Microbiologics

CORPORATE HEADQUARTERS
200 COOPER AVE N, ST. CLOUD, MN 56303
320.253.1640 | MICROBIOLOGICS.COM

EU Declaration of Conformity (N° TD01)

Manufacturer	**European Authorized Representative**
Microbiologics, Inc.	MediMark® Europe Sarl
200 Cooper Avenue North	11, rue Emile Zola – BP 2332
Saint Cloud, MN 56303 USA	38033 Grenoble Cedex 2 – France
+1 (320) 253-1640	+33 (0)4 76 86 43 22
SRN: US-MF-000000466	SRN: FR-AR-000000182

Basic UDI-DI: See Annexes

Device Name: KWIK-STIK™, LYFO DISK™, QC Sets and Panels: KWIK-STIK™

Catalog Number(s): See Annexes

Intended Purpose: KWIK-STIK™ and LYFO DISK™ are intended to be external clinically-relevant viable control materials (as specified in Annex 1 by catalog number) to assist in identification of cultured microorganism isolates, and to verify the performance of assays, reagents, or media that are intended to be used in microbial testing. These products have no qualitative or quantitative assigned value. These control materials are nonautomated and not intended to be used for screening, monitoring, or diagnosis. These controls are not intended for any specific patient population or specimen.

Classification: Class B; rule 7

Notified Body: BSI Group The Netherlands B.V., Say Building, John M. Keynesplein 9, 1066 EP Amsterdam, The Netherlands

Notified Body ID: 2797

Certificate(s) Issued: EU Quality Management System, with IVDR 734881

Expiration Date(s) : 15/12/2026 (DD/MM/YYYY)

Conformity Assessment Route: EU Declaration of Conformity via Annex IX Chapter I Quality Management System, Chapter II Assessment of Technical Documentation, and Chapter III Administrative Provisions.

This declaration of conformity is issued under the sole responsibility of Microbiologics, Inc. and declares that the IVD medical device(s) specified above and listed in Annex 1 and Annex 2 are in conformity with the requirements of the Regulation (EU) IVDR 2017/746 on in vitro diagnostic medical devices. This declaration is valid for an undetermined length of time.

Name: Brian Carter

Function: Senior Regulatory Specialist

Signature: _____

Place and date of issue: Saint Cloud, MN, USA 14/08/2024 DD/MM/YYYY

STATE OF MINNESOTA
COUNTY OF STEARNS
Signed or attested before me on 14-Aug-24 (date) by Jennifer L. Rowan (names(s) of individual(s))
(stamp)

Jennifer Lynn Rowan
Notary Public
Minnesota
My Commission Expires Jan. 31, 2029

(Signature of notarial officer)

Notary Public
Title (and Rank)
My Commission Expires: 31-Jan-2029
dd/mm/yyyy

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Annex 1 to Declaration of Conformity

KWIK-STIK AND LYFO DISK CATALOG NUMBERS

KWIK-STIK		LYFO DISK	Product Description	Basic UDI-DI
2-Pack	6-Pack			
0101P	0101K	0101L	Streptococcus equi subsp. zooepidemicus derived from ATCC® 43079™	0845357001018T
0104P	0104K	0104L	Streptococcus agalactiae (B,3) derived from ATCC® 12403™	0845357001048Z
0105P	0105K	0105L	Streptococcus agalactiae derived from ATCC® BAA-611™	08453570010593
0106P	0106K	0106L	Citrobacter koseri derived from ATCC® 27156™	08453570010695
0110P	0110K	0110L	Prevotella melaninogenica derived from ATCC® 25845™	0845357001108U
0111P	0111K	0111L	Campylobacter jejuni subsp. jejuni derived from ATCC® 33560™	0845357001118W
0112P	0112K	0112L	Mycobacterium tuberculosis derived from ATCC® 25177™	0845357001128Y
0113P	0113K	0113L	Staphylococcus aureus subsp. aureus derived from ATCC® 9144™	08453570011392
0114P	0114K	0114L	Mycobacterium smegmatis derived from ATCC® 607™	08453570011494
0116P	0116K	0116L	Priestia megaterium derived from ATCC® 9885™	08453570011698
0119P	0119K	0119L	Acinetobacter baumannii derived from ATCC® BAA-747™	0845357001199E
0120P	0120K	0120L	Bacillus subtilis derived from ATCC® 6051™	0845357001208X
0121P	0121K	0121L	Campylobacter coli derived from ATCC® 43478™	0845357001218Z
0122P	0122K	0122L	Candida glabrata derived from ATCC® MYA-2950™	08453570012293
0123P	0123K	0123L	Clostridium perfringens derived from ATCC® 12916™	08453570012395
0126P	0126K	0126L	Kocuria kristinae derived from ATCC® BAA-752™	0845357001269B
0127P	0127K	0127L	Lactobacillus gasseri derived from ATCC® 19992™	0845357001279D
0128P	0128K	0128L	Lactobacillus sakei subsp. sakei derived from ATCC® 15521™	0845357001289F
0129P	0129K	0129L	Listeria monocytogenes (4b) derived from ATCC® 13932™	0845357001299H
0130P	0130K	0130L	Listeria monocytogenes derived from ATCC® BAA-751™	08453570013092
0131P	0131K	0131L	Neisseria mucosa derived from ATCC® 49233™	08453570013194
0132P	0132K	0132L	Ochrobactrum anthropi derived from ATCC® BAA-749™	08453570013296
0134P	0134K	0134L	Staphylococcus saprophyticus derived from ATCC® BAA-750™	0845357001349A
0135P	0135K	0135L	Stenotrophomonas maltophilia derived from ATCC® 49130™	0845357001359C
0136P	0136K	0136L	Streptococcus thermophilus derived from ATCC® 19258™	0845357001369E
0139P	0139K	0139L	Brevibacillus agri derived from ATCC® 51663™	0845357001399L
0140P	0140K	0140L	Niallia circulans derived from ATCC® 61™	08453570014095
0141P	0141K	0141L	Aneurinibacillus aneurinilyticus derived from ATCC® 11376™	08453570014197
0142P	0142K	0142L	Paenibacillus macerans derived from ATCC® 8509™	08453570014299
0143P	0143K	0143L	Rhodotorula mucilaginosa derived from ATCC® 66034™	0845357001439B
0144P	0144K	0144L	Brevibacillus laterosporus derived from ATCC® 64™	0845357001449D
0146P	0146K	0146L	Staphylococcus aureus subsp. aureus derived from ATCC® BAA-976™	0845357001469H

0147P	0147K	0147L	Staphylococcus aureus subsp. aureus derived from ATCC® BAA-977™	0845357001479K
0149P	0149K	0149L	Lactococcus lactis subsp. lactis derived from ATCC® 19435™	0845357001499P
N/A	N/A	0151L	Ureaplasma parvum derived from ATCC® 27813™	0845357001519A
0152P	0152K	0152L	Lactococcus cremoris derived from ATCC® 19257™	0845357001529C
0153P	0153K	0153L	Corynebacterium xerosis derived from ATCC® 373™	0845357001539E
0154P	0154K	0154L	Listeria monocytogenes derived from ATCC® 19118™	0845357001549G
N/A	N/A	0156L	Metamycoplasma hominis derived from ATCC® 15488™	0845357001569L
0157P	0157K	0157L	Mycobacterium intracellulare derived from ATCC® 13950™	0845357001579N
0158P	0158K	0158L	Staphylococcus aureus subsp. aureus derived from ATCC® 700699™	0845357001589Q
0159P	0159K	0159L	Staphylococcus aureus subsp. aureus derived from ATCC® 51740™	0845357001599S
0164P	0164K	0164L	Chryseobacterium shigense derived from ATCC® 51823™	0845357001649K
0165P	0165K	0165L	Hafnia alvei derived from ATCC® 51815™	0845357001659M
0167P	0167K	0167L	Klebsiella oxytoca derived from ATCC® 51817™	0845357001679R
0168P	0168K	0168L	Pseudomonas brenneri derived from ATCC® 49642™	0845357001689T
0169P	0169K	0169L	Herminiimonas species derived from ATCC® 49643™	0845357001699V
0170P	0170K	0170L	Cutibacterium acnes derived from ATCC® 6919™	0845357001709E
0172P	0172K	0172L	Geobacillus stearothermophilus derived from ATCC® 10149™	0845357001729J
0173P	0173K	0173L	Staphylococcus aureus subsp. aureus derived from ATCC® 12600™	0845357001739L
0175P	0175K	0175L	Bifidobacterium breve derived from ATCC® 15700™	0845357001759Q
0176P	0176K	0176L	Lacticaseibacillus casei derived from ATCC® 393™	0845357001769S
0177P	0177K	0177L	Aspergillus oryzae derived from ATCC® 10124™	0845357001779U
0178P	0178K	0178L	Penicillium chrysogenum derived from ATCC® 10106™	0845357001789W
0179P	0179K	0179L	Staphylococcus aureus derived from ATCC® BAA-1026™	0845357001799Y
0180P	0180K	0180L	Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 51812™	0845357001809H
0181P	0181K	0181L	Enterococcus faecalis derived from ATCC® 14506™	0845357001819K
0182P	0182K	0182L	Pseudomonas aeruginosa derived from ATCC® 35554™	0845357001829M
0184P	0184K	0184L	Aggregatibacter aphrophilus derived from ATCC® 33389™	0845357001849R
0185P	0185K	0185L	Haemophilus influenzae derived from ATCC® 9007™	0845357001859T
0188P	0188K	0188L	Campylobacter jejuni subsp. jejuni derived from ATCC® BAA-1153™	0845357001889Z
0189P	0189K	0189L	Eikenella corrodens derived from ATCC® BAA-1152™	084535700189A3
0191P	0191K	0191L	Raoultella terrigena derived from ATCC® 33257™	0845357001919N
0196P	0196K	0196L	Klebsiella aerogenes derived from ATCC® 51697™	0845357001969Y
0197P	0197K	0197L	Enterococcus faecalis derived from ATCC® 33186™	084535700197A2
0198P	0198K	0198L	Bacillus cereus derived from ATCC® 33019™	084535700198A4
0199P	0199K	0199L	Pseudomonas aeruginosa derived from ATCC® 19429™	084535700199A6
0200P	0200K	0200L	Bacillus cereus derived from ATCC® 14579™	0845357002008W
0201P	0201K	0201L	Priestia megaterium derived from ATCC® 14581™	0845357002018Y
0203P	0203K	0203L	Escherichia coli derived from ATCC® 23848™	08453570020394

