SEQUENCE LISTING

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<120> COMPOSITIONS COMPRISING BACTERIAL STRAINS

<130> P068687WO

<141> 2017-03-06

<150> GB 1612190.7
<151> 2016-07-13

<150> GB 1616018.6
<151> 2016-09-20

<150> GB 1616016.0
<151> 2016-09-20

<150> GB 1703548.6
<151> 2017-03-06

<150> GB 1703552.8
<151> 2017-03-06

<160> 5

<170> SeqWin2010, version 1.0

<210> 1
<211> 1372
<212> DNA
<213> Blautia stercoris

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ggctgctaat accgcataag cgcacggtat cgcatgatac agtgtgaaaa actccggtgg 180
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acctgactaa gaagccccgg ctaactacgt gccagcagcc gcggtaatac gtaggggcga 480
agcgttatcc ggatttactg ggtgtaaagg gagcgtagac ggaagagcaa gtctgatgtg 540
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<210> 2
<211> 1438
<212> DNA
<213> Blautia wexlerae

<220>
<221> modified base
<222> 19
<223> 'n' is a, c, g or t

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tgctaatacc gcataagcgc acagagctgc atggctcagt gtgaaaaact ccggtggtat 180
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gttatccgga tttactgggt gtaaagggag cgtagacggt gtggcaagtc tgatgtgaaa 540
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<210> 3

<211> 1481
<212> DNA
<213> Blautia stercoris

<220>
<221> modified base
<222> 4
<223> 'k' is g or t

<400> 3
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gcttacgaca gaaccttcgg gggaagatgt aagggactga gcggcggacg ggtgagtaac 120
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cggggacccg cacaagcggt ggagcatgtg gtttattcga agcaacgcga agaaccttac 960
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<210> 4
<211> 1384
<212> DNA
<213> Blautia wexlerae

<220>
<221> modified base
<222> 749
<223> 'n' is a, c, g or t

<400> 4
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<210> 5
<211> 1458
<212> DNA
<213> Blautia hydrogenotrophica

<220>
<221> modified base
<222> 1416
<223> 'n' is a, c, g or t

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gccctataca gggggataac agttagaaat gactgctaat accgcataag cgcacagctt 180
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gccagcagcc gcggtaatac gtaaggggca agcgttatcc ggatttactg ggtgtaaagg 540
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caagcggtgg agcatgtggt ttaattcgaa gcaacgcgaa gaaccttacc aaatcttgac  960
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