

Secuencias

SEQ ID NO:1 (*Blautia stercoris* cepa GAM6-1 gen de ARN ribosomal 16S, secuencia parcial - HM626177)

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1 tgcaagtcga gcgaagcgct tacgacagaa ccttcggggg aagatgtaag ggactgagcg
5 61 gcggacgggt gagtaacgcg tgggtaacct gcctcataca gggggataac agttggaaac
121 ggctgctaata accgcataag cgcacggtat cgcatagatac agtgtgaaaa actccggtgg
181 tatgagatgg acccgctct gattagctag ttggaggggt aacggccac caaggcgacg
241 atcagtagcc ggctgagag ggtgaacggc cacattggga ctgagacacg gccagactc
301 ctacgggagg cagcagtggg gaatattgca caatggggga aaccctgatg cagcgacgcc
10 361 gcgtgaagga agaagtatct cggtatgtaa acttctatca gcagggaaga aaatgacggt
421 acctgactaa gaagccccgg ctaactacgt gccagcagcc gcgtaatac gtagggggca
481 agcgttatcc ggatttactg ggtgtaaagg gagcgtagac ggaagagcaa gtctgatgtg
541 aaaggctggg gcttaacccc aggactgcat tggaaactgt tttcttgag tgccggagag
601 gtaagcggaa ttctagtgt agcggtgaaa tgcgtagata ttaggaggaa caccagtggc
15 661 gaaggcggct tactggacgg taactgacgt tgaggctcga aagcgtgggg agcaaacagg
721 attagatacc ctggtagtcc acgccgtaaa cgatgaatac taggtgttg ggagcaaagc
781 tcttcggtgc cgcagcaaac gcaataagta ttccacctgg ggagtacgtt cgcaagaatg
841 aaactcaaag gaattgacgg ggacccgcac aagcggtgga gcatgtggtt taattcgaag
901 caacgcgaag aaccttacca agtcttgaca tcgatctgac cggttcgtaa tggaacctt
20 961 ccttcgggac agagaagaca ggtggtgcat ggtgtcgtc agctcgtgc gtgagatgtt
1021 gggtaagtc ccgcaacgag cgcaaccct atctcagta gccagcaggt gaagctgggc
1081 actctgtgga gactgccagg gataacctgg aggaaggcgg ggacgacgtc aaatcatcat
1141 gcccttatg atttgggcta cacacgtgct acaatggcgt aaacaaaggg aagcgagccc
1201 gcgaggggga gcaaatccca aaaataacgt ccagttcgg actgcagtct gcaactcgac
25 1261 tgcacgaagc tggaatcgct agtaatcgcg aatcagaatg tcgcggtgaa tacgttcccg
1321 ggtctgtac acaccgccg tcacaccatg ggagtcagta acgcccgaag tc

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SEQ ID NO:2 (*Blautia wexlerae* cepa WAL 14507 gen de ARN ribosomal 16S, secuencia parcial - EF036467)

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30 1 caagtcgaac gggaattant ttattgaaac ttcggtcgat ttaatttaat tctagtggcg
61 gacgggtgag taacgcgtgg gtaacctgcc ttatacaggg ggataacagt cagaaatggc

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121 tgctaatacc gcataagcgc acagagctgc atggctcagt gtgaaaaact ccggtggtat
 181 aagatggacc cgcgttggat tagcttgttg gtggggtaac ggcccaccaa ggcgacgac
 241 catagccggc ctgagagggg gaacggccac attgggactg agacacggcc cagactccta
 301 cgggaggcag cagtggggaa tattgcacaa tgggggaaac cctgatgcag cgacgccgcg
 5 361 tgaaggaaga agtatctcgg tatgtaaact tctatcagca gggaagatag tgacggtacc
 421 tgactaagaa gccccggcta actacgtgcc agcagccgcg gtaatacgta gggggcaagc
 481 gttatccgga ttactgggt gtaaaggag cgtagacggg gtggcaagtc tgatgtgaa
 541 ggcatgggct caacctgtgg actgcattgg aaactgtcat acttgatgc cggaggggta
 601 agcgggaatt ctagttagc ggtgaaatgc gtagatatta ggaggaacac cagtggcgaa
 10 661 ggcggttac tggacggtaa ctgacgttga ggctcgaaag cgtggggagc aaacaggatt
 721 agataccctg gtagtcacg ccgtaaacga tgaataacta ggtgtcgggt ggcaaagcca
 781 ttcggtgccg tcgcaaacgc agtaagtatt ccacctgggg agtacgttcg caagaatgaa
 841 actcaaagga attgacgggg acccgacaaa gcggtggagc atgtggttta attcgaagca
 901 acggaagaa ccttaccag tctgacatc cgcctgaccg atccttaacc ggatcttcc
 15 961 ttcgggacag gcgagacagg tgggtcatgg ttgtcgtcag ctcgtgtcgt gagatgttg
 1021 gtaagtccc gcaacgagcg caaccctat cctcagtagc cagcatttaa ggtgggcact
 1081 ctggggagac tgccagggat aacctggagg aaggcgggga tgacgtcaaa tcatcatgcc
 1141 cttatgatt tgggctacac acgtgctaca atggcgtaaa caaagggaag cgagattgtg
 1201 agatggagca aatcccaaaa ataacgtccc agttcggact gtagtctgca acccgactac
 20 1261 acgaagctgg aatcgctagt aatcgcggat cagaatgccg cgtgaatac gttcccggt
 1321 cttgtacaca ccgcccgtca caccatggga gtcagtaacg cccgaagtca gtgacctaac
 1381 tgcaaagaag gagctgccga aggcgggacc gatgactggg gtgaagtcgt aacaaggt