0204P	0204K	0204L	Escherichia coli (JM101) derived from ATCC® 33876™	08453570020496
0205P	0205K	0205L	Lactococcus lactis subsp. lactis derived from ATCC® 11454™	08453570020598
0207P	0207K	0207L	Penicillium rubens derived from ATCC® 9179™	0845357002079C
0208P	0208K	0208L	Rhizopus stolonifer (-) derived from ATCC® 6227a™	0845357002089E
0209P	0209K	0209L	Rhizopus stolonifer (+) derived from ATCC® 6227b™	0845357002099G
0211P	0211K	0211L	Legionella pneumophila subsp. pneumophila derived from ATCC® 33152™	08453570021193
0212P	0212K	0212L	Legionella bozemanae derived from ATCC® 33217™	08453570021295
0215P	0215K	0215L	Morganella morganii subsp. morganii derived from ATCC® 25829™	0845357002159B
0217P	0217K	0217L	Staphylococcus lugdunensis derived from ATCC® 49576™	0845357002179F
0218P	0218K	0218L	Micrococcus yunnanensis derived from ATCC® 7468™	0845357002189H
0221P	0221K	0221L	Streptococcus equi subsp. equi derived from ATCC® 33398™	08453570022196
0223P	0223K	0223L	Enterococcus saccharolyticus derived from ATCC® 43076™	0845357002239A
0227P	0227K	0227L	Issatchenkia orientalis derived from ATCC® 6258™	0845357002279J
0228P	0228K	0228L	Paenibacillus polymyxa derived from ATCC® 43865™	0845357002289L
0229P	0229K	0229L	Citrobacter freundii derived from ATCC® 43864™	0845357002299N
0231P	0231K	0231L	Escherichia coli derived from ATCC® 700728™	08453570023199
0232P	0232K	0232L	Listeria monocytogenes derived from ATCC® 19112™	0845357002329B
0233P	0233K	0233L	Lactobacillus rhamnosus derived from ATCC® 7469™	0845357002339D
0234P	0234K	0234L	Lactiplantibacillus plantarum derived from ATCC® 8014™	0845357002349F
0235P	0235K	0235L	Lactobacillus delbrueckii subsp. lactis derived from ATCC® 7830™	0845357002359H
0237P	0237K	0237L	Streptococcus salivarius derived from ATCC® 13419™	0845357002379M
0241P	0241K	0241L	Pseudomonas fluorescens derived from ATCC® 13525™	0845357002419C
0242P	0242K	0242L	Micrococcus luteus derived from ATCC® 4698™	0845357002429E
0243P	0243K	0243L	Lactobacillus acidophilus derived from ATCC® 4356™	0845357002439G
0245P	0245K	0245L	Aspergillus niger derived from ATCC® 16888™	0845357002459L
0246P	0246K	0246L	Staphylococcus haemolyticus derived from ATCC® 29970™	0845357002469N
0247P	0247K	0247L	Serratia marcescens derived from ATCC® 13880™	0845357002479Q
0248P	0248K	0248L	Moraxella catarrhalis derived from ATCC® 23246™	0845357002489S
0249P	0249K	0249L	Aggregatibacter aphrophilus derived from ATCC® 49917™	0845357002499U
0250P	0250K	0250L	Candida albicans derived from ATCC® 90029™	0845357002509D
0251P	0251K	0251L	Campylobacter jejuni subsp. jejuni derived from ATCC® 49943™	0845357002519F
0253P	0253K	0253L	Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 25241™	0845357002539K
0254P	0254K	0254L	Listeria monocytogenes Cornell University derived from Silliker® SLR2249	0845357002549M
0255P	0255K	0255L	Zygosaccharomyces rouxii derived from NCYC 381	0845357002559P
0256P	0256K	0256L	Bacillus cereus derived from ATCC® 11778™	0845357002569R
0257P	0257K	0257L	Clostridium perfringens derived from ATCC® 3624™	0845357002579T
0258P	0258K	0258L	Bacillus pumilus derived from ATCC® 14884™	0845357002589V

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0259P	0259K	0259L	Pediococcus acidilactici derived from ATCC® 8042™	0845357002599X
0260P	0260K	0260L	Arthrobacter psychrolactophilus derived from ATCC® 700733™	0845357002609G
0261P	0261K	0261L	Klebsiella variicola derived from ATCC® 31488™	0845357002619J
0262P	0262K	0262L	Serratia marcescens derived from ATCC® 43862™	0845357002629L
0263P	0263K	0263L	Aerococcus viridans derived from ATCC® 10400™	0845357002639N
0264P	0264K	0264L	Candida albicans derived from ATCC® 90028™	0845357002649Q
0265P	0265K	0265L	Alicyclobacillus acidoterrestris derived from ATCC® 49025™	0845357002659S
0266P	0266K	0266L	Streptococcus mutans derived from ATCC® 25175™	0845357002669U
0267P	0267K	0267L	Streptococcus pneumoniae derived from ATCC® 700677™	0845357002679W
0269P	0269K	0269L	Bacillus subtilis derived from ATCC® 11774™	084535700269A2
0270P	0270K	0270L	Bacillus thuringiensis derived from ATCC® 10792™	0845357002709K
0272P	0272K	0272L	Mycobacterium terrae derived from ATCC® 15755™	0845357002729P
0274P	0274K	0274L	Sphingomonas paucimobilis derived from ATCC® 29837™	0845357002749T
0276P	0276K	0276L	Aerococcus viridans derived from ATCC® 11563™	0845357002769X
0277P	0277K	0277L	Listeria monocytogenes derived from ATCC® 19111™	0845357002779Z
0290P	0290K	0290L	Aeromonas salmonicida derived from ATCC® 7965™	0845357002909R
0291P	0291K	0291L	Cryptococcus neoformans derived from ATCC® 14116™	0845357002919T
0293P	0293K	0293L	Corynebacterium renale derived from ATCC® 19412™	0845357002939X
0294P	0294K	0294L	Cellulosimicrobium cellulans derived from ATCC® 27402™	0845357002949Z
0295P	0295K	0295L	Microbacterium testaceum derived from ATCC® 15829™	084535700295A3
0296P	0296K	0296L	Paenibacillus polymyxa derived from ATCC® 7070™	084535700296A5
0299P	0299K	0299L	Listeria ivanovii subsp. londoniensis derived from ATCC® BAA-139™	084535700299AB
0300P	0300K	0300L	Proteus vulgaris derived from ATCC® 33420™	08453570030093
0303P	0303K	0303L	Shigella sonnei derived from ATCC® 25931™	08453570030399
0306P	0306K	0306L	Klebsiella aerogenes derived from ATCC® 13048™	0845357003069F
0310P	0310K	0310L	Proteus mirabilis derived from NCIMB 13283	08453570031096
0312P	0312K	0312L	Staphylococcus aureus subsp. aureus derived from NCIMB 12702	0845357003129A
0313P	0313K	0313L	Enterobacter cloacae subsp. cloacae derived from ATCC® 23355™	0845357003139C
0314P	N/A	N/A	Streptococcus pyogenes derived from NCIMB 13285	0845357003149E
0315P	0315K	0315L	Citrobacter freundii derived from ATCC® 8090™	0845357003159G
0316P	0316K	0316L	Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 23715™	0845357003169J
0317P	0317K	0317L	Clostridium sporogenes derived from ATCC® 19404™	0845357003179L
0318P	0318K	0318L	Clostridium perfringens derived from ATCC® 13124™	0845357003189N
0319P	0319K	0319L	Bacteroides thetaiotaomicron derived from ATCC® 29741™	0845357003199Q
0320P	0320K	0320L	Bacteroides fragilis derived from ATCC® 25285™	08453570032099
0321P	0321K	0321L	Proteus mirabilis derived from ATCC® 29245™	0845357003219B
0322P	0322K	0322L	Peptostreptococcus anaerobius derived from ATCC® 27337™	0845357003229D

