

KWIK-STIK and LYFO DISK

| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|------------------------|------------------|------------------|--|------------|-----|--|
| Acetobacter | | | | | | |
| 0511L | 0511K | 0511P | Acetobacter aceti derived from ATCC® 15973™* | C | 1 | |
| Achromobacter | | | | | | |
| 0516L | 0516K | 0516P | Achromobacter xylosoxidans derived from ATCC® 27061™* | D | 2 | |
| Acinetobacter | | | | | | |
| 0357L | 0357K | 0357P | Acinetobacter baumannii derived from ATCC® 19606™* | B | 2 | |
| 01057L | 01057K | 01057P | Acinetobacter baumannii derived from ATCC® BAA-1605™* | E | 2 | Multi-drug resistant |
| 0119L | 0119K | 0119P | Acinetobacter baumannii derived from ATCC® BAA-747™* | D | 2 | |
| 0599L | 0599K | 0599P | Acinetobacter baumannii derived from NCIMB 12457 | C | 1 | |
| 01266L | 01266K | 01266P | Acinetobacter baumannii derived from NCTC 13304 | C | 2 | OXA-27 |
| 0468L | 0468K | 0468P | Acinetobacter lwoffii derived from ATCC® 15309™* | C | 2 | |
| 0973L | 0973K | 0973P | Acinetobacter lwoffii derived from ATCC® 17925™* | D | 2 | |
| 0635L | 0635K | 0635P | Acinetobacter species derived from ATCC® 49139™* | C | 2 | |
| Actinobacillus | | | | | | |
| 0658L | 0658K | 0658P | Actinobacillus pleuropneumoniae derived from ATCC® 27088™* | D | 2 | |
| Actinomyces | | | | | | |
| 0939L | 0939K | 0939P | Actinomyces odontolyticus derived from ATCC® 17929™* | D | 2 | |
| 0750L | 0750K | 0750P | Actinomyces viscosus derived from ATCC® 15987™* | D | 2 | |
| Aerococcus | | | | | | |
| 0263L | 0263K | 0263P | Aerococcus viridans derived from ATCC® 10400™* | C | 1 | |
| 0276L | 0276K | 0276P | Aerococcus viridans derived from ATCC® 11563™* | C | 1 | |
| 0746L | 0746K | 0746P | Aerococcus viridans derived from ATCC® 700406™* | D | 1 | |
| Aeromonas | | | | | | |
| 0910L | 0910K | 0910P | Aeromonas hydrophila derived from ATCC® 35654™* | C | 2 | |
| 0637L | 0637K | 0637P | Aeromonas hydrophila derived from ATCC® 49140™* | C | 2 | |
| 0870L | 0870K | 0870P | Aeromonas hydrophila derived from ATCC® 7966™* | C | 2 | |
| 0290L | 0290K | 0290P | Aeromonas salmonicida derived from ATCC® 7965™* | C | 2 | Formerly Aeromonas hydrophila |
| Aggregatibacter | | | | | | |
| 01175L | 01175K | 01175P | Aggregatibacter actinomycetemcomitans (b) derived from ATCC® 29522™* | D | 2 | Serotype b |
| 0467L | 0467K | 0467P | Aggregatibacter aphrophilus derived from ATCC® 29241™* | E | 2 | V-dependant; Formerly Haemophilus paraphrophilus |
| 0184L | 0184K | 0184P | Aggregatibacter aphrophilus derived from ATCC® 33389™* | D | 1 | |
| 0645L | 0645K | 0645P | Aggregatibacter aphrophilus derived from ATCC® 49146™* | D | 2 | Formerly Haemophilus paraphrophilus |



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| 0249L | 0249K | 0249P | Aggregatibacter aphrophilus derived from ATCC® 49917™* | D | 2 | Formerly Haemophilus paraphrophilus |
| 0411L | 0411K | 0411P | Aggregatibacter aphrophilus derived from ATCC® 7901™* | C | 2 | Formerly Haemophilus parainfluenzae |
| Alcaligenes | | | | | | |
| 0911L | 0911K | 0911P | Alcaligenes faecalis subsp. faecalis derived from ATCC® 35655™* | C | 1 | |
| 0402L | 0402K | 0402P | Alcaligenes faecalis subsp. faecalis derived from ATCC® 8750™* | C | 1 | |
| Alicyclobacillus | | | | | | |
| 01059L | 01059K | 01059P | Alicyclobacillus acidocaldarius subsp. acidocaldarius derived from NCIMB 11725 | D | 1 | |
| 0265L | 0265K | 0265P | Alicyclobacillus acidoterrestris derived from ATCC® 49025™* | E | 1 | |
| Alternaria | | | | | | |
| 01134L | 01134K | 01134P | Alternaria species derived from ATCC® 20084™* | D | 1 | Mold |
| Aneurinibacillus | | | | | | |
| 0141L | 0141K | 0141P | Aneurinibacillus aneurinilyticus derived from ATCC® 11376™* | C | 1 | Formerly Aneurinibacillus aneurinolyticus |
| Arcanobacterium | | | | | | |
| 01038L | 01038K | 01038P | Arcanobacterium haemolyticum derived from ATCC® BAA-1784™* | D | 2 | |
| Arthrobacter | | | | | | |
| 0260L | 0260K | 0260P | Arthrobacter psychrolactophilus derived from ATCC® 700733™* | E | 1 | Psychrotroph |
| Aspergillus | | | | | | |
| 0392L | 0392K | 0392P | Aspergillus brasiliensis derived from ATCC® 16404™* | E | 1 | Mold |
| 01130L | 01130K | 01130P | Aspergillus brasiliensis derived from ATCC® 9642™* | E | 1 | Mold |
| 01140L | 01140K | 01140P | Aspergillus caesiellus derived from ATCC® 42693™* | E | 1 | Mold |
| 01182L | 01182K | 01182P | Aspergillus flavus derived from ATCC® 9643™* | E | 1 | Mold |
| 01021L | 01021K | 01021P | Aspergillus fumigatus derived from ATCC® 204305™* | E | 2 | Mold |
| 0245L | 0245K | 0245P | Aspergillus niger derived from ATCC® 16888™* | E | 1 | Mold |
| 0500L | 0500K | 0500P | Aspergillus niger derived from ATCC® 6275™* | E | 1 | Mold |
| 0177L | 0177K | 0177P | Aspergillus oryzae derived from ATCC® 10124™* | E | 1 | Mold |
| 01264L | 01264K | 01264P | Aspergillus terreus derived from ATCC® 1012™* | D | 1 | |
| Aureobasidium | | | | | | |
| 01253L | 01253K | 01253P | Aureobasidium pullulans derived from ATCC® 11942™* | E | 1 | Mold |
| 01081L | 01081K | 01081P | Aureobasidium pullulans var. melanigenum derived from ATCC® 15233™* | E | 1 | Mold |
| Bacillus | | | | | | |
| 0953L | 0953K | 0953P | Bacillus atrophaeus derived from ATCC® 9372™* | B | 1 | |



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| 01008L | 01008K | 01008P | Bacillus badius derived from ATCC® 14574™* | B | 1 | |
| 0998L | 0998K | 0998P | Bacillus cereus derived from ATCC® 10876™* | B | 1 | |
| 0256L | 0256K | 0256P | Bacillus cereus derived from ATCC® 11778™* | C | 1 | |
| 0999L | 0999K | 0999P | Bacillus cereus derived from ATCC® 13061™* | B | 1 | |
| 0200L | 0200K | 0200P | Bacillus cereus derived from ATCC® 14579™* | D | 1 | |
| 0198L | 0198K | 0198P | Bacillus cereus derived from ATCC® 33019™* | E | 2 | |
| 0330L | 0330K | 0330P | Bacillus cereus derived from NCIMB 7464 | B | 2 | |
| 0507L | 0507K | 0507P | Bacillus circulans derived from ATCC® 4516™* | E | 1 | |
| 0140L | 0140K | 0140P | Bacillus circulans derived from ATCC® 61™* | C | 1 | |
| 0596L | 0596K | 0596P | Bacillus coagulans derived from ATCC® 7050™* | C | 1 | |
| 0812L | 0812K | 0812P | Bacillus licheniformis derived from ATCC® 12759™* | B | 1 | |
| 0799L | 0799K | 0799P | Bacillus licheniformis derived from ATCC® 14580™* | D | 1 | |
| 0201L | 0201K | 0201P | Bacillus megaterium derived from ATCC® 14581™* | C | 1 | |
| 0116L | 0116K | 0116P | Bacillus megaterium derived from ATCC® 9885™* | C | 1 | |
| 0258L | 0258K | 0258P | Bacillus pumilus derived from ATCC® 14884™* | C | 1 | |
| 0577L | 0577K | 0577P | Bacillus pumilus derived from ATCC® 700814™* | D | 1 | |
| 0474L | 0474K | 0474P | Bacillus pumilus derived from ATCC® BAA-1434™* | C | 1 | |
| 0269L | 0269K | 0269P | Bacillus subtilis derived from ATCC® 11774™* | C | 1 | |
| 0540L | 0540K | 0540P | Bacillus subtilis derived from ATCC® 19659™* | C | 1 | |
| 0486L | 0486K | 0486P | Bacillus subtilis subsp. spizizenii derived from ATCC® 6633™* | B | 1 | |
| 0120L | 0120K | 0120P | Bacillus subtilis subsp. subtilis derived from ATCC® 6051™* | C | 1 | |
| 0270L | 0270K | 0270P | Bacillus thuringiensis derived from ATCC® 10792™* | C | 1 | |
| 0539L | 0539K | 0539P | Bacillus thuringiensis derived from ATCC® 33679™* | D | 1 | Serotype H3:3a,3b |
| Bacteroides | | | | | | |
| 0940L | 0940K | 0940P | Bacteroides fragilis derived from ATCC® 23745™* | D | 2 | |
| 0320L | 0320K | 0320P | Bacteroides fragilis derived from ATCC® 25285™* | D | 2 | Beta-lactamase positive; CLSI QC strain for Anaerobic Antimicrobial Susceptibility Testing |
| 0358L | 0358K | 0358P | Bacteroides fragilis derived from NCTC 9343 | D | 2 | |
| 0400L | 0400K | 0400P | Bacteroides ovatus derived from ATCC® 8483™* | D | 2 | |
| 0585L | 0585K | 0585P | Bacteroides ovatus derived from ATCC® BAA-1296™* | D | 2 | |
| 0587L | 0587K | 0587P | Bacteroides ovatus derived from ATCC® BAA-1304™* | D | 2 | |
| 0319L | 0319K | 0319P | Bacteroides thetaiotaomicron derived from ATCC® 29741™* | D | 2 | Beta-lactamase positive; CLSI QC strain for Anaerobic Antimicrobial Susceptibility Testing |



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| 0619L | 0619K | 0619P | Bacteroides uniformis derived from ATCC® 8492™* | D | 2 | |
| 0445L | 0445K | 0445P | Bacteroides vulgatus derived from ATCC® 8482™* | D | 2 | |
| Bifidobacterium | | | | | | |
| 01092L | 01092K | 01092P | Bifidobacterium animalis subsp. animalis derived from ATCC® 25527™* | E | 1 | |
| 01025L | 01025K | 01025P | Bifidobacterium bifidum derived from ATCC® 11863™* | E | 1 | |
| 0175L | 0175K | 0175P | Bifidobacterium breve derived from ATCC® 15700™* | D | 1 | |
| Bordetella | | | | | | |
| 0655L | 0655K | 0655P | Bordetella bronchiseptica derived from ATCC® 10580™* | D | 2 | |
| 0671L | 0671K | 0671P | Bordetella bronchiseptica derived from ATCC® 4617™* | D | 2 | |
| 0842L | 0842K | 0842P | Bordetella parapertussis derived from ATCC® 15311™* | D | 2 | |
| 0489L | 0489K | 0489P | Bordetella pertussis derived from ATCC® 8467™* | D | 2 | |
| 0843L | 0843K | 0843P | Bordetella pertussis derived from ATCC® 9797™* | D | 2 | |
| Brevibacillus | | | | | | |
| 0139L | 0139K | 0139P | Brevibacillus agri derived from ATCC® 51663™* | C | 1 | |
| 0144L | 0144K | 0144P | Brevibacillus laterosporus derived from ATCC® 64™* | C | 1 | |
| Brevundimonas | | | | | | |
| 0754L | 0754K | 0754P | Brevundimonas diminuta derived from ATCC® 11568™* | B | 1 | |
| 0805L | 0805K | 0805P | Brevundimonas diminuta derived from ATCC® 19146™* | D | 1 | |
| 01167L | 01167K | 01167P | Brevundimonas vesicularis derived from ATCC® 11426™* | D | 1 | |
| Brochothrix | | | | | | |
| 0150L | 0150K | 0150P | Brochothrix thermosphacta derived from ATCC® 11509™* | D | 1 | |
| Burkholderia | | | | | | |
| 01269L | 01269K | 01269P | Burkholderia cenocepacia derived from ATCC® BAA-245™* | C | 2 | |
| 0488L | 0488K | 0488P | Burkholderia cepacia derived from ATCC® 25416™* | C | 2 | |
| 0836L | 0836K | 0836P | Burkholderia cepacia derived from ATCC® 25608™* | C | 2 | |
| 01270L | 01270K | 01270P | Burkholderia multivorans derived from ATCC® BAA-247™* | C | 2 | |
| Campylobacter | | | | | | |
| 01023L | 01023K | 01023P | Campylobacter coli derived from ATCC® 33559™* | E | 2 | |
| 0121L | 0121K | 0121P | Campylobacter coli derived from ATCC® 43478™* | E | 2 | |
| 01237L | 01237K | 01237P | Campylobacter fetus subsp. fetus derived from ATCC® 27374™* | D | 2 | Quality Control Strain for FDA Bacteriological Analytical Manual |
| 0325L | 0325K | 0325P | Campylobacter jejuni subsp. jejuni derived from ATCC® 29428™* | E | 2 | |



