Supplementary Data

Recent Advances in Aptamer-Based Biosensors for Bacterial Detection

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Summary

- Table S1. Sequence of aptamers used for bacterial detection
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Table S1. Sequences of aptamer used in biosensors for bacterial detection.

Bacteria	Aptamer	Sequence	Ref
S. aureus	SA20	GCGCCCTCTCACGTGGCATCAGAGTGCCGGAAGTTCTGCGTTAT	[1]
	SA23	GGGCTGGCCAGATCAGACCCCGGATGATCATCCTTGTGAGAACCA	
	SA34	CACAGTCACTCAGACGGCCGCTATTGTTGCCAGATTGCCTTTGGC	
	SA31	TCCCACGATCTCATTAGTCTGTGGATAAGCGTGGGACGTCTATGA	
	SA43	TCGGCACGTTCTCAGTAGCGCTCGCTGGTCATCCCACAGCTACGTC	
S. aureus	T1	ACTGTCrGrCrGrCrArCrGrCrGrUrGrUrGrUrArGrUrArCrArCrArCrGrArUrCrGrCrGrCrGrCrArCrArArUrArU	[2]
	T2	ACTGTCrArArUrUrUrGrArArUrArUrArUrUrUrArGrUrGrCrGrCrGrCrCrGrUrArGrUrGrUrGrUrArArArArArUrU	
	T3	ACTGTCrArArUrUrUrGrArGrUrGrUrGrUrGrArUrCrArUrArUrArUrCrGrUrArGrCrGrCrGrCrUrArCrArArCrC	
	A14	ACTGTCCACACCGCAGCAGTGGGAACGTTTCAGCCATGCAAGCATCACGCCCGT	
S. aureus	H1	GCAATGGTACGGTACTTCCTCGGCACGTTCTCAGTAGCGCTCGCTGGTCATCCCACAGCTACGTCAAAAGTGCACGCTACT TTGCTAA	[3]
S. aureus	H1	GCAATGGTACGGTACCCCTATGCGCATGTACCATTGCAGTTGTCAGAGAGCGA	[4]
	H2	GGTACA/Dabcyl/TGCGCATAGGGGTACCGTACCATACCCCTATGCGCA/FAM/	
	cApt	GTACCGTACCATTGCTAGCGTCTTCCCGTCCTT	
S. aureus	Apt1	TCCCTACGGCGCTAACCCCCCCCAGTCCGTCCTCCCAGCCTCACACCGCCACCGTGCTACAAC	[5,6]
	Apt2	TCCCTACGGCGCTAACCTCCCAACCGCTCCACCCTGCCTCCGCCTCGCCACCGTGCTACAAC	
S. aureus	SH-Apt$_2$	GCAATGGTACGGTACTTCCTCGGCACGTTCTCAGTAGCGCTCGCTGGTCATCCCACAGCTACGTCAAAAGTGCACGCTACT TTGCTAA	[7,8]
S. aureus	A15	AGCAG CACAGAGGTCAGATG(N40)CCTATGCGTGCTACCGTGAA-	[9]
S. aureus	APT^{seb1}	GGTATTGAGGGTCGCATCCACTGGTCGTTGTTGTCTGTTGTCTGTTATGTTGTTTCGTGATGGCTCTAACTCTCCTCT	[10]
S. aureus	A-SEB	TGCAGGATCCGGTATCCGTGCACACACACCCAACAACCAGCTGCCGCACCGGAGGAATTCTCGT	[11]
S. aureus	G1	UCCGAACAGCGGAAGGUGGUUCGAAGUUGGGGCUUUGGA	[12]
	#2	GGGAGUUUUGAUACGGCUUCAUGCAGUAAUGUUUUUAU	
	#18	UCCGAACAGCGGAAGGUGGUUCGAAGUUGGGGCUUUGGA	