0323P	0323K	0323L	Enterobacter cloacae subsp. cloacae derived from ATCC® 13047™	0845357003239F
0324P	0324K	0324L	Myroides odoratus derived from ATCC® 4651™	0845357003249H
0325P	0325K	0325L	Campylobacter jejuni subsp. jejuni derived from ATCC® 29428™	0845357003259K
0327P	0327K	0327L	Clostridium histolyticum derived from ATCC® 19401™	0845357003279P
0328P	0328K	0328L	Fusobacterium nucleatum subsp. nucleatum derived from ATCC® 25586™	0845357003289R
0329P	0329K	0329L	Clostridioides difficile derived from ATCC® 9689™	0845357003299T
0330P	0330K	0330L	Bacillus cereus derived from NCIMB 7464	0845357003309C
0331P	0331K	0331L	Paeniclostridium sordellii derived from ATCC® 9714™	0845357003319E
0332P	0332K	0332L	Candida albicans derived from ATCC® 14053™	0845357003329G
0333P	0333K	0333L	Papiliotrema laurentii derived from ATCC® 18803™	0845357003339J
0334P	0334K	0334L	Cryptococcus neoformans derived from ATCC® 32045™	0845357003349L
0335P	0335K	0335L	Escherichia coli derived from ATCC® 25922™	0845357003359N
0336P	0336K	0336L	Enterococcus faecalis derived from NCIMB 13280	0845357003369Q
0338P	0338K	0338L	Haemophilus influenzae derived from ATCC® 33533™	0845357003389U
0341P	0341K	0341L	Salmonella enterica subsp. enterica serovar Paratyphi A derived from ATCC® 9150™	0845357003419H
0343P	0343K	0343L	Salmonella enterica subsp. enterica serovar Choleraesuis derived from ATCC® 7001™	0845357003439M
0345P	0345K	0345L	Salmonella enterica subsp. enterica serovar Enteritidis derived from ATCC® 13076™	0845357003459R
0346P	0346K	0346L	Salmonella enterica subsp. enterica serovar Anatum derived from ATCC® 9270™	0845357003469T
0348P	0348K	0348L	Shigella flexneri derived from ATCC® 9199™	0845357003489X
0349P	0349K	0349L	Shigella boydii derived from ATCC® 9207™	0845357003499Z
0350P	0350K	0350L	Shigella sonnei derived from ATCC® 29930™	0845357003509J
0351P	0351K	0351L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 13883™	0845357003519L
0352P	0352K	0352L	Staphylococcus aureus subsp. aureus derived from ATCC® 33862™	0845357003529N
0353P	0353K	0353L	Pseudomonas aeruginosa derived from ATCC® 27853™	0845357003539Q
0354P	0354K	0354L	Serratia marcescens derived from ATCC® 8100™	0845357003549S
0355P	0355K	0355L	Proteus hauseri derived from ATCC® 13315™	0845357003559U
0356P	0356K	0356L	Shigella flexneri derived from ATCC® 12022™	0845357003569W
0357P	0357K	0357L	Acinetobacter baumannii derived from ATCC® 19606™	0845357003579Y
0358P	0358K	0358L	Bacteroides fragilis derived from NCTC 9343	084535700358A2
0360P	0360K	0360L	Staphylococcus aureus subsp. aureus derived from ATCC® 25923™	0845357003609M
0363P	0363K	0363L	Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 14028™	0845357003639T
0365P	0365K	0365L	Staphylococcus aureus subsp. aureus derived from ATCC® 29213™	0845357003659X
0366P	0366K	0366L	Enterococcus faecalis derived from ATCC® 29212™	0845357003669Z
0367P	0367K	0367L	Enterococcus faecalis derived from ATCC® 19433™	084535700367A3
0369P	0369K	0369L	Stenotrophomonas maltophilia derived from ATCC® 13637™	084535700369A7
0370P	0370K	0370L	Streptococcus agalactiae derived from ATCC® 13813™	0845357003709Q

0371P	0371K	0371L	Staphylococcus epidermidis derived from ATCC® 12228™	0845357003719S
0375P	0375K	0375L	Neisseria gonorrhoeae derived from ATCC® 31426™	084535700375A2
0376P	0376K	0376L	Haemophilus influenzae derived from ATCC® 19418™	084535700376A4
0377P	0377K	0377L	Haemophilus parahaemolyticus derived from ATCC® 10014™	084535700377A6
0378P	0378K	0378L	Neisseria gonorrhoeae derived from ATCC® 19424™	084535700378A8
0379P	0379K	0379L	Candida albicans derived from NCYC 1363	084535700379AA
0380P	0380K	0380L	Streptococcus pneumoniae derived from ATCC® 6303™	0845357003809T
0383P	0383K	0383L	Streptococcus mitis derived from NCIMB 13770	0845357003839Z
0384P	0384K	0384L	Providencia stuartii derived from ATCC® 33672™	084535700384A3
0385P	0385K	0385L	Streptococcus pyogenes derived from ATCC® 19615™	084535700385A5
0388P	0388K	0388L	Enterobacter cloacae subsp. cloacae derived from ATCC® 35030™	084535700388AB
0389P	0389K	0389L	Streptococcus anginosus derived from NCTC 10713	084535700389AD
0391P	0391K	0391L	Streptococcus gallolyticus derived from ATCC® 9809™	0845357003919Y
0392P	0392K	0392L	Aspergillus brasiliensis derived from ATCC® 16404™	084535700392A2
0393P	0393K	0393L	Proteus vulgaris derived from NCTC 4636	084535700393A4
0398P	0398K	0398L	Listeria monocytogenes derived from ATCC® 7644™	084535700398AE
0399P	0399K	0399L	Klebsiella aerogenes derived from ATCC® 35029™	084535700399AG
0400P	0400K	0400L	Bacteroides ovatus derived from ATCC® 8483™	08453570040098
0402P	0402K	0402L	Alcaligenes faecalis subsp. faecalis derived from ATCC® 8750™	0845357004029C
0403P	0403K	0403L	Moraxella catarrhalis derived from ATCC® 25240™	0845357004039E
0404P	0404K	0404L	Neisseria meningitidis derived from ATCC® 13102™	0845357004049G
0405P	0405K	0405L	Neisseria lactamica derived from ATCC® 23970™	0845357004059J
0406P	0406K	0406L	Neisseria sicca derived from ATCC® 9913™	0845357004069L
0407P	0407K	0407L	Fusobacterium necrophorum subsp. necrophorum derived from ATCC® 25286™	0845357004079N
0409P	0409K	0409L	Finegoldia magna derived from ATCC® 29328™	0845357004099S
0410P	0410K	0410L	Gardnerella vaginalis derived from ATCC® 14018™	0845357004109B
0411P	0411K	0411L	Aggregatibacter aphrophilus derived from ATCC® 7901™	0845357004119D
0412P	0412K	0412L	Staphylococcus epidermidis derived from ATCC® 14990™	0845357004129F
0414P	0414K	0414L	Listeria innocua derived from NCTC 11288	0845357004149K
0416P	0416K	0416L	Pseudomonas aeruginosa derived from ATCC® 10145™	0845357004169P
0419P	0419K	0419L	Cutibacterium acnes derived from ATCC® 11827™	0845357004199V
0421P	0421K	0421L	Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 13311™	0845357004219G
0422P	0422K	0422L	Escherichia coli derived from ATCC® 35421™	0845357004229J
0423P	0423K	0423L	Streptococcus oralis derived from ATCC® 6249™	0845357004239L
0425P	0425K	0425L	Candida albicans derived from ATCC® 60193™	0845357004259Q
0426P	0426K	0426L	Neisseria gonorrhoeae derived from ATCC® 43070™	0845357004269S
0429P	0429K	0429L	Neisseria gonorrhoeae derived from ATCC® 43069™	0845357004299Y
0431P	0431K	0431L	Paenibacillus larvae subsp. larvae derived from ATCC® 9545™	0845357004319K
0432P	0432K	0432L	Proteus mirabilis derived from ATCC® 43071™	0845357004329M

0433P	0433K	0433L	Escherichia coli derived from ATCC® 4157™	0845357004339P
0435P	0435K	0435L	Streptococcus pneumoniae derived from ATCC® 27336™	0845357004359T
0436P	0436K	0436L	Streptococcus agalactiae derived from ATCC® 27956™	0845357004369V
0439P	0439K	0439L	Streptococcus agalactiae derived from ATCC® 12386™	084535700439A3
0440P	0440K	0440L	Proteus mirabilis derived from ATCC® 12453™	0845357004409L
0441P	0441K	0441L	Haemophilus influenzae derived from ATCC® 10211™	0845357004419N
0442P	0442K	0442L	Trichophyton interdigitale derived from ATCC® 9533™	0845357004429Q
0443P	0443K	0443L	Candida albicans derived from ATCC® 10231™	0845357004439S
0444P	0444K	0444L	Trichophyton rubrum derived from ATCC® 28188™	0845357004449U
0445P	0445K	0445L	Phocaeicola vulgatus derived from ATCC® 8482™	0845357004459W
0446P	0446K	0446L	Shigella sonnei derived from ATCC® 9290™	0845357004469Y
0447P	0447K	0447L	Streptococcus pneumoniae derived from ATCC® 6305™	084535700447A2
0450P	0450K	0450L	Candida tropicalis derived from ATCC® 13803™	0845357004509P
0453P	0453K	0453L	Neisseria meningitidis derived from ATCC® 13077™	0845357004539V
0454P	0454K	0454L	Neisseria meningitidis derived from ATCC® 13090™	0845357004549X
0458P	0458K	0458L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 13882™	084535700458A7
0459P	0459K	0459L	Proteus vulgaris derived from ATCC® 6380™	084535700459A9
0462P	0462K	0462L	Staphylococcus aureus subsp. aureus derived from ATCC® 11632™	0845357004629W
0463P	0463K	0463L	Streptococcus bovis derived from ATCC® 33317™	0845357004639Y
0464P	0464K	0464L	Neisseria sicca derived from ATCC® 29193™	084535700464A2
0465P	0465K	0465L	Escherichia coli derived from ATCC® 11775™	084535700465A4
0468P	0468K	0468L	Acinetobacter lwoffii derived from ATCC® 15309™	084535700468AA
0469P	0469K	0469L	Streptococcus pneumoniae derived from NCIMB 13286	084535700469AC
0472P	0472K	0472L	Enterococcus faecalis derived from NCTC 775	0845357004729Z
0473P	0473K	0473L	Paenibacillus gordonae derived from ATCC® 29948™	084535700473A3
0474P	0474K	0474L	Bacillus pumilus derived from ATCC® BAA-1434™	084535700474A5
0475P	0475K	0475L	Moraxella osloensis derived from ATCC® 10973™	084535700475A7
0476P	0476K	0476L	Haemophilus influenzae derived from ATCC® 43163™	084535700476A9
0478P	0478K	0478L	Citrobacter braakii derived from ATCC® 43162™	084535700478AD
0480P	N/A	N/A	Streptococcus agalactiae derived from NCTC 8017	0845357004809Y
0481P	0481K	0481L	Campylobacter jejuni subsp. jejuni derived from ATCC® 33291™	084535700481A2
0483P	0483K	0483L	Escherichia coli derived from ATCC® 8739™	084535700483A6
0484P	0484K	0484L	Pseudomonas paraeruginosa derived from ATCC® 9027™	084535700484A8
0485P	0485K	0485L	Staphylococcus aureus subsp. aureus derived from ATCC® 6538™	084535700485AA
0486P	0486K	0486L	Bacillus spizizenii derived from ATCC® 6633™	084535700486AC
0487P	0487K	0487L	Clostridium sporogenes derived from ATCC® 11437™	084535700487AE
0488P	0488K	0488L	Burkholderia cepacia derived from ATCC® 25416™	084535700488AG
0489P	0489K	0489L	Bordetella pertussis derived from ATCC® 8467™	084535700489AJ
0494P	0494K	0494L	Staphylococcus saprophyticus derived from ATCC® 15305™	084535700494AB