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| 0481L | 0481K | 0481P | Campylobacter jejuni subsp. jejuni derived from ATCC® 33291™* | E | 2 | |
| 0111L | 0111K | 0111P | Campylobacter jejuni subsp. jejuni derived from ATCC® 33560™* | E | 2 | EUCAST QC strain for Disk Diffusion and MIC determination |
| 0525L | 0525K | 0525P | Campylobacter jejuni subsp. jejuni derived from ATCC® 43430™* | E | 2 | Serotype O:2 |
| 0251L | 0251K | 0251P | Campylobacter jejuni subsp. jejuni derived from ATCC® 49943™* | E | 2 | |
| 0188L | 0188K | 0188P | Campylobacter jejuni subsp. jejuni derived from ATCC® BAA-1153™* | E | 2 | |
| 0718L | 0718K | 0718P | Campylobacter jejuni subsp. jejuni derived from NCTC 11322 | E | 2 | |
| 0712L | 0712K | 0712P | Campylobacter jejuni subsp. jejuni derived from NCTC 13367 | E | 2 | |
| 01132L | 01132K | 01132P | Campylobacter lari derived from ATCC® 35221™* | E | 1 | |
| Candida | | | | | | |
| 0443L | 0443K | 0443P | Candida albicans derived from ATCC® 10231™* | D | 1 | Yeast cells |
| 0332L | 0332K | 0332P | Candida albicans derived from ATCC® 14053™* | D | 1 | Yeast cells |
| 0896L | 0896K | 0896P | Candida albicans derived from ATCC® 2091™* | D | 1 | Yeast cells |
| 0800L | 0800K | 0800P | Candida albicans derived from ATCC® 24433™* | D | 1 | Yeast cells |
| 0425L | 0425K | 0425P | Candida albicans derived from ATCC® 60193™* | D | 1 | Yeast cells |
| 0981L | 0981K | 0981P | Candida albicans derived from ATCC® 66027™* | D | 1 | Yeast cells |
| 0264L | 0264K | 0264P | Candida albicans derived from ATCC® 90028™* | D | 1 | CLSI QC strain for Antimicrobial Susceptibility Testing using the Disk Diffusion Method; yeast cells |
| 0250L | 0250K | 0250P | Candida albicans derived from ATCC® 90029™* | D | 1 | Yeast cells |
| 0379L | 0379K | 0379P | Candida albicans derived from NCYC 1363 | D | 1 | Yeast cells |
| 01256L | 01256K | 01256P | Candida auris derived from CDC B11903 | E | 2 | Yeast cells |
| 01139L | 01139K | 01139P | Candida dubliniensis derived from NCPF 3949 | D | 1 | Yeast cells |
| 0520L | 0520K | 0520P | Candida geochares derived from ATCC® 36852™* | D | 1 | Yeast cells |
| 0737L | 0737K | 0737P | Candida glabrata derived from ATCC® 15126™* | D | 1 | Yeast cells |
| 0992L | 0992K | 0992P | Candida glabrata derived from ATCC® 2001™* | D | 1 | Yeast cells |
| 0986L | 0986K | 0986P | Candida glabrata derived from ATCC® 66032™* | D | 1 | Yeast cells |
| 0122L | 0122K | 0122P | Candida glabrata derived from ATCC® MYA-2950™* | E | 1 | Yeast cells |
| 0990L | 0990K | 0990P | Candida kefyr derived from ATCC® 2512™* | D | 1 | Yeast cells |
| 0982L | 0982K | 0982P | Candida kefyr derived from ATCC® 66028™* | D | 1 | Yeast cells |
| 01185L | 01185K | 01185P | Candida kefyr derived from ATCC® 8553™* | D | 1 | Yeast cells |
| 0809L | 0809K | 0809P | Candida krusei derived from ATCC® 14243™* | D | 1 | Yeast cells |
| 0510L | 0510K | 0510P | Candida krusei derived from ATCC® 34135™* | E | 1 | Yeast cells |



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| 0774L | 0774K | 0774P | <i>Candida lusitanae</i> derived from ATCC® 34449™* | D | 1 | Yeast cells |
| 0726L | 0726K | 0726P | <i>Candida parapsilosis</i> derived from ATCC® 22019™* | D | 1 | CLSI QC strain for Disk Diffusion and MIC determination; yeast cells |
| 01036L | 01036K | 01036P | <i>Candida tropicalis</i> derived from ATCC® 1369™* | E | 1 | Yeast cells |
| 0450L | 0450K | 0450P | <i>Candida tropicalis</i> derived from ATCC® 13803™* | D | 1 | Yeast cells |
| 0983L | 0983K | 0983P | <i>Candida tropicalis</i> derived from ATCC® 66029™* | D | 1 | Yeast cells |
| 0847L | 0847K | 0847P | <i>Candida tropicalis</i> derived from ATCC® 750™* | D | 1 | CLSI QC strain for Disk Diffusion; yeast cells |
| 0897L | 0897K | 0897P | <i>Candida tropicalis</i> derived from ATCC® 9968™* | D | 1 | Yeast cells |
| 0779L | 0779K | 0779P | <i>Candida utilis</i> derived from ATCC® 9950™* | D | 1 | Yeast cells |
| Capnocytophaga | | | | | | |
| 0749L | 0749K | 0749P | <i>Capnocytophaga sputigena</i> derived from ATCC® 33612™* | C | 2 | |
| Cedecea | | | | | | |
| 0517L | 0517K | 0517P | <i>Cedecea neteri</i> derived from ATCC® 33855™* | E | 2 | |
| Cellulosimicrobium | | | | | | |
| 01046L | 01046K | 01046P | <i>Cellulosimicrobium cellulans</i> derived from ATCC® BAA-1816™* | D | 1 | |
| 01047L | 01047K | 01047P | <i>Cellulosimicrobium cellulans</i> derived from ATCC® BAA-1817™* | D | 1 | |
| 0294L | 0294K | 0294P | <i>Cellulosimicrobium cellulans</i> derived from ATCC® 27402™* | D | 1 | |
| Chaetomium | | | | | | |
| 01094L | 01094K | 01094P | <i>Chaetomium globosum</i> derived from ATCC® 6205™* | E | 1 | Mold |
| Chryseobacterium | | | | | | |
| 01186L | 01186K | 01186P | <i>Chryseobacterium indologenes</i> derived from ATCC® 29897™* | E | 2 | |
| 0164L | 0164K | 0164P | <i>Chryseobacterium shigense</i> derived from ATCC® 51823™* | E | 1 | |
| Citrobacter | | | | | | |
| 0478L | 0478K | 0478P | <i>Citrobacter braakii</i> derived from ATCC® 43162™* | C | 1 | |
| 0578L | 0578K | 0578P | <i>Citrobacter braakii</i> derived from ATCC® 51113™* | D | 1 | |
| 0229L | 0229K | 0229P | <i>Citrobacter freundii</i> derived from ATCC® 43864™* | B | 1 | |
| 0315L | 0315K | 0315P | <i>Citrobacter freundii</i> derived from ATCC® 8090™* | C | 1 | |
| 01169L | 01169K | 01169P | <i>Citrobacter freundii</i> derived from ATCC® 8454™* | C | 1 | |
| 0574L | 0574K | 0574P | <i>Citrobacter freundii</i> derived from NCTC 9750 | C | 1 | |
| 0106L | 0106K | 0106P | <i>Citrobacter koseri</i> derived from ATCC® 27156™* | E | 1 | |
| Cladosporium | | | | | | |
| 0537L | 0537K | 0537P | <i>Cladosporium cladosporioides</i> derived from ATCC® 16022™* | E | 1 | Mold |



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| 01254L | 01254K | 01254P | Cladosporium sphaerospermum derived from ATCC® 11289™* | E | 1 | Mold |
| Clostridioides | | | | | | |
| 0329L | 0329K | 0329P | Clostridioides difficile derived from ATCC® 9689™* | D | 2 | Genotype: tcdA positive, tcdB positive; cdtB negative; Toxinotype 0; Ribotype 001; Produces cytotoxin |
| 01206L | 01206K | 01206P | Clostridioides difficile derived from ATCC® 43255™* | D | 2 | Formerly Clostridium difficile; tcdA positive, tcdB positive; Toxinotype 0; Ribotype 087 |
| 0833L | 0833K | 0833P | Clostridioides difficile derived from ATCC® 43593™* | D | 2 | Formerly Clostridium difficile; Genotype: tcdA negative, tcdB negative; cdtB negative; Toxin A and B negative; Ribotype: 060; Serogroup B |
| 0527L | 0527K | 0527P | Clostridioides difficile derived from ATCC® 700057™* | D | 2 | Formerly Clostridium difficile; Beta-lactamase positive; nontoxigenic; CLSI QC strain for Anaerobic Antimicrobial Susceptibility Testing; Genotype: tcdA negative, tcdB negative; cdtB negative; Ribotype 038 |
| 01048L | 01048K | 01048P | Clostridioides difficile derived from ATCC® BAA-1870™* | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive; tcdA positive, tcdB positive; cdtB positive; Binary Toxin positive; Toxinotype IIIb; Ribotype 027; PFGE Type NAP1; REA type BI 8 |
| 01161L | 01161K | 01161P | Clostridioides difficile derived from CDC 20110736 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA negative, tcdB negative, tcdC _deletion negative; cdtA negative, cdtB negative PFGE Type: Unnamed Isolation: NY 2011 |
| 01162L | 01162K | 01162P | Clostridioides difficile derived from CDC 20110995 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive, tcdC _deletion 0 bp; cdtA negative, cdtB negative PFGE Type: NAP11 Isolation: MN 2011 |
| 01163L | 01163K | 01163P | Clostridioides difficile derived from CDC 20120296 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive, tcdC _deletion 39 bp; cdtA positive, cdtB positive PFGE Type: NAP8 Isolation: GA 2011 |
| 01164L | 01164K | 01164P | Clostridioides difficile derived from CDC 20120905 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive, tcdC _deletion 0 bp; cdtA negative, cdtB negative; PFGE; Isolation: MN 2011 |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|------------------------|------------------|------------------|---|------------|-----|--|
| 01165L | 01165K | 01165P | Clostridioides difficile derived from CDC 20121308 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive, tcdC_deletion 18 bp; cdtA positive, cdtB positive PFGE Type: NAP1 Isolation: NY 2011 |
| 01166L | 01166K | 01166P | Clostridioides difficile derived from CDC 20131122 | D | 2 | Formerly Clostridium difficile; Genotype: tcdA positive, tcdB positive, tcdC_deletion 39 bp; cdtA positive, cdtB positive PFGE Type: NAP7 Isolation: CO 2011 |
| Clostridium | | | | | | |
| 0828L | 0828K | 0828P | Clostridium bifermentans derived from ATCC® 638™* | D | 1 | |
| 01196L | 01196K | 01196P | Clostridium bifermentans derived from NCTC 506 | D | 2 | |
| 0327L | 0327K | 0327P | Clostridium histolyticum derived from ATCC® 19401™* | D | 2 | |
| 0801L | 0801K | 0801P | Clostridium perfringens derived from ATCC® 12915™* | E | 2 | |
| 0123L | 0123K | 0123P | Clostridium perfringens derived from ATCC® 12916™* | D | 2 | |
| 0674L | 0674K | 0674P | Clostridium perfringens derived from ATCC® 12919™* | D | 2 | Export license required for shipping outside the U.S. |
| 0318L | 0318K | 0318P | Clostridium perfringens derived from ATCC® 13124™* | D | 2 | |
| 0257L | 0257K | 0257P | Clostridium perfringens derived from ATCC® 3624™* | D | 2 | |
| 0572L | 0572K | 0572P | Clostridium perfringens derived from NCTC 8237 | D | 2 | Type A |
| 0547L | 0547K | 0547P | Clostridium perfringens derived from NCTC 8678 | D | 2 | Non-epsilon toxin producing; heat resistant |
| 0586L | 0586K | 0586P | Clostridium septicum derived from ATCC® 12464™* | D | 2 | |
| 0331L | 0331K | 0331P | Clostridium sordellii derived from ATCC® 9714™* | D | 2 | |
| 0487L | 0487K | 0487P | Clostridium sporogenes derived from ATCC® 11437™* | D | 1 | |
| 0317L | 0317K | 0317P | Clostridium sporogenes derived from ATCC® 19404™* | D | 1 | |
| 0676L | 0676K | 0676P | Clostridium sporogenes derived from ATCC® 3584™* | D | 1 | |
| 01188L | 01188K | 01188P | Clostridium sporogenes derived from ATCC® 9690™* | D | 2 | |
| 0580L | 0580K | 0580P | Clostridium sporogenes derived from NCIMB 12343 | D | 1 | |
| Corynebacterium | | | | | | |
| 0844L | 0844K | 0844P | Corynebacterium diphtheriae derived from ATCC® 13812™* | D | 2 | |
| 01108L | 01108K | 01108P | Corynebacterium jeikeium derived from ATCC® 43734™* | D | 2 | |
| 0965L | 0965K | 0965P | Corynebacterium pseudodiphtheriticum derived from ATCC® 10700™* | D | 1 | |
| 0884L | 0884K | 0884P | Corynebacterium pseudodiphtheriticum derived from ATCC® 10701™* | D | 1 | |
| 0293L | 0293K | 0293P | Corynebacterium renale derived from ATCC® 19412™* | D | 2 | |
| 01040L | 01040K | 01040P | Corynebacterium renale derived from ATCC® BAA-1785™* | D | 2 | |



