

1 **tggtgatac** gtcagttctgg gtgttagtc acagcaatcg cagccaaaat gggctcatct
 61 **gccacatgg** tctttcgct ggcgcagacg gcccttaatt tcattccagaa acttccatat
 121 **tgc**ccctgca atacaattga gtccttggat ttttttctc gcctccgtac ggcgtactgc
 181 **cgtcgagtac** gctctgcgac acatccogat ctacaatttt tcacagccat aattcaggga
 241 **ctttttgt**a tactggtcta tttcaaatga gacaactcg~~c~~ tttccaaccc catataaccg
 301 **tctatctact** tcatgggtca tcatgcattt gtttccaaact ttctacacct atttgttaaa
 361 **caat**gtact aatattgtca taggtttaga tcagagaact ctccagcgt~~a~~ aattcagaa
 421 **cctat**gagaa attttttttc aggaaaaattt ttctatgaga ctcatctcg ggcccttgt
 481 **tggat**ttta tataaggaac actagaatcc cttat~~ttttt~~ aattgaatta caaaaatttt
 541 **tttctat**aca tatatctcat atattaatta atcaaa**atgg** gtaaaagaaaa gactcacg~~tt~~
 601 **aatctcg**ttg tcattggtca cg~~t~~cgattct ggttaatcta ctaccacccg tcacttaatt
 661 **taca**agtgtg gtggtatcg~~a~~ taagagaacc attgaaaagt tcgaaaagga agccgctgaa
 721 **ttaggt**taagg gtttttcaa gtacgcctgg gtttttagaca agttaaaggc tgaaagagaa
 781 **agagg**tatca ccattgat~~at~~ cgctttatgg aagttcgaaa ctccaaagtt ccacgtcacc
 841 **attatt**gatg ctccagg~~t~~ca cagagattc atcaagaaca tgattactgg tac~~tt~~cccaa
 901 **gctg**attgtg ctat~~ttt~~gat tattgtgg~~t~~ ggtattgg~~t~~ aattcgaagc cggtatctcc
 961 **aaggatgg**tc aaaccagaga acacgcttta ttagcttaca ccttaggtgt taagcaattt
 1021 **attgt**tgcta tcaacaagat ggactctgtc aaatggaca aggccagatt cgacgaaata
 1081 **gtca**aggaaa cctctaaact cgtcaagaag gtcggttca accccaaagag t~~gtt~~ccattc
 1141 **gtccc**aatct ctgggtggaa cggtgacaac atgattgagc catcctctaa ctgtccatgg
 1201 **taca**agggtt gggaaaagga aaccaaggct ggttaagtctt ctggtaagac tttgttagaa
 1261 **gctatt**gatg ctatcgaaacc accaaccaga ccaactgaca aggcttaag attaccattt
 1321 **caagatgt**tct acaagatcg~~t~~ tggtat~~gg~~ actgtgccag t~~cg~~gttagagt tgaaaccggt
 1381 **atcat**caagg cccgtatgg~~t~~ tgcac~~tt~~c gccccagctg g~~t~~gttaccac tgaagtcaag
 1441 **tccgtt**gaaa tgcac~~t~~ac~~g~~ acaattaacc gaagg~~t~~gttc caggtgacaa cgttgg~~tt~~c
 1501 **aacgt**caaga atgttccgt taaggaaatc agaagaggta acgtctgtgg tgactccaaag
 1561 **aacgaccc**ac caaagggtgc tgactt~~tt~~cc accgctcaag ttattgtctt aaaccatcca
 1621 **ggtca**aatct cctctggta ctctccagtc ttagattgtc acactgtc~~a~~ cattgtt~~gt~~
