

CR382121.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome A complete sequence
1062590
CR382122.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome B complete sequence
1320834
CR382123.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome C complete sequence
1753957
CR382124.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome D complete sequence
1715506
CR382125.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome E complete sequence
2234072
CR382126.1 *Kluyveromyces lactis* strain NRRL Y-1140 chromosome F complete sequence
2602197
CR382127.1 *Yarrowia lipolytica* CLIB122 chromosome A complete sequence
2303261
CR382128.1 *Yarrowia lipolytica* CLIB122 chromosome B complete sequence
3066374
CR382129.1 *Yarrowia lipolytica* CLIB122 chromosome C complete sequence
3272609
CR382130.1 *Yarrowia lipolytica* CLIB122 chromosome D complete sequence
3633272
CR382131.1 *Yarrowia lipolytica* CLIB122 chromosome E complete sequence
4224103
CR382132.1 *Yarrowia lipolytica* CLIB122 chromosome F complete sequence
4003362
CR380947.2 *Candida glabrata* strain CBS138 chromosome A complete sequence
491328
CR380948.1 *Candida glabrata* strain CBS138 chromosome B complete sequence
502101
CR380949.1 *Candida glabrata* strain CBS138 chromosome C complete sequence
558804
CR380950.1 *Candida glabrata* strain CBS138 chromosome D complete sequence
651701
CR380951.2 *Candida glabrata* strain CBS138 chromosome E complete sequence
687738
CR380952.1 *Candida glabrata* strain CBS138 chromosome F complete sequence
927101
CR380953.1 *Candida glabrata* strain CBS138 chromosome G complete sequence
992211
CR380954.1 *Candida glabrata* strain CBS138 chromosome H complete sequence
1050361
CR380955.2 *Candida glabrata* strain CBS138 chromosome I complete sequence
1100349
CR380956.2 *Candida glabrata* strain CBS138 chromosome J complete sequence
1195132
CR380957.2 *Candida glabrata* strain CBS138 chromosome K complete sequence
1302831
CR380958.2 *Candida glabrata* strain CBS138 chromosome L complete sequence
1455689
CR380959.2 *Candida glabrata* strain CBS138 chromosome M complete sequence
1402899
CU329670.1 *Schizosaccharomyces pombe* chromosome I, complete sequence
5579133
CU329671.1 *Schizosaccharomyces pombe* chromosome II, complete sequence
4539804
CU329672.1 *Schizosaccharomyces pombe* chromosome III, complete sequence
2452883
X54421.1 *Schizosaccharomyces pombe* complete mitochondrial genome
19431
CR382133.2 *Debaryomyces hansenii* CBS767 chromosome A complete sequence
1249940

CR382134.2 *Debaryomyces hansenii* CBS767 chromosome B complete sequence
1344482
CR382135.2 *Debaryomyces hansenii* CBS767 chromosome C complete sequence
1587442
CR382136.2 *Debaryomyces hansenii* CBS767 chromosome A complete sequence
1606296
CR382137.2 *Debaryomyces hansenii* CBS767 chromosome E complete sequence
2007515
CR382138.2 *Debaryomyces hansenii* CBS767 chromosome F complete sequence
2305761
CR382139.2 *Debaryomyces hansenii* CBS767 chromosome G complete sequence
2051050
CU928173.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome A complete sequence
1114666
CU928174.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome B complete sequence
1388208
CU928175.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome C complete sequence
1464093
CU928176.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome D complete sequence
1496342
CU928181.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome E complete sequence
881646
CU928178.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome F complete sequence
1554288
CU928179.1 *Zygosaccharomyces rouxii* strain CBS732 chromosome G complete sequence
1865392
FM992695.1 *Candida dubliniensis* CD36 chromosome R, complete sequence
2267510
FM992688.1 *Candida dubliniensis* CD36 chromosome 1, complete sequence
3214061
FM992689.1 *Candida dubliniensis* CD36 chromosome 2, complete sequence
2289089
FM992690.1 *Candida dubliniensis* CD36 chromosome 3, complete sequence
1863824
FM992691.1 *Candida dubliniensis* CD36 chromosome 4, complete sequence
1641709
FM992692.1 *Candida dubliniensis* CD36 chromosome 5, complete sequence
1245899
FM992693.1 *Candida dubliniensis* CD36 chromosome 6, complete sequence
1073895
FM992694.1 *Candida dubliniensis* CD36 chromosome 7, complete sequence
1022435
FN392319.1 *Pichia pastoris* GS115 chromosome 1, complete sequence
2798491
FN392320.1 *Pichia pastoris* GS115 chromosome 2, complete sequence
2394163
FN392321.1 *Pichia pastoris* GS115 chromosome 3, complete sequence
2245428
FN392322.1 *Pichia pastoris* GS115 chromosome 4, complete sequence
1778296
AE016814.2 *Ashbya gossypii* ATCC 10895 chromosome I, complete sequence
693414
AE016815.5 *Ashbya gossypii* ATCC 10895 chromosome II, complete sequence
870771
AE016816.3 *Ashbya gossypii* ATCC 10895 chromosome III, complete sequence
907494
AE016817.6 *Ashbya gossypii* ATCC 10895 chromosome IV, complete sequence
1467287
AE016818.2 *Ashbya gossypii* ATCC 10895 chromosome V, complete sequence
1519140

AE016819.5 *Ashbya gossypii* ATCC 10895 chromosome VI, complete sequence
1836693
AE016820.4 *Ashbya gossypii* ATCC 10895 chromosome VII, complete sequence
1800949
AE016821.1 *Ashbya gossypii* (= *Eremothecium gossypii*) ATCC 10895 mitochondrion, complete
genome
23564
AE017341.1 *Cryptococcus neoformans* var. *neoformans* JEC21 chromosome 1, complete
sequence
2300533
X59720.2 *S.cerevisiae* chromosome III complete DNA sequence
316613
D50617.1 *Saccharomyces cerevisiae* DNA, chromosome VI, complete genome
270148
CU928165.1 *Lachancea thermotolerans* CBS 6340 chromosome A complete sequence
687718
CU928166.1 *Lachancea thermotolerans* CBS 6340 chromosome B complete sequence
893706
CU928167.1 *Lachancea thermotolerans* CBS 6340 chromosome C complete sequence
999246
CU928168.1 *Lachancea thermotolerans* CBS 6340 chromosome D complete sequence
1513537
CU928169.1 *Lachancea thermotolerans* CBS 6340 chromosome E complete sequence
1521774
CU928170.1 *Lachancea thermotolerans* CBS 6340 chromosome F complete sequence
1632914
CU928171.1 *Lachancea thermotolerans* CBS 6340 chromosome G complete sequence
1720065
CU928180.1 *Lachancea thermotolerans* CBS 6340 chromosome H complete sequence
1423902
CP009805.1 *Botrytis cinerea* B05.10 chromosome BCIN01, complete sequence
4109373
CP009806.1 *Botrytis cinerea* B05.10 chromosome BCIN02, complete sequence
3341473
CP009807.1 *Botrytis cinerea* B05.10 chromosome BCIN03, complete sequence
3226611
CP009808.1 *Botrytis cinerea* B05.10 chromosome BCIN04, complete sequence
2468882
CP009809.1 *Botrytis cinerea* B05.10 chromosome BCIN05, complete sequence
2959378
CP009810.1 *Botrytis cinerea* B05.10 chromosome BCIN06, complete sequence
2725906
CP009811.1 *Botrytis cinerea* B05.10 chromosome BCIN07, complete sequence
2652353
CP009812.1 *Botrytis cinerea* B05.10 chromosome BCIN08, complete sequence
2617329
CP009813.1 *Botrytis cinerea* B05.10 chromosome BCIN09, complete sequence
2547566
CP009814.1 *Botrytis cinerea* B05.10 chromosome BCIN10, complete sequence
2419276
CP009815.1 *Botrytis cinerea* B05.10 chromosome BCIN11, complete sequence
2359939
CP009816.1 *Botrytis cinerea* B05.10 chromosome BCIN12, complete sequence
2352958
CP009817.1 *Botrytis cinerea* B05.10 chromosome BCIN13, complete sequence
2257609
CP009818.1 *Botrytis cinerea* B05.10 chromosome BCIN14, complete sequence
2138025
CP009819.1 *Botrytis cinerea* B05.10 chromosome BCIN15, complete sequence
2027721

CP009820.1 *Botrytis cinerea* B05.10 chromosome BCIN16, complete sequence
1969743
CP009821.1 *Botrytis cinerea* B05.10 chromosome BCIN17, complete sequence
247158
CP009822.1 *Botrytis cinerea* B05.10 chromosome BCIN18, complete sequence
208766
BK006935.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome I, complete sequence
230218
BK006936.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome II, complete sequence
813184
BK006937.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome III, complete sequence
316620
BK006938.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome IV, complete sequence
1531933
BK006939.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome V, complete sequence
576874
BK006940.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome VI, complete sequence
270161
BK006941.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome VII, complete sequence
1090940
BK006934.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome VIII, complete sequence
562643
BK006942.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome IX, complete sequence
439888
BK006943.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome X, complete sequence
745751
BK006944.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XI, complete sequence
666816
BK006945.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XII, complete sequence
1078177
BK006946.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XIII, complete sequence
924431
BK006947.3 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XIV, complete sequence
784333
BK006948.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XV, complete sequence
1091291
BK006949.2 TPA_inf: *Saccharomyces cerevisiae* S288C chromosome XVI, complete sequence
948066
CP001942.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome I, complete sequence
160332
CP001943.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome II, complete sequence
175776
CP001944.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome III, complete sequence
176815
CP001945.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome IV, complete sequence
193740
CP001946.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome V, complete sequence
196642
CP001947.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VI, complete sequence
198217
CP001948.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VII, complete sequence
205935
CP001949.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VIII, complete sequence
204910
CP001951.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome X, complete sequence
234890
CP001952.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome XI, complete sequence
236244
CP003820.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 1, complete sequence
2291499

CP003821.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 2, complete sequence
1621675
CP003822.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 3, complete sequence
1575141
CP003823.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 4, complete sequence
1084805
CP003824.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 5, complete sequence
1814975
CP003825.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 6, complete sequence
1422463
CP003826.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 7, complete sequence
1399503
CP003827.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 8, complete sequence
1398693
CP003828.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 9, complete sequence
1186808
CP003829.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 10, complete sequence
1059964
CP003830.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 11, complete sequence
1561994
CP003831.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 12, complete sequence
774062
CP003832.1 *Cryptococcus neoformans* var. *grubii* H99 chromosome 13, complete sequence
756744
CP003833.2 *Cryptococcus neoformans* var. *grubii* H99 chromosome 14, complete sequence
942867
CP003834.1 *Cryptococcus neoformans* var. *grubii* H99 mitochondrion, complete genome
24919
KC683708.1 *Neurospora crassa* OR74A mitochondrion, complete genome
64840
AP007176.1 *Aspergillus oryzae* RIB40 mitochondrial DNA, complete genome, AOmito_02s
29202
CP000286.1 *Cryptococcus gattii* WM276 chromosome A, complete sequence
1984823
CP000287.1 *Cryptococcus gattii* WM276 chromosome B, complete sequence
2187695
CP000288.1 *Cryptococcus gattii* WM276 chromosome C, complete sequence
1961512
CP000289.1 *Cryptococcus gattii* WM276 chromosome D, complete sequence
2233618
CP000290.1 *Cryptococcus gattii* WM276 chromosome E, complete sequence
1333124
CP000291.1 *Cryptococcus gattii* WM276 chromosome F, complete sequence
1325755
CP000292.1 *Cryptococcus gattii* WM276 chromosome G, complete sequence
1324677
CP000293.1 *Cryptococcus gattii* WM276 chromosome H, complete sequence
1265488
CP000294.1 *Cryptococcus gattii* WM276 chromosome I, complete sequence
989306
CP000295.1 *Cryptococcus gattii* WM276 chromosome J, complete sequence
522727
CP000296.1 *Cryptococcus gattii* WM276 chromosome K, complete sequence
1040760
CP000297.1 *Cryptococcus gattii* WM276 chromosome L, complete sequence
820445
CP000298.1 *Cryptococcus gattii* WM276 chromosome M, complete sequence
705823
CP000299.1 *Cryptococcus gattii* WM276 chromosome N, complete sequence
679007

CP000496.1 *Scheffersomyces stipitis* CBS 6054 chromosome 2, complete sequence
2740984
CP000497.1 *Scheffersomyces stipitis* CBS 6054 chromosome 3, complete sequence
1841851
CP000498.1 *Scheffersomyces stipitis* CBS 6054 chromosome 4, complete sequence
1803401
CP000499.1 *Scheffersomyces stipitis* CBS 6054 chromosome 5, complete sequence
1725948
CP000500.1 *Scheffersomyces stipitis* CBS 6054 chromosome 6, complete sequence
1724953
CP000501.1 *Scheffersomyces stipitis* CBS 6054 chromosome 7, complete sequence
1114415
CP000502.1 *Scheffersomyces stipitis* CBS 6054 chromosome 8, complete sequence
979380
FR839628.1 *Pichia pastoris* CBS 7435 chromosome 1, complete replicon sequence
2891190
FR839629.1 *Pichia pastoris* CBS 7435 chromosome 2, complete replicon sequence
2399323
FR839630.1 *Pichia pastoris* CBS 7435 chromosome 3, complete replicon sequence
2256069
FR839631.1 *Pichia pastoris* CBS 7435 chromosome 4, complete replicon sequence
1820458
FR839632.1 *Pichia pastoris* CBS 7435 mitochondrion, complete replicon sequence
35683
CP003002.1 *Myceliophthora thermophila* ATCC 42464 chromosome 1, complete sequence
10931058
CP003003.1 *Myceliophthora thermophila* ATCC 42464 chromosome 2, complete sequence
5454242
CP003004.1 *Myceliophthora thermophila* ATCC 42464 chromosome 3, complete sequence
5062665
CP003005.1 *Myceliophthora thermophila* ATCC 42464 chromosome 4, complete sequence
4710208
CP003006.1 *Myceliophthora thermophila* ATCC 42464 chromosome 5, complete sequence
4342007
CP003007.1 *Myceliophthora thermophila* ATCC 42464 chromosome 6, complete sequence
4127677
CP003008.1 *Myceliophthora thermophila* ATCC 42464 chromosome 7, complete sequence
4116359
CP003009.1 *Thielavia terrestris* NRRL 8126 chromosome 1, complete sequence
10101509
CP003010.1 *Thielavia terrestris* NRRL 8126 chromosome 2, complete sequence
9477512
CP003011.1 *Thielavia terrestris* NRRL 8126 chromosome 3, complete sequence
4786945
CP003012.1 *Thielavia terrestris* NRRL 8126 chromosome 4, complete sequence
4578922
CP003013.1 *Thielavia terrestris* NRRL 8126 chromosome 5, complete sequence
4396881
CP003014.1 *Thielavia terrestris* NRRL 8126 chromosome 6, complete sequence
3570487
HE580267.1 *Naumovozya dairenensis* CBS 421 chromosome 1, complete genome
2054635
HE580268.1 *Naumovozya dairenensis* CBS 421 chromosome 2, complete genome
1549475
HE580269.1 *Naumovozya dairenensis* CBS 421 chromosome 3, complete genome
1536206
HE580270.1 *Naumovozya dairenensis* CBS 421 chromosome 4, complete genome
1230053
HE580271.1 *Naumovozya dairenensis* CBS 421 chromosome 5, complete genome
1184459