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|-------------------------|------------------|------------------|--|------------|-----|--|
| 0583L | 0583K | 0583P | Corynebacterium striatum derived from ATCC® BAA-1293™* | D | 1 | |
| 01039L | 01039K | 01039P | Corynebacterium urealyticum derived from ATCC® 43044™* | D | 2 | |
| 0153L | 0153K | 0153P | Corynebacterium xerosis derived from ATCC® 373™* | D | 1 | |
| Cronobacter | | | | | | |
| 0756L | 0756K | 0756P | Cronobacter muytjensii derived from ATCC® 51329™* | D | 1 | |
| 01088L | 01088K | 01088P | Cronobacter sakazakii derived from ATCC® 29544™* | E | 1 | |
| Cryptococcus | | | | | | |
| 0984L | 0984K | 0984P | Cryptococcus albidus derived from ATCC® 66030™* | D | 2 | Yeast cells |
| 01051L | 01051K | 01051P | Cryptococcus gattii derived from ATCC® MYA-4560™* | E | 2 | Yeast cells |
| 0333L | 0333K | 0333P | Cryptococcus laurentii derived from ATCC® 18803™* | D | 1 | Yeast cells |
| 0991L | 0991K | 0991P | Cryptococcus laurentii derived from ATCC® 66036™* | D | 2 | Yeast cells |
| 0451L | 0451K | 0451P | Cryptococcus liquefaciens derived from ATCC® 34140™* | D | 2 | Formerly Cryptococcus albidus; yeast cells |
| 01173L | 01173K | 01173P | Cryptococcus neoformans (D) derived from ATCC® 36556™* | D | 2 | Serotype D |
| 01172L | 01172K | 01172P | Cryptococcus neoformans derived from ATCC® 13690™* | D | 2 | Yeast cells |
| 0291L | 0291K | 0291P | Cryptococcus neoformans derived from ATCC® 14116™* | D | 2 | Yeast cells |
| 0781L | 0781K | 0781P | Cryptococcus neoformans derived from ATCC® 204092™* | D | 2 | Yeast cells |
| 0334L | 0334K | 0334P | Cryptococcus neoformans derived from ATCC® 32045™* | D | 2 | Yeast cells |
| 01063L | 01063K | 01063P | Cryptococcus neoformans derived from ATCC® 34877™* | D | 2 | Serotype B; yeast cells |
| 0985L | 0985K | 0985P | Cryptococcus neoformans derived from ATCC® 66031™* | D | 2 | Yeast cells |
| 0987L | 0987K | 0987P | Cryptococcus uniguttulatus derived from ATCC® 66033™* | D | 2 | Yeast cells |
| Curtobacterium | | | | | | |
| 01044L | 01044K | 01044P | Curtobacterium pusillum derived from ATCC® 19096™* | D | 1 | |
| Cutibacterium | | | | | | |
| 0170L | 0170K | 0170P | Cutibacterium acnes derived from ATCC® 6919™* | E | 1 | |
| 0419L | 0419K | 0419P | Cutibacterium acnes derived from ATCC® 11827™* | D | 1 | |
| Deinococcus | | | | | | |
| 01184L | 01184K | 01184P | Deinococcus radiophilus derived from ATCC® 27603™* | E | 1 | |
| Desulfotomaculum | | | | | | |
| 01026L | 01026K | 01026P | Desulfotomaculum nigrificans derived from ATCC® 7946™* | E | 1 | |
| Edwardsiella | | | | | | |
| 0845L | 0845K | 0845P | Edwardsiella tarda derived from ATCC® 15947™* | C | 2 | |



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|------------------------|------------------|------------------|---|------------|-----|--|
| Eggerthella | | | | | | |
| 0936L | 0936K | 0936P | Eggerthella lenta derived from ATCC® 43055™* | D | 1 | CLSI QC strain for Anaerobic Antimicrobial Susceptibility Testing |
| Eikenella | | | | | | |
| 0189L | 0189K | 0189P | Eikenella corrodens derived from ATCC® BAA-1152™* | C | 2 | |
| Elizabethkingia | | | | | | |
| 0971L | 0971K | 0971P | Elizabethkingia meningoseptica derived from ATCC® 13253™* | C | 2 | |
| Enterobacter | | | | | | |
| 01202L | 01202K | 01202P | Enterobacter cloacae derived from ATCC® BAA-2341™* | C | 2 | blaKPC positive; produces carbapenemase KPC; carbapenem-resistant (Imipenem and Ertapenem) |
| 01111L | 01111K | 01111P | Enterobacter cloacae derived from NCTC 13406 | D | 2 | Control for AmpC detection tests |
| 01105L | 01105K | 01105P | Enterobacter cloacae derived from NCTC 13464 | D | 2 | blaCTX-M group 9 gene; CTX-M extended-spectrum (beta)-lactamases |
| 0323L | 0323K | 0323P | Enterobacter cloacae subsp. cloacae derived from ATCC® 13047™* | B | 1 | |
| 0313L | 0313K | 0313P | Enterobacter cloacae subsp. cloacae derived from ATCC® 23355™* | B | 1 | |
| 0388L | 0388K | 0388P | Enterobacter cloacae subsp. cloacae derived from ATCC® 35030™* | B | 1 | |
| 0639L | 0639K | 0639P | Enterobacter cloacae subsp. cloacae derived from ATCC® 49141™* | C | 1 | |
| 01018L | 01018K | 01018P | Enterobacter cloacae subsp. cloacae derived from ATCC® BAA-1143™* | D | 2 | High level producer of chromosomal Amp C beta-lactamase; strong positive control strain for the AmpC disk test |
| 0755L | 0755K | 0755P | Enterobacter hormaechei derived from ATCC® 700323™* | C | 1 | Formerly Enterobacter cloacae subsp. cloacae |
| Enterococcus | | | | | | |
| 0761L | 0761K | 0761P | Enterococcus casseliflavus derived from ATCC® 700327™* | B | 1 | |
| 0623L | 0623K | 0623P | Enterococcus durans derived from ATCC® 11576™* | C | 1 | |
| 0651L | 0651K | 0651P | Enterococcus durans derived from ATCC® 6056™* | C | 1 | |
| 0181L | 0181K | 0181P | Enterococcus faecalis derived from ATCC® 14506™* | C | 2 | |
| 0367L | 0367K | 0367P | Enterococcus faecalis derived from ATCC® 19433™* | C | 2 | |
| 0366L | 0366K | 0366P | Enterococcus faecalis derived from ATCC® 29212™* | C | 2 | EUCAST QC strain for Disk Diffusion; CLSI and EUCAST QC strain for MIC Determination |
| 0197L | 0197K | 0197P | Enterococcus faecalis derived from ATCC® 33186™* | D | 2 | |



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|-----------------------|------------------|------------------|---|------------|-----|--|
| 0753L | 0753K | 0753P | Enterococcus faecalis derived from ATCC® 49532™* | E | 2 | High level Gentamicin-resistant and Streptomycin-sensitive |
| 0752L | 0752K | 0752P | Enterococcus faecalis derived from ATCC® 49533™* | C | 2 | High level Gentamicin-resistant and Streptomycin-sensitive |
| 0959L | 0959K | 0959P | Enterococcus faecalis derived from ATCC® 51299™* | C | 2 | vanB positive; Low level vancomycin resistant; resistant to high level aminoglycosides (gentamicin and streptomycin); Sensitive to teichoplanin; ant(6)-1 aac(6') aph(2''); control for antimicrobial susceptibility testing |
| 01089L | 01089K | 01089P | Enterococcus faecalis derived from ATCC® 51575™* | E | 2 | Resistant to gentamicin, streptomycin and vancomycin; vanB positive |
| 0497L | 0497K | 0497P | Enterococcus faecalis derived from ATCC® 7080™* | C | 2 | |
| 0336L | 0336K | 0336P | Enterococcus faecalis derived from NCIMB 13280 | C | 2 | |
| 0714L | 0714K | 0714P | Enterococcus faecalis derived from NCTC 13379 | E | 2 | |
| 0472L | 0472K | 0472P | Enterococcus faecalis derived from NCTC 775 | C | 2 | |
| 01052L | 01052K | 01052P | Enterococcus faecium derived from ATCC® 6057™* | E | 2 | |
| 0677L | 0677K | 0677P | Enterococcus faecium derived from ATCC® 6569™* | D | 2 | |
| 01260L | 01260K | 01260P | Enterococcus faecium derived from ATCC® 8459™* | D | 1 | Used as a surrogate microorganism in the pasteurization of nuts |
| 0679L | 0679K | 0679P | Enterococcus faecium derived from ATCC® 27270™* | D | 2 | |
| 0968L | 0968K | 0968P | Enterococcus faecium derived from ATCC® 35667™* | C | 2 | |
| 01000L | 01000K | 01000P | Enterococcus faecium derived from ATCC® 700221™* | C | 2 | vanA positive; IS16 and esp positive; resistant to vancomycin and teicoplanin |
| 01143L | 01143K | 01143P | Enterococcus faecium derived from NCTC 12204 | C | 2 | vanA-type glycopeptide resistance; vancomycin resistant |
| 0895L | 0895K | 0895P | Enterococcus gallinarum derived from ATCC® 49573™* | C | 1 | |
| 0678L | 0678K | 0678P | Enterococcus hirae derived from ATCC® 10541™* | D | 1 | |
| 0650L | 0650K | 0650P | Enterococcus hirae derived from ATCC® 8043™* | C | 1 | |
| 0857L | 0857K | 0857P | Enterococcus raffinosus derived from ATCC® 49464™* | C | 1 | |
| 0223L | 0223K | 0223P | Enterococcus saccharolyticus derived from ATCC® 43076™* | C | 1 | |
| Erysipelothrix | | | | | | |
| 0661L | 0661K | 0661P | Erysipelothrix rhusiopathiae derived from ATCC® 19414™* | C | 2 | |
| Escherichia | | | | | | |
| 0204L | 0204K | 0204P | Escherichia coli (JM101) derived from ATCC® 33876™* | C | 1 | |



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|-----------|------------------|------------------|--|------------|-----|---|
| 01101L | 01101K | 01101P | Escherichia coli (O103:H11) derived from CDC 06-3008 | F | 2 | Serotype O103:H11; eae positive; stx 1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 01104L | 01104K | 01104P | Escherichia coli (O104:H4) derived from ATCC® BAA-2326™* | F | 2 | Serotype O104:H4; aggR positive; stx 2 positive; export license required for shipping outside the U.S. |
| 01102L | 01102K | 01102P | Escherichia coli (O111:H8) derived from CDC 2010C-3114 | F | 2 | Serotype O111:H8; eae positive; stx 1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 01099L | 01099K | 01099P | Escherichia coli (O121:H19) derived from CDC 02-3211 | F | 2 | Serotype O121:H19; eae positive; stx1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 01097L | 01097K | 01097P | Escherichia coli (O145:NM) derived from CDC 99-3311 | F | 2 | Serotype O145:NM; eae positive; stx1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 0617L | 0617K | 0617P | Escherichia coli (O157:H7) derived from ATCC® 35150™* | D | 2 | Serotype O157:H7; export license required for shipping outside the U.S. |
| 0795L | 0795K | 0795P | Escherichia coli (O157:H7) derived from ATCC® 43888™* | C | 2 | Serotype O157:H7; does not produce Shiga-Like Toxin I or II; export license required for shipping outside the U.S. |
| 01204L | 01204K | 01204P | Escherichia coli (O157:H7) derived from ATCC® 43890™* | D | 2 | Serotype O157:H7; Produces Shiga-like toxin 1; export license required for shipping outside the U.S. |
| 01100L | 01100K | 01100P | Escherichia coli (O26:H11) derived from CDC 03-3014 | F | 2 | Serotype O26:H11; eae positive; stx 1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 01098L | 01098K | 01098P | Escherichia coli (O45:H2) derived from CDC 00-3039 | F | 2 | Serotype O45:H2; eae positive; stx 1 and/or stx 2 positive; export license required for shipping outside the U.S. |
| 0680L | 0680K | 0680P | Escherichia coli derived from ATCC® 10536™* | D | 1 | |
| 01050L | 01050K | 01050P | Escherichia coli derived from ATCC® 10799™* | E | 1 | |
| 0681L | 0681K | 0681P | Escherichia coli derived from ATCC® 11229™* | C | 1 | |
| 0502L | 0502K | 0502P | Escherichia coli derived from ATCC® 11303™* | D | 1 | |
| 0465L | 0465K | 0465P | Escherichia coli derived from ATCC® 11775™* | C | 1 | Serovar O1:K1:H7 |
| 0747L | 0747K | 0747P | Escherichia coli derived from ATCC® 13706™* | C | 1 | |



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|-----------|------------------|------------------|--|------------|-----|--|
| 0682L | 0682K | 0682P | Escherichia coli derived from ATCC® 13762™* | C | 1 | |
| 0543L | 0543K | 0543P | Escherichia coli derived from ATCC® 14169™* | C | 1 | |
| 01037L | 01037K | 01037P | Escherichia coli derived from ATCC® 15597™* | E | 1 | Host for MS2 Bacteriophage (ATCC® 15597™-B1) |
| 0203L | 0203K | 0203P | Escherichia coli derived from ATCC® 23848™* | C | 1 | Genotype B |
| 0335L | 0335K | 0335P | Escherichia coli derived from ATCC® 25922™* | A | 1 | CLSI and EUCAST control for antimicrobial susceptibility testing |
| 0495L | 0495K | 0495P | Escherichia coli derived from ATCC® 35218™* | C | 1 | TEM-1 beta-lactamase producer (non-ESBL); CLSI and EUCAST control for antimicrobial susceptibility testing |
| 0422L | 0422K | 0422P | Escherichia coli derived from ATCC® 35421™* | C | 1 | |
| 0433L | 0433K | 0433P | Escherichia coli derived from ATCC® 4157™* | C | 1 | |
| 0869L | 0869K | 0869P | Escherichia coli derived from ATCC® 51446™* | C | 2 | |
| 01233L | 01233K | 01233P | Escherichia coli derived from ATCC® 51739™* | C | 1 | |
| 0860L | 0860K | 0860P | Escherichia coli derived from ATCC® 51755™* | D | 1 | |
| 0791L | 0791K | 0791P | Escherichia coli derived from ATCC® 51813™* | D | 1 | |
| 01137L | 01137K | 01137P | Escherichia coli derived from ATCC® 700609™* | E | 1 | Designation CN13; Bacteriophage host |
| 0231L | 0231K | 0231P | Escherichia coli derived from ATCC® 700728™* | C | 1 | Serotype O157:H7; nontoxicogenic |
| 0483L | 0483K | 0483P | Escherichia coli derived from ATCC® 8739™* | C | 1 | |
| 0115L | 0115K | 0115P | Escherichia coli derived from ATCC® 9637™* | D | 1 | |
| 01261L | 01261K | 01261P | Escherichia coli derived from ATCC® BAA-1429™* | D | 1 | This is a non-pathogenic E. coli strain recommended by the USDA as a surrogate indicator organism to measure changes in microbial counts |
| 01242L | 01242K | 01242P | Escherichia coli derived from ATCC® BAA-2452™* | C | 2 | New Delhi metallo-beta-lactamase (NDM-1) positive; blaNDM positive by PCR; Carbapenem-resistant (Imipenem and Ertapenem) |
| 01113L | 01113K | 01113P | Escherichia coli derived from ATCC® BAA-2469™* | E | 2 | New Delhi metallo-beta-lactamase (NDM-1) positive; blaKPC negative by PCR; blaNDM positive by PCR; Carbapenem-resistant (Imipenem and Ertapenem) |
| 01229L | 01229K | 01229P | Escherichia coli ATCC® BAA-2523™* | C | 2 | Resistant to Ertapenem |
| 01259L | 01259K | 01259P | Escherichia coli derived from CDC AR-0346 | D | 2 | Extended-spectrum beta-lactamase (ESBL) positive; contains mcr-1 gene |