SEQ ID NO:3 (secuencia de consenso de ARNr 16S para *Blautia stercoris* cepa 830)

25 TTTKGTCTGGCTCAGGATGAACGCTGGCGGCGTGCTTAACACATGCAAGTCGAGCGAAG
 CGCTTACGACAGAACCTTCGGGGGAAGATGTAAGGGACTGAGCGGCGGACGGGTGAGT
 AACGCGTGGGTAACCTGCCTCATACAGGGGGATAACAGTTGGAAACGGCTGCTAATACC
 GCATAAGCGCACAGTATCGCATGATACAGTGTGAAAACTCCGGTGGTATGAGATGGACC
 CGCGTCTGATTAGCTAGTTGGAGGGGTAACGGCCCAACCAAGGCGACGATCAGTAGCCGG
 30 CCTGAGAGGGTGAACGGCCACATTGGGACTGAGACACGGCCCAGACTCCTACGGGAGG
 CAGCAGTGGGGAATATTGCACAATGGGGGAAACCCTGATGCAGCGACGCCGCGTGAAG

GAAGAAGTATCTCGGTATGTAACTTCTATCAGCAGGGAAGAAAATGACGGTACCTGACT
 AAGAAGCCCCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGGGCAAGCGTTA
 TCCGGATTTACTGGGTGTAAAGGGAGCGTAGACGGAAGAGCAAGTCTGATGTGAAAGGC
 TGGGGCTTAACCCCAGGACTGCATTGGAACTGTTTTTCTTGAGTGCCGGAGAGGTAAGC
 5 GGAATTCCTAGTGTAGCGGTGAAATGCGTAGATATTAGGAGGAACACCAGTGGCGAAGG
 CGGCTTACTGGACGGTAACTGACGTTGAGGCTCGAAAGCGTGGGGAGCAAACAGGATTA
 GATACCCTGGTAGTCCACGCCGTAAACGATGAATACTAGGTGTTGGGGAGCAAAGCTCTT
 CGGTGCCGCAGCAAACGCAATAAGTATTCCACCTGGGGAGTACGTTTCGCAAGAATGAAA
 CTCAAAGGAATTGACGGGGACCCGCACAAGCGGTGGAGCATGTGGTTTATTCTGAAGCAA
 10 CGCGAAGAACCTTACCAAGTCTTGACATCGATCTGACCGGTTTCGTAATGGAACCTTTCCTT
 CGGGACAGAGAAGACAGGTGGTGCATGGTTGTCGTCAGCTCGTGTCGTGAGATGTTGGG
 TTAAGTCCCGCAACGAGCGCAACCCCTATCGTCAGTAGCCAGCAGGTAAAGCTGGGCAC
 TCTGAGGAGACTGCCAGGGATAACCTGGAGGAAGGCGGGGACGACGTCAAATCATCATG
 CCCCTTATGATTTGGGCTACACACGTGCTACAATGGCGTAAACAAAGGGAAGCGAGCCC
 15 GCGAGGGGGGAGCAAATCCCCAAAATAACGTCCCAGTTCGGACTGCAGTCTGCAACTCGA
 CTGCACGAAGCTGGAATCGCTAGTAATCGCGAATCAGAATGTCGCGGTGAATACGTTCCC
 GGGTCTTGTACACACCGCCCGTCACACCATGGGAGTCAGTAACGCCCGAAGTCAGTGAC
 CCAACCTTAGGGAGGGAGCTGCCGAAGGCGGGATTGATAACTGGGGTGAAGTCTAGGG
 GGT

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SEQ ID NO:4 (secuencia de consenso de ARNr 16S para *Blautia wexlerae* cepa MRX008)
 TTCATTGAGACTTCGGTGGATTTAGATTCTATTTCTAGTGGCGGACGGGTGAGTAACGCG
 TGGGTAACCTGCCTTATACAGGGGGATAACAGTCAGAAATGGCTGCTAATACCGCATAAG
 CGCACAGAGCTGCATGGCTCAGTGTGAAAACTCCGGTGGTATAAGATGGACCCGCGTT
 25 GGATTAGCTTGTTGGTGGGGTAACGGCCACCAAGGCGACGATCCATAGCCGGCCTGAG
 AGGGTGAACGGCCACATTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAG
 TGGGGAATATTGCACAATGGGGGAAACCCTGATGCAGCGACGCCGCGTGAAGGAAGAAG
 TATCTCGGTATGTAACTTCTATCAGCAGGGAAGATAGTGACGGTACCTGACTAAGAAGC
 CCCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGGGCAAGCGTTATCCGGAT
 30 TTAAGTGGGTGTAAAGGGAGCGTAGACGGTGTGGCAAGTCTGATGTGAAAGGCATGGGCT
 CAACCTGTGGACTGCATTGGAACTGTCATACTTGAGTGCCGGAGGGGGTAAGCGGAATTC