HE580272.1 *Naumovozya dairenensis* CBS 421 chromosome 6, complete genome
1138463
HE580273.2 *Naumovozya dairenensis* CBS 421 chromosome 7, complete genome
1565278
HE580274.1 *Naumovozya dairenensis* CBS 421 chromosome 8, complete genome
1004037
HE580275.1 *Naumovozya dairenensis* CBS 421 chromosome 9, complete genome
786484
HE580276.1 *Naumovozya dairenensis* CBS 421 chromosome 10, complete genome
782356
HE580277.1 *Naumovozya dairenensis* CBS 421 chromosome 11, complete genome
696134
FQ311430.1 *Sporisorium reilianum* SRZ2 chromosome 1 complete DNA sequence
2448206
FQ311441.1 *Sporisorium reilianum* SRZ2 chromosome 2 complete DNA sequence
1791267
FQ311452.1 *Sporisorium reilianum* SRZ2 chromosome 3 complete DNA sequence
1534569
FQ311463.1 *Sporisorium reilianum* SRZ2 chromosome 4 complete DNA sequence
772363
FQ311470.1 *Sporisorium reilianum* SRZ2 chromosome 5 complete DNA sequence
730625
FQ311471.1 *Sporisorium reilianum* SRZ2 chromosome 6 complete DNA sequence
976264
FQ311472.1 *Sporisorium reilianum* SRZ2 chromosome 7 complete DNA sequence
880029
FQ311473.1 *Sporisorium reilianum* SRZ2 chromosome 8 complete DNA sequence
760034
FQ311474.1 *Sporisorium reilianum* SRZ2 chromosome 9 complete DNA sequence
670125
FQ311431.1 *Sporisorium reilianum* SRZ2 chromosome 10 complete DNA sequence
642761
FQ311432.1 *Sporisorium reilianum* SRZ2 chromosome 11 complete DNA sequence
636629
FQ311433.1 *Sporisorium reilianum* SRZ2 chromosome 12 complete DNA sequence
565009
FQ311434.1 *Sporisorium reilianum* SRZ2 chromosome 13 complete DNA sequence
555666
FQ311435.1 *Sporisorium reilianum* SRZ2 chromosome 14 complete DNA sequence
562274
FQ311436.1 *Sporisorium reilianum* SRZ2 chromosome 15 complete DNA sequence
562189
FQ311437.1 *Sporisorium reilianum* SRZ2 chromosome 16 complete DNA sequence
478111
FQ311438.1 *Sporisorium reilianum* SRZ2 chromosome 17 complete DNA sequence
534508
FQ311439.1 *Sporisorium reilianum* SRZ2 chromosome 18 complete DNA sequence
518890
FQ311440.1 *Sporisorium reilianum* SRZ2 chromosome 19 complete DNA sequence
521053
FQ311442.1 *Sporisorium reilianum* SRZ2 chromosome 20 complete DNA sequence
1115737
FQ311443.1 *Sporisorium reilianum* SRZ2 chromosome 21 complete DNA sequence
429375
FQ311444.1 *Sporisorium reilianum* SRZ2 chromosome 22 complete DNA sequence
385084
FQ311445.1 *Sporisorium reilianum* SRZ2 chromosome 23 complete DNA sequence
263978
FQ311446.1 *Sporisorium reilianum* SRZ2 contig_5.00010 complete DNA sequence
1151

FQ311447.1 Sporisorium reilianum SRZ2 contig_5.00013 complete DNA sequence
1431
FQ311448.1 Sporisorium reilianum SRZ2 contig_5.00023 complete DNA sequence
1758
FQ311449.1 Sporisorium reilianum SRZ2 contig_5.00029 complete DNA sequence
1581
FQ311450.1 Sporisorium reilianum SRZ2 contig_5.00032 complete DNA sequence
1796
FQ311451.1 Sporisorium reilianum SRZ2 contig_5.00045 complete DNA sequence
1309
FQ311453.1 Sporisorium reilianum SRZ2 contig_5.00046 complete DNA sequence
721
FQ311454.1 Sporisorium reilianum SRZ2 contig_5.00050 complete DNA sequence
2378
FQ311455.1 Sporisorium reilianum SRZ2 contig_5.00055 complete DNA sequence
4673
FQ311456.1 Sporisorium reilianum SRZ2 contig_5.00068 complete DNA sequence
1537
FQ311457.1 Sporisorium reilianum SRZ2 contig_5.00069 complete DNA sequence
864
FQ311458.1 Sporisorium reilianum SRZ2 contig_5.00075 complete DNA sequence
19032
FQ311459.1 Sporisorium reilianum SRZ2 contig_5.00080 complete DNA sequence
1970
FQ311460.1 Sporisorium reilianum SRZ2 contig_5.00081 complete DNA sequence
1618
FQ311461.1 Sporisorium reilianum SRZ2 contig_5.00088 complete DNA sequence
1446
FQ311462.1 Sporisorium reilianum SRZ2 sr_contig_5.00092 complete DNA sequence
759
FQ311464.1 Sporisorium reilianum SRZ2 sr_contig_5.00094 complete DNA sequence
796
FQ311465.1 Sporisorium reilianum SRZ2 contig_5.00095 complete DNA sequence
776
FQ311466.1 Sporisorium reilianum SRZ2 contig_5.00096 complete DNA sequence
3131
FQ311467.1 Sporisorium reilianum SRZ2 contig_5.00104 complete DNA sequence
1823
FQ311468.1 Sporisorium reilianum SRZ2 contig_5.00115 complete DNA sequence
1082
CP002497.1 Eremothecium cymbalariae DBVPG#7215 chromosome 1, complete sequence
1110245
CP002498.1 Eremothecium cymbalariae DBVPG#7215 chromosome 2, complete sequence
1601921
CP002499.1 Eremothecium cymbalariae DBVPG#7215 chromosome 3, complete sequence
1193613
CP002500.1 Eremothecium cymbalariae DBVPG#7215 chromosome 4, complete sequence
1540972
CP002501.1 Eremothecium cymbalariae DBVPG#7215 chromosome 5, complete sequence
1385851
CP002502.1 Eremothecium cymbalariae DBVPG#7215 chromosome 6, complete sequence
959278
CP002503.1 Eremothecium cymbalariae DBVPG#7215 chromosome 7, complete sequence
980088
CP002504.1 Eremothecium cymbalariae DBVPG#7215 chromosome 8, complete sequence
897456
HE612856.1 Tetrapisispora phaffii CBS 4417 chromosome 1, complete genome
1377933
HE612857.1 Tetrapisispora phaffii CBS 4417 chromosome 2, complete genome
1164785

HE612858.1 *Tetrapisispora phaffii* CBS 4417 chromosome 3, complete genome
1096211
HE612859.1 *Tetrapisispora phaffii* CBS 4417 chromosome 4, complete genome
1042069
HE612860.1 *Tetrapisispora phaffii* CBS 4417 chromosome 5, complete genome
854350
HE612861.1 *Tetrapisispora phaffii* CBS 4417 chromosome 6, complete genome
815984
HE612862.1 *Tetrapisispora phaffii* CBS 4417 chromosome 7, complete genome
808783
HE612863.1 *Tetrapisispora phaffii* CBS 4417 chromosome 8, complete genome
752451
HE612864.1 *Tetrapisispora phaffii* CBS 4417 chromosome 9, complete genome
740259
HE612865.1 *Tetrapisispora phaffii* CBS 4417 chromosome 10, complete genome
713392
HE612866.1 *Tetrapisispora phaffii* CBS 4417 chromosome 11, complete genome
536385
HE612867.1 *Tetrapisispora phaffii* CBS 4417 chromosome 12, complete genome
504740
HE612868.1 *Tetrapisispora phaffii* CBS 4417 chromosome 13, complete genome
452480
HE612869.1 *Tetrapisispora phaffii* CBS 4417 chromosome 14, complete genome
432466
HE612870.1 *Tetrapisispora phaffii* CBS 4417 chromosome 15, complete genome
413379
HE612871.1 *Tetrapisispora phaffii* CBS 4417 chromosome 16, complete genome
394523
HE576752.1 *Naumovozya castellii* CBS 4309 chromosome 1, complete genome
3019861
HE576753.1 *Naumovozya castellii* CBS 4309 chromosome 2, complete genome
1756361
HE576754.1 *Naumovozya castellii* CBS 4309 chromosome 3, complete genome
1245273
HE576755.1 *Naumovozya castellii* CBS 4309 chromosome 4, complete genome
958728
HE576756.1 *Naumovozya castellii* CBS 4309 chromosome 5, complete genome
835708
HE576757.1 *Naumovozya castellii* CBS 4309 chromosome 6, complete genome
828437
HE576758.1 *Naumovozya castellii* CBS 4309 chromosome 7, complete genome
804126
HE576759.1 *Naumovozya castellii* CBS 4309 chromosome 8, complete genome
699326
HE576760.1 *Naumovozya castellii* CBS 4309 chromosome 9, complete genome
603245
HE576761.1 *Naumovozya castellii* CBS 4309 chromosome 10, complete genome
468474
HE616742.1 *Torulaspora delbrueckii* CBS 1146 chromosome 1, complete genome
1418021
HE616743.1 *Torulaspora delbrueckii* CBS 1146 chromosome 2, complete genome
1354397
HE616744.1 *Torulaspora delbrueckii* CBS 1146 chromosome 3, complete genome
1292049
HE616745.1 *Torulaspora delbrueckii* CBS 1146 chromosome 4, complete genome
1218070
HE616746.1 *Torulaspora delbrueckii* CBS 1146 chromosome 5, complete genome
1086982
HE616747.1 *Torulaspora delbrueckii* CBS 1146 chromosome 6, complete genome
1070623

HE616748.1 *Torulaspora delbrueckii* CBS 1146 chromosome 7, complete genome
946563
HE616749.1 *Torulaspora delbrueckii* CBS 1146 chromosome 8, complete genome
833973
CP002713.1 *Encephalitozoon hellem* ATCC 50504 chromosome I, complete sequence
161584
CP002714.1 *Encephalitozoon hellem* ATCC 50504 chromosome II, complete sequence
179949
CP002715.1 *Encephalitozoon hellem* ATCC 50504 chromosome III, complete sequence
182562
CP002716.1 *Encephalitozoon hellem* ATCC 50504 chromosome IV, complete sequence
196397
CP002717.1 *Encephalitozoon hellem* ATCC 50504 chromosome V, complete sequence
201478
CP002718.1 *Encephalitozoon hellem* ATCC 50504 chromosome VI, complete sequence
200132
CP002719.1 *Encephalitozoon hellem* ATCC 50504 chromosome VII, complete sequence
213233
CP002720.1 *Encephalitozoon hellem* ATCC 50504 chromosome VIII, complete sequence
203699
CP002721.1 *Encephalitozoon hellem* ATCC 50504 chromosome IXa, complete sequence
167643
CP002722.1 *Encephalitozoon hellem* ATCC 50504 chromosome IXb, complete sequence
66626
CP002723.1 *Encephalitozoon hellem* ATCC 50504 chromosome X, complete sequence
240865
CP002724.1 *Encephalitozoon hellem* ATCC 50504 chromosome XI, complete sequence
237616
CP003518.1 *Encephalitozoon romaleae* SJ-2008 chromosome I, complete sequence
146158
CP003519.1 *Encephalitozoon romaleae* SJ-2008 chromosome II, complete sequence
173763
CP003520.1 *Encephalitozoon romaleae* SJ-2008 chromosome III, complete sequence
179400
CP003521.1 *Encephalitozoon romaleae* SJ-2008 chromosome IV, complete sequence
197308
CP003522.1 *Encephalitozoon romaleae* SJ-2008 chromosome V, complete sequence
191809
CP003523.1 *Encephalitozoon romaleae* SJ-2008 chromosome VI, complete sequence
194799
CP003524.1 *Encephalitozoon romaleae* SJ-2008 chromosome VII, complete sequence
204533
CP003525.1 *Encephalitozoon romaleae* SJ-2008 chromosome VIII, complete sequence
207682
CP003529.1 *Encephalitozoon romaleae* SJ-2008 chromosome X, complete sequence
230122
CP003530.1 *Encephalitozoon romaleae* SJ-2008 chromosome XI, complete sequence
231554
CP015461.1 *Agaricus bisporus* var. *bisporus* H97 chromosome 5, complete sequence
2550681
CP015462.1 *Agaricus bisporus* var. *bisporus* H97 chromosome 6, complete sequence
2329815
HE650821.1 *Kazachstania africana* CBS 2517 chromosome 1, complete genome
1757399
HE650822.1 *Kazachstania africana* CBS 2517 chromosome 2, complete genome
1485491
HE650823.1 *Kazachstania africana* CBS 2517 chromosome 3, complete genome
1340033
HE650824.1 *Kazachstania africana* CBS 2517 chromosome 4, complete genome
1026673