 1681 **aagtt**cgaca cttaatcg~~a~~ aaagattgac agaagaactg gtaagaagtt agaagacaac
 1741 **ccta**agttca tcaagtctgg tgatgtct~~gt~~ attgtcaaga tggteccatc taagccaaatg
 1801 **tgtgtt**gaaag ctttca~~t~~ctga ctacccacca ttaggttagat t~~cg~~ctgtgag agatatgaga
 1861 **caaactgtt**g ctgtcggtgt tatcaagtct~~t~~ gttaaaag~~t~~ ctgacaagtc cggtaagg~~tt~~
 1921 **acta**agg~~t~~gtc ctcaaaaggc tgctaa~~g~~aaa taagctc~~tt~~ aattt~~g~~aaat t~~gt~~ccggag
 1981 **aagctact**tg aattt~~g~~aaat~~t~~ acgtaaaat~~t~~ c~~g~~agttttt~~t~~ c~~o~~ctacagtt agaattctgt
 2041 **ttat**ttt~~g~~at a~~g~~taat~~ttt~~ atactt~~ta~~at g~~t~~taat~~at~~gt tgatctaaac g~~c~~tcc~~ttt~~gt
 2101 **agga**acctat ccaattt~~g~~aa c~~g~~caaactct attcg~~g~~actg c~~g~~atatacac agt~~tt~~ccgtt
 2161 **gac**ctcaato gtcggatatt atgtcoaaca tcagtcgat~~g~~ c~~at~~gaacac~~a~~ aaaagg~~gt~~tc
 2221 **cgaaaa**at~~t~~ttt ccttat~~ca~~ag ccacat~~gt~~ga aaagtc~~gt~~cc tcaat~~g~~ac~~g~~a g~~t~~caaagatc
 2281 **tgata**agccc accaaactt~~a~~ tattaagatc cctgttata~~t~~ gaagttccat tttt~~cacata~~
 2341 **ttctt**taaca cagctgg~~t~~ac cctcaactgt ttcaaaactt~~t~~ gagatggatt agtattgcgt
 2401 **tattt**gat~~gt~~ agaaaag~~t~~ta g~~t~~aaaaatc~~t~~ atcattgaac c~~g~~agtcataa tctggggaaa
 2461 **cagagtc**agt aaaataccgt g~~t~~taat~~t~~ttt ttcatt~~g~~ac tctgag~~t~~gtt acaactttcg
 2521 **tatt**tgct~~a~~ tgaagt~~t~~ttt aaaccat~~cg~~a gaat~~c~~ttttt~~t~~aaaat~~g~~acaa taaat~~g~~at~~t~~
 2581 **ttaat**atcat t~~c~~at~~ttt~~ttg ataaat~~aca~~aa tataat~~cata~~ gaattaat~~t~~ ttttt~~gtttt~~
 2641 **ataa**acccaa aaaaat~~aga~~ga agaggat~~gt~~at ttcaag~~g~~at~~t~~ gtt~~g~~caagat cggctcg~~gt~~tc
 2701 **taat**agatt~~t~~ ttac~~tt~~ga ccagat~~ct~~gt tccaaagg~~t~~ta tacaattt~~g~~a acttggaaaat
 2761 **tgc**atcg~~gt~~ agcagat~~ttt~~ tagctt~~ca~~aa g~~cc~~gcacaca t~~c~~at~~c~~agc~~g~~at~~c~~ attatgaaaa
 2821 **agttt**ac~~gt~~at cca~~g~~tgcc~~a~~ agactgt~~gt~~tc tttgg~~g~~caat ttggata~~acgt~~ t~~g~~cc~~c~~agac~~g~~
 2881 **tc**at~~tt~~gt~~gt~~ cctacccctg~~t~~ acaat~~gt~~ca~~a~~ gac~~g~~at~~gt~~tg agtac~~ctt~~gg gata~~c~~aa~~g~~ga
 2941 **tttgg**ac~~g~~ag tttt~~g~~ag~~ca~~ aagccat~~ccc~~ agagcac~~gt~~t~~t~~ ttatac~~aaa~~ gggcattaca
 3001 **gatt**caac~~cc~~ caacaagg~~t~~tt acact~~g~~ag~~c~~t g~~g~~aaaat~~gt~~tt~~g~~ g~~a~~ac~~at~~tt~~g~~c aaaaatt~~gg~~c