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|----------------------|------------------|------------------|---|------------|-----|---|
| 01062L | 01062K | 01062P | Escherichia coli derived from NCTC 8622 | C | 2 | Serovar O126:K7(B16):H2; export license required for shipping outside the U.S. |
| 01109L | 01109K | 01109P | Escherichia coli derived from NCTC 9001 | C | 2 | |
| 0706L | 0706K | 0706P | Escherichia coli derived from NCTC 10538 | B | 1 | K12 |
| 0861L | 0861K | 0861P | Escherichia coli derived from NCTC 12900 | C | 1 | Serotype O157:H7; non toxigenic |
| 01193L | 01193K | 01193P | Escherichia coli derived from NCTC 13167 | C | 2 | |
| 01192L | 01192K | 01192P | Escherichia coli derived from NCTC 13216 | C | 2 | |
| 01085L | 01085K | 01085P | Escherichia coli derived from NCTC 13351 | D | 1 | TEM-3 ESBL |
| 01136L | 01136K | 01136P | Escherichia coli derived from NCTC 13476 | D | 2 | IMP-type |
| 01265L | 01265K | 01265P | Escherichia coli derived from NCTC 13353 | C | 2 | Routine QC strain for Disk Diffusion and MIC Testing of Beta-Lactam Combination Agents per CLSI M100-28Ed; blaCTX-M-15 gene; susceptible to Cefepime-tazobactam combination |
| 01244L | 01244K | 01244P | Escherichia coli derived from NCTC 13846 | D | 2 | Colistin resistant; MCR-1 positive; harbors multiple other resistance genes (visit product page at microbiologics.com to view all genes) |
| Eurotium | | | | | | |
| 01160L | 01160K | 01160P | Eurotium rubrum derived from ATCC® 42690™* | E | 1 | Mold |
| Finegoldia | | | | | | |
| 0409L | 0409K | 0409P | Finegoldia magna derived from ATCC® 29328™* | D | 1 | |
| Fluoribacter | | | | | | |
| 0212L | 0212K | 0212P | Fluoribacter bozemanai derived from ATCC® 33217™* | E | 2 | |
| Fusarium | | | | | | |
| 0531L | 0531K | 0531P | Fusarium solani derived from ATCC® 36031™* | E | 2 | Mold |
| Fusobacterium | | | | | | |
| 01191L | 01191K | 01191P | Fusobacterium mortiferum derived from ATCC® 25557™* | D | 1 | |
| 0407L | 0407K | 0407P | Fusobacterium necrophorum subsp. necrophorum derived from ATCC® 25286™* | D | 2 | |
| 0328L | 0328K | 0328P | Fusobacterium nucleatum subsp. nucleatum derived from ATCC® 25586™* | D | 2 | |
| Gardnerella | | | | | | |
| 0410L | 0410K | 0410P | Gardnerella vaginalis derived from ATCC® 14018™* | C | 2 | |



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|----------------------|------------------|------------------|---|------------|-----|---|
| 0643L | 0643K | 0643P | Gardnerella vaginalis derived from ATCC® 49145™* | C | 2 | |
| Geobacillus | | | | | | |
| 0172L | 0172K | 0172P | Geobacillus stearothermophilus derived from ATCC® 10149™* | C | 1 | |
| 0872L | 0872K | 0872P | Geobacillus stearothermophilus derived from ATCC® 12980™* | B | 1 | |
| 0871L | 0871K | 0871P | Geobacillus stearothermophilus derived from ATCC® 7953™* | D | 1 | |
| Geotrichum | | | | | | |
| 0519L | 0519K | 0519P | Geotrichum candidum derived from ATCC® 34614™* | D | 1 | Yeast cells |
| Haemophilus | | | | | | |
| 0441L | 0441K | 0441P | Haemophilus influenzae derived from ATCC® 10211™* | C | 2 | Type b; beta lactamase negative |
| 0376L | 0376K | 0376P | Haemophilus influenzae derived from ATCC® 19418™* | C | 2 | |
| 0338L | 0338K | 0338P | Haemophilus influenzae derived from ATCC® 33533™* | C | 2 | Type b; beta lactamase producer |
| 0993L | 0993K | 0993P | Haemophilus influenzae derived from ATCC® 35056™* | D | 2 | beta lactamase producer |
| 0438L | 0438K | 0438P | Haemophilus influenzae derived from ATCC® 43065™* | C | 2 | |
| 0476L | 0476K | 0476P | Haemophilus influenzae derived from ATCC® 43163™* | C | 2 | Beta lactamase producer |
| 0644L | 0644K | 0644P | Haemophilus influenzae derived from ATCC® 49144™* | C | 2 | Beta lactamase producer |
| 0647L | 0647K | 0647P | Haemophilus influenzae derived from ATCC® 49247™* | C | 2 | Reduced susceptibility to beta-lactam agents due to PBP mutations in Haemophilus influenzae; Ampicillin resistant; CLSI and EUCAST control for antimicrobial susceptibility testing |
| 0919L | 0919K | 0919P | Haemophilus influenzae derived from ATCC® 49766™* | C | 2 | CLSI and EUCAST antimicrobial susceptibility control strain |
| 0620L | 0620K | 0620P | Haemophilus influenzae derived from ATCC® 9006™* | C | 2 | Type A |
| 0185L | 0185K | 0185P | Haemophilus influenzae derived from ATCC® 9007™* | D | 2 | Type C |
| 01024L | 01024K | 01024P | Haemophilus influenzae derived from NCTC 8468 | C | 2 | |
| 0377L | 0377K | 0377P | Haemophilus parahaemolyticus derived from ATCC® 10014™* | C | 2 | |
| 01171L | 01171K | 01171P | Haemophilus parainfluenzae derived from ATCC® 9796™* | C | 2 | |
| Hafnia | | | | | | |
| 0165L | 0165K | 0165P | Hafnia alvei derived from ATCC® 51815™* | E | 1 | |
| Hanseniaspora | | | | | | |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|----------------------|------------------|------------------|---|------------|-----|---|
| 01012L | 01012K | 01012P | Hanseniaspora valbyensis derived from ATCC® 58370™* | C | 1 | Formerly Kloeckera japonica; yeast cells |
| Herminiimonas | | | | | | |
| 0169L | 0169K | 0169P | Herminiimonas species derived from ATCC® 49643™* | E | 1 | Formerly Aquaspirillum species |
| Issatchenkia | | | | | | |
| 0227L | 0227K | 0227P | Issatchenkia orientalis derived from ATCC® 6258™* | D | 2 | CLSI QC strain for Disk Diffusion and MIC determination; yeast cells |
| Klebsiella | | | | | | |
| 0306L | 0306K | 0306P | Klebsiella aerogenes derived from ATCC® 13048™* | B | 1 | Quality control for DuPont RiboPrinter® Microbial Characterization System; Formerly Enterobacter aerogenes |
| 0399L | 0399K | 0399P | Klebsiella aerogenes derived from ATCC® 35029™* | B | 1 | Formerly Enterobacter aerogenes |
| 0196L | 0196K | 0196P | Klebsiella aerogenes derived from ATCC® 51697™* | D | 1 | Formerly Enterobacter aerogenes |
| 0597L | 0597K | 0597P | Klebsiella aerogenes derived from NCIMB 10102 | B | 2 | Formerly Enterobacter aerogenes |
| 0530L | 0530K | 0530P | Klebsiella oxytoca derived from ATCC® 13182™* | B | 2 | |
| 0626L | 0626K | 0626P | Klebsiella oxytoca derived from ATCC® 49131™* | B | 2 | |
| 0167L | 0167K | 0167P | Klebsiella oxytoca derived from ATCC® 51817™* | E | 2 | |
| 0757L | 0757K | 0757P | Klebsiella oxytoca derived from ATCC® 700324™* | D | 2 | |
| 0840L | 0840K | 0840P | Klebsiella oxytoca derived from ATCC® 8724™* | B | 2 | |
| 01147L | 01147K | 01147P | Klebsiella oxytoca derived from NCTC 11686 | D | 2 | |
| 01019L | 01019K | 01019P | Klebsiella pneumoniae derived from ATCC® BAA-1144™* | D | 2 | Control strain for the AmpC disk test; weak positive |
| 01005L | 01005K | 01005P | Klebsiella pneumoniae derived from ATCC® BAA-1705™* | B | 2 | blaKPC positive; blaNDM negative; Produces carbapenemase KPC; Carbapenem-resistant (imipenem and ertapenem); Modified Hodge Test (MHT) positive control |
| 01006L | 01006K | 01006P | Klebsiella pneumoniae derived from ATCC® BAA-1706™* | B | 2 | Modified Hodge Test (MHT) negative control |
| 01060L | 01060K | 01060P | Klebsiella pneumoniae derived from ATCC® BAA-2146™* | E | 2 | New Delhi metallo-beta-lactamase (NDM-1) positive; blaNDM positive; blaKPC negative |
| 01263L | 01263K | 01263P | Klebsiella pneumoniae derived from ATCC® BAA-2814™* | D | 2 | KPC-3 |
| 01153L | 01153K | 01153P | Klebsiella pneumoniae derived from CDC 1100192 | D | 2 | blaNDM-1 metallo-beta-lactamase gene |
| | 01241K | 01241P | Klebsiella pneumoniae derived from CDC AR-0039 | C | 2 | Produces OXA-181 carbapenemase |



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|----------------------|------------------|------------------|---|------------|-----|--|
| | 01240K | 01240P | Klebsiella pneumoniae derived from CDC AR-0066 | C | 2 | Produces OXA-232 carbapenemase |
| 01117L | 01117K | 01117P | Klebsiella pneumoniae derived from NCTC 13438 | D | 2 | Produces carbapenemase KPC-3 |
| | | 01245P | Klebsiella pneumoniae derived from NCTC 13439 | C | 2 | blaVIM-1 gene; Metallo-beta-lactamase positive |
| 01112L | 01112K | 01112P | Klebsiella pneumoniae derived from NCTC 13440 | D | 2 | Metallo-beta-lactamase positive; VIM-1 |
| 01148L | 01148K | 01148P | Klebsiella pneumoniae derived from NCTC 13442 | D | 2 | OXA 48 |
| 01145L | 01145K | 01145P | Klebsiella pneumoniae derived from NCTC 13443 | D | 2 | NDM-1 positive; New Delhi metallo-beta-lactamase |
| 0684L | 0684K | 0684P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 10031™* | C | 2 | Methyl Red positive, Voges-Proskauer negative |
| 0458L | 0458K | 0458P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 13882™* | B | 2 | |
| 0351L | 0351K | 0351P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 13883™* | A | 2 | Methyl Red positive; Vog; Not thermotolerant: does not grow at 44.5°C |
| 0685L | 0685K | 0685P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 27736™* | B | 2 | |
| 0957L | 0957K | 0957P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 33495™* | C | 2 | |
| 0942L | 0942K | 0942P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 35657™* | D | 2 | |
| 0683L | 0683K | 0683P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 4352™* | B | 2 | |
| 0784L | 0784K | 0784P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® 700603™* | D | 2 | Produces SHV-18; Extended-spectrum beta-lactamase (ESBL) positive; CLSI and EUCAST positive control for ESBL tests |
| 01247L | 01247K | 01247P | Klebsiella pneumoniae subsp. pneumoniae derived from ATCC® BAA-2524™* | C | 2 | QC strain for Biomerieux Chrom-ID OXA 48 agar (bMX1103199) |
| 0261L | 0261K | 0261P | Klebsiella variicola derived from ATCC® 31488™* | D | 2 | Formerly Klebsiella pneumoniae subsp. pneumoniae; thermotolerant |
| Kocuria | | | | | | |
| 0126L | 0126K | 0126P | Kocuria kristinae derived from ATCC® BAA-752™* | C | 1 | |
| 0670L | 0670K | 0670P | Kocuria rhizophila derived from ATCC® 15957™* | E | 1 | |
| 0669L | 0669K | 0669P | Kocuria rhizophila derived from ATCC® 9341a™* | C | 1 | |
| 0688L | 0688K | 0688P | Kocuria rhizophila derived from ATCC® 9341™* | D | 1 | |
| 0766L | 0766K | 0766P | Kocuria rosea derived from ATCC® 186™* | D | 1 | |
| Lactobacillus | | | | | | |
| 0885L | 0885K | 0885P | Lactobacillus acidophilus derived from ATCC® 314™* | C | 1 | |
| 0243L | 0243K | 0243P | Lactobacillus acidophilus derived from ATCC® 4356™* | D | 1 | |