CTAGTGTAGCGGTGAAATGCGTAGATATTAGGAGGAACACCAGTGGCGAAGGCGGCTTA
 CTGGACGGTAACTGACGTTGAGGCTCGAAAGCGTGGGGAGCAAACAGGATTAGATACCC
 TGGTAGTCCACGCCGTAAACGATGAATACTAGGTGTCNNGGGAGCATGGCTCTTCGGTG
 CCGTCGCAAACGCAGTAAGTATTCCACCTGGGGAGTACGTTGCGAAGAATGAAACTCAA
 5 GGAATTGACGGGGACCCGCACAAGCGGTGGAGCATGTGGTTTAATTGGAAGCAACGCGA
 AGAACCTTACCAAGTCTTGACATCCGCCTGACCGATCCTTAACCGGATCTTTCCTTCGGG
 ACAGGCGAGACAGGTGGTGCATGGTTGTCGTCAGCTCGTGTGTCGTGAGATGTTGGGTAA
 GTCCCGCAACGAGCGCAACCCCTATCCTCAGTAGCCAGCATTAAAGGTGGGCACTCTGG
 GGAGACTGCCAGGGATAACCTGGAGGAAGGCGGGGATGACGTCAAATCATCATGCCCT
 10 TATGATTTGGGCTACACACGTGCTACAATGGCGTAAACAAAGGGAAGCGAGATCGTGAGA
 TGGAGCAAATCCCAAAAATAACGTCCCAGTTCGGACTGTAGTCTGCAACCCGACTACACG
 AAGCTGGAATCGCTAGTAATCGCGGATCAGAATGCCGCGGTGAATACGTTCCCGGGTCTT
 GTACACACCGCCCGTACACCATGGGAGTCAGTAACGCCCGAAGTCAGTGACCTAACTG
 CAAAGAAGGAGCTGCCGAA

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SEQ ID NO:5 (*Blautia hydrogenotrophica* cepa S5a36 gen de ARN ribosomal 16S, secuencia parcial - X95624.1)

1 gatgaacgct ggcggcgtgc ttaacacatg caagtcgaac gaagcgatag agaacggaga
 61 ttccggttga agttttctat tgactgagtg gcggacgggt gagtaacgcg tgggtaacct
 20 121 gccctataca gggggataac agttagaaat gactgctaata accgcataag cgcacagctt
 181 cgcatagaag ggtgtgaaaa actgaggtgg tataggtatg acccgcggtg gattagctag
 241 ttggtgaggt aacggccac caaggcgacg atccatagcc ggcctgagag ggtgaacggc
 301 cacattggga ctgagacacg gcccacaaact ctacgggagg cagcagtgga gaattatgca
 361 caatggggga aacctgatg cagcgacgcc gcgtgaagga agaagtatct cggatatgaa
 25 421 acttctatca gcaggaaga aagtacggt acctgactaa gaagccccgg ctaattacgt
 481 gccagcagcc gcggaatac gtaaggggca agcggtatcc ggatttactg ggtgtaaagg
 541 gagcgtagac ggtttggcaa gtctgatgtg aaaggcatgg gctcaacctg tggactgcat
 601 tggaaactgt cagacttgag tgccggagag gcaagcggaa ttctagtgt agcggtgaaa
 661 tgcgtagata ttaggaggaa caccagtggc gaaggcggcc tgctggacgg taactgacgt
 30 721 tgaggctcga aagcgtgggg agcaaacagg attagatacc ctggtagtcc acgctgtaa
 781 cgatgaatac taggtgtcgg gtggcaaagc cattcgggtc cgcagcaaac gcaataagta

841 ttccacctg gggagtagt tcgcaagaat gaaactcaaa ggaattgacg gggacccgca
 901 caagcgggtg agcatgtgtt ttaattcgaa gcaacgcgaa gaaccttacc aaatcttgac
 961 atccctctga ccgggaagta atgtccctt ttctcgga cagaggagac aggtggtgca
 1021 tggttgctgt cagctcgtgt cgtgagatgt tgggttaagt cccgcaacga gcgcaaccct
 5 1081 tattcttagt agccagcagg tagagctggg cactctaggg agactgccag ggataacctg
 1141 gaggaagggtg gggatgacgt caaatcatca tgcccttat gatttgggct acacacgtgc
 1201 tacaatggcg taaacaaagg gaagcgaagg ggtgacctgg agcaaattc aaaaataacg
 1261 tctcagttcg gattgtagtc tgcaactcga ctacatgaag ctggaatcgc tagtaatcgc
 1321 gaatcagaat gtcgcggtga atacgttccc gggctctgta cacaccgccc gtcacaccat
 10 1381 gggagtcagt aacgcccga gtcagtgacc caaccnaag gagggagctg ccgaagggtg
 1441 gactgataac tggggtga

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