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0495P	0495K	0495L	Escherichia coli derived from ATCC® 35218™	084535700495AD
0496P	0496K	0496L	Staphylococcus aureus subsp. aureus derived from ATCC® 33591™	084535700496AF
0497P	0497K	0497L	Enterococcus faecalis derived from ATCC® 7080™	084535700497AH
0500P	0500K	0500L	Aspergillus niger derived from ATCC® 6275™	0845357005009D
0501P	0501K	0501L	Salmonella enterica subsp. enterica derived from ATCC® 51741™	0845357005019F
0502P	0502K	0502L	Escherichia coli derived from ATCC® 11303™	0845357005029H
N/A	N/A	0503L	Mycoplasma pneumoniae derived from ATCC® 15531™	0845357005039K
0506P	0506K	0506L	Serratia marcescens derived from ATCC® 14041™	0845357005069R
0508P	0508K	0508L	Streptococcus pyogenes derived from ATCC® 12344™	0845357005089V
0510P	0510K	0510L	Candida krusei derived from ATCC® 34135™	0845357005109G
0511P	0511K	0511L	Acetobacter aceti derived from ATCC® 15973™	0845357005119J
0513P	0513K	0513L	Mycobacterium fortuitum subsp. fortuitum derived from ATCC® 6841™	0845357005139N
0514P	0514K	0514L	Mycobacterium smegmatis derived from ATCC® 19420™	0845357005149Q
0519P	0519K	0519L	Galactomyces candidus derived from ATCC® 34614™	084535700519A2
0520P	0520K	0520L	Candida geochares derived from ATCC® 36852™	0845357005209K
0522P	0522K	0522L	Mycobacterium avium subsp. avium derived from ATCC® 15769™	0845357005229P
0524P	0524K	0524L	Pseudomonas protegens (G) derived from ATCC® 17386™	0845357005249T
0527P	0527K	0527L	Clostridioides difficile derived from ATCC® 700057™	0845357005279Z
0530P	0530K	0530L	Klebsiella oxytoca derived from ATCC® 13182™	0845357005309N
0531P	0531K	0531L	Fusarium keratoplasticum derived from ATCC® 36031™™	0845357005319Q
0533P	0533K	0533L	Staphylococcus capitis derived from ATCC® 146™	0845357005339U
0534P	0534K	0534L	Saccharomyces cerevisiae derived from ATCC® 4098™	0845357005349W
0537P	0537K	0537L	Cladosporium cladosporioides derived from ATCC® 16022™	084535700537A4
0539P	0539K	0539L	Bacillus thuringiensis derived from ATCC® 33679™	084535700539A8
0540P	0540K	0540L	Bacillus spizizenii derived from ATCC® 19659™	0845357005409R
0543P	0543K	0543L	Escherichia coli derived from ATCC® 14169™	0845357005439X
0544P	0544K	0544L	Mycobacterium avium subsp. avium derived from ATCC® 25291™	0845357005449Z
0545P	0545K	0545L	Mycobacterium kansasii derived from ATCC® 12478™	084535700545A3
0546P	0546K	0546L	Lactobacillus paracasei derived from ATCC® 334™	084535700546A5
0547P	0547K	0547L	Clostridium perfringens derived from NCTC 8678	084535700547A7
0571P	0571K	0571L	Staphylococcus epidermidis derived from NCIMB 8853	084535700571A4
0572P	0572K	0572L	Clostridium perfringens derived from NCTC 8237	084535700572A6
0574P	0574K	0574L	Citrobacter freundii derived from NCTC 9750	084535700574AA
0576P	0576K	0576L	Pseudomonas aeruginosa derived from NCIMB 8626	084535700576AE
0577P	0577K	0577L	Bacillus pumilus derived from ATCC® 700814™	084535700577AG
0578P	0578K	0578L	Citrobacter braakii derived from ATCC® 51113™	084535700578AJ
0580P	0580K	0580L	Clostridium sporogenes derived from NCIMB 12343	084535700580A5
0583P	0583K	0583L	Corynebacterium striatum derived from ATCC® BAA-1293™	084535700583AB

0584P	0584K	0584L	Parabacteroides distasonis derived from ATCC® BAA-1295™	084535700584AD
0585P	0585K	0585L	Bacteroides ovatus derived from ATCC® BAA-1296™	084535700585AF
0586P	0586K	0586L	Clostridium septicum derived from ATCC® 12464™	084535700586AH
0587P	0587K	0587L	Bacteroides ovatus derived from ATCC® BAA-1304™	084535700587AK
0595P	0595K	0595L	Salmonella bongori derived from ATCC® 43975™	084535700595AJ
0596P	0596K	0596L	Weizmannia coagulans derived from ATCC® 7050™	084535700596AL
0597P	0597K	0597L	Klebsiella aerogenes derived from NCIMB 10102	084535700597AN
0598P	0598K	0598L	Pseudomonas aeruginosa derived from NCIMB 12469	084535700598AQ
0599P	0599K	0599L	Acinetobacter baumannii derived from NCIMB 12457	084535700599AS
0602P	0602K	0602L	Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 12394™	0845357006029N
0603P	0603K	0603L	Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 12388™	0845357006039Q
0604P	0604K	0604L	Salmonella enterica subsp. enterica serovar Pullorum derived from ATCC® 13036™	0845357006049S
0605P	0605K	0605L	Staphylococcus xylosus derived from ATCC® 29971™	0845357006059U
0607P	0607K	0607L	Proteus mirabilis derived from ATCC® 7002™	0845357006079Y
0617P	0617K	0617L	Escherichia coli (O157:H7) derived from ATCC® 35150™	084535700617A3
0618P	0618K	0618L	Parabacteroides distasonis derived from ATCC® 8503™	084535700618A5
0619P	0619K	0619L	Bacteroides uniformis derived from ATCC® 8492™	084535700619A7
0620P	0620K	0620L	Haemophilus influenzae derived from ATCC® 9006™	0845357006209Q
0621P	0621K	0621L	Oligella urethralis derived from ATCC® 17960™	0845357006219S
0622P	0622K	0622L	Moraxella catarrhalis derived from ATCC® 8176™	0845357006229U
0626P	0626K	0626L	Klebsiella oxytoca derived from ATCC® 49131™	084535700626A4
0627P	0627K	0627L	Pseudomonas putida derived from ATCC® 49128™	084535700627A6
0628P	0628K	0628L	Staphylococcus epidermidis derived from ATCC® 49134™	084535700628A8
0629P	0629K	0629L	Staphylococcus gallinarum derived from ATCC® 49148™	084535700629AA
0630P	0630K	0630L	Streptococcus pasteurianus derived from ATCC® 49133™	0845357006309T
0631P	0631K	0631L	Streptococcus gallolyticus derived from ATCC® 49147™	0845357006319V
0632P	0632K	0632L	Streptococcus pneumoniae derived from ATCC® 49136™	0845357006329X
0633P	0633K	0633L	Streptococcus pneumoniae derived from ATCC® 49150™	0845357006339Z
0635P	0635K	0635L	Acinetobacter species derived from ATCC® 49139™	084535700635A5
0637P	0637K	0637L	Aeromonas hydrophila derived from ATCC® 49140™	084535700637A9
0638P	0638K	0638L	Shewanella haliotis derived from ATCC® 49138™	084535700638AB
0639P	0639K	0639L	Enterobacter ludwigii derived from ATCC® 49141™	084535700639AD
0640P	0640K	0640L	Proteus vulgaris derived from ATCC® 49132™	0845357006409W
0641P	0641K	0641L	Ralstonia insidiosa derived from ATCC® 49129™	0845357006419Y
0642P	0642K	0642L	Moraxella catarrhalis derived from ATCC® 49143™	084535700642A2
0643P	0643K	0643L	Gardnerella vaginalis derived from ATCC® 49145™	084535700643A4
0644P	0644K	0644L	Haemophilus influenzae derived from ATCC® 49144™	084535700644A6
0645P	0645K	0645L	Aggregatibacter aphrophilus derived from ATCC® 49146™	084535700645A8
0646P	0646K	0646L	Neisseria lactamica derived from ATCC® 49142™	084535700646AA
0647P	0647K	0647L	Haemophilus influenzae derived from ATCC® 49247™	084535700647AC