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|--------------------|------------------|------------------|--|------------|-----|--|
| 01262L | 01262K | 01262P | Lactobacillus brevis derived from ATCC® 14869™* | E | 1 | |
| 0546L | 0546K | 0546P | Lactobacillus casei derived from ATCC® 334™* | E | 1 | |
| 0176L | 0176K | 0176P | Lactobacillus casei derived from ATCC® 393™* | D | 1 | |
| 0813L | 0813K | 0813P | Lactobacillus fermentum derived from ATCC® 9338™* | C | 1 | |
| 0127L | 0127K | 0127P | Lactobacillus gasseri derived from ATCC® 19992™* | C | 1 | |
| 0235L | 0235K | 0235P | Lactobacillus leichmannii derived from ATCC® 7830™* | C | 1 | Formerly Lactobacillus delbrueckii subsp. lactis |
| 0526L | 0526K | 0526P | Lactobacillus paracasei subsp. paracasei derived from ATCC® BAA-52™* | D | 1 | |
| 0234L | 0234K | 0234P | Lactobacillus plantarum derived from ATCC® 8014™* | D | 1 | |
| 01144L | 01144K | 01144P | Lactobacillus plantarum subsp. plantarum derived from ATCC® 14917™* | C | 1 | |
| 01090L | 01090K | 01090P | Lactobacillus rhamnosus derived from ATCC® 53103™* | E | 1 | |
| 0233L | 0233K | 0233P | Lactobacillus rhamnosus derived from ATCC® 7469™* | D | 1 | |
| 0989L | 0989K | 0989P | Lactobacillus rhamnosus derived from ATCC® 9595™* | C | 1 | |
| 0128L | 0128K | 0128P | Lactobacillus sakei subsp. sakei derived from ATCC® 15521™* | D | 1 | |
| Lactococcus | | | | | | |
| 0980L | 0980K | 0980P | Lactococcus lactis derived from ATCC® 49032™* | D | 1 | |
| 0152L | 0152K | 0152P | Lactococcus lactis subsp. cremoris derived from ATCC® 19257™* | D | 1 | |
| 0205L | 0205K | 0205P | Lactococcus lactis subsp. lactis derived from ATCC® 11454™* | D | 1 | |
| 0149L | 0149K | 0149P | Lactococcus lactis subsp. lactis derived from ATCC® 19435™* | D | 1 | |
| Legionella | | | | | | |
| 01195L | 01195K | 01195P | Legionella anisa derived from ATCC® 35292™* | E | 2 | |
| 01003L | 01003K | 01003P | Legionella longbeachae derived from ATCC® 33462™* | E | 2 | |
| 0211L | 0211K | 0211P | Legionella pneumophila subsp. pneumophila derived from ATCC® 33152™* | E | 2 | |
| Listeria | | | | | | |
| 0856L | 0856K | 0856P | Listeria grayi derived from ATCC® 25401™* | D | 1 | |
| 0222L | 0222K | 0222P | Listeria grayi derived from ATCC® 700545™* | D | 1 | |
| 0814L | 0814K | 0814P | Listeria innocua derived from ATCC® 33090™* | D | 1 | Serotype 6a |
| 01243L | 01243K | 01243P | Listeria innocua derived from ATCC® 51742™* | D | 1 | |
| 0414L | 0414K | 0414P | Listeria innocua derived from NCTC 11288 | D | 1 | Serotype 6a |
| 0815L | 0815K | 0815P | Listeria ivanovii subsp. ivanovii derived from ATCC® 19119™* | D | 2 | |
| 0299L | 0299K | 0299P | Listeria ivanovii subsp. londoniensis derived from ATCC® BAA-139™* | E | 1 | Formerly Listeria ivanovii subsp. ivanovii |



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|-------------------------|------------------|------------------|--|------------|-----|-----------------------------------|
| 0254L | 0254K | 0254P | Listeria monocytogenes Cornell University derived from Silliker® SLR2249 | D | 2 | Act A gene removed |
| 0129L | 0129K | 0129P | Listeria monocytogenes derived from ATCC® 13932™* | C | 2 | Serotype 4b |
| 0727L | 0727K | 0727P | Listeria monocytogenes derived from ATCC® 15313™* | D | 2 | Non-hemolytic on sheep blood agar |
| 0277L | 0277K | 0277P | Listeria monocytogenes derived from ATCC® 19111™* | D | 2 | Serotype 1 |
| 0232L | 0232K | 0232P | Listeria monocytogenes derived from ATCC® 19112™* | D | 2 | Serotype 2 |
| 0686L | 0686K | 0686P | Listeria monocytogenes derived from ATCC® 19114™* | D | 2 | Serotype 4a |
| 0687L | 0687K | 0687P | Listeria monocytogenes derived from ATCC® 19115™* | D | 2 | Serotype 4b |
| 0154L | 0154K | 0154P | Listeria monocytogenes derived from ATCC® 19118™* | D | 2 | Serotype 4e |
| 01190L | 01190K | 01190P | Listeria monocytogenes derived from ATCC® 35152™* | C | 2 | Non-hemolytic variation |
| 0398L | 0398K | 0398P | Listeria monocytogenes derived from ATCC® 7644™* | D | 2 | |
| 0130L | 0130K | 0130P | Listeria monocytogenes derived from ATCC® BAA-751™* | C | 2 | |
| 0783L | 0783K | 0783P | Listeria monocytogenes derived from NCTC 10890 | D | 2 | Serotype 7 |
| 0802L | 0802K | 0802P | Listeria seeligeri derived from ATCC® 35967™* | D | 1 | |
| 0816L | 0816K | 0816P | Listeria welshimeri derived from ATCC® 35897™* | D | 1 | |
| Malassezia | | | | | | |
| 0701L | 0701K | 0701P | Malassezia furfur derived from ATCC® 14521™* | D | 2 | Yeast cells |
| Mannheimia | | | | | | |
| 0664L | 0664K | 0664P | Mannheimia haemolytica derived from ATCC® 33396™* | E | 2 | |
| Methylobacterium | | | | | | |
| 01110L | 01110K | 01110P | Methylobacterium extorquens derived from ATCC® BAA-2500™* | E | 1 | |
| 01181L | 01181K | 01181P | Methylobacterium organophilum derived from ATCC® 27886™* | E | 1 | |
| Meyerozyma | | | | | | |
| 0738L | 0738K | 0738P | Meyerozyma guilliermondii derived from ATCC® 6260™* | D | 1 | Yeast cells |
| Microbacterium | | | | | | |
| 01042L | 01042K | 01042P | Microbacterium liquefaciens derived from ATCC® BAA-1819™* | D | 1 | |
| 01041L | 01041K | 01041P | Microbacterium paraoxydans derived from ATCC® BAA-1818™* | D | 1 | |
| 0295L | 0295K | 0295P | Microbacterium testaceum derived from ATCC® 15829™* | D | 1 | |
| Micrococcus | | | | | | |
| 0689L | 0689K | 0689P | Micrococcus luteus derived from ATCC® 10240™* | C | 1 | |
| 0242L | 0242K | 0242P | Micrococcus luteus derived from ATCC® 4698™* | D | 1 | |
| 0804L | 0804K | 0804P | Micrococcus luteus derived from ATCC® 49732™* | C | 1 | |
| 0740L | 0740K | 0740P | Micrococcus species derived from ATCC® 700405™* | D | 1 | |



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|----------------------|------------------|------------------|--|------------|-----|--|
| 0218L | 0218K | 0218P | Micrococcus yunnanensis derived from ATCC® 7468™* | C | 1 | Formerly Micrococcus luteus |
| Microsporium | | | | | | |
| 0894L | 0894K | 0894P | Microsporium canis derived from ATCC® 36299™* | E | 2 | Mold |
| 0893L | 0893K | 0893P | Microsporium gypseum derived from ATCC® 24102™* | E | 2 | Mold |
| Moraxella | | | | | | |
| 0248L | 0248K | 0248P | Moraxella catarrhalis derived from ATCC® 23246™* | D | 1 | Formerly Moraxella (Branhamella) catarrhalis |
| 0951L | 0951K | 0951P | Moraxella catarrhalis derived from ATCC® 25238™* | C | 1 | |
| 0403L | 0403K | 0403P | Moraxella catarrhalis derived from ATCC® 25240™* | C | 1 | Formerly Moraxella (Branhamella) catarrhalis |
| 0642L | 0642K | 0642P | Moraxella catarrhalis derived from ATCC® 49143™* | C | 1 | Formerly Moraxella (Branhamella) catarrhalis |
| 0622L | 0622K | 0622P | Moraxella catarrhalis derived from ATCC® 8176™* | C | 1 | Formerly Moraxella (Branhamella) catarrhalis |
| 0475L | 0475K | 0475P | Moraxella osloensis derived from ATCC® 10973™* | C | 2 | Formerly Moraxella (Moraxella) osloensis |
| Morganella | | | | | | |
| 0215L | 0215K | 0215P | Morganella morganii subsp. morganii derived from ATCC® 25829™* | C | 2 | |
| 0839L | 0839K | 0839P | Morganella morganii subsp. morganii derived from ATCC® 25830™* | D | 2 | |
| Mucor | | | | | | |
| 01142L | 01142K | 01142P | Mucor racemosus derived from ATCC® 42647™* | D | 1 | Mold |
| 01255L | 01255K | 01255P | Mucor racemosus f. racemosus derived from ATCC® 22365™* | D | 1 | Mold |
| Mycobacterium | | | | | | |
| 01149L | 01149K | 01149P | Mycobacterium avium derived from ATCC® 700898™* | E | 2 | |
| 0522L | 0522K | 0522P | Mycobacterium avium subsp. avium derived from ATCC® 15769™* | E | 2 | Serotype 1 |
| 0544L | 0544K | 0544P | Mycobacterium avium subsp. avium derived from ATCC® 25291™* | E | 2 | Serotype 2 |
| 0513L | 0513K | 0513P | Mycobacterium fortuitum subsp. fortuitum derived from ATCC® 6841™* | E | 2 | |
| 0995L | 0995K | 0995P | Mycobacterium gordonae derived from ATCC® 14470™* | E | 2 | |
| 01049L | 01049K | 01049P | Mycobacterium haemophilum derived from ATCC® 29548™* | E | 2 | Requires hemin (x factor) for growth |
| 0157L | 0157K | 0157P | Mycobacterium intracellulare derived from ATCC® 13950™* | E | 2 | |
| 0545L | 0545K | 0545P | Mycobacterium kansasii derived from ATCC® 12478™* | E | 2 | |
| 0721L | 0721K | 0721P | Mycobacterium smegmatis derived from ATCC® 14468™* | E | 1 | |
| 0514L | 0514K | 0514P | Mycobacterium smegmatis derived from ATCC® 19420™* | E | 1 | |



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|-------------------|------------------|------------------|---|------------|-----|--|
| 0114L | 0114K | 0114P | Mycobacterium smegmatis derived from ATCC® 607™* | E | 1 | |
| 0272L | 0272K | 0272P | Mycobacterium terrae derived from ATCC® 15755™* | E | 1 | |
| 0112L | 0112K | 0112P | Mycobacterium tuberculosis derived from ATCC® 25177™* | E | 2 | Attenuated |
| Mycoplasma | | | | | | |
| 01053L | 01053K | 01053P | Mycoplasma bovis derived from ATCC® 25025™* | E | 2 | |
| 0156L | 0156K | 0156P | Mycoplasma hominis derived from ATCC® 15488™* | E | 2 | |
| 0503L | 0503K | 0503P | Mycoplasma pneumoniae derived from ATCC® 15531™* | E | 2 | |
| Myroides | | | | | | |
| 0324L | 0324K | 0324P | Myroides odoratus derived from ATCC® 4651™* | D | 1 | |
| Neisseria | | | | | | |
| 0378L | 0378K | 0378P | Neisseria gonorrhoeae derived from ATCC® 19424™* | C | 2 | |
| 0375L | 0375K | 0375P | Neisseria gonorrhoeae derived from ATCC® 31426™* | C | 2 | Beta lactamase producer |
| 0429L | 0429K | 0429P | Neisseria gonorrhoeae derived from ATCC® 43069™* | C | 2 | |
| 0426L | 0426K | 0426P | Neisseria gonorrhoeae derived from ATCC® 43070™* | C | 2 | |
| 0648L | 0648K | 0648P | Neisseria gonorrhoeae derived from ATCC® 49226™* | C | 2 | Chromosomally mediated penicillin-resistant Neisseria gonorrhoeae; CLSI control for antimicrobial susceptibility testing |
| 0552L | 0552K | 0552P | Neisseria gonorrhoeae derived from ATCC® 49498™* | C | 2 | |
| 0952L | 0952K | 0952P | Neisseria gonorrhoeae derived from ATCC® 49981™* | D | 2 | Penicillin resistant Neisseria |
| 01239L | 01239K | 01239P | Neisseria gonorrhoeae derived from CDC 10328 | C | 2 | According to the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing," this strain is intermediate to ciprofloxacin (CipI) |
| 01222L | 01222K | 01222P | Neisseria gonorrhoeae derived from CDC 10329 | C | 2 | According to the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing," this isolate is resistant to ciprofloxacin (CipR) |
| 01216L | 01216K | 01216P | Neisseria gonorrhoeae derived from CDC F-18 | C | 2 | This strain is susceptible to relevant antimicrobial agents per the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing" |