HE650825.1 *Kazachstania africana* CBS 2517 chromosome 5, complete genome
925516
HE650826.1 *Kazachstania africana* CBS 2517 chromosome 6, complete genome
888590
HE650827.1 *Kazachstania africana* CBS 2517 chromosome 7, complete genome
812636
HE650828.1 *Kazachstania africana* CBS 2517 chromosome 8, complete genome
738805
HE650829.1 *Kazachstania africana* CBS 2517 chromosome 9, complete genome
607727
HE650830.1 *Kazachstania africana* CBS 2517 chromosome 10, complete genome
578118
HE650831.1 *Kazachstania africana* CBS 2517 chromosome 11, complete genome
547686
HE650832.1 *Kazachstania africana* CBS 2517 chromosome 12, complete genome
421466
FO082059.1 *Millerozyma farinosa* CBS 7064 chromosome A complete sequence
1055225
FO082058.1 *Millerozyma farinosa* CBS 7064 chromosome B complete sequence
1063002
FO082057.1 *Pichia sorbitophila* strain CBS 7064 chromosome C complete sequence
1095816
FO082056.1 *Pichia sorbitophila* strain CBS 7064 chromosome D complete sequence
1102865
FO082055.1 *Pichia sorbitophila* strain CBS 7064 chromosome E complete sequence
1307235
FO082054.1 *Pichia sorbitophila* strain CBS 7064 chromosome F complete sequence
1407903
FO082053.1 *Pichia sorbitophila* strain CBS 7064 chromosome G complete sequence
1423303
FO082052.1 *Pichia sorbitophila* strain CBS 7064 chromosome H complete sequence
1423303
FO082051.1 *Pichia sorbitophila* strain CBS 7064 chromosome I complete sequence
1666063
FO082050.1 *Pichia sorbitophila* strain CBS 7064 chromosome J complete sequence
1803284
FO082049.1 *Pichia sorbitophila* strain CBS 7064 chromosome K complete sequence
1933849
FO082048.1 *Pichia sorbitophila* strain CBS 7064 chromosome L complete sequence
1933849
FO082047.1 *Pichia sorbitophila* strain CBS 7064 chromosome M complete sequence
2121241
FO082046.1 *Pichia sorbitophila* strain CBS 7064 chromosome N complete sequence
2122704
HE806316.1 *Tetrapisispora blattae* CBS 6284 chromosome 1, complete genome
2697332
HE806317.1 *Tetrapisispora blattae* CBS 6284 chromosome 2, complete genome
2450908
HE806318.1 *Tetrapisispora blattae* CBS 6284 chromosome 3, complete genome
1779678
HE806319.1 *Tetrapisispora blattae* CBS 6284 chromosome 4, complete genome
1449145
HE806320.1 *Tetrapisispora blattae* CBS 6284 chromosome 5, complete genome
1324051
HE806321.1 *Tetrapisispora blattae* CBS 6284 chromosome 6, complete genome
1053862
HE806322.1 *Tetrapisispora blattae* CBS 6284 chromosome 7, complete genome
986391
HE806323.1 *Tetrapisispora blattae* CBS 6284 chromosome 8, complete genome
956377

HE806324.1 *Tetrapisispora blattae* CBS 6284 chromosome 9, complete genome
857919
HE806325.1 *Tetrapisispora blattae* CBS 6284 chromosome 10, complete genome
492930
HE978314.1 *Kazachstania naganishii* CBS 8797 chromosome 1, complete genome
1290777
HE978315.1 *Kazachstania naganishii* CBS 8797 chromosome 2, complete genome
1410756
HE978316.1 *Kazachstania naganishii* CBS 8797 chromosome 3, complete genome
1310278
HE978317.1 *Kazachstania naganishii* CBS 8797 chromosome 4, complete genome
993246
HE978318.1 *Kazachstania naganishii* CBS 8797 chromosome 5, complete genome
856010
HE978319.1 *Kazachstania naganishii* CBS 8797 chromosome 6, complete genome
770551
HE978320.1 *Kazachstania naganishii* CBS 8797 chromosome 7, complete genome
780220
HE978321.1 *Kazachstania naganishii* CBS 8797 chromosome 8, complete genome
724298
HE978322.1 *Kazachstania naganishii* CBS 8797 chromosome 9, complete genome
628580
HE978323.1 *Kazachstania naganishii* CBS 8797 chromosome 10, complete genome
586734
HE978324.1 *Kazachstania naganishii* CBS 8797 chromosome 11, complete genome
538562
HE978325.1 *Kazachstania naganishii* CBS 8797 chromosome 12, complete genome
450952
HE978326.1 *Kazachstania naganishii* CBS 8797 chromosome 13, complete genome
504857
CP009075.1 *Verticillium dahliae* JR2 chromosome 1, complete sequence
9275483
CP009079.1 *Verticillium dahliae* JR2 chromosome 2, complete sequence
4277765
CP009077.1 *Verticillium dahliae* JR2 chromosome 3, complete sequence
4168633
CP009080.1 *Verticillium dahliae* JR2 chromosome 4, complete sequence
4086908
CP009078.1 *Verticillium dahliae* JR2 chromosome 5, complete sequence
4171808
CP009076.1 *Verticillium dahliae* JR2 chromosome 6, complete sequence
3530890
CP009082.1 *Verticillium dahliae* JR2 chromosome 7, complete sequence
3277570
CP009081.1 *Verticillium dahliae* JR2 chromosome 8, complete sequence
3361230
CP006020.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome I, complete sequence
809179
CP006021.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome II, complete sequence
1019532
CP006022.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome III, complete sequence
1493473
CP006023.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome IV, complete sequence
1099572
CP006024.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome V, complete sequence
1513917
CP006025.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome VI, complete sequence
1334226
CP006026.1 *Saccharomycetaceae* sp. '*Ashbya aceri*' chromosome VII, complete sequence
1597628

CP006258.1 *Saccharomycetaceae* sp. 'Ashbya aceri' mitochondrion, complete genome
26996
HG934059.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOA
2210998
HG934060.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOB
3023969
HG934061.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOC
3189950
HG934062.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOD
3555057
HG934063.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOE
4142257
HG934064.1 *Yarrowia lipolytica* WSH-Z06 complete genome, chromosome YALIOF
3968352
CP004528.1 *Saccharomyces cerevisiae* YJM993 plasmid 2 micron, complete sequence
6303
CP006506.1 *Saccharomyces cerevisiae* YJM993 mitochondrion, complete genome
78907
CM002802.1 *Penicillium chrysogenum* strain P2niaD18 mitochondrion, complete sequence, whole
genome shotgun sequence
27017
CP010980.1 *Verticillium dahliae* VdLs.17 chromosome 1, complete sequence
5989981
CP010981.1 *Verticillium dahliae* VdLs.17 chromosome 2, complete sequence
6210300
CP010982.1 *Verticillium dahliae* VdLs.17 chromosome 3, complete sequence
5894008
CP010983.1 *Verticillium dahliae* VdLs.17 chromosome 4, complete sequence
4331027
CP010984.1 *Verticillium dahliae* VdLs.17 chromosome 5, complete sequence
3581864
CP010985.1 *Verticillium dahliae* VdLs.17 chromosome 6, complete sequence
3402923
CP010986.1 *Verticillium dahliae* VdLs.17 chromosome 7, complete sequence
3272870
CP010987.1 *Verticillium dahliae* VdLs.17 chromosome 8, complete sequence
3290897
CP002705.1 *Ashbya gossypii* FDAG1 chromosome I, complete sequence
693310
CP002706.1 *Ashbya gossypii* FDAG1 chromosome II, complete sequence
874947
CP002707.1 *Ashbya gossypii* FDAG1 chromosome III, complete sequence
911446
CP002708.1 *Ashbya gossypii* FDAG1 chromosome IV, complete sequence
1465717
CP002709.1 *Ashbya gossypii* FDAG1 chromosome V, complete sequence
1527143
CP002710.1 *Ashbya gossypii* FDAG1 chromosome VI, complete sequence
1837132
CP002711.1 *Ashbya gossypii* FDAG1 chromosome VII, complete sequence
1823962
CP002712.1 *Ashbya gossypii* FDAG1 mitochondrion, complete sequence
23497
CP006476.1 *Saccharomyces cerevisiae* YJM195 mitochondrion, complete genome
78961
CP004504.1 *Saccharomyces cerevisiae* YJM244 plasmid 2 micron, complete sequence
6303
CP006477.1 *Saccharomyces cerevisiae* YJM244 mitochondrion, complete genome
80737
CP004116.1 *Saccharomyces cerevisiae* YJM1078 plasmid 2micron, complete sequence

6303
CP004115.1 *Saccharomyces cerevisiae* YJM1078 mitochondrion, complete genome
82025
CP004530.1 *Saccharomyces cerevisiae* YJM1083 plasmid 2 micron, complete sequence
6118
CP006508.1 *Saccharomyces cerevisiae* YJM1083 mitochondrion, complete genome
89444
CP004531.1 *Saccharomyces cerevisiae* YJM1129 plasmid 2 micron, complete sequence
6178
CP006509.1 *Saccharomyces cerevisiae* YJM1129 mitochondrion, complete genome
81706
CP004502.1 *Saccharomyces cerevisiae* YJM189 plasmid 2 micron, complete sequence
6178
CP006474.1 *Saccharomyces cerevisiae* YJM189 mitochondrion, complete genome
80811
CP004503.1 *Saccharomyces cerevisiae* YJM193 plasmid 2 micron, complete sequence
6303
CP006475.1 *Saccharomyces cerevisiae* YJM193 mitochondrion, complete genome
84644
CP004505.1 *Saccharomyces cerevisiae* YJM248 plasmid 2 micron, complete sequence
6178
CP006478.1 *Saccharomyces cerevisiae* YJM248 mitochondrion, complete genome
86593
CP006479.1 *Saccharomyces cerevisiae* YJM270 mitochondrion, complete genome
82331
CP004506.1 *Saccharomyces cerevisiae* YJM271 plasmid 2 micron, complete sequence
6178
CP006480.1 *Saccharomyces cerevisiae* YJM271 mitochondrion, complete genome
84530
CP004507.1 *Saccharomyces cerevisiae* YJM320 plasmid 2 micron, complete sequence
6118
CP006481.1 *Saccharomyces cerevisiae* YJM320 mitochondrion, complete genome
86949
CP004508.1 *Saccharomyces cerevisiae* YJM326 plasmid 2 micron, complete sequence
6119
CP006482.1 *Saccharomyces cerevisiae* YJM326 mitochondrion, complete genome
91172
CP004509.1 *Saccharomyces cerevisiae* YJM428 plasmid 2 micron, complete sequence
6118
CP006483.1 *Saccharomyces cerevisiae* YJM428 mitochondrion, complete genome
85782
CP004510.1 *Saccharomyces cerevisiae* YJM450 plasmid 2 micron, complete sequence
6178
CP006484.1 *Saccharomyces cerevisiae* YJM450 mitochondrion, complete genome
86522
CP004511.1 *Saccharomyces cerevisiae* YJM451 plasmid 2 micron, complete sequence
6106
CP006485.1 *Saccharomyces cerevisiae* YJM451 mitochondrion, complete genome
84234
CP004512.1 *Saccharomyces cerevisiae* YJM453 plasmid 2 micron, complete sequence
6178
CP006486.1 *Saccharomyces cerevisiae* YJM453 mitochondrion, complete genome
82494
CP004513.1 *Saccharomyces cerevisiae* YJM456 plasmid 2 micron, complete sequence
6347
CP006487.1 *Saccharomyces cerevisiae* YJM456 mitochondrion, complete genome
79631
CP006488.1 *Saccharomyces cerevisiae* YJM470 mitochondrion, complete genome
83571
CP004514.1 *Saccharomyces cerevisiae* YJM541 plasmid 2 micron, complete sequence

6118
CP006489.1 *Saccharomyces cerevisiae* YJM541 mitochondrion, complete genome
87003
CP004515.1 *Saccharomyces cerevisiae* YJM554 plasmid 2 micron, complete sequence
6118
CP006490.1 *Saccharomyces cerevisiae* YJM554 mitochondrion, complete genome
87008
CP004516.1 *Saccharomyces cerevisiae* YJM555 plasmid 2 micron, complete sequence
6117
CP006491.1 *Saccharomyces cerevisiae* YJM555 mitochondrion, complete genome
84193
CP004517.1 *Saccharomyces cerevisiae* YJM627 plasmid 2 micron, complete sequence
6303
CP006492.1 *Saccharomyces cerevisiae* YJM627 mitochondrion, complete genome
80791
CP004518.1 *Saccharomyces cerevisiae* YJM681 plasmid 2 micron, complete sequence
6118
CP006493.1 *Saccharomyces cerevisiae* YJM681 mitochondrion, complete genome
81045
CP004519.1 *Saccharomyces cerevisiae* YJM682 plasmid 2 micron, complete sequence
6119
CP006494.1 *Saccharomyces cerevisiae* YJM682 mitochondrion, complete genome
88887
CP006495.1 *Saccharomyces cerevisiae* YJM683 mitochondrion, complete genome
88953
CP006496.1 *Saccharomyces cerevisiae* YJM689 mitochondrion, complete genome
84098
CP006497.1 *Saccharomyces cerevisiae* YJM693 mitochondrion, complete genome
89161
CP004520.1 *Saccharomyces cerevisiae* YJM969 plasmid 2 micron, complete sequence
6178
CP006498.1 *Saccharomyces cerevisiae* YJM969 mitochondrion, complete genome
79066
CP004521.1 *Saccharomyces cerevisiae* YJM972 plasmid 2 micron, complete sequence
6303
CP006499.1 *Saccharomyces cerevisiae* YJM972 mitochondrion, complete genome
78907
CP004522.1 *Saccharomyces cerevisiae* YJM975 plasmid 2 micron, complete sequence
6303
CP006500.1 *Saccharomyces cerevisiae* YJM975 mitochondrion, complete genome
78950
CP004523.1 *Saccharomyces cerevisiae* YJM978 plasmid 2 micron, complete sequence
6303
CP006501.1 *Saccharomyces cerevisiae* YJM978 mitochondrion, complete genome
78916
CP004524.1 *Saccharomyces cerevisiae* YJM981 plasmid 2 micron, complete sequence
6303
CP006502.1 *Saccharomyces cerevisiae* YJM981 mitochondrion, complete genome
78916
CP004525.1 *Saccharomyces cerevisiae* YJM984 plasmid 2 micron, complete sequence
6303
CP006503.1 *Saccharomyces cerevisiae* YJM984 mitochondrion, complete genome
78907
CP004526.1 *Saccharomyces cerevisiae* YJM987 plasmid 2 micron, complete sequence
6303
CP006504.1 *Saccharomyces cerevisiae* YJM987 mitochondrion, complete genome
78974
CP004527.1 *Saccharomyces cerevisiae* YJM990 plasmid 2 micron, complete sequence
6303
CP006505.1 *Saccharomyces cerevisiae* YJM990 mitochondrion, complete genome