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0648P	0648K	0648L	Neisseria gonorrhoeae derived from ATCC® 49226™	084535700648AE
0650P	0650K	0650L	Enterococcus hirae derived from ATCC® 8043™	0845357006509Z
0651P	0651K	0651L	Enterococcus durans derived from ATCC® 6056™	084535700651A3
0655P	0655K	0655L	Bordetella bronchiseptica derived from ATCC® 10580™	084535700655AB
0656P	0656K	0656L	Streptococcus equi subsp. equi derived from ATCC® 9528™	084535700656AD
0660P	0660K	0660L	Trueperella pyogenes derived from ATCC® 19411™	084535700660A4
0661P	0661K	0661L	Erysipelothrix rhusiopathiae derived from ATCC® 19414™	084535700661A6
0668P	0668K	0668L	Pasteurella multocida subsp. multocida derived from ATCC® 12945™	084535700668AL
0669P	0669K	0669L	Kocuria rhizophila derived from ATCC® 9341a™	084535700669AN
0670P	0670K	0670L	Kocuria rhizophila derived from ATCC® 15957™	084535700670A7
0671P	0671K	0671L	Bordetella bronchiseptica derived from ATCC® 4617™	084535700671A9
0674P	0674K	0674L	Clostridium perfringens derived from ATCC® 12919™	084535700674AF
0676P	0676K	0676L	Clostridium sporogenes derived from ATCC® 3584™	084535700676AK
0677P	0677K	0677L	Enterococcus faecium derived from ATCC® 6569™	084535700677AM
0678P	0678K	0678L	Enterococcus hirae derived from ATCC® 10541™	084535700678AP
0679P	0679K	0679L	Enterococcus faecium derived from ATCC® 27270™	084535700679AR
0680P	0680K	0680L	Escherichia coli derived from ATCC® 10536™	084535700680AA
0681P	0681K	0681L	Escherichia coli derived from ATCC® 11229™	084535700681AC
0683P	0683K	0683L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 4352™	084535700683AG
0684P	0684K	0684L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 10031™	084535700684AJ
0685P	0685K	0685L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 27736™	084535700685AL
0686P	0686K	0686L	Listeria monocytogenes derived from ATCC® 19114™	084535700686AN
0687P	0687K	0687L	Listeria monocytogenes derived from ATCC® 19115™	084535700687AQ
0688P	0688K	0688L	Kocuria rhizophila derived from ATCC® 9341™	084535700688AS
0689P	0689K	0689L	Micrococcus luteus derived from ATCC® 10240™	084535700689AU
0690P	0690K	0690L	Proteus mirabilis derived from ATCC® 25933™	084535700690AD
0691P	0691K	0691L	Proteus vulgaris derived from ATCC® 8427™	084535700691AF
0693P	0693K	0693L	Pseudomonas aeruginosa derived from ATCC® 15442™	084535700693AK
0695P	0695K	0695L	Pseudomonas aeruginosa derived from ATCC® 25619™	084535700695AP
0697P	0697K	0697L	Rhodococcus equi derived from ATCC® 6939™	084535700697AT
0698P	0698K	0698L	Saccharomyces kudriavzevii derived from ATCC® 2601™	084535700698AV
0699P	0699K	0699L	Saccharomyces cerevisiae derived from ATCC® 9763™	084535700699AX
0701P	0701K	0701L	Malassezia furfur derived from ATCC® 14521™	0845357007019R
0702P	0702K	0702L	Pseudomonas sp. derived from ATCC® 31483™	0845357007029T
0703P	0703K	0703L	Pluralibacter gergoviae derived from ATCC® 33028™	0845357007039V
0706P	0706K	0706L	Escherichia coli derived from NCTC 10538	084535700706A3
0709P	0709K	0709L	Streptococcus agalactiae (B) derived from NCIMB 701348	084535700709A9
0710P	0710K	0710L	Streptococcus agalactiae derived from NCTC 9993	0845357007109S
0712P	0712K	0712L	Campylobacter jejuni subsp. jejuni derived from NCTC 13367	0845357007129W

0713P	0713K	0713L	Staphylococcus aureus subsp. aureus derived from NCTC 12973	0845357007139Y
0714P	0714K	0714L	Enterococcus faecalis derived from NCTC 13379	084535700714A2
0718P	0718K	0718L	Campylobacter jejuni subsp. jejuni derived from NCTC 11322	084535700718AA
0720P	0720K	0720L	Vibrio furnissii derived from NCTC 11218	0845357007209V
0721P	0721K	0721L	Mycobacterium smegmatis derived from ATCC® 14468™	0845357007219X
0726P	0726K	0726L	Candida parapsilosis derived from ATCC® 22019™	084535700726A9
0727P	0727K	0727L	Listeria monocytogenes derived from ATCC® 15313™	084535700727AB
0728P	0728K	0728L	Thermoanaerobacterium thermosaccharolyticum derived from ATCC® 7956™	084535700728AD
0736P	0736K	0736L	Saccharomyces cerevisiae derived from NCYC 79	084535700736AC
0737P	0737K	0737L	Candida glabrata derived from ATCC® 15126™	084535700737AE
0738P	0738K	0738L	Meyerozyma guilliermondii derived from ATCC® 6260™	084535700738AG
0739P	0739K	0739L	Staphylococcus lentus derived from ATCC® 700403™	084535700739AJ
0740P	0740K	0740L	Micrococcus species derived from ATCC® 700405™	084535700740A3
0741P	0741K	0741L	Staphylococcus xylosus derived from ATCC® 700404™	084535700741A5
0742P	0742K	0742L	Stenotrophomonas maltophilia derived from ATCC® 51331™	084535700742A7
0743P	0743K	0743L	Streptococcus equi subsp. zooepidemicus derived from ATCC® 700400™	084535700743A9
0744P	0744K	0744L	Streptococcus uberis derived from ATCC® 700407™	084535700744AB
0747P	0747K	0747L	Escherichia coli derived from ATCC® 13706™	084535700747AH
0749P	0749K	0749L	Capnocytophaga sputigena derived from ATCC® 33612™	084535700749AM
0750P	0750K	0750L	Actinomyces viscosus derived from ATCC® 15987™	084535700750A6
0754P	0754K	0754L	Brevundimonas diminuta derived from ATCC® 11568™	084535700754AE
0755P	0755K	0755L	Enterobacter hormaechei derived from ATCC® 700323™	084535700755AG
0756P	0756K	0756L	Cronobacter muytjensii derived from ATCC® 51329™	084535700756AJ
0757P	0757K	0757L	Klebsiella oxytoca derived from ATCC® 700324™	084535700757AL
0759P	0759K	0759L	Stenotrophomonas maltophilia derived from ATCC® 17666™	084535700759AQ
0761P	0761K	0761L	Enterococcus casseliflavus derived from ATCC® 700327™	084535700761AB
0763P	0763K	0763L	Streptococcus pneumoniae derived from ATCC® 6301™	084535700763AF
0764P	0764K	0764L	Mammaliicoccus sciuri derived from ATCC® 29061™	084535700764AH
0766P	0766K	0766L	Kocuria rosea derived from ATCC® 186™	084535700766AM
0774P	0774K	0774L	Candida lusitaniae derived from ATCC® 34449™	084535700774AL
0778P	0778K	0778L	Cutaneotrichosporon dermatis derived from ATCC® 204094™	084535700778AU
0779P	0779K	0779L	Candida utilis derived from ATCC® 9950™	084535700779AW
0780P	0780K	0780L	Prototheca wickerhamii derived from ATCC® 16529™	084535700780AF
0781P	0781K	0781L	Cryptococcus neoformans derived from ATCC® 204092™	084535700781AH
0783P	0783K	0783L	Listeria monocytogenes derived from NCTC 10890	084535700783AM
0784P	0784K	0784L	Klebsiella quasipneumoniae derived from ATCC® 700603™	084535700784AP
0791P	0791K	0791L	Escherichia coli derived from ATCC® 51813™	084535700791AL
0794P	0794K	0794L	Penicillium venetum derived from ATCC® 16025™	084535700794AS

0795P	0795K	0795L	Escherichia coli (O157:H7) derived from ATCC® 43888™	084535700795AU
0799P	0799K	0799L	Bacillus licheniformis derived from ATCC® 14580™	084535700799B4
0800P	0800K	0800L	Candida albicans derived from ATCC® 24433™	0845357008009U
0801P	0801K	0801L	Clostridium perfringens derived from ATCC® 12915™	0845357008019W
0802P	0802K	0802L	Listeria seeligeri derived from ATCC® 35967™	0845357008029Y
0803P	0803K	0803L	Zygosaccharomyces rouxii derived from ATCC® 28253™	084535700803A2
0804P	0804K	0804L	Micrococcus luteus derived from ATCC® 49732™	084535700804A4
0805P	0805K	0805L	Brevundimonas diminuta derived from ATCC® 19146™	084535700805A6
0806P	0806K	0806L	Serratia marcescens derived from ATCC® 14756™	084535700806A8
0809P	0809K	0809L	Candida krusei derived from ATCC® 14243™	084535700809AE
0812P	0812K	0812L	Bacillus licheniformis derived from ATCC® 12759™	084535700812A3
0813P	0813K	0813L	Lactobacillus fermentum derived from ATCC® 9338™	084535700813A5
0814P	0814K	0814L	Listeria innocua (6a) derived from ATCC® 33090™	084535700814A7
0815P	0815K	0815L	Listeria ivanovii subsp. ivanovii derived from ATCC® 19119™	084535700815A9
0816P	0816K	0816L	Listeria welshimeri derived from ATCC® 35897™	084535700816AB
0817P	0817K	0817L	Salmonella enterica subsp. enterica serovar Abaetetuba derived from ATCC® 35640™	084535700817AD
0818P	0818K	0818L	Vibrio parahaemolyticus derived from ATCC® 17802™	084535700818AF
0819P	0819K	0819L	Vibrio alginolyticus derived from ATCC® 17749™	084535700819AH
0826P	0826K	0826L	Salmonella enterica subsp. enterica serovar Abaetetuba derived from Silliker® SLR156	084535700826AE
0827P	0827K	0827L	Staphylococcus aureus subsp. aureus derived from ATCC® 6538P™	084535700827AG
0828P	0828K	0828L	Clostridium bifermentans derived from ATCC® 638™	084535700828AJ
0830P	0830K	0830L	Pseudomonas aeruginosa derived from NCTC 10662	084535700830A5
0831P	0831K	0831L	Staphylococcus aureus derived from NCTC 6571	084535700831A7
0832P	0832K	0832L	Staphylococcus aureus subsp. aureus derived from ATCC® 51153™	084535700832A9
0833P	0833K	0833L	Clostridioides difficile derived from ATCC® 43593™	084535700833AB
0836P	0836K	0836L	Burkholderia cenocepacia derived from ATCC® 25608™	084535700836AH
0838P	0838K	0838L	Serratia liquefaciens derived from ATCC® 27592™	084535700838AM
0839P	0839K	0839L	Morganella morganii subsp. morganii derived from ATCC® 25830™	084535700839AP
0840P	0840K	0840L	Klebsiella michiganensis derived from ATCC® 8724™	084535700840A8
0841P	0841K	0841L	Proteus vulgaris derived from ATCC® 6896™	084535700841AA
0842P	0842K	0842L	Bordetella parapertussis derived from ATCC® 15311™	084535700842AC
0843P	0843K	0843L	Bordetella pertussis derived from ATCC® 9797™	084535700843AE
0844P	0844K	0844L	Corynebacterium diphtheriae derived from ATCC® 13812™	084535700844AG
0845P	0845K	0845L	Edwardsiella tarda derived from ATCC® 15947™	084535700845AJ
0847P	0847K	0847L	Candida tropicalis derived from ATCC® 750™	084535700847AN
0851P	0851K	0851L	Salmonella enterica subsp. enterica serovar Poona derived from NCTC 4840	084535700851AD
0852P	0852K	0852L	Staphylococcus aureus subsp. aureus derived from ATCC® 43300™	084535700852AF
0853P	0853K	0853L	Pseudomonas stutzeri derived from ATCC® 17588™	084535700853AH

