

Staphylococcus aureus* Fluorescent Reporter Plasmid pSFRFPS1, Recombinant in *Staphylococcus aureus

Catalog No. NR-51165

Product Description: NR-51165 is a glycerol stock of *Staphylococcus aureus* (*S. aureus*), strain RN4220 containing the eqFP650 far-red fluorescent protein (FRFP) reporter plasmid pSFRFPS1, a derivative of the *Escherichia coli* (*E. coli*) - staphylococcal shuttle vector pKK30.

Lot¹: 70010752

Manufacturing Date: 22NOV2017

TEST	SPECIFICATIONS	RESULTS
Phenotypic Analysis Cellular morphology Colony morphology ² Motility (wet mount) Hemolysis ³ Biochemical characterization Catalase VITEK [®] MS (MALDI-TOF)	Gram-positive cocci Report results Report results Report results Positive <i>S. aureus</i>	Gram-positive cocci Circular, convex, entire, smooth and cream to slight lavender (Figure 1) Non-motile Non-hemolytic ⁴ Positive <i>S. aureus</i> (99.9%)
Genotypic Analysis Sequencing of 16S ribosomal RNA gene (~ 770 base pairs)	≥ 99% sequence identity to <i>S. aureus</i> , strain RN4220 (GenBank: AFGU01000017.1)	100% sequence identity to <i>S. aureus</i> , strain RN4220 (GenBank: AFGU01000017.1) ⁵
Confirmation of pSFRFPS1 plasmid	Report results	Consistent with pSFRFPS1 plasmid description (Figure 2, Table 1) ^{6,7}
Functional Activity of Antibiotic Resistance Gene in <i>S. aureus</i> Trimethoprim ²	Growth	Growth
Purity (post-freeze)⁸	Growth consistent with expected colony morphology	Growth consistent with expected colony morphology
Viability (post-freeze)²	Growth	Growth

¹NR-51165 was produced by inoculation of the deposited material in Tryptic Soy broth containing 10 µg/mL trimethoprim and incubated for 1 day at 37°C in an aerobic atmosphere. Broth inoculum was used to inoculate Tryptic Soy agar with 10 µg/mL trimethoprim kolles, which were grown for 1 day at 37°C in an aerobic atmosphere to produce this lot.

²1 day at 37°C in an aerobic atmosphere on Tryptic Soy agar with 10 µg/mL trimethoprim

³1 day at 37°C in an aerobic atmosphere on Tryptic Soy agar with 5% defibrinated sheep blood.

⁴Limited β-hemolysis was observed after 2 days

⁵Also consistent with *S. argenteus* and *S. simiae*

⁶Illumina[®] MiSeq[®] sequence was analyzed with CLC Genomics Workbench Version 7.0.2.

⁷pSFRFPS1 was sequenced and annotated by BEI Resources and is consistent with the vector described in Rodriguez, M. D., et al. "Construction of Stable Fluorescent Reporter Plasmids for Use in *Staphylococcus aureus*." *Front. Microbiol.* 8 (2017): 2491. PubMed: 29312199.

⁸Purity of this lot was assessed for 7 days at 37°C in an aerobic atmosphere with 5% CO₂ on Tryptic Soy agar with 5% defibrinated sheep blood.

