 3 mannosyltransferase EEQ89360.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH48246.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein partial OAC98443.1
Rhizopus delemar RA 99 880 RO3G 09130 EIE84420.1
Rhizopus delemar RA 99 880 RO3G 14430 EIE89719.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH57823.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein partial OAC98339.1
Rhizopus delemar RA 99 880 RO3G 04986 EIE80281.1
Lichtheimia corymbifera JMRC FSU 9682 glycolipid 2 alpha mannosyltransferase CDH56281.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC99636.1
Rhizopus delemar RA 99 880 RO3G 02104 EIE77400.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC99919.1
Lichtheimia corymbifera JMRC FSU 9682 ktr1p CDH57617.1
Rhizopus delemar RA 99 880 RO3G 15769 EIE91058.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAD04767.1
Lichtheimia corymbifera JMRC FSU 9682 predicted protein CDH57821.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC98341.1
Rhizopus delemar RA 99 880 glycolipid 2 alpha mannosyltransferase EIE76986.1
Rhizopus delemar RA 99 880 RO3G 04985 EIE80280.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAC99972.1
Lichtheimia corymbifera JMRC FSU 9682 alpha mannosyltransferase CDH49609.1
Lichtheimia corymbifera JMRC FSU 9682 glycolipid 2 alpha mannosyltransferase CDH56586.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAD01519.1
Rhizopus delemar RA 99 880 RO3G 09375 EIE84665.1
Rhizopus delemar RA 99 880 RO3G 14697 EIE89986.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH49115.1
Lichtheimia corymbifera JMRC FSU 9682 glycosyltransferase family 15 protein CDH56989.1
Rhizopus delemar RA 99 880 RO3G 11315 EIE86604.1
Mucor circinelloides f. lusitanicus CBS 277.49 glycosyltransferase family 15 protein OAD01494.1
Cryptococcus neoformans var. neoformans JEC21 alpha 1 2 mannosyltransferase putative AAW41592.1
Cryptococcus gattii CA1873 alpha 1 2 mannosyltransferase KIR59963.1
Candida albicans SC5314 Mnt3p AOW31254.1
Candida tropicalis MYA 3404 CTRG 00566 EER35827.1
Candida albicans SC5314 alpha 1 2 mannosyltransferase KTR1 AOW28222.1
Candida tropicalis MYA 3404 glycolipid 2 alpha mannosyltransferase EER33315.1
Candida albicans SC5314 alpha 1 2 mannosyltransferase KRE2 AOW28224.1
Candida tropicalis MYA 3404 glycolipid 2 alpha mannosyltransferase EER33313.1
Verruconis gallopava PV09 06761 KIW01920.1
Scedosporium apiospermum SAPIO CDS7578 KEZ41447.1
Sporothrix schenckii 1099 18 alpha 1 2 mannosyltransferase KJR84335.1
Purpureocillium lilacinum glycolipid 2 alpha mannosyltransferase OAQ90126.1
Exophiala oligosperma PV06 08651 KIW38812.1
Rhinocladiella mackenziei CBS 650.93 Z518 03292 KIX08635.1
Cladophialophora bantiana CBS 173.52 Z519 08726 KIW90943.1
Aspergillus fumigatus Af293 alpha 1 2 mannosyltransferase Kre2 putative EAL91532.1
Aspergillus lentulus glycolipid 2 alpha mannosyltransferase 1 GAQ09571.1
Talaromyces marneffei ATCC 18224 alpha 1 2 mannosyltransferase Kre2 putative EEA26261.1
Trichophyton rubrum CBS 118892 TERG 05005 KFL61807.1
Trichophyton rubrum CBS 118892 TERG 05005 EGD88758.1
Trichophyton rubrum CBS 118892 TERG 05005 KFL61808.1
Trichophyton rubrum CBS 118892 TERG 05005 KFL61809.1
Coccidioides immitis RS glycolipid 2 alpha mannosyltransferase EAS35364.3
Histoplasma capsulatum NAm1 glycolipid 2 alpha mannosyltransferase EDN03089.1
Paracoccidioides brasiliensis Pb18 PADG 05426 EEH49347.2