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|----------------------|------------------|------------------|--|------------|-----|---|
| 01217L | 01217K | 01217P | Neisseria gonorrhoeae derived from CDC F-28 | C | 2 | According to the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing," this strain is resistant to spectinomycin (SpCR) |
| 01219L | 01219K | 01219P | Neisseria gonorrhoeae derived from CDC SPJ-15 | C | 2 | According to the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing," this is an isolate with a critical MIC greater than or equal to 1.0 ug/ml of azithromycin (AznC) |
| 01218L | 01218K | 01218P | Neisseria gonorrhoeae derived from CDC SPL-4 | C | 2 | According to the CDC document, "Neisseria gonorrhoeae Reference Strains for Susceptibility Testing," this strain exhibits decreased sensitivity to cefixime (CfxDS) |
| 0405L | 0405K | 0405P | Neisseria lactamica derived from ATCC® 23970™* | C | 2 | |
| 0943L | 0943K | 0943P | Neisseria lactamica derived from ATCC® 23971™* | C | 2 | |
| 0646L | 0646K | 0646P | Neisseria lactamica derived from ATCC® 49142™* | C | 2 | |
| 0453L | 0453K | 0453P | Neisseria meningitidis derived from ATCC® 13077™* | C | 2 | Serogroup A |
| 0454L | 0454K | 0454P | Neisseria meningitidis derived from ATCC® 13090™* | C | 2 | Serogroup B |
| 0404L | 0404K | 0404P | Neisseria meningitidis derived from ATCC® 13102™* | C | 2 | Serogroup C |
| 0649L | 0649K | 0649P | Neisseria meningitidis derived from NCTC 10026 | E | 2 | Serogroup B |
| 0131L | 0131K | 0131P | Neisseria mucosa derived from ATCC® 49233™* | D | 2 | |
| 0464L | 0464K | 0464P | Neisseria sicca derived from ATCC® 29193™* | D | 1 | |
| 0406L | 0406K | 0406P | Neisseria sicca derived from ATCC® 9913™* | C | 1 | |
| Nocardia | | | | | | |
| 0866L | 0866K | 0866P | Nocardia brasiliensis derived from ATCC® 19296™* | D | 2 | |
| Ochrobactrum | | | | | | |
| 0132L | 0132K | 0132P | Ochrobactrum anthropi derived from ATCC® BAA-749™* | C | 1 | |
| Oligella | | | | | | |
| 0868L | 0868K | 0868P | Oligella ureolytica derived from ATCC® 43534™* | C | 1 | |
| 0621L | 0621K | 0621P | Oligella urethralis derived from ATCC® 17960™* | C | 2 | |
| Paenibacillus | | | | | | |
| 0473L | 0473K | 0473P | Paenibacillus gordonae derived from ATCC® 29948™* | C | 1 | |
| 0431L | 0431K | 0431P | Paenibacillus larvae subsp. larvae derived from ATCC® 9545™* | E | 1 | |
| 0142L | 0142K | 0142P | Paenibacillus macerans derived from ATCC® 8509™* | C | 1 | |
| 0228L | 0228K | 0228P | Paenibacillus polymyxa derived from ATCC® 43865™* | B | 1 | |



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|---------------------------|------------------|------------------|---|------------|-----|---|
| 0296L | 0296K | 0296P | Paenibacillus polymyxa derived from ATCC® 7070™* | D | 1 | |
| 0883L | 0883K | 0883P | Paenibacillus polymyxa derived from ATCC® 842™* | D | 1 | |
| Parabacteroides | | | | | | |
| 0618L | 0618K | 0618P | Parabacteroides distasonis derived from ATCC® 8503™* | D | 2 | |
| 0584L | 0584K | 0584P | Parabacteroides distasonis derived from ATCC® BAA-1295™* | D | 2 | |
| Parvimonas | | | | | | |
| 0958L | 0958K | 0958P | Parvimonas micra derived from ATCC® 33270™* | D | 1 | |
| Pasteurella | | | | | | |
| 0668L | 0668K | 0668P | Pasteurella multocida subsp. multocida derived from ATCC® 12945™* | C | 2 | |
| Pediococcus | | | | | | |
| 0259L | 0259K | 0259P | Pediococcus acidilactici derived from ATCC® 8042™* | C | 1 | |
| 01075L | 01075K | 01075P | Pediococcus pentosaceus derived from ATCC® 33316™* | E | 1 | |
| Penicillium | | | | | | |
| 0178L | 0178K | 0178P | Penicillium chrysogenum derived from ATCC® 10106™* | E | 1 | Mold |
| 01133L | 01133K | 01133P | Penicillium citrinum derived from ATCC® 9849™* | E | 1 | Mold |
| 01252L | 01252K | 01252P | Penicillium corylophilum derived from ATCC® 9784™* | E | 1 | Mold |
| 01200L | 01200K | 01200P | Penicillium rubens derived from ATCC® 11709™* | E | 1 | Mold; produces penicillin; formerly Penicillium chrysogenum |
| 0207L | 0207K | 0207P | Penicillium rubens derived from ATCC® 9179™* | D | 1 | Mold |
| 0794L | 0794K | 0794P | Penicillium venetum derived from ATCC® 16025™* | E | 1 | Mold |
| Peptostreptococcus | | | | | | |
| 0322L | 0322K | 0322P | Peptostreptococcus anaerobius derived from ATCC® 27337™* | D | 1 | |
| Pluralibacter | | | | | | |
| 0703L | 0703K | 0703P | Pluralibacter gergoviae derived from ATCC® 33028™* | D | 2 | Formerly Enterobacter gergoviae |
| Porphyromonas | | | | | | |
| 0912L | 0912K | 0912P | Porphyromonas gingivalis derived from ATCC® 33277™* | D | 2 | |
| Prevotella | | | | | | |
| 01187L | 01187K | 01187P | Prevotella intermedia derived from ATCC® 25611™* | E | 2 | |
| 0110L | 0110K | 0110P | Prevotella melaninogenica derived from ATCC® 25845™* | D | 2 | |
| Proteus | | | | | | |
| 0355L | 0355K | 0355P | Proteus hauseri derived from ATCC® 13315™* | B | 2 | |
| 0440L | 0440K | 0440P | Proteus mirabilis derived from ATCC® 12453™* | B | 2 | |
| 01116L | 01116K | 01116P | Proteus mirabilis derived from ATCC® 21100™* | D | 2 | |
| 0690L | 0690K | 0690P | Proteus mirabilis derived from ATCC® 25933™* | D | 2 | |



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|--------------------|------------------|------------------|---|------------|-----|------------------------|
| 0321L | 0321K | 0321P | Proteus mirabilis derived from ATCC® 29245™* | B | 2 | |
| 01215L | 01215K | 01215P | Proteus mirabilis derived from ATCC® 29906™* | C | 2 | |
| 0190L | 0190K | 0190P | Proteus mirabilis derived from ATCC® 33583™* | D | 2 | |
| 0944L | 0944K | 0944P | Proteus mirabilis derived from ATCC® 35659™* | D | 2 | |
| 0432L | 0432K | 0432P | Proteus mirabilis derived from ATCC® 43071™* | B | 2 | |
| 0607L | 0607K | 0607P | Proteus mirabilis derived from ATCC® 7002™* | B | 2 | |
| 0310L | 0310K | 0310P | Proteus mirabilis derived from NCIMB 13283 | D | 2 | |
| 01152L | 01152K | 01152P | Proteus vulgaris derived from ATCC® 29905™* | D | 2 | |
| 0300L | 0300K | 0300P | Proteus vulgaris derived from ATCC® 33420™* | B | 2 | |
| 0640L | 0640K | 0640P | Proteus vulgaris derived from ATCC® 49132™* | B | 2 | |
| 0459L | 0459K | 0459P | Proteus vulgaris derived from ATCC® 6380™* | B | 2 | |
| 0841L | 0841K | 0841P | Proteus vulgaris derived from ATCC® 6896™* | B | 2 | |
| 0691L | 0691K | 0691P | Proteus vulgaris derived from ATCC® 8427™* | B | 2 | |
| 0393L | 0393K | 0393P | Proteus vulgaris derived from NCTC 4636 | C | 2 | |
| Prototheca | | | | | | |
| 0780L | 0780K | 0780P | Prototheca wickerhamii derived from ATCC® 16529™* | D | 1 | Yeast cells |
| Providencia | | | | | | |
| 0692L | 0692K | 0692P | Providencia alcalifaciens derived from ATCC® 9886™* | C | 1 | |
| 0997L | 0997K | 0997P | Providencia rettgeri derived from ATCC® 9250™* | D | 2 | |
| 0384L | 0384K | 0384P | Providencia stuartii derived from ATCC® 33672™* | C | 1 | |
| 0879L | 0879K | 0879P | Providencia stuartii derived from ATCC® 49809™* | C | 1 | |
| Pseudomonas | | | | | | |
| 0416L | 0416K | 0416P | Pseudomonas aeruginosa derived from ATCC® 10145™** | B | 2 | |
| 01078L | 01078K | 01078P | Pseudomonas aeruginosa derived from ATCC® 13388™* | E | 2 | |
| 0693L | 0693K | 0693P | Pseudomonas aeruginosa derived from ATCC® 15442™* | C | 2 | Pyocyanin not produced |
| 0199L | 0199K | 0199P | Pseudomonas aeruginosa derived from ATCC® 19429™* | D | 2 | |
| 0695L | 0695K | 0695P | Pseudomonas aeruginosa derived from ATCC® 25619™* | C | 2 | |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|-------------------|------------------|------------------|---|------------|-----|---|
| 0353L | 0353K | 0353P | <i>Pseudomonas aeruginosa</i> derived from ATCC® 27853™* | A | 2 | Contains inducible AmpC beta-lactamase; CLSI and EUCAST control for antimicrobial susceptibility testing; Microbiologics does not recommend the use of this strain for testing Ceftriaxone using the disk diffusion method. We have found that this strain is displaying resistance to Ceftriaxone using this method - this strain may not give consistent test results for the Ceftriaxone MIC well. |
| 0975L | 0975K | 0975P | <i>Pseudomonas aeruginosa</i> derived from ATCC® 35032™* | C | 2 | |
| 0182L | 0182K | 0182P | <i>Pseudomonas aeruginosa</i> derived from ATCC® 35554™* | D | 2 | |
| 0484L | 0484K | 0484P | <i>Pseudomonas aeruginosa</i> derived from ATCC® 9027™** | B | 2 | |
| 01009L | 01009K | 01009P | <i>Pseudomonas aeruginosa</i> derived from ATCC® 9721™* | B | 2 | |
| 01010L | 01010K | 01010P | <i>Pseudomonas aeruginosa</i> derived from ATCC® BAA-1744™* | B | 2 | |
| 0598L | 0598K | 0598P | <i>Pseudomonas aeruginosa</i> derived from NCIMB 12469 | C | 2 | Microbiologics does not recommend the use of this strain for testing Ceftriaxone using the disk diffusion method. We have found that this strain is displaying resistance to Ceftriaxone using this method. |
| 0304L | 0304K | 0304P | <i>Pseudomonas aeruginosa</i> derived from NCIMB 8295 | C | 2 | |
| 0576L | 0576K | 0576P | <i>Pseudomonas aeruginosa</i> derived from NCIMB 8626 | C | 2 | |
| 0830L | 0830K | 0830P | <i>Pseudomonas aeruginosa</i> derived from NCTC 10662 | D | 2 | |
| 0168L | 0168K | 0168P | <i>Pseudomonas brenneri</i> derived from ATCC® 49642™* | E | 1 | Formerly <i>Pseudomonas fluorescens</i> |
| 0241L | 0241K | 0241P | <i>Pseudomonas fluorescens</i> derived from ATCC® 13525™* | D | 1 | |
| 0880L | 0880K | 0880P | <i>Pseudomonas mosselii</i> derived from ATCC® 49838™* | C | 1 | Formerly <i>Pseudomonas fluorescens</i> |
| 0524L | 0524K | 0524P | <i>Pseudomonas protegens</i> (G) derived from ATCC® 17386™* | D | 1 | |
| 0702L | 0702K | 0702P | <i>Pseudomonas putida</i> derived from ATCC® 31483™* | E | 1 | |
| 0627L | 0627K | 0627P | <i>Pseudomonas putida</i> derived from ATCC® 49128™* | C | 1 | |
| 0853L | 0853K | 0853P | <i>Pseudomonas stutzeri</i> derived from ATCC® 17588™* | D | 1 | |
| Ralstonia | | | | | | |
| 0641L | 0641K | 0641P | <i>Ralstonia insidiosa</i> derived from ATCC® 49129™* | C | 1 | Formerly <i>Ralstonia pickettii</i> |
| 01159L | 01159K | 01159P | <i>Ralstonia pickettii</i> derived from ATCC® 27511™* | D | 1 | |
| 01197L | 01197K | 01197P | <i>Ralstonia pickettii</i> derived from ATCC® 700591™* | D | 1 | |
| Raoultella | | | | | | |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|----------------------|------------------|------------------|--|------------|-----|---|
| 0191L | 0191K | 0191P | Raoultella terrigena derived from ATCC® 33257™* | E | 2 | |
| Rhizopus | | | | | | |
| 0208L | 0208K | 0208P | Rhizopus stolonifer (-) derived from ATCC® 6227a™* | E | 1 | Mold |
| 0209L | 0209K | 0209P | Rhizopus stolonifer (+) derived from ATCC® 6227b™* | E | 1 | Mold |
| Rhodococcus | | | | | | |
| 0697L | 0697K | 0697P | Rhodococcus equi derived from ATCC® 6939™* | D | 2 | |
| Rhodotorula | | | | | | |
| 0143L | 0143K | 0143P | Rhodotorula mucilaginosa derived from ATCC® 66034™* | D | 2 | Yeast cells |
| Saccharomyces | | | | | | |
| 01066L | 01066K | 01066P | Saccharomyces cerevisiae derived from ATCC® 18824™* | E | 1 | Yeast cells |
| 0534L | 0534K | 0534P | Saccharomyces cerevisiae derived from ATCC® 4098™* | D | 1 | Yeast cells |
| 0186L | 0186K | 0186P | Saccharomyces cerevisiae derived from ATCC® 7754™* | E | 1 | Yeast cells |
| 0900L | 0900K | 0900P | Saccharomyces cerevisiae derived from ATCC® 9080™* | D | 1 | Yeast cells |
| 0699L | 0699K | 0699P | Saccharomyces cerevisiae derived from ATCC® 9763™* | D | 1 | Yeast cells |
| 0736L | 0736K | 0736P | Saccharomyces cerevisiae derived from NCYC 79 | D | 1 | Yeast cells |
| 0374L | 0374K | 0374P | Saccharomyces cerevisiae derived from NCYC 853 | D | 1 | Yeast cells |
| 0698L | 0698K | 0698P | Saccharomyces kudriavzevii derived from ATCC® 2601™* | D | 1 | Formerly Saccharomyces cerevisiae; yeast cells |
| Salmonella | | | | | | |
| 0595L | 0595K | 0595P | Salmonella bongori derived from ATCC® 43975™* | E | 2 | |
| 0901L | 0901K | 0901P | Salmonella enterica subsp. arizonae derived from ATCC® 13314™* | D | 2 | |
| 01054L | 01054K | 01054P | Salmonella enterica subsp. diarizonae derived from ATCC® 12325™* | E | 2 | Lactose broth positive; H2S positive |
| 01045L | 01045K | 01045P | Salmonella enterica subsp. diarizonae derived from ATCC® 29934™* | E | 2 | Lactose broth positive; H2S negative |
| 0501L | 0501K | 0501P | Salmonella enterica subsp. enterica derived from ATCC® 51741™* | D | 2 | H2S negative; formerly Salmonella choleraesuis subsp. choleraesuis serovar Infantis |
| 0817L | 0817K | 0817P | Salmonella enterica subsp. enterica serovar Abaetetuba derived from ATCC® 35640™* | D | 2 | |
| 0826L | 0826K | 0826P | Salmonella enterica subsp. enterica serovar Abaetetuba derived from Silliker® SLR156 | C | 2 | |
| 0890L | 0890K | 0890P | Salmonella enterica subsp. enterica serovar Abony derived from NCTC 6017 | C | 2 | Serovar Abony |
| 0346L | 0346K | 0346P | Salmonella enterica subsp. enterica serovar Anatum derived from ATCC® 9270™* | C | 2 | Group E |
| 01056L | 01056K | 01056P | Salmonella enterica subsp. enterica serovar Bispebjerg derived from ATCC® 9842™* | E | 2 | H2S negative; formerly Salmonella Abortusequi |
| 0902L | 0902K | 0902P | Salmonella enterica subsp. enterica serovar Choleraesuis derived from ATCC® 10708™* | C | 2 | H2S negative |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|-------------------|------------------|------------------|--|------------|-----|----------------------------------|
| 0343L | 0343K | 0343P | Salmonella enterica subsp. enterica serovar Choleraesuis derived from ATCC® 7001™* | C | 2 | Group C; H2S negative |
| 0345L | 0345K | 0345P | Salmonella enterica subsp. enterica serovar Enteritidis derived from ATCC® 13076™* | C | 2 | Group D |
| 01103L | 01103K | 01103P | Salmonella enterica subsp. enterica serovar Enteritidis derived from ATCC® 49223™* | E | 2 | |
| 01095L | 01095K | 01095P | Salmonella enterica subsp. enterica serovar Newport derived from ATCC® 6962™* | D | 2 | |
| 0341L | 0341K | 0341P | Salmonella enterica subsp. enterica serovar Paratyphi A derived from ATCC® 9150™* | C | 2 | Group A; H2S negative |
| 01170L | 01170K | 01170P | Salmonella enterica subsp. enterica serovar Paratyphi B derived from ATCC® 8759™* | C | 2 | |
| 0851L | 0851K | 0851P | Salmonella enterica subsp. enterica serovar Poona derived from NCTC 4840 | C | 2 | |
| 0604L | 0604K | 0604P | Salmonella enterica subsp. enterica serovar Pullorum derived from ATCC® 13036™* | D | 2 | H2S negative |
| 0344L | 0344K | 0344P | Salmonella enterica subsp. enterica serovar Tallahassee derived from ATCC® 12002™* | D | 2 | |
| 0421L | 0421K | 0421P | Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 13311™* | D | 2 | |
| 0363L | 0363K | 0363P | Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 14028™* | C | 2 | |
| 0253L | 0253K | 0253P | Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 25241™* | C | 2 | |
| 0180L | 0180K | 0180P | Salmonella enterica subsp. enterica serovar Typhimurium derived from ATCC® 51812™* | D | 2 | |
| 0340L | 0340K | 0340P | Salmonella enterica subsp. enterica serovar Typhimurium derived from NCTC 74 | D | 2 | |
| 0342L | 0342K | 0342P | Salmonella enterica subsp. enterica serovar Vellore derived from ATCC® 15611™* | D | 2 | Group B |
| 01087L | 01087K | 01087P | Salmonella enterica subsp. salamae serotype Tranaroa derived from NCTC 10252 | D | 2 | Serotype Tranaroa |
| Serratia | | | | | | |
| 0838L | 0838K | 0838P | Serratia liquefaciens derived from ATCC® 27592™* | C | 1 | |
| 0247L | 0247K | 0247P | Serratia marcescens derived from ATCC® 13880™* | D | 1 | |
| 0506L | 0506K | 0506P | Serratia marcescens derived from ATCC® 14041™* | D | 1 | Pigmented |
| 0806L | 0806K | 0806P | Serratia marcescens derived from ATCC® 14756™* | D | 1 | |
| 0216L | 0216K | 0216P | Serratia marcescens derived from ATCC® 43861™* | C | 1 | |
| 0262L | 0262K | 0262P | Serratia marcescens derived from ATCC® 43862™* | C | 1 | |
| 0354L | 0354K | 0354P | Serratia marcescens derived from ATCC® 8100™* | C | 1 | |
| 01146L | 01146K | 01146P | Serratia marcescens derived from NCTC 9743 | C | 2 | |
| Shewanella | | | | | | |
| 0638L | 0638K | 0638P | Shewanella haliotis derived from ATCC® 49138™* | C | 1 | Formerly Shewanella putrefaciens |