S. aureus	AT-27 AT-33 AT-36 AT-49	ACCCCTGCAGGATCCTTTGCTGGTACC-(N42)-AGTATCGCTAATCAGTCTAGAGGGCCCCAGAAT	[13]
S. aureus	H1, P1	GCAATGGTACGGTACTTCCTCGGCACGTTCTCAGTAGCGCTCGCTGGTCATCCCACAGCTACGTCAAAAGTGCACGCTACT TTGCTAA	[14]
S. aureus	Antibac1 Antibac2	GGGACAGGGAGTGCGCTGCTCCCC GGGGACTAGAGGACTTGTGCGGCC	[15,16]
E. coli	SH-Apt1	GCAATGGTACGGTACTTCCCCATGAGTGTTGTGAAATGTTGGGACACTAGGTGGCATAGAGCCGCAAAAGTGCAGCTACT TTGCTAA	[17,18]
E. coli	/	ATCCGTCACACCTGCTCTATCAAATGTGCAGATATCAAGACGATTTGTACAAGATGGTGTTGGCTCCCGTAT	[19]
E. coli	GN6 GN12	ATACCAGCTTATTCAATTGGGTGAGGGGGGGTTCACAACGTTAAAGATAGACGGGGGAAGATAGTAAGTGCAATCT	[20]
E. coli	6-3 8-1 8-7 8-8 8-12 8-13 8-19 8-35	UGGUUUCAGCGACAGGAGGGGUGUAGGUGGAUUGCUGUCCUUUGCGUGU UGCUAGUGUUGUAUGCACGUGGAGGAGGAGGCGUACACUUGCUUUGUGGU GAUUGACCGUAUGGAGGAUGCAAAGGGAGGGAGGUCACUUGAGUUAGUUA GCAGGAUGUGGAGGAGGCAUCUGCUGCAAUCGGGACUUGUGUCGAGUAUC GCAUUGUCUGCGUGUGGAGGCAGGAGGCAAGAUAAGAGGUGAUGCGGUUG CAUGUUGGCGAUACGUCUAAACGGUGGGUUGUGGAGGAUUGAUUUAUACG AGUAGUGUCAGCGUGUGGUGGAGGUUGGCGACAUAUGUAGGGUGCGAUUG UGCGCAAUACACGGUGAGGAGGUGGAGAGAUGUAGGUGCUUAGCAGUUGA	[21]
E. coli	ECA I ECA II	GTCTGCGAGCGGGGCGCGGGCCCGGCGGGGGATGCGC ACGGCGCTCCCAACAGGCCTCTCCTTACGGCATATTA	[22,23]
E. coli	/	GCAATGGTACGGTACTTCCCCATGAGTGTTGTGAAATGTTGGGACACTAGGTGGCATAGAGCCGCAAAAGTGCACGCTAC TTTGCTAA	[24]
E. coli	Stx1 stx2	ATCCAGAGTGACGCAGCAGTAGTTTGTTGGTTATTACGGCGGGTTGCGATGGGTGCGAATCGGTGGACACGGTGGCTTAG T ATCCAGAGTGACGCAGCAGGAAAGGACGTCAAATTAGGGGCCGGGACAACGAAAGCCCACAACTGGACGGTGGCTTAG T	[25]

E.coli	E1 E2 E3	GCAATGGTACGGTACTTCCACTTAGGTCGAGGTTAGTTTGTCTTGCTGGCGCATCCACTGAGCGCAAAAGTGCACGCTACTTTGCTAA GCAATGGTACGGTACTTCCCCATGAGTGTTGTGAAATGTTGGGACACTAGGTGGCATAGAGCCGCAAAAGTGCACGCTACTTTGCTA GCAATGGTACGGTACTTCCGTTGCACTGTGCGGCCGAGCTGCCCCCTGGTTTGTGAATACCCTGGGCAAAAGTGCACGCTACTTTGCTAA	[26]
P. aeroginosa	F23	CCCCCGTTGCTTTCGCTTTTCCTTTCGCTTTTGTTCGTTTCGTCCCTGCTTCCTTTCTTG	[27]
B. cereus	/	AGCAGCACAGAGGTCAGATGCCCCCCTTTTATCCGTCGGCATGATGTCTCCCGATCCGGTCCTATGCGTGCTA	[28]
B. cereus	B15 B16	AGCAGCACAGAGGTCAGATGGGCGGGTTTGGATCTTTGGTTGGCGCCTGTTTCTTTATGACCTATGCGTGCTACCGTGAA AGCAGCACAGAGGTCAGATGATATGTTTACGCCAGTGGTATTATTGGGGTTGATATGTCACCTATGCGTGCTACCGTGAA	[29]
B. cereus	13-18 13-24	AGCACAGAGGTCAGATGGGCTACTGGAGCATCTGGTAACGAAGTACCCTCGGGGCGG AGCACAGAGGTCAGATGATCGAGGGCGCAGACCGAACCCGCGTGCGCAGTACAAGGGC	[30]
Acinetobacter baumannii	AB	AATCAGGCTCAGCATGGAGTTGCGAGGCCAATATCCGGTTAAGCG	[31,32]
Acinetobacterer baumanni	K2	ACAGCACCACAGACCACATATCACATGCTGTCGCCTTGCGATATCAATTCCAGTGATGTTTGTCTTCCTGCC	[31]
Leptospira interrogans	LAP3	TGGCGTTAGAGATACCGGAACCGGTGTCGGGCGTCTGAAGAATCC	[33]
Bacillus cytotoxicus	BAS6R	ATCCGTCACACCTGCTCTGCACGGGCTCAGTTTGGCTTTGTATCCTAAGAGGATGGTGTTGGCTCCCGTAT	[34]
Vibrio cholerae	CT916	GGCAAAAAGGATTGCCCAGGTCTGCTGTCTAGCCGGATTC	[35]
Clostridium difficile	/	TAGTGATGCCTTTGTTGAGA-[N40]-TCTTCATCGTCCACTAAATT	[36]
S. typhimurium	H2	TATGGCGGCGTCACCCGACGGGGACTTGACATTATGACAG	[14]
S. Typhimurium	Apt S.T	TGGCTAGCTCAGTCATATGGCGGCGTCACCCGACGGGGACTTGACATTATGACAGCCGCG	[37]

Vibrio parahaemolyticus	Apt VP	TGGCTAGCTCAGTCATCTAAAAATGGGCAAAGAAACAGTGACTCGTTGAGATACTCCGCG	[37]
Clostridium perfringens	/	TCAACGGCAGTAACATTAGC	[38]

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