0856P	0856K	0856L	Listeria grayi derived from ATCC® 25401™	084535700856AP
0857P	0857K	0857L	Enterococcus raffinosus derived from ATCC® 49464™	084535700857AR
0858P	0858K	0858L	Streptococcus sanguinis derived from ATCC® 10556™	084535700858AT
0859P	0859K	0859L	Streptomyces griseus subsp. griseus derived from ATCC® 10137™	084535700859AV
0860P	0860K	0860L	Escherichia coli derived from ATCC® 51755™	084535700860AE
0861P	0861K	0861L	Escherichia coli derived from NCTC 12900	084535700861AG
0864P	0864K	0864L	Streptococcus agalactiae derived from ATCC® 12401™	084535700864AN
0865P	0865K	0865L	Streptococcus pneumoniae derived from ATCC® 10015™	084535700865AQ
0866P	0866K	0866L	Nocardia brasiliensis derived from ATCC® 19296™	084535700866AS
0867P	0867K	0867L	Veillonella parvula derived from ATCC® 10790™	084535700867AU
0868P	0868K	0868L	Oligella ureolytica derived from ATCC® 43534™	084535700868AW
0869P	0869K	0869L	Escherichia coli derived from ATCC® 51446™	084535700869AY
0870P	0870K	0870L	Aeromonas hydrophila derived from ATCC® 7966™	084535700870AH
0871P	0871K	0871L	Geobacillus stearothermophilus derived from ATCC® 7953™	084535700871AK
0872P	0872K	0872L	Geobacillus stearothermophilus derived from ATCC® 12980™	084535700872AM
0879P	0879K	0879L	Providencia stuartii derived from ATCC® 49809™	084535700879B3
0880P	0880K	0880L	Pseudomonas mosselii derived from ATCC® 49838™	084535700880AL
0881P	0881K	0881L	Staphylococcus saprophyticus derived from ATCC® 49907™	084535700881AN
0883P	0883K	0883L	Paenibacillus polymyxa derived from ATCC® 842™	084535700883AS
0884P	0884K	0884L	Corynebacterium pseudodiphtheriticum derived from ATCC® 10701™	084535700884AU
0885P	0885K	0885L	Lactobacillus acidophilus derived from ATCC® 314™	084535700885AW
0889P	0889K	0889L	Staphylococcus aureus subsp. aureus derived from ATCC® 33592™	084535700889B6
0890P	0890K	0890L	Salmonella enterica subsp. enterica serovar Abony derived from NCTC 6017	084535700890AP
0891P	0891K	0891L	Trichophyton tonsurans derived from ATCC® 28942™	084535700891AR
0893P	0893K	0893L	Microsporum gypseum derived from ATCC® 24102™	084535700893AV
0894P	0894K	0894L	Microsporum canis derived from ATCC® 36299™	084535700894AX
0895P	0895K	0895L	Enterococcus gallinarum derived from ATCC® 49573™	084535700895AZ
0896P	0896K	0896L	Candida albicans derived from ATCC® 2091™	084535700896B3
0897P	0897K	0897L	Candida tropicalis derived from ATCC® 9968™	084535700897B5
0900P	0900K	0900L	Saccharomyces cerevisiae derived from ATCC® 9080™	0845357009009Z
0901P	0901K	0901L	Salmonella enterica subsp. arizonae derived from ATCC® 13314™	084535700901A3
0902P	0902K	0902L	Salmonella enterica subsp. enterica serovar Choleraesuis derived from ATCC® 10708™	084535700902A5
0906P	0906K	0906L	Staphylococcus aureus subsp. aureus derived from ATCC® 29737™	084535700906AD
0907P	0907K	0907L	Staphylococcus pseudintermedius derived from ATCC® 49444™	084535700907AF
0909P	0909K	0909L	Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 27729™	084535700909AK
0910P	0910K	0910L	Aeromonas hydrophila derived from ATCC® 35654™	084535700910A4

0911P	0911K	0911L	Alcaligenes faecalis subsp. faecalis derived from ATCC® 35655™	084535700911A6
0912P	0912K	0912L	Porphyromonas gingivalis derived from ATCC® 33277™	084535700912A8
0919P	0919K	0919L	Haemophilus influenzae derived from ATCC® 49766™	084535700919AN
0936P	0936K	0936L	Eggerthella lenta derived from ATCC® 43055™	084535700936AN
0937P	0937K	0937L	Staphylococcus aureus subsp. aureus derived from ATCC® 49476™	084535700937AQ
0938P	0938K	0938L	Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 9610™	084535700938AS
0939P	0939K	0939L	Schaalia odontolytica derived from ATCC® 17929™	084535700939AU
0940P	0940K	0940L	Bacteroides fragilis derived from ATCC® 23745™	084535700940AD
0942P	0942K	0942L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 35657™	084535700942AH
0944P	0944K	0944L	Proteus mirabilis derived from ATCC® 35659™	084535700944AM
0945P	0945K	0945L	Staphylococcus saprophyticus derived from ATCC® 49453™	084535700945AP
0946P	0946K	0946L	Staphylococcus warneri derived from ATCC® 49454™	084535700946AR
0947P	0947K	0947L	Streptococcus pneumoniae derived from ATCC® 49619™	084535700947AT
0948P	0948K	0948L	Sphingobacterium multivorum derived from ATCC® 35656™	084535700948AV
0950P	0950K	0950L	Yarrowia lipolytica derived from ATCC® 9773™	084535700950AG
0951P	0951K	0951L	Moraxella catarrhalis derived from ATCC® 25238™	084535700951AJ
0952P	0952K	0952L	Neisseria gonorrhoeae derived from ATCC® 49981™	084535700952AL
0953P	0953K	0953L	Bacillus atrophaeus derived from ATCC® 9372™	084535700953AN
0957P	0957K	0957L	Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 33495™	084535700957AW
0958P	0958K	0958L	Parvimonas micra derived from ATCC® 33270™	084535700958AY
0959P	0959K	0959L	Enterococcus faecalis derived from ATCC® 51299™	084535700959B2
0963P	0963K	0963L	Staphylococcus capitis subsp. capitis derived from ATCC® 35661™	084535700963AR
0965P	0965K	0965L	Corynebacterium pseudodiphtheriticum derived from ATCC® 10700™	084535700965AV
0967P	0967K	0967L	Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 35666™	084535700967AZ
0968P	0968K	0968L	Enterococcus faecium derived from ATCC® 35667™	084535700968B3
0969P	0969K	0969L	Streptococcus mutans derived from ATCC® 35668™	084535700969B5
0971P	0971K	0971L	Elizabethkingia meningoseptica derived from ATCC® 13253™	084535700971AQ
0973P	0973K	0973L	Acinetobacter lwoffii derived from ATCC® 17925™	084535700973AU
0975P	0975K	0975L	Pseudomonas aeruginosa derived from ATCC® 35032™	084535700975AY
0976P	0976K	0976L	Staphylococcus epidermidis derived from ATCC® 49461™	084535700976B2
0978P	0978K	0978L	Streptococcus species derived from ATCC® 12392™	084535700978B6
0979P	0979K	0979L	Streptococcus pyogenes, (A,3) derived from ATCC® 12384™	084535700979B8
0980P	0980K	0980L	Lactococcus lactis derived from ATCC® 49032™	084535700980AR
0981P	0981K	0981L	Candida albicans derived from ATCC® 66027™	084535700981AT
0982P	0982K	0982L	Candida kefyr derived from ATCC® 66028™	084535700982AV
0983P	0983K	0983L	Candida tropicalis derived from ATCC® 66029™	084535700983AX
0984P	0984K	0984L	Cryptococcus albidus derived from ATCC® 66030™	084535700984AZ