Figure S1: Nucleotide sequence of the *D. macquariensis* strain D50 *TEF1* gene (bold text) with 5' and 3'-flanking regions. The putative promoter sequence is red.

1 cactactcag acttaccacc gcatatacag acattaacca taatagtgcc atctacaggg
61 cacaatgcat ttaatacaca atctaataata cagcttaagt agagccatc tgtaattaa
121 cccacacaaa gagtaatcc gteacatata ctaatgtgaa cgccattcat ttgacctagt
181 gtcacatgg caaaagctt cgccgcgtt tcgcaatttc actacatgtc gcctgtcatt
241 agtacgaaac tgctatgtaa gatcaactgc aaaatgtcc actggccat aatcgaaagg
301 tctaggagtt tgccaaaaaa tttagtccca tactatccca accaaatgtc ctcaaataac
361 taccatcaca cctatttgc caagtttcc cttatctactata tagctact tcatgctgt
421 tgcataatccg tagtctacgg ccatagoact tggattctga atacttagag gaaataggg
481 aaagccctat cgggtaaat taaatatttg tctgtccaa aaaaaaaaaatt tggccacgac
541 accctctgtat gaatggaaacg caattggctc tgcaaaaatt tataaggaat taaattgaag
601 aacctttttt tccctctt ttttcaata cagtatacta ttattataca taaaacaaaaa
661 tggtaaaga aaagactcac gttaatcttg ttgtcattgg tcacgtcgat tccggtaaat
721 ctaccactac cggtcaactt attacaagt gtgggttat cgataagaga actattgaaa
781 agttcgaaaa ggaagccgct gaatttaggtt agggttctt caagtacgct tgggttttag
841 acaagttaaa ggctgaaaga gaaagaggtt tcaccattga tatcgcttta tggaaagttc
901 aaactccaaa gttccacgtc accattattt atgctccagg tcacagagat ttcatcaaga
961 acatgattac tggtaattcc caagctgatt gtgttattt gattattgtt ggtggattt
1021 gtgaattcga agctggtatac tccaaaggatg gtcaaaaccag agaacacgct ttattagctt
1081 acaccttagg tgtaagcaa ttgattgtt cttatcaacaa gatggactct gtcaaatggg
1141 acaaggccag attcgacgaa atcgtaagg aaacctctaa cttcgtaag aaggtcggtt
1201 tcaacccaaa gagtgttcca ttctgtccaa tctctgggtt gaacggtgac aacatgattt
1261 aaccatcatc taactgtcca tggtacaagg gttggggaaaa ggaacccaag gctggtaagt
1321 cctctggtaa gactttgttta gaagctattt atgctatcga accaccaacc agaccaactg
1381 acaaggctt aagattacca ttgcaagatg tctacaagat cgggtgttatt ggtactgtgc
1441 cagtcggtag agttgaaacc ggttattatca aggccggat ggttgcacc ttggccccag
1501 ctgggtgtcac cactgaagtc aagtccgtt aatgcatca cgaacaattt accgaagggt
1561 ttccaggtga caacgttggt ttcaacgtca agaacgttcc cgttaaggaa atcagaagag
1621 gtaacgtctg tggtaactcc aagaatgacc caccaaaaggg tgctgactct ttcaccgctc
1681 aagttattgt cttaaaccat ccaggtcaaa tctctctgg ttactctcca gtcttagatt
1741 gtcacacacgc tcacattgtt tgtaagttcg acactttat cggaaaggatt gacagaagaaa
1801 ctggtaagaa gttagaagac aaccctaaat tcatcaagtc tgggtatgtt gctattgtca
1861 agatggttcc atctaagcca atgtgtgtt aagcttccac tgactacccca ccatttaggt
1921 gattcgctgt cagagatatg agacaaaactg ttggcgtccg tgttatcaag tctgttgaga
1981 agtctgacaa gtccggtaag gttaccaagg ctgctaaaaa ggctgctaaag aaataaagctg
2041 attaatgttag aattttat tggtaacttg ttttatatta ggcttttga atacatatga
2101 actttaaaaa tgattgttgc ctccgataag cactagctt ttgtctctgt aaatcggtt
2161 gttggcgtggc caagtatctg gecgtatgcac attttagtgcgtaat aagactatt taatagaatt
2221 tgaacttcac actggcgaac cctacttagtt aacgcactt tatacttttcccgagtaat
2281 aacaaagaac gtacgttgc ttataatcaa catatgcacat atcacccttccatggctttag
2341 gtattaaatta atagcatgtt gaaaattttt ttagcacaat agatactgtt tacaaccaga
2401 tatcatgttc aactggaaat agatacgtt aatgaactt tggtatgatgaa gggcttattt
2461 gacatagaac cgtgttgcg ataaac

Figure S2: Nucleotide sequence of the *D. macquariensis* strain D50 *IEF2* gene (bold text) with 5' and 3'-flanking regions. The putative promoter sequence is pink.