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|-------------------------|------------------|------------------|--|------------|-----|---|
| 0888L | 0888K | 0888P | Shewanella putrefaciens derived from ATCC® 8071™* | D | 1 | |
| Shigella | | | | | | |
| 01209L | 01209K | 01209P | Shigella boydii (2) derived from ATCC® 8700™* | C | 2 | Serotype 2 |
| 0349L | 0349K | 0349P | Shigella boydii derived from ATCC® 9207™* | C | 2 | Serotype 1 |
| 01235L | 01235K | 01235P | Shigella dysenteriae derived from ATCC® 13313™* | D | 2 | Export license required for shipping outside the U.S. |
| 0356L | 0356K | 0356P | Shigella flexneri derived from ATCC® 12022™* | C | 2 | Serotype 2b |
| 0348L | 0348K | 0348P | Shigella flexneri derived from ATCC® 9199™* | C | 2 | Serotype 1a |
| 0303L | 0303K | 0303P | Shigella sonnei derived from ATCC® 25931™* | C | 2 | |
| 0350L | 0350K | 0350P | Shigella sonnei derived from ATCC® 29930™* | D | 2 | |
| 0446L | 0446K | 0446P | Shigella sonnei derived from ATCC® 9290™* | B | 2 | |
| 0705L | 0705K | 0705P | Shigella sonnei derived from NCTC 12984 | C | 2 | |
| Sphingobacterium | | | | | | |
| 0948L | 0948K | 0948P | Sphingobacterium multivorum derived from ATCC® 35656™* | C | 2 | |
| Sphingomonas | | | | | | |
| 0274L | 0274K | 0274P | Sphingomonas paucimobilis derived from ATCC® 29837™* | C | 2 | |
| Sporidiobolus | | | | | | |
| 01013L | 01013K | 01013P | Sporidiobolus salmonicolor derived from ATCC® MYA-4550™* | D | 1 | Yeast cells |
| Staphylococcus | | | | | | |
| 0179L | 0179K | 0179P | Staphylococcus aureus derived from ATCC® BAA-1026™* | D | 2 | |
| 01007L | 01007K | 01007P | Staphylococcus aureus derived from ATCC® BAA-1708™* | D | 2 | mecA positive; SCCmec type II positive; Methicillin resistant; mupA positive; CLSI screening control for high level mupirocin resistance |
| 01122L | 01122K | 01122P | Staphylococcus aureus derived from ATCC® BAA-2312™* | D | 2 | mecC positive; SCCmec type XI positive; Methicillin resistant; this strain contains a divergent mecA gene that is not detected by standard PCR kits |
| 01065L | 01065K | 01065P | Staphylococcus aureus derived from NCTC 12493 | C | 2 | mecA positive; Methicillin resistant; EUCAST QC strain for cefoxitin |
| 0831L | 0831K | 0831P | Staphylococcus aureus derived from NCTC 6571 | B | 2 | |
| 0462L | 0462K | 0462P | Staphylococcus aureus subsp. aureus derived from ATCC® 11632™* | C | 2 | Beta lactamase producer |
| 01246L | 01246K | 01246P | Staphylococcus aureus subsp. aureus derived from ATCC® 12598™* | C | 2 | Cowan's serotype 1; contains a protein A |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|-----------|------------------|------------------|---|------------|-----|--|
| 0173L | 0173K | 0173P | Staphylococcus aureus subsp. aureus derived from ATCC® 12600™* | D | 2 | |
| 0360L | 0360K | 0360P | Staphylococcus aureus subsp. aureus derived from ATCC® 25923™* | A | 2 | mecA negative; CAMP test control; CLSI control for antimicrobial susceptibility testing using disk diffusion method |
| 0365L | 0365K | 0365P | Staphylococcus aureus subsp. aureus derived from ATCC® 29213™* | A | 2 | mecA negative; weak beta-lactamase producing strain; Oxacillin sensitive; EUCAST QC strain for Disk Diffusion; CLSI and EUCAST QC strain for MIC determination |
| 0906L | 0906K | 0906P | Staphylococcus aureus subsp. aureus derived from ATCC® 29737™* | C | 2 | |
| 0496L | 0496K | 0496P | Staphylococcus aureus subsp. aureus derived from ATCC® 33591™* | C | 2 | Methicillin resistant |
| 0889L | 0889K | 0889P | Staphylococcus aureus subsp. aureus derived from ATCC® 33592™* | B | 2 | mecA positive; SCCmec type III positive; Methicillin and gentamicin resistant |
| 0352L | 0352K | 0352P | Staphylococcus aureus subsp. aureus derived from ATCC® 33862™* | B | 2 | Recommended for CAMP Test |
| 0852L | 0852K | 0852P | Staphylococcus aureus subsp. aureus derived from ATCC® 43300™* | C | 2 | mecA positive; SCCmec type II positive; Methicillin and oxacillin resistant; CLSI control for cefoxitin disk diffusion, oxacillin agar, and antimicrobial susceptibility testing |
| 0937L | 0937K | 0937P | Staphylococcus aureus subsp. aureus derived from ATCC® 49476™* | D | 2 | |
| 0832L | 0832K | 0832P | Staphylococcus aureus subsp. aureus derived from ATCC® 51153™* | D | 2 | |
| 0159L | 0159K | 0159P | Staphylococcus aureus subsp. aureus derived from ATCC® 51740™* | D | 2 | |
| 0827L | 0827K | 0827P | Staphylococcus aureus subsp. aureus derived from ATCC® 6538F™* | C | 2 | |
| 0485L | 0485K | 0485P | Staphylococcus aureus subsp. aureus derived from ATCC® 6538™* | C | 2 | |
| 01022L | 01022K | 01022P | Staphylococcus aureus subsp. aureus derived from ATCC® 700698™* | E | 2 | mecA positive; SCCmec type II positive; Methicillin resistant; Heterogeneous susceptibility to vancomycin; GRD Etest® control |
| 0158L | 0158K | 0158P | Staphylococcus aureus subsp. aureus derived from ATCC® 700699™* | D | 2 | mecA positive; SCCmec type II positive; Methicillin resistant; reduced vancomycin susceptibility, Mu50; absence of pvl gene; propagate on BHI with 4 µg/l vancomycin in order to retain reduced susceptibility |



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KWIK-STIK and LYFO DISK

| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|-----------|------------------|------------------|--|------------|-----|---|
| 0113L | 0113K | 0113P | Staphylococcus aureus subsp. aureus derived from ATCC® 9144™* | D | 2 | |
| 01055L | 01055K | 01055P | Staphylococcus aureus subsp. aureus derived from ATCC® BAA-44™* | D | 2 | Methicillin-resistant |
| 0146L | 0146K | 0146P | Staphylococcus aureus subsp. aureus derived from ATCC® BAA-976™* | C | 2 | Clindamycin sensitive; negative control for inducible clindamycin resistant D-zone test; positive for msaA-mediated macrolide-only resistance |
| 0147L | 0147K | 0147P | Staphylococcus aureus subsp. aureus derived from ATCC® BAA-977™* | C | 2 | Positive control for inducible clindamycin resistant D-zone test; positive for ermA-mediated resistance |
| 0312L | 0312K | 0312P | Staphylococcus aureus subsp. aureus derived from NCIMB 12702 | A | 2 | |
| 0713L | 0713K | 0713P | Staphylococcus aureus subsp. aureus derived from NCTC 12973 | C | 2 | |
| 0533L | 0533K | 0533P | Staphylococcus capitis derived from ATCC® 146™* | E | 1 | Formerly Staphylococcus epidermidis |
| 0963L | 0963K | 0963P | Staphylococcus capitis subsp. capitis derived from ATCC® 35661™* | C | 1 | |
| 0371L | 0371K | 0371P | Staphylococcus epidermidis derived from ATCC® 12228™* | B | 1 | |
| 0412L | 0412K | 0412P | Staphylococcus epidermidis derived from ATCC® 14990™* | B | 1 | |
| 01189L | 01189K | 01189P | Staphylococcus epidermidis derived from ATCC® 35984™* | E | 1 | |
| 0628L | 0628K | 0628P | Staphylococcus epidermidis derived from ATCC® 49134™* | C | 2 | |
| 0976L | 0976K | 0976P | Staphylococcus epidermidis derived from ATCC® 49461™* | C | 2 | |
| 01068L | 01068K | 01068P | Staphylococcus epidermidis derived from ATCC® 51625™* | E | 2 | Methicillin resistant |
| 0571L | 0571K | 0571P | Staphylococcus epidermidis derived from NCIMB 8853 | C | 1 | |
| 0629L | 0629K | 0629P | Staphylococcus gallinarum derived from ATCC® 49148™* | D | 2 | Formerly Staphylococcus xylosum |
| 0246L | 0246K | 0246P | Staphylococcus haemolyticus derived from ATCC® 29970™* | D | 2 | |
| 0739L | 0739K | 0739P | Staphylococcus lentus derived from ATCC® 700403™* | D | 1 | |
| 0217L | 0217K | 0217P | Staphylococcus lugdunensis derived from ATCC® 49576™* | B | 1 | |
| 0762L | 0762K | 0762P | Staphylococcus lugdunensis derived from ATCC® 700328™* | D | 2 | |
| 01150L | 01150K | 01150P | Staphylococcus lugdunensis derived from NCTC 7990 | D | 2 | |
| 0907L | 0907K | 0907P | Staphylococcus pseudintermedius derived from ATCC® 49444™* | C | 2 | Formerly Staphylococcus aureus subsp. aureus; recommended for CAMP test for Listeria monocytogenes |