0985P	0985K	0985L	Cryptococcus neoformans derived from ATCC® 66031™	084535700985B3
0986P	0986K	0986L	Candida glabrata derived from ATCC® 66032™	084535700986B5
0987P	0987K	0987L	Cryptococcus uniguttulatus derived from ATCC® 66033™	084535700987B7
0989P	0989K	0989L	Lactobacillus rhamnosus derived from ATCC® 9595™	084535700989BB
0990P	0990K	0990L	Candida kefyr derived from ATCC® 2512™	084535700990AU
0991P	0991K	0991L	Papiliotrema laurentii derived from ATCC® 66036™	084535700991AW
0992P	0992K	0992L	Candida glabrata derived from ATCC® 2001™	084535700992AY
0993P	0993K	0993L	Haemophilus influenzae derived from ATCC® 35056™	084535700993B2
0994P	0994K	0994L	Streptococcus pyogenes derived from ATCC® 49399™	084535700994B4
0995P	0995K	0995L	Mycobacterium gordonae derived from ATCC® 14470™	084535700995B6
0998P	0998K	0998L	Bacillus cereus derived from ATCC® 10876™	084535700998BC
0999P	0999K	0999L	Bacillus paranthracis derived from ATCC® 13061™	084535700999BE
01000P	01000K	01000L	Enterococcus faecium derived from ATCC® 700221™	0845357001000VQ
01003P	01003K	01003L	Legionella longbeachae derived from ATCC® 33462™	0845357001003VW
01005P	01005K	01005L	Klebsiella pneumoniae derived from ATCC® BAA-1705™	0845357001005W2
01006P	01006K	01006L	Klebsiella pneumoniae derived from ATCC® BAA-1706™	0845357001006W4
01007P	01007K	01007L	Staphylococcus aureus derived from ATCC® BAA-1708™	0845357001007W6
01008P	01008K	01008L	Bacillus badius derived from ATCC® 14574™	0845357001008W8
01009P	01009K	01009L	Pseudomonas aeruginosa derived from ATCC® 9721™	0845357001009WA
01010P	01010K	01010L	Pseudomonas aeruginosa derived from ATCC® BAA-1744™	0845357001010VT
01011P	01011K	01011L	Zygosaccharomyces parabailii derived from ATCC® MYA-4549™	0845357001011VV
01012P	01012K	01012L	Hanseniaspora valbyensis derived from ATCC® 58370™	0845357001012VX
01013P	01013K	01013L	Sporidiobolus salmonicolor derived from ATCC® MYA-4550™	0845357001013VZ
01018P	01018K	01018L	Enterobacter cloacae subsp. cloacae derived from ATCC® BAA-1143™	0845357001018WB
01019P	01019K	01019L	Klebsiella pneumoniae derived from ATCC® BAA-1144™	0845357001019WD
01020P	01020K	01020L	Stenotrophomonas maltophilia derived from ATCC® 13636™	0845357001020VW
01021P	01021K	01021L	Aspergillus fumigatus derived from ATCC® 204305™	0845357001021VY
01022P	01022K	01022L	Staphylococcus aureus subsp. aureus derived from ATCC® 700698™	0845357001022W2
01023P	01023K	01023L	Campylobacter coli derived from ATCC® 33559™	0845357001023W4
01024P	01024K	01024L	Haemophilus influenzae derived from NCTC 8468	0845357001024W6
01025P	01025K	01025L	Bifidobacterium bifidum derived from ATCC® 11863™	0845357001025W8
01035P	01035K	01035L	Staphylococcus saprophyticus derived from ATCC® 35552™	0845357001035WB
01036P	01036K	01036L	Candida tropicalis derived from ATCC® 1369™	0845357001036WD
01037P	01037K	01037L	Escherichia coli derived from ATCC® 15597™	0845357001037WF
01038P	01038K	01038L	Arcanobacterium haemolyticum derived from ATCC® BAA-1784™	0845357001038WH
01039P	01039K	01039L	Corynebacterium urealyticum derived from ATCC® 43044™	0845357001039WK
01040P	01040K	01040L	Corynebacterium renale derived from ATCC® BAA-1785™	0845357001040W4
01041P	N/A	N/A	Microbacterium paraoxydans derived from ATCC® BAA-1818™	0845357001041W6

01042P	01042K	01042L	Microbacterium liquefaciens derived from ATCC® BAA-1819™	0845357001042W8
01044P	01044K	01044L	Curtobacterium pusillum derived from ATCC® 19096™	0845357001044WC
01045P	01045K	01045L	Salmonella enterica subsp. diarizonae derived from ATCC® 29934™	0845357001045WE
01046P	01046K	01046L	Cellulosimicrobium cellulans derived from ATCC® BAA-1816™	0845357001046WG
01047P	01047K	01047L	Cellulosimicrobium cellulans derived from ATCC® BAA-1817™	0845357001047WJ
01048P	01048K	01048L	Clostridioides difficile derived from ATCC® BAA-1870™	0845357001048WL
01049P	01049K	01049L	Mycobacterium haemophilum derived from ATCC® 29548™	0845357001049WN
01050P	01050K	01050L	Escherichia coli derived from ATCC® 10799™	0845357001050W7
01051P	01051K	01051L	Cryptococcus gattii derived from ATCC® MYA-4560™	0845357001051W9
01052P	01052K	01052L	Enterococcus faecium derived from ATCC® 6057™	0845357001052WB
N/A	N/A	01053L	Mycoplasma bovis derived from ATCC® 25025™	0845357001053WD
01054P	01054K	01054L	Salmonella enterica subsp. diarizonae derived from ATCC® 12325™	0845357001054WF
01055P	01055K	01055L	Staphylococcus aureus subsp. aureus derived from ATCC® BAA-44™	0845357001055WH
01056P	01056K	01056L	Salmonella enterica subsp. enterica serovar Bispebjerg derived from ATCC® 9842™	0845357001056WK
01057P	01057K	01057L	Acinetobacter baumannii derived from ATCC® BAA-1605™	0845357001057WM
01060P	01060K	01060L	Klebsiella pneumoniae derived from ATCC® BAA-2146™	0845357001060WA
01062P	01062K	01062L	Escherichia coli derived from NCTC 8622	0845357001062WE
01063P	01063K	01063L	Cryptococcus gattii (B) derived from ATCC® 34877™	0845357001063WG
01065P	01065K	01065L	Staphylococcus aureus derived from NCTC 12493	0845357001065WL
01066P	01066K	01066L	Saccharomyces cerevisiae derived from ATCC® 18824™	0845357001066WN
01068P	01068K	01068L	Staphylococcus epidermidis derived from ATCC® 51625™	0845357001068WS
01075P	01075K	01075L	Pediococcus pentosaceus derived from ATCC® 33316™	0845357001075WP
01076P	01076K	01076L	Vibrio vulnificus derived from ATCC® 27562™	0845357001076WR
01077P	01077K	01077L	Streptococcus anginosus derived from ATCC® 33397™	0845357001077WT
01078P	01078K	01078L	Pseudomonas aeruginosa derived from ATCC® 13388™	0845357001078WV
01081P	01081K	01081L	Aureobasidium pullulans var. melanigenum derived from ATCC® 15233™	0845357001081WJ
01085P	01085K	01085L	Escherichia coli derived from NCTC 13351	0845357001085WS
01087P	01087K	01087L	Salmonella enterica subsp. salamae serotype Tranaroa derived from NCTC 10252	0845357001087WW
01088P	01088K	01088L	Cronobacter sakazakii derived from ATCC® 29544™	0845357001088WY
01089P	01089K	01089L	Enterococcus faecalis derived from ATCC® 51575™	0845357001089X2
01090P	01090K	01090L	Lactobacillus rhamnosus derived from ATCC® 53103™	0845357001090WK
01092P	01092K	01092L	Bifidobacterium animalis subsp. animalis derived from ATCC® 25527™	0845357001092WP
01094P	01094K	01094L	Chaetomium globosum derived from ATCC® 6205™	0845357001094WT
01095P	01095K	01095L	Salmonella enterica subsp. enterica serovar Newport derived from ATCC® 6962™	0845357001095WV
01097P	01097K	01097L	Escherichia coli (O145:NM) derived from CDC 99-3311	0845357001097WZ
01098P	01098K	01098L	Escherichia coli (O45:H2) derived from CDC 00-3039	0845357001098X3

01099P	01099K	01099L	Escherichia coli (O121:H19) derived from CDC 02-3211	0845357001099X5
01100P	01100K	01100L	Escherichia coli (O26:H11) derived from CDC 03-3014	0845357001100VV
01101P	01101K	01101L	Escherichia coli (O103:H11) derived from CDC 06-3008	0845357001101VX
01102P	01102K	01102L	Escherichia coli (O111:H8) derived from CDC 2010C-3114	0845357001102VZ
01103P	01103K	01103L	Salmonella enterica subsp. enterica serovar Enteritidis derived from ATCC® 49223™	0845357001103W3
01104P	01104K	01104L	Escherichia coli (O104:H4) derived from ATCC® BAA-2326™	0845357001104W5
01105P	01105K	01105L	Enterobacter cloacae derived from NCTC 13464	0845357001105W7
01108P	01108K	01108L	Corynebacterium jeikeium derived from ATCC® 43734™	0845357001108WD
01110P	01110K	01110L	Methylobacterium extorquens derived from ATCC® BAA-2500™	0845357001110VY
01111P	01111K	01111L	Enterobacter cloacae derived from NCTC 13406	0845357001111W2
01112P	01112K	01112L	Klebsiella pneumoniae derived from NCTC 13440	0845357001112W4
01113P	01113K	01113L	Escherichia coli derived from ATCC® BAA-2469™	0845357001113W6
01117P	01117K	01117L	Klebsiella pneumoniae derived from NCTC 13438	0845357001117WE
01122P	01122K	01122L	Staphylococcus aureus derived from ATCC® BAA-2312™	0845357001122W7
01130P	01130K	01130L	Aspergillus brasiliensis derived from ATCC® 9642™	0845357001130W6
01132P	01132K	01132L	Campylobacter lari derived from ATCC® 35221™	0845357001132WA
01133P	01133K	01133L	Penicillium citrinum derived from ATCC® 9849™	0845357001133WC
01134P	01134K	01134L	Alternaria species derived from ATCC® 20084™	0845357001134WE
01136P	01136K	01136L	Escherichia coli derived from NCTC 13476	0845357001136WJ
01139P	01139K	01139L	Candida dubliniensis derived from NCPF 3949	0845357001139WQ
01140P	01140K	01140L	Aspergillus caesiellus derived from ATCC® 42693™	0845357001140W9
01141P	01141K	01141L	Wallemia mellicola derived from ATCC® 42694™	0845357001141WB
01142P	01142K	01142L	Mucor racemosus derived from ATCC® 42647™	0845357001142WD
01143P	01143K	01143L	Enterococcus faecium derived from NCTC 12204	0845357001143WF
01144P	01144K	01144L	Lactobacillus plantarum subsp. plantarum derived from ATCC® 14917™	0845357001144WH
01148P	01148K	01148L	Klebsiella pneumoniae derived from NCTC 13442	0845357001148WR
01149P	01149K	01149L	Mycobacterium avium derived from ATCC® 700898™	0845357001149WT
01150P	01150K	01150L	Staphylococcus lugdunensis derived from NCTC 7990	0845357001150WC
01152P	01152K	01152L	Proteus vulgaris derived from ATCC® 29905™	0845357001152WG
01159P	01159K	01159L	Ralstonia pickettii derived from ATCC® 27511™	0845357001159WW
01160P	01160K	01160L	Eurotium rubrum derived from ATCC® 42690™	0845357001160WF
01167P	01167K	01167L	Brevundimonas vesicularis derived from ATCC® 11426™	0845357001167WV
01169P	01169K	01169L	Citrobacter freundii derived from ATCC® 8454™	0845357001169WZ
01170P	01170K	01170L	Salmonella enterica subsp. enterica serovar Paratyphi B derived from ATCC® 8759™	0845357001170WJ
01171P	01171K	01171L	Haemophilus parainfluenzae derived from ATCC® 9796™	0845357001171WL
01172P	01172K	01172L	Cryptococcus neoformans derived from ATCC® 13690™	0845357001172WN
01173P	01173K	01173L	Cryptococcus neoformans (D) derived from ATCC® 36556™	0845357001173WQ
01175P	01175K	01175L	Aggregatibacter actinomycetemcomitans (b) derived from ATCC® 29522™	0845357001175WU