78916
CP004529.1 *Saccharomyces cerevisiae* YJM996 plasmid 2 micron, complete sequence
6303
CP006507.1 *Saccharomyces cerevisiae* YJM996 mitochondrion, complete genome
78907
CP004532.1 *Saccharomyces cerevisiae* YJM1133 plasmid 2 micron, complete sequence
6118
CP006510.1 *Saccharomyces cerevisiae* YJM1133 mitochondrion, complete genome
84963
CP004533.1 *Saccharomyces cerevisiae* YJM1190 plasmid 2 micron, complete sequence
6118
CP006511.2 *Saccharomyces cerevisiae* YJM1190 mitochondrion, complete genome
88996
CP004534.1 *Saccharomyces cerevisiae* YJM1199 plasmid 2 micron, complete sequence
6118
CP006512.1 *Saccharomyces cerevisiae* YJM1199 mitochondrion, complete genome
90633
CP004535.1 *Saccharomyces cerevisiae* YJM1202 plasmid 2 micron, complete sequence
6118
CP006513.1 *Saccharomyces cerevisiae* YJM1202 mitochondrion, complete genome
90638
CP004536.1 *Saccharomyces cerevisiae* YJM1208 plasmid 2 micron, complete sequence
6118
CP006514.1 *Saccharomyces cerevisiae* YJM1208 mitochondrion, complete genome
84252
CP004537.1 *Saccharomyces cerevisiae* YJM1242 plasmid 2 micron, complete sequence
6303
CP006515.1 *Saccharomyces cerevisiae* YJM1242 mitochondrion, complete genome
95658
CP004538.1 *Saccharomyces cerevisiae* YJM1244 plasmid 2 micron, complete sequence
6303
CP006516.1 *Saccharomyces cerevisiae* YJM1244 mitochondrion, complete genome
83285
CP006517.1 *Saccharomyces cerevisiae* YJM1248 mitochondrion, complete genome
81013
CP004539.1 *Saccharomyces cerevisiae* YJM1250 plasmid 2 micron, complete sequence
6300
CP006518.1 *Saccharomyces cerevisiae* YJM1250 mitochondrion, complete genome
88746
CP006519.1 *Saccharomyces cerevisiae* YJM1252 mitochondrion, complete genome
83856
CP006520.1 *Saccharomyces cerevisiae* YJM1273 mitochondrion, complete genome
77475
CP004540.1 *Saccharomyces cerevisiae* YJM1304 plasmid 2 micron, complete sequence
6166
CP006521.1 *Saccharomyces cerevisiae* YJM1304 mitochondrion, complete genome
81083
CP004541.1 *Saccharomyces cerevisiae* YJM1307 plasmid 2 micron, complete sequence
6178
CP006522.1 *Saccharomyces cerevisiae* YJM1307 mitochondrion, complete genome
79733
CP004542.1 *Saccharomyces cerevisiae* YJM1311 plasmid 2 micron, complete sequence
6118
CP006523.1 *Saccharomyces cerevisiae* YJM1311 mitochondrion, complete genome
87022
CP004543.1 *Saccharomyces cerevisiae* YJM1326 plasmid 2 micron, complete sequence
6178
CP006524.1 *Saccharomyces cerevisiae* YJM1326 mitochondrion, complete genome
85887
CP004544.1 *Saccharomyces cerevisiae* YJM1332 plasmid 2 micron, complete sequence

6303
CP006525.1 *Saccharomyces cerevisiae* YJM1332 mitochondrion, complete genome
81501
CP004545.1 *Saccharomyces cerevisiae* YJM1336 plasmid 2 micron, complete sequence
6178
CP006526.1 *Saccharomyces cerevisiae* YJM1336 mitochondrion, complete genome
82967
CP004546.1 *Saccharomyces cerevisiae* YJM1338 plasmid 2 micron, complete sequence
6303
CP006527.1 *Saccharomyces cerevisiae* YJM1338 mitochondrion, complete genome
80970
CP004547.1 *Saccharomyces cerevisiae* YJM1341 plasmid 2 micron, complete sequence
6178
CP006528.1 *Saccharomyces cerevisiae* YJM1341 mitochondrion, complete genome
84119
CP006529.1 *Saccharomyces cerevisiae* YJM1342 mitochondrion, complete genome
85762
CP004548.1 *Saccharomyces cerevisiae* YJM1355 plasmid 2 micron, complete sequence
6117
CP006530.1 *Saccharomyces cerevisiae* YJM1355 mitochondrion, complete genome
79418
CP004549.1 *Saccharomyces cerevisiae* YJM1356 plasmid 2 micron, complete sequence
6303
CP006531.1 *Saccharomyces cerevisiae* YJM1356 mitochondrion, complete genome
84849
CP004550.1 *Saccharomyces cerevisiae* YJM1381 plasmid 2 micron, complete sequence
6119
CP006532.1 *Saccharomyces cerevisiae* YJM1381 mitochondrion, complete genome
83874
CP004551.1 *Saccharomyces cerevisiae* YJM1383 plasmid 2 micron, complete sequence
6194
CP006533.1 *Saccharomyces cerevisiae* YJM1383 mitochondrion, complete genome
86820
CP006534.1 *Saccharomyces cerevisiae* YJM1385 mitochondrion, complete genome
92176
CP004552.1 *Saccharomyces cerevisiae* YJM1386 plasmid 2 micron, complete sequence
6118
CP006535.1 *Saccharomyces cerevisiae* YJM1386 mitochondrion, complete genome
85554
CP006536.1 *Saccharomyces cerevisiae* YJM1387 mitochondrion, complete genome
80666
CP006537.1 *Saccharomyces cerevisiae* YJM1388 mitochondrion, complete genome
77572
CP006538.1 *Saccharomyces cerevisiae* YJM1389 mitochondrion, complete genome
80794
CP006539.1 *Saccharomyces cerevisiae* YJM1399 mitochondrion, complete genome
73450
CP004553.1 *Saccharomyces cerevisiae* YJM1400 plasmid 2 micron, complete sequence
6094
CP006540.1 *Saccharomyces cerevisiae* YJM1400 mitochondrion, complete genome
88306
CP004554.1 *Saccharomyces cerevisiae* YJM1401 plasmid 2 micron, complete sequence
6095
CP006541.1 *Saccharomyces cerevisiae* YJM1401 mitochondrion, complete genome
81306
CP006542.1 *Saccharomyces cerevisiae* YJM1402 mitochondrion, complete genome
78309
CP004555.1 *Saccharomyces cerevisiae* YJM1415 plasmid 2 micron, complete sequence
6178
CP006543.1 *Saccharomyces cerevisiae* YJM1415 mitochondrion, complete genome

83950
CP004556.1 *Saccharomyces cerevisiae* YJM1417 plasmid 2 micron, complete sequence
6303
CP006544.1 *Saccharomyces cerevisiae* YJM1417 mitochondrion, complete genome
82284
CP006545.1 *Saccharomyces cerevisiae* YJM1418 mitochondrion, complete genome
76794
CP004557.1 *Saccharomyces cerevisiae* YJM1419 plasmid 2 micron, complete sequence
6118
CP006546.1 *Saccharomyces cerevisiae* YJM1419 mitochondrion, complete genome
84604
CP004558.1 *Saccharomyces cerevisiae* YJM1433 plasmid 2 micron, complete sequence
6346
CP006547.1 *Saccharomyces cerevisiae* YJM1433 mitochondrion, complete genome
81976
CP006548.1 *Saccharomyces cerevisiae* YJM1434 mitochondrion, complete genome
77515
CP006549.1 *Saccharomyces cerevisiae* YJM1439 mitochondrion, complete genome
80882
CP006550.1 *Saccharomyces cerevisiae* YJM1443 mitochondrion, complete genome
77956
CP006551.1 *Saccharomyces cerevisiae* YJM1444 mitochondrion, complete genome
83736
CP006552.1 *Saccharomyces cerevisiae* YJM1447 mitochondrion, complete genome
74237
CP004559.1 *Saccharomyces cerevisiae* YJM1450 plasmid 2 micron, complete sequence
6162
CP006553.1 *Saccharomyces cerevisiae* YJM1450 mitochondrion, complete genome
76603
CP006554.1 *Saccharomyces cerevisiae* YJM1460 mitochondrion, complete genome
82773
CP004560.1 *Saccharomyces cerevisiae* YJM1463 plasmid 2 micron, complete sequence
6179
CP006555.1 *Saccharomyces cerevisiae* YJM1463 mitochondrion, complete genome
79676
CP004561.1 *Saccharomyces cerevisiae* YJM1477 plasmid 2 micron, complete sequence
6303
CP006556.1 *Saccharomyces cerevisiae* YJM1477 mitochondrion, complete genome
79244
CP004562.1 *Saccharomyces cerevisiae* YJM1478 plasmid 2 micron, complete sequence
6303
CP006557.1 *Saccharomyces cerevisiae* YJM1478 mitochondrion, complete genome
83765
CP004563.1 *Saccharomyces cerevisiae* YJM1479 plasmid 2 micron, complete sequence
6094
CP006558.1 *Saccharomyces cerevisiae* YJM1479 mitochondrion, complete genome
88309
CP004564.1 *Saccharomyces cerevisiae* YJM1526 plasmid 2 micron, complete sequence
6178
CP006559.1 *Saccharomyces cerevisiae* YJM1526 mitochondrion, complete genome
81825
CP004565.1 *Saccharomyces cerevisiae* YJM1527 plasmid 2 micron, complete sequence
6303
CP006560.1 *Saccharomyces cerevisiae* YJM1527 mitochondrion, complete genome
78592
CP006561.1 *Saccharomyces cerevisiae* YJM1549 mitochondrion, complete genome
78275
CP006562.1 *Saccharomyces cerevisiae* YJM1573 mitochondrion, complete genome
78385
CP004566.1 *Saccharomyces cerevisiae* YJM1574 plasmid 2 micron, complete sequence

6178
CP006563.1 *Saccharomyces cerevisiae* YJM1574 mitochondrion, complete genome
79923
CP006564.1 *Saccharomyces cerevisiae* YJM1592 mitochondrion, complete genome
80788
CP004567.1 *Saccharomyces cerevisiae* YJM1615 plasmid 2 micron, complete sequence
6118
CP006565.1 *Saccharomyces cerevisiae* YJM1615 mitochondrion, complete genome
84325
CP010913.1 *Sporisorium scitamineum* strain SSC39 chromosome 1, complete sequence
2009762
CP010914.1 *Sporisorium scitamineum* strain SSC39 chromosome 2, complete sequence
1671606
CP010915.1 *Sporisorium scitamineum* strain SSC39 chromosome 3, complete sequence
1566833
CP010916.1 *Sporisorium scitamineum* strain SSC39 chromosome 4, complete sequence
1271980
CP010917.1 *Sporisorium scitamineum* strain SSC39 chromosome 5, complete sequence
1055339
CP010918.1 *Sporisorium scitamineum* strain SSC39 chromosome 6, complete sequence
1032954
CP010919.1 *Sporisorium scitamineum* strain SSC39 chromosome 7, complete sequence
938453
CP010920.1 *Sporisorium scitamineum* strain SSC39 chromosome 8, complete sequence
837966
CP010921.1 *Sporisorium scitamineum* strain SSC39 chromosome 9, complete sequence
808176
CP010922.1 *Sporisorium scitamineum* strain SSC39 chromosome 10, complete sequence
875830
CP010923.1 *Sporisorium scitamineum* strain SSC39 chromosome 11, complete sequence
790822
CP010924.1 *Sporisorium scitamineum* strain SSC39 chromosome 12, complete sequence
716888
CP010925.1 *Sporisorium scitamineum* strain SSC39 chromosome 13, complete sequence
681008
CP010926.1 *Sporisorium scitamineum* strain SSC39 chromosome 14, complete sequence
652314
CP010927.1 *Sporisorium scitamineum* strain SSC39 chromosome 15, complete sequence
634981
CP010928.1 *Sporisorium scitamineum* strain SSC39 chromosome 16, complete sequence
653278
CP010929.1 *Sporisorium scitamineum* strain SSC39 chromosome 17, complete sequence
605135
CP010930.1 *Sporisorium scitamineum* strain SSC39 chromosome 18, complete sequence
598089
CP010931.1 *Sporisorium scitamineum* strain SSC39 chromosome 19, complete sequence
580969
CP010932.1 *Sporisorium scitamineum* strain SSC39 chromosome 20, complete sequence
573042
CP010933.1 *Sporisorium scitamineum* strain SSC39 chromosome 21, complete sequence
539152
CP010934.1 *Sporisorium scitamineum* strain SSC39 chromosome 22, complete sequence
475902
CP010935.1 *Sporisorium scitamineum* strain SSC39 chromosome 23, complete sequence
140422
CP010939.1 *Sporisorium scitamineum* strain SSC39 mitochondrion, complete sequence
88018
CP011547.1 *Saccharomyces cerevisiae* strain ySR127 chromosome I, complete sequence
224154
CP011548.1 *Saccharomyces cerevisiae* strain ySR127 chromosome II, complete sequence