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|-------------------------|------------------|------------------|--|------------|-----|--|
| 0494L | 0494K | 0494P | Staphylococcus saprophyticus derived from ATCC® 15305™* | B | 1 | |
| 01035L | 01035K | 01035P | Staphylococcus saprophyticus derived from ATCC® 35552™* | D | 1 | |
| 0945L | 0945K | 0945P | Staphylococcus saprophyticus derived from ATCC® 49453™* | D | 1 | |
| 0881L | 0881K | 0881P | Staphylococcus saprophyticus derived from ATCC® 49907™* | D | 1 | |
| 0134L | 0134K | 0134P | Staphylococcus saprophyticus derived from ATCC® BAA-750™* | B | 1 | |
| 0764L | 0764K | 0764P | Staphylococcus sciuri subsp. sciuri derived from ATCC® 29061™* | D | 1 | |
| 01067L | 01067K | 01067P | Staphylococcus species derived from ATCC® 27626™* | E | 1 | |
| 0946L | 0946K | 0946P | Staphylococcus warneri derived from ATCC® 49454™* | C | 1 | |
| 0605L | 0605K | 0605P | Staphylococcus xylosus derived from ATCC® 29971™* | D | 2 | |
| 0741L | 0741K | 0741P | Staphylococcus xylosus derived from ATCC® 700404™* | D | 1 | |
| Stenotrophomonas | | | | | | |
| 01020L | 01020K | 01020P | Stenotrophomonas maltophilia derived from ATCC® 13636™* | D | 1 | Positive control for Etest® metallo Beta-Lactamase strip |
| 0369L | 0369K | 0369P | Stenotrophomonas maltophilia derived from ATCC® 13637™* | C | 1 | |
| 0759L | 0759K | 0759P | Stenotrophomonas maltophilia derived from ATCC® 17666™* | C | 1 | |
| 0135L | 0135K | 0135P | Stenotrophomonas maltophilia derived from ATCC® 49130™* | B | 1 | |
| 0742L | 0742K | 0742P | Stenotrophomonas maltophilia derived from ATCC® 51331™* | D | 1 | |
| Streptococcus | | | | | | |
| 01234L | 01234K | 01234P | Streptococcus agalactiae (III) derived from ATCC® BAA-22™* | C | 2 | Serotype III |
| 0439L | 0439K | 0439P | Streptococcus agalactiae derived from ATCC® 12386™* | C | 2 | Group B |
| 0104L | 0104K | 0104P | Streptococcus agalactiae derived from ATCC® 12403™* | E | 2 | Group B; Type III |
| 0370L | 0370K | 0370P | Streptococcus agalactiae derived from ATCC® 13813™* | B | 2 | Group B; nonhemolytic in absence of CAMP Factor |
| 0436L | 0436K | 0436P | Streptococcus agalactiae derived from ATCC® 27956™* | B | 2 | Group B |
| 0105L | 0105K | 0105P | Streptococcus agalactiae derived from ATCC® BAA-611™* | D | 2 | Group B; serotype V |
| 0709L | 0709K | 0709P | Streptococcus agalactiae derived from NCIMB 701348 | C | 2 | Group B |
| 0480L | 0480K | 0480P | Streptococcus agalactiae derived from NCTC 8017 | C | 2 | Group B |
| 0710L | 0710K | 0710P | Streptococcus agalactiae derived from NCTC 9993 | E | 2 | Group B |
| 01077L | 01077K | 01077P | Streptococcus anginosus derived from ATCC® 33397™* | C | 2 | Group G |
| 0389L | 0389K | 0389P | Streptococcus anginosus derived from NCTC 10713 | D | 2 | Group G |
| 0463L | 0463K | 0463P | Streptococcus bovis derived from ATCC® 33317™* | C | 1 | |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|-----------|------------------|------------------|---|------------|-----|--|
| 0603L | 0603K | 0603P | Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 12388™* | C | 2 | Group C |
| 0602L | 0602K | 0602P | Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 12394™* | C | 2 | Group G |
| 0967L | 0967K | 0967P | Streptococcus dysgalactiae subsp. equisimilis derived from ATCC® 35666™* | E | 2 | Group C |
| 0373L | 0373K | 0373P | Streptococcus dysgalactiae subsp. equisimilis derived from NCTC 8543 | E | 2 | Group C |
| 0221L | 0221K | 0221P | Streptococcus equi subsp. equi derived from ATCC® 33398™* | C | 2 | Group C |
| 0656L | 0656K | 0656P | Streptococcus equi subsp. equi derived from ATCC® 9528™* | C | 2 | Group C |
| 0101L | 0101K | 0101P | Streptococcus equi subsp. zooepidemicus derived from ATCC® 43079™* | C | 2 | |
| 0743L | 0743K | 0743P | Streptococcus equi subsp. zooepidemicus derived from ATCC® 700400™* | C | 2 | |
| 0631L | 0631K | 0631P | Streptococcus gallolyticus derived from ATCC® 49147™* | C | 1 | |
| 0391L | 0391K | 0391P | Streptococcus gallolyticus derived from ATCC® 9809™* | C | 1 | |
| 0949L | 0949K | 0949P | Streptococcus gallolyticus subsp. gallolyticus derived from ATCC® 49475™* | D | 1 | |
| 0383L | 0383K | 0383P | Streptococcus mitis derived from NCIMB 13770 | C | 2 | |
| 0266L | 0266K | 0266P | Streptococcus mutans derived from ATCC® 25175™* | C | 1 | |
| 0969L | 0969K | 0969P | Streptococcus mutans derived from ATCC® 35668™* | C | 1 | |
| 0423L | 0423K | 0423P | Streptococcus oralis derived from ATCC® 6249™* | C | 2 | Formerly Streptococcus mitis |
| 0630L | 0630K | 0630P | Streptococcus pasteurianus derived from ATCC® 49133™* | C | 1 | |
| 0865L | 0865K | 0865P | Streptococcus pneumoniae derived from ATCC® 10015™* | C | 2 | |
| 0435L | 0435K | 0435P | Streptococcus pneumoniae derived from ATCC® 27336™* | C | 2 | |
| 0632L | 0632K | 0632P | Streptococcus pneumoniae derived from ATCC® 49136™* | C | 2 | |
| 0633L | 0633K | 0633P | Streptococcus pneumoniae derived from ATCC® 49150™* | C | 2 | |
| 0947L | 0947K | 0947P | Streptococcus pneumoniae derived from ATCC® 49619™* | C | 2 | Penicillin intermediate by altered penicillin-binding protein; serotype 19F, type 19; CLSI and EUCAST QC strain for Disk Diffusion and MIC Determination |
| 0763L | 0763K | 0763P | Streptococcus pneumoniae derived from ATCC® 6301™* | C | 2 | |
| 0380L | 0380K | 0380P | Streptococcus pneumoniae derived from ATCC® 6303™* | C | 2 | Mucoid colonies |
| 0447L | 0447K | 0447P | Streptococcus pneumoniae derived from ATCC® 6305™* | C | 2 | |
| 0267L | 0267K | 0267P | Streptococcus pneumoniae derived from ATCC® 700677™* | D | 2 | |



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|------------------------------|------------------|------------------|---|------------|-----|---|
| 0469L | 0469K | 0469P | Streptococcus pneumoniae derived from NCIMB 13286 | E | 2 | |
| 0508L | 0508K | 0508P | Streptococcus pyogenes derived from ATCC® 12344™* | D | 2 | Group A; type 1 |
| 0979L | 0979K | 0979P | Streptococcus pyogenes derived from ATCC® 12384™* | B | 2 | Group A; type 3 |
| 0385L | 0385K | 0385P | Streptococcus pyogenes derived from ATCC® 19615™* | B | 2 | Group A |
| 0994L | 0994K | 0994P | Streptococcus pyogenes derived from ATCC® 49399™* | B | 2 | Group A |
| | | 0314P | Streptococcus pyogenes derived from NCIMB 13285 | C | 2 | Group A |
| 0237L | 0237K | 0237P | Streptococcus salivarius derived from ATCC® 13419™* | D | 1 | |
| 0136L | 0136K | 0136P | Streptococcus salivarius subsp. thermophilus derived from ATCC® 19258™* | C | 1 | |
| 0858L | 0858K | 0858P | Streptococcus sanguinis derived from ATCC® 10556™* | C | 2 | |
| 0978L | 0978K | 0978P | Streptococcus species derived from ATCC® 12392™* | D | 1 | Group F; type 2 |
| 0864L | 0864K | 0864P | Streptococcus species derived from ATCC® 12401™* | C | 2 | Group B; type 1b |
| 0744L | 0744K | 0744P | Streptococcus uberis derived from ATCC® 700407™* | D | 1 | |
| Streptomyces | | | | | | |
| 0859L | 0859K | 0859P | Streptomyces griseus subsp. griseus derived from ATCC® 10137™* | C | 1 | |
| Talaromyces | | | | | | |
| 01238L | 01238K | 01238P | Talaromyces pinophilus derived from ATCC® 11797™* | D | 1 | Mold; fungus resistance testing; formerly Penicillium funiculosum |
| 0535L | 0535K | 0535P | Talaromyces wortmannii derived from ATCC® 32333™* | E | 1 | Mold; formerly Penicillium variabile |
| Thermoanaerobacterium | | | | | | |
| 0728L | 0728K | 0728P | Thermoanaerobacterium thermosaccharolyticum derived from ATCC® 7956™* | E | 1 | |
| Trichophyton | | | | | | |
| 0442L | 0442K | 0442P | Trichophyton interdigitale derived from ATCC® 9533™* | E | 2 | Mold; formerly Trichophyton mentagrophytes |
| 0444L | 0444K | 0444P | Trichophyton rubrum derived from ATCC® 28188™* | E | 2 | Mold |
| 0891L | 0891K | 0891P | Trichophyton tonsurans derived from ATCC® 28942™* | E | 2 | Mold |
| Trichosporon | | | | | | |
| 0778L | 0778K | 0778P | Trichosporon dermatis derived from ATCC® 204094™* | E | 2 | Formerly Trichosporon mucoides; yeast cells |
| Trueperella | | | | | | |
| 0660L | 0660K | 0660P | Trueperella pyogenes derived from ATCC® 19411™* | D | 2 | Formerly Arcanobacterium pyogenes |
| Ureaplasma | | | | | | |
| 0151L | 0151K | 0151P | Ureaplasma parvum derived from ATCC® 27813™* | E | 2 | |
| Veillonella | | | | | | |
| 0867L | 0867K | 0867P | Veillonella parvula derived from ATCC® 10790™* | E | 1 | |



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| LYFO DISK | KWIK-STIK 6-Pack | KWIK-STIK 2-Pack | Product Description | Price Code | BSL | Comment |
|--------------------------|------------------|------------------|--|------------|-----|---|
| Vibrio | | | | | | |
| 0819L | 0819K | 0819P | Vibrio alginolyticus derived from ATCC® 17749™* | C | 1 | |
| 0720L | 0720K | 0720P | Vibrio furnissii derived from NCTC 11218 | C | 1 | |
| 01236L | 01236K | 01236P | Vibrio parahaemolyticus (o2:k3) derived from NCTC 10885 | E | 2 | Serovar o2:k3 |
| 0818L | 0818K | 0818P | Vibrio parahaemolyticus derived from ATCC® 17802™* | D | 2 | |
| 01076L | 01076K | 01076P | Vibrio vulnificus derived from ATCC® 27562™* | D | 2 | |
| 01214L | 01214K | 01214P | Vibrio vulnificus derived from ATCC® 29307™* | E | 2 | Halophilic; recommended by ISO 11133 for performance testing of culture media |
| Walleimia | | | | | | |
| 01141L | 01141K | 01141P | Walleimia mellicola derived from ATCC® 42694™* | D | 1 | Mold; formerly Walleimia sebi |
| Yarrowia | | | | | | |
| 0950L | 0950K | 0950P | Yarrowia lipolytica derived from ATCC® 9773™* | D | 1 | Yeast cells |
| Yersinia | | | | | | |
| 0316L | 0316K | 0316P | Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 23715™* | C | 2 | Biotype 1; serotype 8 |
| 0909L | 0909K | 0909P | Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 27729™* | C | 2 | Biotype 1; serotype 8 |
| 0938L | 0938K | 0938P | Yersinia enterocolitica subsp. enterocolitica derived from ATCC® 9610™* | D | 2 | Biovar 1; serogroup O:8 |
| 0974L | 0974K | 0974P | Yersinia kristensenii derived from ATCC® 33639™* | D | 2 | |
| 0785L | 0785K | 0785P | Yersinia ruckeri derived from ATCC® 29473™* | E | 1 | |
| Zygosaccharomyces | | | | | | |
| 01011L | 01011K | 01011P | Zygosaccharomyces parabailii derived from ATCC® MYA-4549™* | E | 1 | Yeast cells |
| 0803L | 0803K | 0803P | Zygosaccharomyces rouxii derived from ATCC® 28253™* | E | 1 | Yeast cells |
| 0255L | 0255K | 0255P | Zygosaccharomyces rouxii derived from NCYC 381 | E | 1 | |



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