01181P	01181K	01181L	Methylobacterium organophilum derived from ATCC® 27886™	0845357001181WP
01182P	01182K	01182L	Aspergillus flavus derived from ATCC® 9643™	0845357001182WR
01184P	01184K	01184L	Deinococcus radiophilus derived from ATCC® 27603™	0845357001184WV
01185P	01185K	01185L	Candida kefyr derived from ATCC® 8553™	0845357001185WX
01186P	01186K	01186L	Chryseobacterium indologenes derived from ATCC® 29897™	0845357001186WZ
01187P	01187K	01187L	Prevotella intermedia derived from ATCC® 25611™	0845357001187X3
01188P	01188K	01188L	Clostridium sporogenes derived from ATCC® 9690™	0845357001188X5
01189P	01189K	01189L	Staphylococcus epidermidis derived from ATCC® 35984™	0845357001189X7
01190P	01190K	01190L	Listeria monocytogenes derived from ATCC® 35152™	0845357001190WQ
01192P	01192K	01192L	Escherichia coli derived from NCTC 13216	0845357001192WU
01193P	01193K	01193L	Escherichia coli derived from NCTC 13167	0845357001193WW
01195P	01195K	01195L	Legionella anisa derived from ATCC® 35292™	0845357001195X2
01196P	01196K	01196L	Clostridium bifermentans derived from NCTC 506	0845357001196X4
01197P	01197K	01197L	Ralstonia pickettii derived from ATCC® 700591™	0845357001197X6
01200P	01200K	01200L	Penicillium rubens derived from ATCC® 11709™	0845357001200W2
01202P	01202K	01202L	Enterobacter cloacae derived from ATCC® BAA-2341™	0845357001202W6
01204P	01204K	01204L	Escherichia coli (O157:H7) derived from ATCC® 43890™	0845357001204WA
01209P	01209K	01209L	Shigella boydii (2) derived from ATCC® 8700™	0845357001209WL
01213P	N/A	N/A	Yersinia enterocolitica derived from NCTC 11174	0845357001213WB
01214P	01214K	01214L	Vibrio vulnificus derived from ATCC® 29307™	0845357001214WD
01215P	01215K	01215L	Proteus mirabilis derived from ATCC® 29906™	0845357001215WF
01229P	01229K	01229L	Escherichia coli derived from ATCC® BAA-2523™	0845357001229WS
01233P	01233K	01233L	Escherichia coli derived from ATCC® 51739™	0845357001233WH
01234P	01234K	01234L	Streptococcus agalactiae (III) derived from ATCC® BAA-22™	0845357001234WK
01235P	01235K	01235L	Shigella dysenteriae derived from ATCC® 13313™	0845357001235WM
01236P	01236K	01236L	Vibrio parahaemolyticus (o2:k3) derived from NCTC 10885	0845357001236WP
01237P	01237K	01237L	Campylobacter fetus subsp. fetus derived from ATCC® 27374™	0845357001237WR
01238P	01238K	01238L	Talaromyces pinophilus derived from ATCC® 11797™	0845357001238WT
01242P	01242K	01242L	Escherichia coli derived from ATCC® BAA-2452™	0845357001242WJ
01244P	01244K	01244L	Escherichia coli derived from NCTC 13846	0845357001244WN
01245P	N/A	N/A	Klebsiella pneumoniae derived from NCTC 13439	0845357001245WQ
01247P	01247K	01247L	Klebsiella pneumonia subsp. pneumoniae derived from ATCC® BAA-2524™	0845357001247WU
01256P	01256K	01256L	Candida auris derived from CDC B11903	0845357001256WV
01257P	01257K	01257L	Salmonella enterica subsp. enterica serotype Nottingham derived from NCTC7832	0845357001257WX
01259P	01259K	01259L	Escherichia coli derived from CDC AR-0346	0845357001259X3
01260P	01260K	01260L	Enterococcus faecium derived from ATCC® 8459™	0845357001260WL
01261P	01261K	01261L	Escherichia coli derived from ATCC® BAA-1429™	0845357001261WN
01262P	01262K	01262L	Levilactobacillus brevis derived from ATCC® 14869™	0845357001262WQ
01263P	01263K	01263L	Klebsiella pneumoniae derived from ATCC® BAA-2814™	0845357001263WS

01264P	01264K	01264L	Aspergillus terreus derived from ATCC® 1012™	0845357001264WU
01265P	01265K	01265L	Escherichia coli derived from NCTC 13353	0845357001265WW
01266P	01266K	01266L	Acinetobacter baumannii derived from NCTC 13304	0845357001266WY
01269P	01269K	01269L	Burkholderia cenocepacia derived from ATCC® BAA-245™	0845357001269X6
01270P	01270K	01270L	Burkholderia multivorans derived from ATCC® BAA-247™	0845357001270WP
01272P	01272K	01272L	Lactobacillus johnsonii derived from ATCC® 11506™	0845357001272WT
0158MRSA	N/A	N/A	Staphylococcus aureus subsp. aureus derived from ATCC® 700699™ (MRSA Live Culture Positive Control)	084535700158MRSAQC
0360MSSA	N/A	N/A	Staphylococcus aureus subsp. aureus derived from ATCC® 25923™ (MSSA Live Culture Positive Control)	084535700360MSSAP8
0371MSSE	N/A	N/A	Staphylococcus epidermidis derived from ATCC® 12228™ (MSSE Live Culture Positive Control)	084535700371MSSEQ8

Microbiologics

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Annex 2 to Declaration of Conformity

QC SETS AND PANELS KWIK-STIK

Catalog #	Product Description	Basic UDI-DI
5002P	CLSI® MIC QC Set without Klebsiella pneumoniae	0845357050029T
5027P	CLSI® M22 Chocolate Agar QC Set	084535705027AB
5062P	Dried Overnight – Gram Negative Panel QC Set	084535705062AD
5063P	Dried Overnight – Gram Positive Panel QC Set	084535705063AF
5065P	ANA (3 Strains) QC Set	084535705065AK
5066P	NH (4 Strains) QC Set	084535705066AM
5107P	Non-Fastidious Gram Negative (5 Strains) QC Set	084535705107AA
5112P	AST-GP (4 Strains) QC Set	084535705112A3
5184P	AST-YS QC Set	084535705184AU
5187P	AST-GP (5 Strains) QC Set	084535705187B2
5189P	AST-GN QC Set with Klebsiella	084535705189B6
5190P	ANC Comprehensive QC Set	084535705190AP
5191P	BCL Comprehensive QC Set	084535705191AR
5192P	GN Comprehensive QC Set	084535705192AT
5193P	GP Comprehensive QC Set	084535705193AV
5194P	NH Comprehensive QC Set	084535705194AX
5195P	YST Comprehensive QC Set	084535705195AZ
5209P	CBC QC Set	084535705209AK
5214P	GN Streamlined QC Set	084535705214AC
5215P	GP Streamlined QC Set	084535705215AE
5216P	ANC Streamlined QC Set	084535705216AG
5220P	AST-GP (6 Strains) QC Set	084535705220A7
5225P	Gram-Positive Blood Culture Control Panel (Live Culture)	084535705225AH
5246P	Anaerobe Identification Panel	084535705246AR
5247P	Gram Negative Identification Panel	084535705247AT
5248P	Gram Positive Identification Panel	084535705248AV
5252P	Yeast Identification Panel	084535705252AL
8164	GBS QC Set (Live Culture)	084535708164BB
8171	C. difficile QC Set (Live Culture)	084535708171B8
8173	GBS Verification Panel (Live Culture)	084535708173BC
8174	MRSA Organism Set (Live Culture)	084535708174BE
8178	SA Organism Set (Live Culture)	084535708178BN
8179	Enteric Bacterial Verification Panel (Live Culture)	084535708179BQ
8243	Cdiff Verification Panel (Live Culture)	084535708243B8