808595
CP011549.1 *Saccharomyces cerevisiae* strain ySR127 chromosome III, complete sequence
313668
CP011550.1 *Saccharomyces cerevisiae* strain ySR127 chromosome IV, complete sequence
1531682
CP011551.1 *Saccharomyces cerevisiae* strain ySR127 chromosome V, complete sequence
552153
CP011552.1 *Saccharomyces cerevisiae* strain ySR127 chromosome VI, complete sequence
270121
CP011553.1 *Saccharomyces cerevisiae* strain ySR127 chromosome VII, complete sequence
1084888
CP011554.1 *Saccharomyces cerevisiae* strain ySR127 chromosome VIII, complete sequence
562640
CP011555.1 *Saccharomyces cerevisiae* strain ySR127 chromosome IX, complete sequence
434899
CP011556.1 *Saccharomyces cerevisiae* strain ySR127 chromosome X, complete sequence
745071
CP011557.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XI, complete sequence
666444
CP011558.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XII, complete sequence
1062294
CP011559.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XIII, complete sequence
922588
CP011560.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XIV, complete sequence
784302
CP011561.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XV, complete sequence
1089058
CP011562.1 *Saccharomyces cerevisiae* strain ySR127 chromosome XVI, complete sequence
948057
CP011563.1 *Saccharomyces cerevisiae* strain ySR127 mitochondrion, complete sequence
85731
CM003593.1 *Saccharomyces eubayanus* strain FM1318 mitochondrion, complete sequence,
whole genome shotgun sequence
63999
CP019371.1 *Trametes hirsuta* strain 072 chromosome 1, complete sequence
4437623
CP019376.1 *Trametes hirsuta* strain 072 chromosome 3, complete sequence
3674663
CP019377.1 *Trametes hirsuta* strain 072 chromosome 4, complete sequence
3112363
CP019378.1 *Trametes hirsuta* strain 072 chromosome 5, complete sequence
3088601
CP019379.1 *Trametes hirsuta* strain 072 chromosome 6, complete sequence
3045029
CP019380.1 *Trametes hirsuta* strain 072 chromosome 7, complete sequence
2738869
CP019381.1 *Trametes hirsuta* strain 072 chromosome 8, complete sequence
2614590
CP019382.1 *Trametes hirsuta* strain 072 chromosome 9, complete sequence
2611572
CP019370.1 *Trametes hirsuta* strain 072 chromosome 10, complete sequence
2280792
CP019372.1 *Trametes hirsuta* strain 072 chromosome 11, complete sequence
2042872
CP019373.1 *Trametes hirsuta* strain 072 chromosome 12, complete sequence
1949064
CP019374.1 *Trametes hirsuta* strain 072 chromosome 13, complete sequence
1948569
AP014599.1 *Kluyveromyces marxianus* DNA, chromosome 1, complete genome, strain: NBRC
1777

1722779
AP014600.1 *Kluyveromyces marxianus* DNA, chromosome 2, complete genome, strain: NBRC 1777
1688883
AP014601.1 *Kluyveromyces marxianus* DNA, chromosome 3, complete genome, strain: NBRC 1777
1564943
AP014602.1 *Kluyveromyces marxianus* DNA, chromosome 4, complete genome, strain: NBRC 1777
1422013
AP014603.1 *Kluyveromyces marxianus* DNA, chromosome 5, complete genome, strain: NBRC 1777
1351988
AP014604.1 *Kluyveromyces marxianus* DNA, chromosome 6, complete genome, strain: NBRC 1777
1212614
AP014605.1 *Kluyveromyces marxianus* DNA, chromosome 7, complete genome, strain: NBRC 1777
940253
AP014606.1 *Kluyveromyces marxianus* DNA, chromosome 8, complete genome, strain: NBRC 1777
945852
AP014607.1 *Kluyveromyces marxianus* mitochondrial DNA, complete genome, strain: NBRC 1777
46256
AP012213.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 1
1745387
AP012214.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 2
1711476
AP012215.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 3
1588169
AP012216.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 4
1421472
AP012217.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 5
1353011
AP012218.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 6
1197921
AP012219.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 7
963005
AP012220.1 *Kluyveromyces marxianus* DMKU3-1042 DNA, complete genome, chromosome 8
939718
AP012221.1 *Kluyveromyces marxianus* DMKU3-1042 mitochondrial DNA, complete genome
46308
CP014242.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome II, complete sequence
2061902
CP014243.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome III, complete sequence
1398029
CP014244.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome IV, complete sequence
1404966
CP014245.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome V, complete sequence
1123606
CP014246.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome VI, complete sequence
912650
CP014247.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome VII, complete sequence
928141
CP014248.1 *Eremothecium sinECAUDUM* strain ATCC 58844 chromosome VIII, complete sequence
1093229
CP014249.1 *Eremothecium sinECAUDUM* strain ATCC 58844 mitochondrion, complete genome
26238
CP017344.1 *Talaromyces pinophilus* strain 1-95 chromosome 1, complete sequence
7684667

CP017345.1 *Talaromyces pinophilus* strain 1-95 chromosome 2, complete sequence
6009755
CP017346.1 *Talaromyces pinophilus* strain 1-95 chromosome 3, complete sequence
4804168
CP017347.1 *Talaromyces pinophilus* strain 1-95 chromosome 4, complete sequence
4274674
CP017348.1 *Talaromyces pinophilus* strain 1-95 chromosome 5, complete sequence
3951088
CP017349.1 *Talaromyces pinophilus* strain 1-95 chromosome 6, complete sequence
3820271
CP017350.1 *Talaromyces pinophilus* strain 1-95 chromosome 7, complete sequence
2993891
CP017351.1 *Talaromyces pinophilus* strain 1-95 chromosome 8, complete sequence
2941929
CP017352.1 *Talaromyces pinophilus* strain 1-95 mitochondrion, complete genome
31729
CM004311.1 *Saccharomyces cerevisiae* strain GLBRCY22-3 mitochondrion, complete sequence,
whole genome shotgun sequence
82897
CP014501.1 *Sugiyamaella lignohabitans* strain CBS 10342 chromosome A, complete sequence
5930846
CP014503.1 *Sugiyamaella lignohabitans* strain CBS 10342 chromosome B, complete sequence
4956242
CP014500.1 *Sugiyamaella lignohabitans* strain CBS 10342 chromosome C, complete sequence
2773601
CP014502.1 *Sugiyamaella lignohabitans* strain CBS 10342 chromosome D, complete sequence
2277569
CP015470.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 1, complete sequence
3779356
CP015471.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 2, complete sequence
3241356
CP015472.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 3, complete sequence
3126264
CP015473.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 4, complete sequence
3115696
CP015474.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 5, complete sequence
2477729
CP015476.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 7, complete sequence
2233015
CP015477.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 8, complete sequence
2155647
CP015479.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 10, complete sequence
1787784
CP015481.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 12, complete sequence
1576767
CP015482.1 *Agaricus bisporus* var. *bisporus* strain H39 chromosome 13, complete sequence
1405511
CM008299.1 *Fusarium oxysporum* f. sp. *radicis-cucumerinum* strain Forc016 mitochondrion,
complete sequence, whole genome shotgun sequence
47541
CP014709.1 *Komagataella phaffii* strain WT chromosome 2, complete sequence
2396783
CP014585.1 *Komagataella pastoris* strain ATCC 28485 chromosome 2, complete sequence
2687137
CP014716.1 *Komagataella phaffii* GS115 chromosome 2, complete sequence
2396528
CP017553.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1A, complete sequence
2257857
CP017554.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1B, complete sequence
3044971

CP017555.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1C, complete sequence
3366276
CP017556.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1D, complete sequence
3629463
CP017557.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1E, complete sequence
4198534
CP017558.1 *Yarrowia lipolytica* strain CLIB89(W29) chromosome 1F, complete sequence
4002965
CP017559.1 *Yarrowia lipolytica* strain CLIB89(W29) mitochondrion, complete genome
47926
CP017814.1 *Sclerotinia sclerotiorum* chromosome 1, complete sequence
3951982
CP017815.1 *Sclerotinia sclerotiorum* chromosome 2, complete sequence
3683506
CP017816.1 *Sclerotinia sclerotiorum* chromosome 3, complete sequence
3351453
CP017817.1 *Sclerotinia sclerotiorum* chromosome 4, complete sequence
2873318
CP017818.1 *Sclerotinia sclerotiorum* chromosome 5, complete sequence
2822964
CP017819.1 *Sclerotinia sclerotiorum* chromosome 6, complete sequence
2483831
CP017821.1 *Sclerotinia sclerotiorum* chromosome 8, complete sequence
2299506
CP017822.1 *Sclerotinia sclerotiorum* chromosome 9, complete sequence
2122865
CP017823.1 *Sclerotinia sclerotiorum* chromosome 10, complete sequence
2105496
CP017824.1 *Sclerotinia sclerotiorum* chromosome 11, complete sequence
2052242
CP017825.1 *Sclerotinia sclerotiorum* chromosome 12, complete sequence
1878461
CP017826.1 *Sclerotinia sclerotiorum* chromosome 13, complete sequence
1845946
CP017827.1 *Sclerotinia sclerotiorum* chromosome 14, complete sequence
1815632
CP017828.1 *Sclerotinia sclerotiorum* chromosome 15, complete sequence
1765292
CP017829.1 *Sclerotinia sclerotiorum* chromosome 16, complete sequence
1419421
CP012809.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A1, complete sequence
4911352
CP012810.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A2, complete sequence
4155011
CP012811.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A3, complete sequence
3014088
CP012812.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A4, complete sequence
2703655
CP012813.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A5, complete sequence
2119959
CP012814.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A6, complete sequence
1448773
CP012815.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome A7, complete sequence

1364045
CP012816.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B1, complete sequence
4331628
CP012817.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B2, complete sequence
4036768
CP012818.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B3, complete sequence
2645872
CP012819.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B4, complete sequence
2651701
CP012820.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B5, complete sequence
2379027
CP012821.1 *Saccharomycopsis fibuligera* x *Saccharomycopsis* cf. *fibuligera* strain KJJ81
chromosome B6, complete sequence
1410247
CP012823.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 1, complete sequence
4889271
CP012824.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 2, complete sequence
4132043
CP012825.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 3, complete sequence
3000806
CP012826.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 4, complete sequence
2649207
CP012827.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 5, complete sequence
2100688
CP012828.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 6, complete sequence
1441802
CP012829.1 *Saccharomycopsis fibuligera* strain KPH12 chromosome 7, complete sequence
1353399
CP015982.1 *Saccharomycopsis fibuligera* strain ATCC 36309 chromosome 5, complete sequence
2406875
CP015984.1 *Saccharomycopsis fibuligera* strain ATCC 36309 chromosome 7, complete sequence
1411284
CP019490.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 1, complete
sequence
2110500
CP019491.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 2, complete
sequence
2005801
CP019492.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 3, complete
sequence
1751495
CP019494.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 5, complete
sequence
1443312
CP019495.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 6, complete
sequence
1315104
CP019496.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 7, complete
sequence
1283838
CP019497.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 8, complete
sequence
1249162
CP019498.1 *Zygosaccharomyces parabailii* strain ATCC 60483 chromosome 9, complete
sequence

1240939
CP019499.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 10, complete sequence
1189704
CP019500.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 11, complete sequence
1115933
CP019501.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 12, complete sequence
1091360
CP019502.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 13, complete sequence
1077716
CP019503.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 14, complete sequence
1007293
CP019504.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 15, complete sequence
858772
CP019505.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 chromosome 16, complete sequence
571967
CP019507.2 *Zygosaccharomyces parabaillii* strain ATCC 60483 plasmid pSB2, complete sequence
5427
CP019506.1 *Zygosaccharomyces parabaillii* strain ATCC 60483 mitochondrion, complete genome
29945
CP016232.1 *Trichoderma reesei* QM6a chromosome I, complete sequence
6835803
CP016233.1 *Trichoderma reesei* QM6a chromosome II, complete sequence
6234656
CP016234.1 *Trichoderma reesei* QM6a chromosome III, complete sequence
5311445
CP016235.1 *Trichoderma reesei* QM6a chromosome IV, complete sequence
4556834
CP016236.1 *Trichoderma reesei* QM6a chromosome V, complete sequence
4159965
CP016237.1 *Trichoderma reesei* QM6a chromosome VI, complete sequence
4000387
CP016238.1 *Trichoderma reesei* QM6a chromosome VII, complete sequence
3823438
CP020123.1 *Saccharomyces cerevisiae* strain S288c chromosome I, complete sequence
219929
CP020124.1 *Saccharomyces cerevisiae* strain S288c chromosome II, complete sequence
813597
CP020125.1 *Saccharomyces cerevisiae* strain S288c chromosome III, complete sequence
341580
CP020126.1 *Saccharomyces cerevisiae* strain S288c chromosome IV, complete sequence
1566853
CP020127.1 *Saccharomyces cerevisiae* strain S288c chromosome V, complete sequence
583092
CP020128.1 *Saccharomyces cerevisiae* strain S288c chromosome VI, complete sequence
271539
CP020129.1 *Saccharomyces cerevisiae* strain S288c chromosome VII, complete sequence
1091538
CP020130.1 *Saccharomyces cerevisiae* strain S288c chromosome VIII, complete sequence
581049
CP020131.1 *Saccharomyces cerevisiae* strain S288c chromosome IX, complete sequence
440036
CP020132.1 *Saccharomyces cerevisiae* strain S288c chromosome X, complete sequence
751611