Figure 1: Colony Morphology

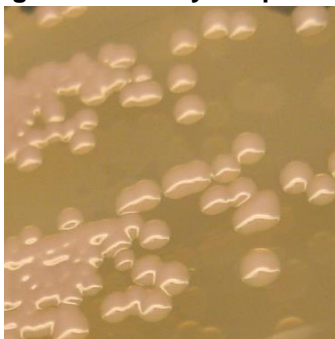


Figure 2: Fluorescent Report Plasmid pSFRFPS1

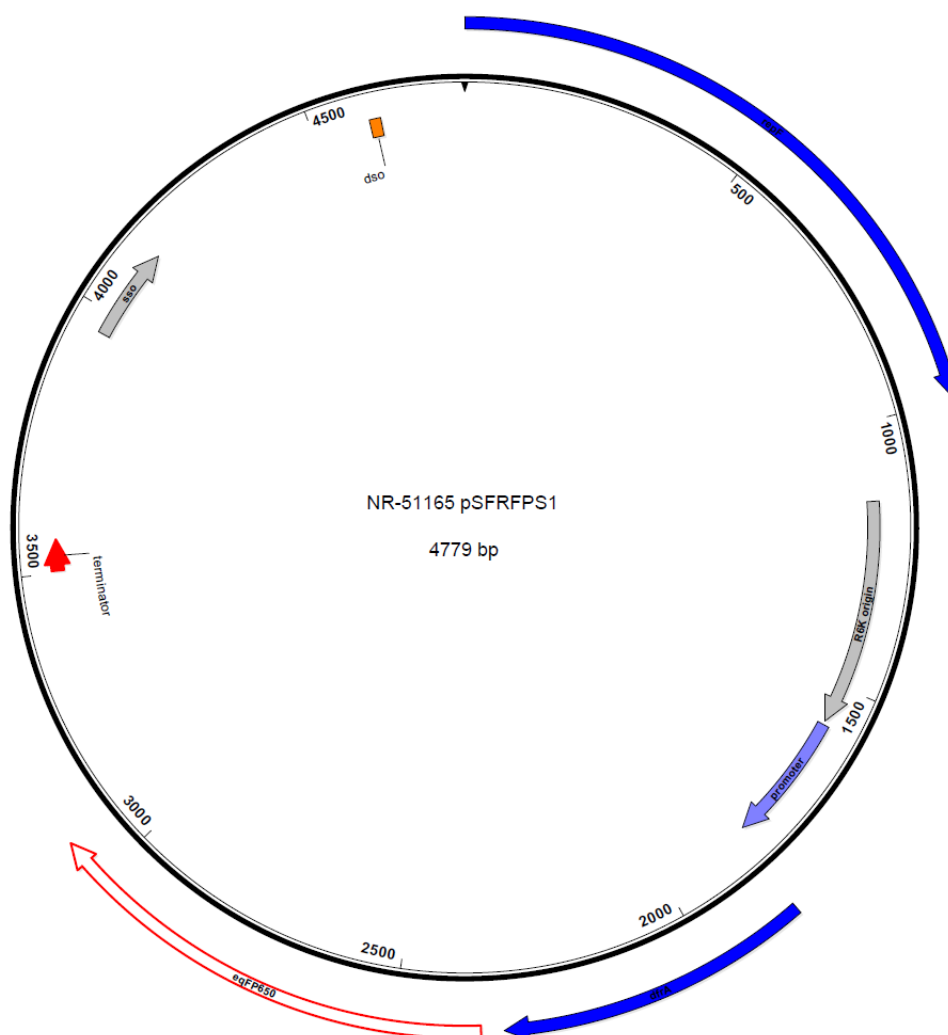


Table 1: Sequence of pSFRFPS1

1	ATGCAATATA	ATACTACTAG	AAGTATAACC	GAAAAATCAAG	ATAATAAAAAC	GTTAAAAAGAT
61	ATGACGAAAA	GTGGGAAACA	ACGCCCATGG	AGAGAAAAGA	AAATAGATAA	TGTAAGCTAT
121	GCAGATATAC	TAGAAATTTT	AAAAATCAAA	AAGGCTTTTA	ATGTAAAACA	ATGTGGTAAT
181	ATTTTAGAAT	TTAAGCCAAC	TGATGAAGGC	TATTTGAAGT	TACATAAGAC	ATGGTTTTGT
241	AAATCAAAAT	TATGTCCGGT	TTGTAATTGG	AGACGTGCTA	TGAAAAATAG	TTATCAAGCT
301	CAAAAAGTGA	TTGAAAAAGT	AATTAAGGAA	AAGCCAAAAG	CACGTTGGTT	GTTTTTAACA
361	CTTTCAACAA	AAAATGCGAT	AGATGGAGAT	ACTTTAGAAC	AAAGTTTGAA	GCATCTAACT
421	AAAGCATTTG	ATAGGTTGAG	TAGATATAAA	AAGGTTAAAC	AAAATCTTGT	TGGATTTATG
481	CGTTCAACAG	AAGTTACCGT	TAATAAAAAAT	GACGGTAGTT	ATAATCAGCA	CATGCATGTT
541	TTGTTATGTG	TTGAAAATGC	ATATTTTAGA	AAAAAAGAGA	ATTATATAAC	TCAAGAAGAA
601	TGGGTTAATT	TATGGCAAAG	AGCATTACAA	GTTGACTATC	GACCTGTTGC	TAATGTTAAA
661	GCGATCAAAC	CGAATAGAAA	AGGCGATAAA	GACATTGAAT	CGGCAATCAA	AGAGACCTCA
721	AAATATTCGG	TTAAATCATC	TGATTTTTTTA	ACTGATGATG	ATGAAAAAAA	TCAAGAAATT
781	GTAAGTGATT	TAGAAAAAGG	TTTGTATCGA	AAACGTATGT	TAAGTTATGG	TGGATTGCTT
841	AAACAAAAAC	ATAAAATTTT	AAACTTAGAC	GATGTGGAAG	ATGGTAATTT	GATTAATGCA
901	AGTGATGAAG	ATAAAACAAC	AGACGAAGAA	GAAAAAGCAC	ATTCAATTAC	CGCAATTTGG
961	AATTTTCGAAA	AGCAAAATTA	TTATTTAAGA	CATTAGTGTT	GACTAATGTC	TTTTTTGTTG
1021	ATTTTTTATA	AAAAAGTACT	CTTATTTTTG	TGACAAATGC	TGTATGTAGT	GTCACAAAAA
1081	TAAGACAAAC	GCAATATATT	GTGTCACAAA	AATAAGACAG	TACAGCTTTG	TATGATCCGT
1141	CGACGAAAGC	CTGGCCACGA	TGCGTCCGGC	GTAGAGGATC	TGAAGATCAG	CAGTTCAACC
1201	TGTTGATAGT	ACGTACTAAG	CTCTCATGTT	TCACGTAATA	AGCTCTCATG	TTTAACGTAC
1261	TAAGCTCTCA	TGTTTAACGA	ACTAAACCCT	CATGGCTAAC	GTAATAAGCT	CTCATGGCTA
1321	ACGTACTAAG	CTCTCATGTT	TCACGTAATA	AGCTCTCATG	TTTGAACAAT	AAAATTAATA
1381	TAAATCAGCA	ACTTAAATAG	CCTCTAAGGT	TTTTAAGTTT	ATAAGAAAAA	AAAGAAATATA
1441	TAAGGCTTTT	AAAGCTTTTA	AGGTTTAACG	GTTGTGGACA	ACAAGCCAGG	GATGTAACGC