1 **tcgtacgtta** cattatatac actccatat atgaagtaga ccaattcttt aacactgtga
 61 ccaacatcg **ttctgtacgt** ctattggcg aattcgccga aatatacata ctccctgtat
 121 atggtcagaa **tcttcacate** ggtgttatta ggcgacaccg aaacccttaa acccccgett
 181 cgcaactcggt cggtcgtgtc tggtaacaa atgcgttacg ctacaccat tcgtaaattc
 241 caaacgtcat tccataggtc gggtccgacg aaagcctcac ttccgoacac cacatccatt
 301 catcaggata tgacttcgcg cagacgtcat cgaccctgg ggctcggtt cggtttcgc
 361 ttcaatccca gggtgccg **cgccatccca** ctctccggga cgaccatca taccacaatt
 421 caagacggcc **cttatgtgc** tggttccaa cgtgtccta tcgatcata atccacccct
 481 gggttgtct caatcgccag atttacgggt tctcttaat tgcgcctaca ttttgttgc
 541 tcacgttcgt tgctctttt ctacatggac gagttgtgc caacgaccat ttttcacggt
 601 atcaagtggt ggtaggcctc cggcgaccc tggacggctc gtttcgtt atccattccat
 661 actaattaac ggcttcggac ttttgttgcg cgaatcatg agaatcaca aactccggc
 721 aactagctcg gtcgtcccc gcaccggggc ctctatggg ttcttcgccc cgggagtcca
 781 gcaataagaa catagtggaa aattataat atgagaatt cggcgtgcaa tgcacatgga
 841 gtatattta aaggttatct tattattcaa ttgaatataa ttaactagta agaatattta
 901 ataattaaac **atggctatca** ctattggat taacggtttc ggtcgattt gacggtttagt
 961 **cttaagagtc** gcttttagaaa gatctgacat caaggttggt ggcgtcaatg atccattcat
 1021 tgccgctgaa tacgtgtttt acatgttcaa gtacgattct actcacggta gatacaaggg
 1081 tgatgtaa **gc**ccaaggacg acaccttagt catcgacggt caatctatca aggttttcgg
 1141 tgaaagagac ccagecctgca ttccatgggg taaggtcgggt gttgactacg tcacgaaatc
 1201 caccgggtgc ttcaccacta tggaaagggtgc taaaaagcac attgacggtg gtggcaagaa
 1261 ggtcattatc actgttccat ctcagatgc tccaatgttc gttgttgggt ttaacgaaaca
 1321 aaaatacacc ccagacttga agattgttcc caacgcttct tgcgtactacta actgtttggc
 1381 tccattagcc aaggttatca acgacacett cgttattgaa gaaggtttaa tgaccactgt
 1441 ccactccatc actgccaccc aaaagacccgt tgatggtcca tcccacaagg actggagagg
 1501 tggtagaacc gcttctggta acattatccc atttccacc ggtgctgcca aggcggtcgg
 1561 taaggttatt ccagaattaa acggtaagtt gaccggatgt tctttgagag ttcctactgt
 1621 cgatgttca gttgttggatt taaccgtcag attgaagaaa cctgtctaccc acgaagaaat
 1681 ctccgaagct atcaagtctg cttctaacgg tgaattaaag ggtgttttag gttacactga
 1741 agatgttgc gtttccaccc atttcttagg ttccagctac tcttctatct tcgatcaaaa
 1801 ggccggtatac ttattatccc caaccttcgt caagttgatc tcttggatc ataaacgaaatt
 1861 cggttactcc accagagttt tcgattttt agaacacgtc gccaaaggctt ctaattaact
 1921 agccgaccct aaactataat ttatttatgg ttatcgcc ctagcatata tatgcgaat
 1981 aaaatgtatt tattttaa **aca**ttttaaaaact tttatgttgc gcgagtttgg catgtatgt
 2041 aattcttagg ctcgtatggg tttccaaacct tagctgggtt aggtgtgtatg tacattttaaa
 2101 ctatattaa **c**gttctaata tcttgcgtt atttattttt aaatattcac atattttttt
 2161 aaattctata catatacatt atcactatca catacattat cactattata tacacagtgc
 2221 atatttcctt attacctaaa tttgacactg ctccctgtt ttacaacggt ttctttcat
 2281 ggacaaattt ttcttagatc aaacatcaa attataaaaa tcaaatgata ctgtacctca
 2341 ctgtactcaa caacaattgg actatatctc aacaataaca tgcataattt tattccaaac
 2401 gttggtaat gatatgatc cagattacag atcatttgag gatttactcc tagattttaa
 2461 aatcactcat agtcgagac agttaatca attacaatc acatataattt ccataagaca
 2521 caaggattt ttgcctcgac aaaccataat attcatcaa gctactgata ctaacgacaa
 2581 cgcacatca ttcaatacgt tggtgtatgaaa aaaaacaacg aactcatct tattaaggat
 2641 tctaaattcag gcgattgaaa atctacatc agaagttccc ataatagtac gggattttaaa
 2701 attcaacgag aaatatctca acttaattat aaataatgtt tctaaacagct tattatctat
 2761 agataattat gctgagggtca taggtgacgt tgaatttgatt tatcaaacta gccacctcg
 2821 atcgaacgt tcattaaagaa acatcatcat caatattccc gaaaaagatg taagtattat
 2881 atcggattcc aaaggactca ttaataacgt caatgatttcc ttgaaacaaa cgacgctgtt
 2941 aaattttat aaccttaccc ttgattcggtt cacttcaagg