CP020133.1 *Saccharomyces cerevisiae* strain S288c chromosome XI, complete sequence
666862
CP020135.1 *Saccharomyces cerevisiae* strain S288c chromosome XIII, complete sequence
930506
CP020136.1 *Saccharomyces cerevisiae* strain S288c chromosome XIV, complete sequence
777615
CP020137.1 *Saccharomyces cerevisiae* strain S288c chromosome XV, complete sequence
1091343
CP020138.1 *Saccharomyces cerevisiae* strain S288c chromosome XVI, complete sequence
954457
CP020139.1 *Saccharomyces cerevisiae* strain S288c mitochondrion, complete genome
85793
CP020157.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome I, complete sequence
215496
CP020158.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome II, complete sequence
795659
CP020159.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome III, complete sequence
327289
CP020160.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome IV, complete sequence
1488087
CP020161.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome V, complete sequence
576784
CP020162.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome VI, complete sequence
257436
CP020163.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome VII, complete sequence
1070236
CP020164.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome VIII, complete sequence
533397
CP020165.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome IX, complete sequence
419821
CP020166.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome X, complete sequence
730957
CP020167.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome XI, complete sequence
657417
CP020169.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome XIII, complete sequence
913017
CP020170.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome XIV, complete sequence
765110
CP020171.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome XV, complete sequence
1120088
CP020172.1 *Saccharomyces cerevisiae* strain DBVPG6765 chromosome XVI, complete sequence
920208
CP020173.1 *Saccharomyces cerevisiae* strain DBVPG6765 mitochondrion, complete genome
81722
CP020174.1 *Saccharomyces cerevisiae* strain SK1 chromosome I, complete sequence
228861
CP020175.1 *Saccharomyces cerevisiae* strain SK1 chromosome II, complete sequence
829469
CP020176.1 *Saccharomyces cerevisiae* strain SK1 chromosome III, complete sequence
340914
CP020177.1 *Saccharomyces cerevisiae* strain SK1 chromosome IV, complete sequence
1486921
CP020178.1 *Saccharomyces cerevisiae* strain SK1 chromosome V, complete sequence
589812
CP020179.1 *Saccharomyces cerevisiae* strain SK1 chromosome VI, complete sequence
299318
CP020180.1 *Saccharomyces cerevisiae* strain SK1 chromosome VII, complete sequence
1080440
CP020181.1 *Saccharomyces cerevisiae* strain SK1 chromosome VIII, complete sequence
542723

CP020182.1 *Saccharomyces cerevisiae* strain SK1 chromosome IX, complete sequence
449612
CP020183.1 *Saccharomyces cerevisiae* strain SK1 chromosome X, complete sequence
753937
CP020184.1 *Saccharomyces cerevisiae* strain SK1 chromosome XI, complete sequence
690901
CP020186.1 *Saccharomyces cerevisiae* strain SK1 chromosome XIII, complete sequence
923535
CP020187.1 *Saccharomyces cerevisiae* strain SK1 chromosome XIV, complete sequence
791982
CP020188.1 *Saccharomyces cerevisiae* strain SK1 chromosome XV, complete sequence
1053869
CP020189.1 *Saccharomyces cerevisiae* strain SK1 chromosome XVI, complete sequence
946846
CP020190.1 *Saccharomyces cerevisiae* strain SK1 mitochondrion, complete genome
84638
CP020208.1 *Saccharomyces cerevisiae* strain YPS128 chromosome I, complete sequence
237661
CP020209.1 *Saccharomyces cerevisiae* strain YPS128 chromosome II, complete sequence
808481
CP020210.1 *Saccharomyces cerevisiae* strain YPS128 chromosome III, complete sequence
319304
CP020211.1 *Saccharomyces cerevisiae* strain YPS128 chromosome IV, complete sequence
1495051
CP020212.1 *Saccharomyces cerevisiae* strain YPS128 chromosome V, complete sequence
575962
CP020213.1 *Saccharomyces cerevisiae* strain YPS128 chromosome VI, complete sequence
289276
CP020214.1 *Saccharomyces cerevisiae* strain YPS128 chromosome VII, complete sequence
1082279
CP020215.1 *Saccharomyces cerevisiae* strain YPS128 chromosome VIII, complete sequence
543240
CP020216.1 *Saccharomyces cerevisiae* strain YPS128 chromosome IX, complete sequence
440919
CP020217.1 *Saccharomyces cerevisiae* strain YPS128 chromosome X, complete sequence
723414
CP020218.1 *Saccharomyces cerevisiae* strain YPS128 chromosome XI, complete sequence
667446
CP020220.1 *Saccharomyces cerevisiae* strain YPS128 chromosome XIII, complete sequence
928527
CP020221.1 *Saccharomyces cerevisiae* strain YPS128 chromosome XIV, complete sequence
766731
CP020222.1 *Saccharomyces cerevisiae* strain YPS128 chromosome XV, complete sequence
1072763
CP020223.1 *Saccharomyces cerevisiae* strain YPS128 chromosome XVI, complete sequence
932263
CP020224.1 *Saccharomyces cerevisiae* strain YPS128 mitochondrion, complete genome
77479
CP020225.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome I, complete
sequence
214332
CP020226.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome II, complete
sequence
815151
CP020227.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome III, complete
sequence
309137
CP020228.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome IV, complete
sequence
1468232

CP020229.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome V, complete sequence
555692

CP020230.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome VI, complete sequence
290867

CP020231.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome VII, complete sequence
632616

CP020232.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome VIII, complete sequence
738767

CP020233.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome IX, complete sequence
428968

CP020234.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome X, complete sequence
1092164

CP020235.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome XI, complete sequence
792116

CP020237.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome XIII, complete sequence
662343

CP020238.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome XIV, complete sequence
765143

CP020239.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome XV, complete sequence
1062502

CP020240.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 chromosome XVI, complete sequence
909189

CP020241.1 *Saccharomyces cerevisiae* strain UWOPS03-461.4 mitochondrion, complete genome
74179

CP020191.1 *Saccharomyces cerevisiae* strain Y12 chromosome I, complete sequence
197190

CP020192.1 *Saccharomyces cerevisiae* strain Y12 chromosome II, complete sequence
800619

CP020193.1 *Saccharomyces cerevisiae* strain Y12 chromosome III, complete sequence
322503

CP020194.1 *Saccharomyces cerevisiae* strain Y12 chromosome IV, complete sequence
1497473

CP020195.1 *Saccharomyces cerevisiae* strain Y12 chromosome V, complete sequence
575802

CP020196.1 *Saccharomyces cerevisiae* strain Y12 chromosome VI, complete sequence
285938

CP020197.1 *Saccharomyces cerevisiae* strain Y12 chromosome VII, complete sequence
1107723

CP020198.1 *Saccharomyces cerevisiae* strain Y12 chromosome VIII, complete sequence
547529

CP020199.1 *Saccharomyces cerevisiae* strain Y12 chromosome IX, complete sequence
431184

CP020200.1 *Saccharomyces cerevisiae* strain Y12 chromosome X, complete sequence
725652

CP020201.1 *Saccharomyces cerevisiae* strain Y12 chromosome XI, complete sequence
692235

CP020203.1 *Saccharomyces cerevisiae* strain Y12 chromosome XIII, complete sequence
903361

CP020204.1 *Saccharomyces cerevisiae* strain Y12 chromosome XIV, complete sequence

800685
CP020205.1 *Saccharomyces cerevisiae* strain Y12 chromosome XV, complete sequence
1048295
CP020206.1 *Saccharomyces cerevisiae* strain Y12 chromosome XVI, complete sequence
901780
CP020207.1 *Saccharomyces cerevisiae* strain Y12 mitochondrion, complete genome
82868
CP020140.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome I, complete sequence
217365
CP020141.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome II, complete sequence
815565
CP020142.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome III, complete sequence
332771
CP020143.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome IV, complete sequence
1486873
CP020144.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome V, complete sequence
572248
CP020145.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome VI, complete sequence
298678
CP020146.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome VII, complete sequence
1078455
CP020147.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome VIII, complete sequence
535633
CP020148.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome IX, complete sequence
443501
CP020149.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome X, complete sequence
728645
CP020150.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome XI, complete sequence
695907
CP020152.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome XIII, complete sequence
930256
CP020153.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome XIV, complete sequence
784343
CP020154.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome XV, complete sequence
1062383
CP020155.1 *Saccharomyces cerevisiae* strain DBVPG6044 chromosome XVI, complete sequence
939911
CP020156.1 *Saccharomyces cerevisiae* strain DBVPG6044 mitochondrion, complete genome
81093
CP020242.1 *Saccharomyces paradoxus* strain CBS432 chromosome I, complete sequence
235665
CP020243.1 *Saccharomyces paradoxus* strain CBS432 chromosome II, complete sequence
835812
CP020244.1 *Saccharomyces paradoxus* strain CBS432 chromosome III, complete sequence
357685
CP020245.1 *Saccharomyces paradoxus* strain CBS432 chromosome IV, complete sequence
1477988
CP020246.1 *Saccharomyces paradoxus* strain CBS432 chromosome V, complete sequence
565208
CP020247.1 *Saccharomyces paradoxus* strain CBS432 chromosome VI, complete sequence
296034
CP020248.1 *Saccharomyces paradoxus* strain CBS432 chromosome VII, complete sequence
1119193
CP020249.1 *Saccharomyces paradoxus* strain CBS432 chromosome VIII, complete sequence
531267
CP020250.1 *Saccharomyces paradoxus* strain CBS432 chromosome IX, complete sequence
443128
CP020251.1 *Saccharomyces paradoxus* strain CBS432 chromosome X, complete sequence
743843
CP020252.1 *Saccharomyces paradoxus* strain CBS432 chromosome XI, complete sequence

690553
CP020254.1 *Saccharomyces paradoxus* strain CBS432 chromosome XIII, complete sequence
941477
CP020255.1 *Saccharomyces paradoxus* strain CBS432 chromosome XIV, complete sequence
778926
CP020256.1 *Saccharomyces paradoxus* strain CBS432 chromosome XV, complete sequence
1078405
CP020257.1 *Saccharomyces paradoxus* strain CBS432 chromosome XVI, complete sequence
903028
CP020258.1 *Saccharomyces paradoxus* strain CBS432 mitochondrion, complete genome
71482
CP020259.1 *Saccharomyces paradoxus* strain N44 chromosome I, complete sequence
224181
CP020260.1 *Saccharomyces paradoxus* strain N44 chromosome II, complete sequence
800085
CP020261.1 *Saccharomyces paradoxus* strain N44 chromosome III, complete sequence
312241
CP020262.1 *Saccharomyces paradoxus* strain N44 chromosome IV, complete sequence
1466838
CP020263.1 *Saccharomyces paradoxus* strain N44 chromosome V, complete sequence
553973
CP020264.1 *Saccharomyces paradoxus* strain N44 chromosome VI, complete sequence
296174
CP020265.1 *Saccharomyces paradoxus* strain N44 chromosome VII, complete sequence
1082435
CP020266.1 *Saccharomyces paradoxus* strain N44 chromosome VIII, complete sequence
529296
CP020267.1 *Saccharomyces paradoxus* strain N44 chromosome IX, complete sequence
442403
CP020268.1 *Saccharomyces paradoxus* strain N44 chromosome X, complete sequence
729272
CP020269.1 *Saccharomyces paradoxus* strain N44 chromosome XI, complete sequence
679893
CP020271.1 *Saccharomyces paradoxus* strain N44 chromosome XIII, complete sequence
926729
CP020272.1 *Saccharomyces paradoxus* strain N44 chromosome XIV, complete sequence
776644
CP020273.1 *Saccharomyces paradoxus* strain N44 chromosome XV, complete sequence
1061158
CP020274.1 *Saccharomyces paradoxus* strain N44 chromosome XVI, complete sequence
904322
CP020275.1 *Saccharomyces paradoxus* strain N44 mitochondrion, complete genome
69948
CP020276.1 *Saccharomyces paradoxus* strain YPS138 chromosome I, complete sequence
254647
CP020277.1 *Saccharomyces paradoxus* strain YPS138 chromosome II, complete sequence
789597
CP020278.1 *Saccharomyces paradoxus* strain YPS138 chromosome III, complete sequence
315836
CP020279.1 *Saccharomyces paradoxus* strain YPS138 chromosome IV, complete sequence
1469832
CP020280.1 *Saccharomyces paradoxus* strain YPS138 chromosome V, complete sequence
571131
CP020281.1 *Saccharomyces paradoxus* strain YPS138 chromosome VI, complete sequence
296071
CP020282.1 *Saccharomyces paradoxus* strain YPS138 chromosome VII, complete sequence
1067218
CP020283.1 *Saccharomyces paradoxus* strain YPS138 chromosome VIII, complete sequence
531566
CP020284.1 *Saccharomyces paradoxus* strain YPS138 chromosome IX, complete sequence

431692
CP020285.1 *Saccharomyces paradoxus* strain YPS138 chromosome X, complete sequence
707288
CP020286.1 *Saccharomyces paradoxus* strain YPS138 chromosome XI, complete sequence
687600
CP020288.1 *Saccharomyces paradoxus* strain YPS138 chromosome XIII, complete sequence
935281
CP020289.1 *Saccharomyces paradoxus* strain YPS138 chromosome XIV, complete sequence
769158
CP020290.1 *Saccharomyces paradoxus* strain YPS138 chromosome XV, complete sequence
1049886
CP020291.1 *Saccharomyces paradoxus* strain YPS138 chromosome XVI, complete sequence
905611
CP020292.1 *Saccharomyces paradoxus* strain YPS138 mitochondrion, complete genome
71396
CP020293.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome I, complete sequence
220177
CP020294.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome II, complete sequence
731301
CP020295.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome III, complete sequence
308064
CP020296.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome IV, complete sequence
1391618
CP020297.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome V, complete sequence
577572
CP020298.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome VI, complete sequence
282957
CP020299.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome VII, complete sequence
1105967
CP020300.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome VIII, complete sequence
546662
CP020301.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome IX, complete sequence
374625
CP020302.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome X, complete sequence
751133
CP020303.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome XI, complete sequence
805821
CP020305.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome XIII, complete sequence
869764
CP020306.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome XIV, complete sequence
872155
CP020307.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome XV, complete sequence
1201643
CP020308.1 *Saccharomyces paradoxus* strain UFRJ50816 chromosome XVI, complete sequence
1029411
CP020309.1 *Saccharomyces paradoxus* strain UFRJ50816 mitochondrion, complete genome
77386
CP020310.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome I, complete
sequence
211120
CP020311.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome II, complete
sequence
801843
CP020312.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome III, complete
sequence
316051
CP020313.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome IV, complete
sequence
1513677

CP020314.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome V, complete sequence
1007496

CP020315.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome VI, complete sequence
306036

CP020316.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome VII, complete sequence
1057931

CP020317.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome VIII, complete sequence
526907

CP020318.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome IX, complete sequence
439147

CP020319.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome X, complete sequence
750185

CP020320.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome XI, complete sequence
681961

CP020322.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome XIII, complete sequence
475121

CP020323.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome XIV, complete sequence
769517