1501	ACTGAGAAGC	CCTTAGAGCC	TCTCAAAGCA	ATTTTGAGTG	ACACAGGAAC	ACTTAACGGC
1561	TGACATGGGA	ATTTCGAGCTG	ATATTTTTGA	CTAAACCAAA	TGCTAACCCA	GAAATACAAT
1621	CACTGTGTCT	AATGAATAAT	TTGTTTTATA	AACACTTTTT	TGTTTACTTC	TCATTTTTAA
1681	TTAGTTATAA	TTAACTAAAT	AATAGAGCAT	TAAATATATT	TAATAAAACT	TATTTAATGC
1741	AAAATTATGA	CTAACATATC	TATAATAAAT	AAAGATTAGA	TATCAATATA	TTATCGGGCA
1801	AATGTATCGA	GCAAGATGCA	TCGGATCGAT	CCAGGAGGTA	TACCATGACA	TTATCAATAA
1861	TTGTCGCTCA	CGATAAACAA	AGAGTCATTG	GGTACCAAAA	TCAATTACCT	TGGCACTTAC
1921	CAAATGATTT	AAAGCATATT	AAACAACCTGA	CCACTGGGAA	TACACTTGTA	ATGGCACGGA
1981	AAACTTTTAA	TTCTATAGGG	AAGCCATTGC	CAAATAGACG	TAACGTCGTA	CTCACTAACC
2041	AAGCTTCATT	TCGCCATGAA	GGGGTAGATG	TTATAAACTC	TCTTGATGAA	ATTAAAGAGT
2101	TATCTGGTCA	TGTTTTTATA	TTTGGAGGAC	AAACGTTATA	CGAAGCAATG	ATTGACCAGG
2161	TAGATGATAT	GTATATCACA	GTAATAGATG	GAAAAGTTTCA	AGGAGACACA	TTCTTTCCAC
2221	CATACACATT	CGAAAACCTGG	GAAGTCGAAT	CTTCAGTAGA	AGGTCAACTA	GATGAAAAAA
2281	ATACTATACC	GCATACATTC	TTACATTTAG	TGCGTAGAAA	AGGGAAATAG	GCGCGCCTGA
2341	TTAACTTTAT	AAGGAGGAAA	AACATATGGG	TGAAGATAGT	GAATTAATTA	GTGAAAATAT
2401	GCACATGAAA	TTATATATGG	AAGGTACTGT	AAATGGTCAT	CATTTTAAAT	GTAATTCAGA
2461	AGGTGAAGGT	AAACCTTATG	AAGTTACACA	AACTGCAAAA	ATTAAAGTTG	TAGAAGGTGG
2521	TCCATTACCT	TTTGCATTTG	ATATTTTAGC	TACATCATTT	ATGTATGGTA	GTAAAACATT
2581	TATTAATCAT	ACTCAAGGTA	TTCCAGATTT	CTTTAAACAA	TCTTTTCCTG	AAGGTTTTAC
2641	ATGGGAACGT	ATTACAACCT	ATGAAGATGG	TGGTGTTTTA	ACAGCAACTC	AAGATACAAG
2701	TTTACAAAAAT	GGTTGTTTAA	TTTATAATGT	TAAAATTAAT	GGTGTAAATT	TTCCATCAAA
2761	TGGTCCTGTA	ATGCAAAAAGA	AAACATTAGG	TTGGGAAGCA	AGTACTGAAA	TGTTATATCC
2821	AGCTGATTCT	GGTTTACGTG	GTCATTACACA	AATGGCTTTA	AAATTAGTTG	GTGGTGGTTA
2881	TTTACATTGT	AGTTTAAAAA	CAACTTATAG	ATCTAAAAAA	CCAGCAAAAA	ATTTAAAAAT
2941	GCCTGGTTTT	TATTTTGTTG	ATCGTAAATT	AGAAAGAATT	AAAGAAGCTG	ATAAAGAAAC
3001	TTATGTTGAA	CAACATGAAA	TGGCTGTAGC	TCGTTATTGT	GATTTACCAT	CAAAATTAGG
3061	TCATAGTTAA	GGCGCGCCTA	TTCTAATGCA	TAATAAATAC	TGATAACATC	TTATATTTTG
3121	TATTATATTT	TGTATTATCG	TTGACATGTA	TAATTTTGAT	ATCAAAAACCT	GATTTTCCCT