Figure S3: Nucleotide sequence of the *D. macquariensis* strain D50 GAP gene (bold text) with 5' and 3'-flanking regions. The putative promoter sequence is green.

1 aggaattaca aatagaagcc ataaatatat tcattctgt atataaatat tagccaataa
61 aattgcattc gtaacaatt atgagaaatt tttcaccttg attggacat atataaaagg
121 aagctttctg gcccattca aaacaagtgc ttaattccca taatataatc ttttatattt
181 taaccagggtt tcttaacata tatcatgcca **gttccatttg** **aaaaagggttc** **cgaaaagaag**
241 **gttgctaact** **tatTTAAAC** **cagatTTA** **caatgtcaca** **ctgttgaaga** **aggtggtcca**
301 **cacaaggTTG** **gtccaaactt** **acacggTATT** **attggtagaa** **catctggtca** **agccgctggT**
361 **tactcttaca** **ctgatGCCAA** **taAAAAGAAG** **ggtgttcaat** **ggactgaaca** **aactttatcc**
421 **gactacttgg** **aaaACCCAAA** **gaaatatacC** **ccaggtacca** **agatggcTT** **cggTggTTA**
481 **aagaaaccta** **aggacagaaa** **cgatttgatc** **acctatatac** **ccgaagotac** **caaataaacc**
541 **atttatttgc** **tgttggaaag** **tttactcgTC** **gttttgtaca** **aacactgttt** **aatgggcctc**
601 **gaatTTtagCT** **agttacacat** **ttatctgtat** **atacatgaat** **aatattaacg** **aacgatgtta**
661 **gattaaatgc** **ttacagttaga** **gtacaaggcag** **tgggggttat** **atTTggatgg** **acattatctg**
721 **ttcacgtgac** **ccagttatgag** **tgaaaagtTC** **gccaagtcaa** **ccactgttatt** **acagacggca**
781 **attatattac** **aatttagtggc** **aac**

Figure S4: Nucleotide sequence of the *D. macquariensis* strain D50 CYC1 gene (bold text) with 5' and 3'-flanking regions. The putative transcriptional terminator sequence is purple.