CP020324.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome XV, complete sequence
1052871

CP020325.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 chromosome XVI, complete sequence
905577

CP020326.1 *Saccharomyces paradoxus* strain UWOPS91-917.1 mitochondrion, complete genome
73171

CP021239.1 *Kluyveromyces lactis* strain GG799 chromosome A, complete sequence
1050866

CP021240.1 *Kluyveromyces lactis* strain GG799 chromosome B, complete sequence
1319333

CP021241.1 *Kluyveromyces lactis* strain GG799 chromosome C, complete sequence
1752996

CP021242.1 *Kluyveromyces lactis* strain GG799 chromosome D, complete sequence
1700866

CP021243.1 *Kluyveromyces lactis* strain GG799 chromosome E, complete sequence
2217437

CP021244.1 *Kluyveromyces lactis* strain GG799 chromosome F, complete sequence
2601250

CP021245.1 *Kluyveromyces lactis* strain GG799 mitochondrion, complete genome
39914

CM007980.1 *Saccharomyces cerevisiae* W303 plasmid p2-micron, complete sequence, whole genome shotgun sequence
6300

CP022321.1 *Cryptococcus neoformans* var. *grubii* chromosome 1, complete sequence
2291500

CP022322.1 *Cryptococcus neoformans* var. *grubii* chromosome 2, complete sequence
1621676

CP022323.1 *Cryptococcus neoformans* var. *grubii* chromosome 3, complete sequence
1574972

CP022324.1 *Cryptococcus neoformans* var. *grubii* chromosome 4, complete sequence
1084805

CP022325.1 *Cryptococcus neoformans* var. *grubii* chromosome 5, complete sequence
1814975
CP022326.1 *Cryptococcus neoformans* var. *grubii* chromosome 6, complete sequence
1422463
CP022327.1 *Cryptococcus neoformans* var. *grubii* chromosome 7, complete sequence
1399209
CP022328.1 *Cryptococcus neoformans* var. *grubii* chromosome 8, complete sequence
1398693
CP022329.1 *Cryptococcus neoformans* var. *grubii* chromosome 9, complete sequence
1186813
CP022330.1 *Cryptococcus neoformans* var. *grubii* chromosome 10, complete sequence
1059962
CP022331.1 *Cryptococcus neoformans* var. *grubii* chromosome 11, complete sequence
1562107
CP022332.1 *Cryptococcus neoformans* var. *grubii* chromosome 12, complete sequence
774060
CP022333.1 *Cryptococcus neoformans* var. *grubii* chromosome 13, complete sequence
756017
CP022334.1 *Cryptococcus neoformans* var. *grubii* chromosome 14, complete sequence
942472
CP022335.1 *Cryptococcus neoformans* var. *grubii* strain KN99 mitochondrion, complete genome
24923
CM008094.1 [*Candida*] *glabrata* strain DSY562 mitochondrion, complete sequence, whole genome
shotgun sequence
20086
CM008107.1 [*Candida*] *glabrata* strain DSY565 mitochondrion, complete sequence, whole genome
shotgun sequence
20086
CP022800.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 1, complete sequence
3532264
CP022801.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 2, complete sequence
2872662
CP022803.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 3, complete sequence
2401307
CP022802.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 4, complete sequence
2335616
CP022804.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 5, complete sequence
2001616
CP022806.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 6, complete sequence
1779389
CP022807.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 7, complete sequence
1731349
CP022808.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 8, complete sequence
1712619
CP022809.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 9, complete sequence
1657153
CP022805.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 10, complete sequence
1651622
CP022812.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 11, complete sequence
1479508
CP022811.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 13, complete sequence
1441076
CP022813.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 14, complete sequence
1377913
CP022814.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 15, complete sequence
1346229
CP022816.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 16, complete sequence
1292678
CP022815.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 17, complete sequence
1279747

CP022817.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 18, complete sequence
1274223

CP022818.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 19, complete sequence
1125282

CP022820.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 20, complete sequence
1090035

CP022819.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 21, complete sequence
1077963

CP022822.1 *Parastagonospora nodorum* isolate LDN03-Sn4 chromosome 23, complete sequence
476058

CP022852.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 1, complete sequence
3434512

CP022853.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 2, complete sequence
2797422

CP022855.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 3, complete sequence
2253675

CP022854.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 4, complete sequence
2256902

CP022856.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 5, complete sequence
1872577

CP022858.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 6, complete sequence
1730842

CP022857.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 7, complete sequence
1705187

CP022860.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 8, complete sequence
1558419

CP022859.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 9, complete sequence
1583228

CP022869.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 10, complete sequence
1130272

CP022861.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 11, complete sequence
1388164

CP022862.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 13, complete sequence
1396077

CP022865.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 14, complete sequence
1299335

CP022864.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 15, complete sequence
1255146

CP022868.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 16, complete sequence
1170298

CP022866.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 17, complete sequence
1211780

CP022873.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 19, complete sequence
995738

CP022872.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 20, complete sequence
1048157

CP022871.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 21, complete sequence
1070356

CP022867.1 *Parastagonospora nodorum* isolate Sn79-1087 chromosome 22, complete sequence
1168050

CM008263.1 *Phellinus noxius* strain FFPRI411160 mitochondrion, complete sequence, whole
genome shotgun sequence
163449

CP023456.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 1, complete sequence
1734405

CP023457.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 2, complete sequence
1699062

CP023458.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 3, complete sequence
1559681

CP023459.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 4, complete sequence

1406771
CP023460.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 5, complete sequence
1373102
CP023461.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 6, complete sequence
1205333
CP023462.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 7, complete sequence
928821
CP023463.1 *Kluyveromyces marxianus* strain NRRL Y-6860 chromosome 8, complete sequence
930443
CM008875.1 *Saccharomyces cerevisiae* strain CEN.PK113-7D plasmid p2uM, complete
sequence, whole genome shotgun sequence
6318
CM008876.1 *Saccharomyces cerevisiae* strain CEN.PK113-7D mitochondrion, complete
sequence, whole genome shotgun sequence
86616
CP022024.1 *Alternaria solani* isolate NL03003 chromosome 1, complete sequence
6940169
CP022025.1 *Alternaria solani* isolate NL03003 chromosome 2, complete sequence
5062150
CP022026.1 *Alternaria solani* isolate NL03003 chromosome 3, complete sequence
3306264
CP022027.1 *Alternaria solani* isolate NL03003 chromosome 4, complete sequence
2866555
CP022028.1 *Alternaria solani* isolate NL03003 chromosome 5, complete sequence
2771896
CP022029.1 *Alternaria solani* isolate NL03003 chromosome 6, complete sequence
2590027
CP022030.1 *Alternaria solani* isolate NL03003 chromosome 7, complete sequence
2546252
CP022031.1 *Alternaria solani* isolate NL03003 chromosome 8, complete sequence
2508373
CP022032.1 *Alternaria solani* isolate NL03003 chromosome 9, complete sequence
2309181
CP022033.1 *Alternaria solani* isolate NL03003 chromosome 10, complete sequence
1878275
CP025759.1 *Cryptococcus gattii* VGII R265 chromosome 1, complete sequence
2207848
CP025760.1 *Cryptococcus gattii* VGII R265 chromosome 2, complete sequence
2171391
CP025761.1 *Cryptococcus gattii* VGII R265 chromosome 3, complete sequence
1968033
CP025762.1 *Cryptococcus gattii* VGII R265 chromosome 4, complete sequence
1750766
CP025763.1 *Cryptococcus gattii* VGII R265 chromosome 5, complete sequence
1316417
CP025764.1 *Cryptococcus gattii* VGII R265 chromosome 6, complete sequence
1307262
CP025765.1 *Cryptococcus gattii* VGII R265 chromosome 7, complete sequence
1243496
CP025766.1 *Cryptococcus gattii* VGII R265 chromosome 8, complete sequence
1233638
CP025767.1 *Cryptococcus gattii* VGII R265 chromosome 9, complete sequence
972356
CP025768.1 *Cryptococcus gattii* VGII R265 chromosome 10, complete sequence
815166
CP025769.1 *Cryptococcus gattii* VGII R265 chromosome 11, complete sequence
733013
CP025770.1 *Cryptococcus gattii* VGII R265 chromosome 12, complete sequence
663894
CP025771.1 *Cryptococcus gattii* VGII R265 chromosome 13, complete sequence

650339
CP025772.1 *Cryptococcus gattii* VGII R265 chromosome 14, complete sequence
516503
CP025773.1 *Cryptococcus gattii* VGII R265 mitochondrion, complete genome
31189
CP027647.1 *Torulaspora delbrueckii* strain COFT1 chromosome 1, complete sequence
1420397
CP027648.1 *Torulaspora delbrueckii* strain COFT1 chromosome 2, complete sequence
1398547
CP027649.1 *Torulaspora delbrueckii* strain COFT1 chromosome 3, complete sequence
1318504
CP027650.1 *Torulaspora delbrueckii* strain COFT1 chromosome 4, complete sequence
1238785
CP027651.1 *Torulaspora delbrueckii* strain COFT1 chromosome 5, complete sequence
1073930
CP027652.1 *Torulaspora delbrueckii* strain COFT1 chromosome 6, complete sequence
1067747
CP027653.1 *Torulaspora delbrueckii* strain COFT1 chromosome 7, complete sequence
927760
CP027654.1 *Torulaspora delbrueckii* strain COFT1 chromosome 8, complete sequence
863982
CP027655.1 *Torulaspora delbrueckii* strain COFT1 mitochondrion, complete genome
47174
CP021098.1 *Pichia kudriavzevii* strain SJP mitochondrion, complete genome
51126
CP028531.1 *Pichia kudriavzevii* strain CBS5147 chromosome 1, complete sequence
2861343
CP028532.1 *Pichia kudriavzevii* strain CBS5147 chromosome 2, complete sequence
2715831
CP028533.1 *Pichia kudriavzevii* strain CBS5147 chromosome 3, complete sequence
2541826
CP028534.1 *Pichia kudriavzevii* strain CBS5147 chromosome 4, complete sequence
1374539
CP028535.1 *Pichia kudriavzevii* strain CBS5147 chromosome 5, complete sequence
1289534
CP028773.1 *Pichia kudriavzevii* strain CBS573 chromosome 1, complete sequence
2852272
CP028774.1 *Pichia kudriavzevii* strain CBS573 chromosome 2, complete sequence
2745854
CP028775.1 *Pichia kudriavzevii* strain CBS573 chromosome 3, complete sequence
2541747
CP028776.1 *Pichia kudriavzevii* strain CBS573 chromosome 4, complete sequence
1383544
CP028777.1 *Pichia kudriavzevii* strain CBS573 chromosome 5, complete sequence
1289138
CP028778.1 *Pichia kudriavzevii* strain CBS573 mitochondrion, complete genome
51340
CP026301.1 *Saccharomyces cerevisiae* strain BY4742 chromosome I, complete sequence
230490
CP026296.1 *Saccharomyces cerevisiae* strain BY4742 chromosome II, complete sequence
822297
CP026297.1 *Saccharomyces cerevisiae* strain BY4742 chromosome III, complete sequence
333764
CP026298.1 *Saccharomyces cerevisiae* strain BY4742 chromosome IV, complete sequence
1545606
CP026299.1 *Saccharomyces cerevisiae* strain BY4742 chromosome V, complete sequence
582048
CP026302.1 *Saccharomyces cerevisiae* strain BY4742 chromosome VI, complete sequence
270887
CP026294.1 *Saccharomyces cerevisiae* strain BY4742 chromosome VII, complete sequence

1091388
CP026287.1 *Saccharomyces cerevisiae* strain BY4742 chromosome VIII, complete sequence
576806
CP026295.1 *Saccharomyces cerevisiae* strain BY4742 chromosome IX, complete sequence
440032
CP026288.1 *Saccharomyces cerevisiae* strain BY4742 chromosome X, complete sequence
745649
CP026289.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XI, complete sequence
666876
CP026300.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XII, complete sequence
1104511
CP026291.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XIII, complete sequence
930523
CP026293.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XIV, complete sequence
784585
CP026303.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XV, complete sequence
1091321
CP026290.1 *Saccharomyces cerevisiae* strain BY4742 chromosome XVI, complete sequence
948685
HG970332.2 *Fusarium graminearum* chromosome 1, complete genome
11760891
HG970333.1 *Fusarium graminearum* chromosome 2, complete genome
8997558
HG970334.1 *Fusarium graminearum* chromosome 3, complete genome
7792947
HG970335.2 *Fusarium graminearum* chromosome 4, complete genome
9395062
HG970331.1 *Fusarium graminearum* mitochondrial complete genome
95638
CM000169.1 *Aspergillus fumigatus* Af293 chromosome 1, whole genome shotgun sequence
4918979
CM000170.1 *Aspergillus fumigatus* Af293 chromosome 2, whole genome shotgun sequence
4844472
CM000171.1 *Aspergillus fumigatus* Af293 chromosome 3, whole genome shotgun sequence
4079167
CM000172.1 *Aspergillus fumigatus* Af293 chromosome 4, whole genome shotgun sequence
3923705
CM000173.1 *Aspergillus fumigatus* Af293 chromosome 5, whole genome shotgun sequence
3948441
CM000174.1 *Aspergillus fumigatus* Af293 chromosome 6, whole genome shotgun sequence
3778736
CM000175.1 *Aspergillus fumigatus* Af293 chromosome 7, whole genome shotgun sequence
2058334
CP003009.1 *Thielavia terrestris* NRRL 8126 chromosome 1, complete sequence
10101509
CP003010.1 *Thielavia terrestris* NRRL 8126 chromosome 2, complete sequence
9477512
CP003011.1 *Thielavia terrestris* NRRL 8126 chromosome 3, complete sequence
4786945
CP003012.1 *Thielavia terrestris* NRRL 8126 chromosome 4, complete sequence
4578922
CP003013.1 *Thielavia terrestris* NRRL 8126 chromosome 5, complete sequence
4396881
CP003014.1 *Thielavia terrestris* NRRL 8126 chromosome 6, complete sequence
3570487
CH672346.1 *Candida albicans* WO-1 chromosome 1 supercont1.1 genomic scaffold, whole
genome shotgun sequence
3194068
GG670278.1 *Candida albicans* WO-1 chromosome 1 supercont1.10 genomic scaffold, whole
genome shotgun sequence