3181 CTATTATTTT CGAGATTTAT TTTCTTAATT CTCTTTAACA AACTAGAAAT ATTGTATATA
 3241 CAAAAAATTA TAAATAATAG ATGAATAGTT TAATTATAGG TGTTTCATCAA TCGAAAAAGC
 3301 AACGTATCTT ATTTAAAGTG CGTTGCTTTT TTCTCATTTA TAAGGTTAAA TAATTCTCAT
 3361 ATATCAAGCA AAGTGACAGG CGATGCGGCC GCTAGCCTAG GAGCTCGGTA CCCGGGGATC
 3421 CGAATCATGA ATTACAAGCA AAAGTAGCGG TGATTGTTAA AATTGATGGT AAACAATCAC
 3481 CGCTATTTTT GCTTGTGTAT GTATAAAAAA GGGATCAAAG GTCATCCCCC ATGATTGATA
 3541 GTGGGGGGAT GACTTTTGAT CCTATGTTCA TGTTGCTTAT TTAATCGCCT TTGATCACTT
 3601 TAAAATACCT TAAAACCCCT GGAATTTCTG GCTTTGCCAG ACCTATCATT TTTGAATGAT
 3661 AGCAAATTCT CCTTATGCTC TTACGGAGTT TTTAGAGAAA AATTAAAAAT TCTCGATTTT
 3721 TGATAAAAAA CGCCCTGCAG GAATTTAGAA AAACATGTGG AAGTTTTAAA GGATTTTATG
 3781 CTAATTTTTTA ATTTGCATGT AACTCGAGGG GAATATTTGA GGGGATTTTG AAACGAGTTT
 3841 CTTCTTGTTT TCACACTGTT TTTTTATTCC TATTGGTGTT GTTGCTTACT TTTTGTTTTT
 3901 CTATAAAGAT GATGCTTTTG ACGAGATGGA AGAAAAATAT GATTATCATG AAGATAATAA
 3961 AAAATAGACG ACGCATTAT GCGGAGAAAA TTTATTGATG TTGAGAAGAA CCCTTAACTA
 4021 AACTTGCAGA CGAATGTCGG CATAGCGTGA GCTATTAAGC CGACCATTCTG ACAAGTTTTG
 4081 GGATTGTTAA GGGTTCCGAG GCTCAACGTC AATAAAGCAA TTGGAATAAA TAGTATCTAA
 4141 AGTAACCCGT ATTTAAGATT ACATTGTACA TCCTTCGAAA CCATAGTAGC TGCGATTCCA
 4201 GCTAGGTTTC CTTTTATCCC AATTTTAGCT AATTTTTTTG CTACGCCCTT TAAGTTTTTG
 4261 GCTTTAACAT CTTCAAAAAC TGCTGTGATT ATGTTTCCAG TTAAAAGTTC TCCATATGAG
 4321 TTTAAAACCT CAGAATAGAA ACAGTCACCG CTATTTTTGT AGTTCCCACC AACTGCTCTA
 4381 GGTGATATTG AACTAGTTTT CTTTTTATTC TCTATGTCGT TTTTTAGTTT TTGGAATCCT
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 4501 AGAAAGTTTT ATTTATTCAT TGGTTTTTAC ATTAATTTAT TTAGGTTTAG AGGGTCATAA
 4561 AAAGAAGAAG GAATAGGTTG TTTTTTGAAA CGAGTGTGAA CGAGTTTCTT CTTGTCTTGA
 4621 TACTATATAG AAATAACTCG ATTTTATATA TATAGCTGTA ACTGTTGATA TTACAGTGTT
 4681 TAAACGTGTT TTTGTGCGTG AAAGGAAAAT TTGACAATAA AAAACCCAG TTATATTATT
 4741 AAGGTGTCGA ATCTTAAATA ATACTGGGGG TCTTTTTTAT

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