1 atatttatta aaagaaaaaaga aataagatcaa gattaatcca aggttgttta aagaagacaa
61 agacatgtaa cgaggtaat ttataataaa aaaaagaaaa tatgtattt aaagtataca
121 taaaattgtt gaaggatgc gtttagaaag gagctgaatg taacgagatg cattcaatt
181 caaagacaga gagttttac atttatcta gacgaattag gatgttcat tgtagtagat
241 ttctctgatt tggtgcattt gttgtttat tggtcaaaa tagtgaaaaa gaggggaaaga
301 gggaaagataa acattatatg caggtcactt aaacacttga taaagtcatc ataaaacaac
361 ctaaatcaac tatttgatc caaaattttt cataaattac ataaattact ttttggaaatt
421 cttcaatatac aggttcaaa taataaaaaat tagggtagc gtagataaaa cgaatattgt
481 aggtttatgt tggcgtaa attacacaaa gttgagaaga taagatgcg tacaaggatg
541 aaaaaataaa aaattattgt tctgaagatt ttttttttgg gagatttcgg aaaaattttg
601 ggtctcatcg acttatcgat ccggccaaagt gtaacctaaa tagtggaaacc taaaacccaa
661 taactactat gttatagtaa agacgattga gtaagtcaagg attggataaa aaaaataatt
721 gaatcaaata ctctacatga atttattttc cctcttaatg ctcatttetc attttatatt
781 tacatttata tttcataacc tctttttaa cttctcttacttactt ctttcccta
841 cacggttatt gtacgtgtt attccggat ttaaatttgc tatgtgaaa attacgadag
901 aatgttttttta ttgecgctg tatttcaat tatataatgat tattttagca tgcgtcgacc
961 gcacaaaaacc ctcatcgat catattttac ccataactatg tgaaaattttt tttttttt
1021 gctctgcggg tgagggggcg ttggccaaata tataaggatg gcccctgtc ttagacaaaa
1081 attttccagg ttataagttt ttttttttcc ttcttctttt acttcatcta aactaaactt
1141 aattaaaaat gggtaaagat aagaatcagc ttaacgtcgt cgtcatttggc cacgtcgatt
1201 cccgttaagtc aaccactact ggtcactttaa ttttcaagtg tggtgttatt gataagagaa
1261 ctattgaaaaa gttcgaaaag gaagctaatg aatttagaaaa aggttcttcc aagtacgctt
1321 gggttttaga caagcttaag gctgaaaagag aaagaggat cactattgtat attgtttat
1381 ggaagttcga aactccaaag tatacgtta ctgttattga tgctccaggt cacagagatt
1441 tcattaagaa tatgattact ggtacttetc aagctgatttgc tgcttttttattattgtct
1501 gtggtattgg tgaattcga gcccgttattt ctaaggatgg tcaaaaccaga gaacatgctt
1561 tattagcttta caccttaggt gttaaacaat taattgttgc tgtcaacaag atggattccg
1621 tttaaatggga taaggctaga tttgaagaaaa tttcaaaagga aacctctaat ttcgttaaga
1681 aggttggta caatccaaag actgttccat tggttcaat ctctgggttgg aatgggtgata
1741 acatgattga accatctgtt aactgttccat ggtacaaggg ttggaaaag gaaaccaagg
1801 cccgttaaggt cactggtaag actttatttag atgttattga tgcttattgaa ccacctcaaa
1861 gaccaactga taagccatta agattaccat tacaagatgt ttacaagatt ggtggattt
1921 gaactgtgcc agttggtaga gttgaaaccgg tatacattaa ggctggatg gttgtgttt
1981 tgcggccagc tggtgttacc actgaagtca agtccgttga aatgtatcat gaaacattag
2041 ttgaagggtgt tccagggtac aatgttggtt tcaatgttaa gaatgtttcc gttaaaggaaa
2101 tttagaagagg taacgtttgt ggtgacaccca aggttgcatttcc accaaaggct gctgtttcat
2161 tcaatgttca agttattgtc ttaaaccatc caggtcaat ttctgttgc tactctccag
2221 ttttagatttgc tcaacccggcc catattgtttt gtaagttcga tcacttagtt gaaaagattt
2281 atagaagaac tggtaagtca atgaaagatg aacccaaattt catcaagtcc ggtgtgtct
2341 ctattgttag aatggtccccca tccaaagccaa tgggtgttgc agtcttactt gactacccac
2401 cattaggttag attcgttgc agagatatga gacaaactgt cgctgtcggt gttatcaagt
2461 ctgttgacaa gtcggacaaag gttggtaagg tcaccaagtc tgctttaag gtcgtcaaga
2521 agtaaacaac ctgttttacc tttgtatgtaa gtatgttttcc tttgactaat gtatgaaatt
2581 aaatataatataaaaaaaag tagaaatgag cgtattatgt aactgtttgc tgccaaaaat
2641 cgtattatag gtgaattact gtgaaatatg tcttaatatt tatgtcaactc gagccacaaa
2701 tggatgtt gattttataa aacaaaagag tataatgttga tatttaggtt aaaaacaaaat
2761 ttggcatgtt aagggttcttca agagggtttat atctatgtcac ggagaaaaaaa gttaaaagaag
2821 ttcatatgtt tgcagcgtt aattttttgc aataaaatca atcgttcaat gtaacccat
2881 ttatgttca gtaattatataa taaagggttac ctacagctga gttggatttac gttggaaatgt

Figure S5: Nucleotide sequence of the *C. santamariae* strain G12 *TEF1* gene (bold text) with 5' and 3'-flanking regions. The putative promoter sequence is blue.