10847
CH672350.1 *Candida albicans* WO-1 chromosome 2 supercont1.5 genomic scaffold, whole genome shotgun sequence
1368965
CH672354.1 *Candida albicans* WO-1 chromosome 2 supercont1.9 genomic scaffold, whole genome shotgun sequence
891553
CM000310.1 *Candida albicans* WO-1 chromosome 3, whole genome shotgun sequence
1768732
GG670282.1 *Candida albicans* WO-1 chromosome 4 supercont1.14 genomic scaffold, whole genome shotgun sequence
6607
CH672349.1 *Candida albicans* WO-1 chromosome 4 supercont1.4 genomic scaffold, whole genome shotgun sequence
1636855
CM000311.1 *Candida albicans* WO-1 chromosome 5, whole genome shotgun sequence
1198695
CM000312.1 *Candida albicans* WO-1 chromosome 6, whole genome shotgun sequence
1043947
CM000313.1 *Candida albicans* WO-1 chromosome 7, whole genome shotgun sequence
958737
CM000309.1 *Candida albicans* WO-1 chromosome R, whole genome shotgun sequence
2299365
GG670279.1 *Candida albicans* WO-1 supercont1.11 genomic scaffold, whole genome shotgun sequence
10615
GG670280.1 *Candida albicans* WO-1 supercont1.12 genomic scaffold, whole genome shotgun sequence
10435
GG670281.1 *Candida albicans* WO-1 supercont1.13 genomic scaffold, whole genome shotgun sequence
7823
GG670283.1 *Candida albicans* WO-1 supercont1.15 genomic scaffold, whole genome shotgun sequence
5386
GG670284.1 *Candida albicans* WO-1 supercont1.16 genomic scaffold, whole genome shotgun sequence
5039
GG670285.1 *Candida albicans* WO-1 supercont1.17 mitochondrial scaffold, whole genome shotgun sequence
55284
CP017623.1 *Candida albicans* SC5314 chromosome 1 sequence
3188341
CP017624.1 *Candida albicans* SC5314 chromosome 2 sequence
2231883
CP017625.1 *Candida albicans* SC5314 chromosome 3 sequence
1799298
CP017626.1 *Candida albicans* SC5314 chromosome 4 sequence
1603259
CP017627.1 *Candida albicans* SC5314 chromosome 5 sequence
1190845
CP017628.1 *Candida albicans* SC5314 chromosome 6 sequence
1033292
CP017629.1 *Candida albicans* SC5314 chromosome 7 sequence
949511
CP017630.1 *Candida albicans* SC5314 chromosome R sequence
2286237
CP025167.1 *Candida albicans* strain SC5314-GTH12 chromosome 1A
3186784
CP025168.1 *Candida albicans* strain SC5314-GTH12 chromosome 2A

2231038
CP025169.1 *Candida albicans* strain SC5314-GTH12 chromosome 3A
1799051
CP025170.1 *Candida albicans* strain SC5314-GTH12 chromosome 4A
1603040
CP025171.1 *Candida albicans* strain SC5314-GTH12 chromosome 5A
1190317
CP025172.1 *Candida albicans* strain SC5314-GTH12 chromosome 6A
1032182
CP025173.1 *Candida albicans* strain SC5314-GTH12 chromosome 7A
948952
CP025174.1 *Candida albicans* strain SC5314-GTH12 chromosome RA
2285431
CP025175.1 *Candida albicans* strain SC5314-GTH12 chromosome 1B
3186276
CP025176.1 *Candida albicans* strain SC5314-GTH12 chromosome 2B
2231079
CP025177.1 *Candida albicans* strain SC5314-GTH12 chromosome 3B
1798977
CP025178.1 *Candida albicans* strain SC5314-GTH12 chromosome 4B
1602746
CP025179.1 *Candida albicans* strain SC5314-GTH12 chromosome 5B
1190508
CP025180.1 *Candida albicans* strain SC5314-GTH12 chromosome 6B
1032226
CP025181.1 *Candida albicans* strain SC5314-GTH12 chromosome 7B
949045
CP025182.1 *Candida albicans* strain SC5314-GTH12 chromosome RB
2285142
CP025183.1 *Candida albicans* strain SC5314-GTH12 mitochondrion
40418
CP025150.1 *Candida albicans* strain SC5314-P0 chromosome 1A
3187190
CP025151.1 *Candida albicans* strain SC5314-P0 chromosome 2A
2231757
CP025152.1 *Candida albicans* strain SC5314-P0 chromosome 3A
1799323
CP025153.1 *Candida albicans* strain SC5314-P0 chromosome 4A
1603391
CP025154.1 *Candida albicans* strain SC5314-P0 chromosome 5A
1190670
CP025155.1 *Candida albicans* strain SC5314-P0 chromosome 6A
1032578
CP025156.1 *Candida albicans* strain SC5314-P0 chromosome 7A
949203
CP025157.1 *Candida albicans* strain SC5314-P0 chromosome RA
2285906
CP025158.1 *Candida albicans* strain SC5314-P0 chromosome 1B
3186810
CP025159.1 *Candida albicans* strain SC5314-P0 chromosome 2B
2231689
CP025160.1 *Candida albicans* strain SC5314-P0 chromosome 3B
1799405
CP025161.1 *Candida albicans* strain SC5314-P0 chromosome 4B
1603204
CP025162.1 *Candida albicans* strain SC5314-P0 chromosome 5B
1190801
CP025163.1 *Candida albicans* strain SC5314-P0 chromosome 6B
1032709
CP025164.1 *Candida albicans* strain SC5314-P0 chromosome 7B

949291
CP025165.1 *Candida albicans* strain SC5314-P0 chromosome RB
2285569
CP025166.1 *Candida albicans* strain SC5314-P0 mitochondrion
40443
CM001231.1 *Magnaporthe oryzae* 70-15 chromosome 1, whole genome shotgun sequence
7978604
CM001232.1 *Magnaporthe oryzae* 70-15 chromosome 2, whole genome shotgun sequence
8319966
CM001233.1 *Magnaporthe oryzae* 70-15 chromosome 3, whole genome shotgun sequence
6606598
CM001234.1 *Magnaporthe oryzae* 70-15 chromosome 4, whole genome shotgun sequence
5546968
CM001235.1 *Magnaporthe oryzae* 70-15 chromosome 5, whole genome shotgun sequence
4490059
CM001236.1 *Magnaporthe oryzae* 70-15 chromosome 6, whole genome shotgun sequence
4133993
CM001237.1 *Magnaporthe oryzae* 70-15 chromosome 7, whole genome shotgun sequence
3415785
AE016814.2 *Ashbya gossypii* ATCC 10895 chromosome I, complete sequence
693414
AE016815.5 *Ashbya gossypii* ATCC 10895 chromosome II, complete sequence
870771
AE016816.3 *Ashbya gossypii* ATCC 10895 chromosome III, complete sequence
907494
AE016817.6 *Ashbya gossypii* ATCC 10895 chromosome IV, complete sequence
1467287
AE016818.2 *Ashbya gossypii* ATCC 10895 chromosome V, complete sequence
1519140
AE016819.5 *Ashbya gossypii* ATCC 10895 chromosome VI, complete sequence
1836693
AE016820.4 *Ashbya gossypii* ATCC 10895 chromosome VII, complete sequence
1800949
AE016821.1 *Ashbya gossypii* (= *Eremothecium gossypii*) ATCC 10895 mitochondrion, complete
genome
23564
CP002705.1 *Ashbya gossypii* FDAG1 chromosome I, complete sequence
693310
CP002706.1 *Ashbya gossypii* FDAG1 chromosome II, complete sequence
874947
CP002707.1 *Ashbya gossypii* FDAG1 chromosome III, complete sequence
911446
CP002708.1 *Ashbya gossypii* FDAG1 chromosome IV, complete sequence
1465717
CP002709.1 *Ashbya gossypii* FDAG1 chromosome V, complete sequence
1527143
CP002710.1 *Ashbya gossypii* FDAG1 chromosome VI, complete sequence
1837132
CP002711.1 *Ashbya gossypii* FDAG1 chromosome VII, complete sequence
1823962
CP002712.1 *Ashbya gossypii* FDAG1 mitochondrion, complete sequence
23497
CM001563.1 *Saccharomyces arboricola* H-6 chromosome I, whole genome shotgun sequence
194368
CM001564.1 *Saccharomyces arboricola* H-6 chromosome II, whole genome shotgun sequence
782499
CM001565.1 *Saccharomyces arboricola* H-6 chromosome III, whole genome shotgun sequence
298424
CM001566.1 *Saccharomyces arboricola* H-6 chromosome IV, whole genome shotgun sequence
1141293

CM001567.1 *Saccharomyces arboricola* H-6 chromosome V, whole genome shotgun sequence
531626

CM001568.1 *Saccharomyces arboricola* H-6 chromosome VI, whole genome shotgun sequence
272605

CM001569.1 *Saccharomyces arboricola* H-6 chromosome VII, whole genome shotgun sequence
1053431

CM001570.1 *Saccharomyces arboricola* H-6 chromosome VIII, whole genome shotgun sequence
527949

CM001571.1 *Saccharomyces arboricola* H-6 chromosome IX, whole genome shotgun sequence
407361

CM001572.1 *Saccharomyces arboricola* H-6 chromosome X, whole genome shotgun sequence
721635

CM001573.1 *Saccharomyces arboricola* H-6 chromosome XI, whole genome shotgun sequence
650640

CM001574.1 *Saccharomyces arboricola* H-6 chromosome XII, whole genome shotgun sequence
984206

CM001575.1 *Saccharomyces arboricola* H-6 chromosome XIII, whole genome shotgun sequence
1246548

CM001576.1 *Saccharomyces arboricola* H-6 chromosome XIV, whole genome shotgun sequence
771513

CM001577.1 *Saccharomyces arboricola* H-6 chromosome XV, whole genome shotgun sequence
1023324

CM001578.1 *Saccharomyces arboricola* H-6 chromosome XVI, whole genome shotgun sequence
879294

CM000687.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome A, whole genome shotgun
sequence
951467

CM000688.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome B, whole genome shotgun
sequence
1119477

CM000689.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome C, whole genome shotgun
sequence
1252455

CM000690.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome D, whole genome shotgun
sequence
1289280

CM000691.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome E, whole genome shotgun
sequence
1295560

CM000692.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome F, whole genome shotgun
sequence
1385275

CM000693.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome G, whole genome shotgun
sequence
1737261

CM000694.1 *Saccharomyces kluyveri* NRRL Y-12651 chromosome H, whole genome shotgun
sequence
2314951

CP001942.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome I, complete sequence
160332

CP001943.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome II, complete sequence
175776

CP001944.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome III, complete sequence
176815

CP001945.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome IV, complete sequence
193740

CP001946.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome V, complete sequence
196642

CP001947.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VI, complete sequence
198217

CP001948.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VII, complete sequence
205935
CP001949.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome VIII, complete sequence
204910
CP001950.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome IX sequence
233397
CP001951.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome X, complete sequence
234890
CP001952.1 *Encephalitozoon intestinalis* ATCC 50506 chromosome XI, complete sequence
236244
CP010913.1 *Sporisorium scitamineum* strain SSC39 chromosome 1, complete sequence
2009762
CP010914.1 *Sporisorium scitamineum* strain SSC39 chromosome 2, complete sequence
1671606
CP010915.1 *Sporisorium scitamineum* strain SSC39 chromosome 3, complete sequence
1566833
CP010916.1 *Sporisorium scitamineum* strain SSC39 chromosome 4, complete sequence
1271980
CP010917.1 *Sporisorium scitamineum* strain SSC39 chromosome 5, complete sequence
1055339
CP010918.1 *Sporisorium scitamineum* strain SSC39 chromosome 6, complete sequence
1032954
CP010919.1 *Sporisorium scitamineum* strain SSC39 chromosome 7, complete sequence
938453
CP010920.1 *Sporisorium scitamineum* strain SSC39 chromosome 8, complete sequence
837966
CP010921.1 *Sporisorium scitamineum* strain SSC39 chromosome 9, complete sequence
808176
CP010922.1 *Sporisorium scitamineum* strain SSC39 chromosome 10, complete sequence
875830
CP010923.1 *Sporisorium scitamineum* strain SSC39 chromosome 11, complete sequence
790822
CP010924.1 *Sporisorium scitamineum* strain SSC39 chromosome 12, complete sequence
716888
CP010925.1 *Sporisorium scitamineum* strain SSC39 chromosome 13, complete sequence
681008
CP010926.1 *Sporisorium scitamineum* strain SSC39 chromosome 14, complete sequence
652314
CP010927.1 *Sporisorium scitamineum* strain SSC39 chromosome 15, complete sequence
634981
CP010928.1 *Sporisorium scitamineum* strain SSC39 chromosome 16, complete sequence
653278
CP010929.1 *Sporisorium scitamineum* strain SSC39 chromosome 17, complete sequence
605135
CP010930.1 *Sporisorium scitamineum* strain SSC39 chromosome 18, complete sequence
598089
CP010931.1 *Sporisorium scitamineum* strain SSC39 chromosome 19, complete sequence
580969
CP010932.1 *Sporisorium scitamineum* strain SSC39 chromosome 20, complete sequence
573042
CP010933.1 *Sporisorium scitamineum* strain SSC39 chromosome 21, complete sequence
539152
CP010934.1 *Sporisorium scitamineum* strain SSC39 chromosome 22, complete sequence
475902
CP010935.1 *Sporisorium scitamineum* strain SSC39 chromosome 23, complete sequence
140422
CP010936.1 *Sporisorium scitamineum* strain SSC39 chromosome 24 sequence
87293
CP010937.1 *Sporisorium scitamineum* strain SSC39 chromosome 25 sequence
82225

CP010938.1 Sporisorium scitamineum strain SSC39 chromosome 26 sequence
99152
CP010939.1 Sporisorium scitamineum strain SSC39 mitochondrion, complete sequence
88018
>NC_031675.1 Fusarium verticillioides 7600 chromosome 1, whole genome shotgun sequence
6219215