1 tttacgtctg ttttatattt agcggactta tcgggttag gcgggcacaa tcggaaactc
 61 gaaccaaaca atttctact cgaccgttc gacgaataag aattaaaaaa atagtaattt
 121 gacgtgtata taaccgttt tagatgatta aaaattcat gttagcgggt aattttgtgt
 181 aataatagat gattctaata tataagtgaa aagttccag aatgttattt tttgcgggt
 241 ttcaaccttc tttttttt cttttttct cctaacatac atacatacaa ttaat**catgc**
 301 **cagctccatt** tgaacaagg tccgaaaaga agggtgtac tcttttaaa actagatgtt
 361 **tacaatgtca** caccgttgaa aaagggtggtc caaacaagg tggtccaaat ttacatggtg
 421 **tctttggtag** aaaatctggt ttagcagatg gtttcttta tactgatgct aacaagaaga
 481 **agggtgttac** ttggtctgaa caaaccatgt ctgattactt agaaaaaccca aagaagtata
 541 **ttccaggtac** caagatggct ttccgggtt taaagaaacc aaaagacaga aacgatttag
 601 **tcacttactt** agcaagtgc accaaataga tttagaacac tctagataag caaaagcaaa
 661 ataaaagcga aatgctctgt atataataat ttatgttttta ttatgtata gagttagag
 721 agtgagagag attgtaaagcc cccagaaaaat gaagaaaaat atttattgtt gaagatgaga
 781 atactttat gagaccctcc atgggttgct ttatgcagg gagatgcagt tttgcattgag
 841 atcgagatta cttatcctag aagttgatt tataataggc gtatttaggg tagaaaaatag
 901 cactttta agaataactt gaataatctc cattaatcaa aaacacaaac attaactaag
 961 cgtgtcatat ttacttttt tctataatca tctctatcgc ctttttcttt ttttctcag
 1021 attctatttc tcaaacacac totctctgac tctacttca actttaactc tttgacttgt
 1081 **ttcatctgaa** aaatcttgc tatgtcaac tagtgtttcg atatatcgat ctgtaccc
 1141 **ttgactaacc** gctctttcac ttcttttcg tcaaacttcg cttacattat aat

Figure S6: Nucleotide sequence of the *C. santamariae* strain G12 CYC1 gene (bold text) with 5' and 3'-flanking regions. The putative transcriptional terminator sequence is brown.

Table S1. Extracellular proteolytic activity and concentration of extracellular proteins of yeast strains isolated from soil samples collected from various sampling sites in Antarctica¹ after 72 h of cultivation.

Yeast strain	Proteolytic activity at pH 7.0 (nmol/mL × min)	Proteolytic activity at pH 4.0 (nmol/mL × min)	Extracellular proteins concentration (mg/mL)
<i>Debaromyces macquariensis</i> D50	6.30	7.50	0.030
<i>Debaromyces hansenii</i> IBT-D1	6.97	3.20	0.032
<i>Goffeauzyma gastrica</i> IBT-D7	21.11	8.68	0.060
<i>Goffeauzyma gilvescens</i> IBT-D13	16.18	8.56	0.096
<i>Goffeauzyma gilvescens</i> IBT-D15	11.45	12.36	0.112
<i>Goffeauzyma gilvescens</i> IBT-D16	11.84	14.54	0.136
<i>Goffeauzyma gilvescens</i> IBT-D20	15.98	24.47	0.122
<i>Goffeauzyma gilvescens</i> IBT-D60	13.02	15.79	0.178
<i>Leuconeurospora</i> sp. IBT-D59	38.63	17.70	0.362
<i>Naganishia globosa</i> IBT-D37	17.76	8.90	0.158
<i>Naganishia adeliensis</i> IBT-D58	11.45	12.43	0.215
<i>Naganishia albida</i> IBT-D62	27.60	18.30	0.329

<i>Rhodotorula mucilaginosa</i>			
IBT-D12	28.22	17.50	0.150
<i>Rhodotorula mucilaginosa</i>			
IBT-D34	18.55	12.17	0.105
<i>Rhodotorula mucilaginosa</i>			
IBT-D38	18.29	8.29	0.099

¹ Białykowska, A.M.; Szulczewska, K.M.; Krysiak, J.; Florczak, T.; Gromek, E.; Kassassir, H.; Kur, J.; Turkiewicz, M. Genetic and biochemical characterization of yeasts isolated from Antarctic soil samples. *Polar Biol.* **2017**, *40*, 1787-1803, doi: 10.1007/s00300-017-2102-7.