

BA_2677 family bacteriocins

BA_2677	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG	RCGGCGG	R
YP_084014.1	6	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG	RCGG	R
ZP_03234428.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG	RCGGCGG	R
YP_036780.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG	RCGGCGG	R
NP_832442.1	6	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG	RCGGCGG	R
YP_001645299.1	1	MNEFQQELQALNLNAYQPGNVVYWD	QQNQYPYYI	QDDARRCGGCGG	GG	RCGGCGG
ZP_03107636.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGG		RCGGCGG
YP_002338693.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGG		RCGGCGG
ZP_00741089.1	6	MNQFQHELQTLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
YP_002446107.1	1	MNQFQHELQTLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
ZP_00240549.1	1	MNQFQHELQTLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
NP_979019.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	QQNQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
YP_002367408.1	1	MNQFQQELQSLNLNDYQTGNVYVWD	PQQSQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
YP_001375144.1	1	MNEFQQELQSLNLNDYQPGNVVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
YP_001645091.1	1	MNNFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGGCGG		RCGGCGG
ZP_03233379.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
NP_978937.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
NP_832311.1	3	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
YP_002367295.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
ZP_00744092.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
ZP_00239435.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
YP_002445982.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
YP_002338620.1	1	MNQFQQELQALSNDYQSGNIVYWD	QQNQYPYYI	QDDARRCGGCGG		RCGGCGG
NT03BL1107.1	1	MQPFHDELQSLNLDYHFGADDMTHWD	PDRHH	ANVKHCASCW	SCGSC	AS

BA_2677	60	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	88
YP_084014.1	62	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	90
ZP_03234428.1	60	CGGCGAG	RCGGCIG	CVGCFG	CFNCWNWJ	II	88
YP_036780.1	60	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	88
NP_832442.1	65	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	93
YP_001645299.1	59	CGGCGG	CGGG	RCGGCVG	CVGCFG	CSNCWNWJ	II
ZP_03107636.1	57	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	85
YP_002338693.1	57	CGGCGAG	RCGGCIG	CAGCFG	CFNCWNWJ	II	85
ZP_00741089.1	66	CGGCGG	R	CVG	CAGCFG	CFNCWNWJ	II
YP_002446107.1	61	CGGCGG	R	CVG	CAGCFG	CFNCWNWJ	VI
ZP_00240549.1	61	CGG		CVG	CAGCFG	CFNCWNWJ	II
NP_979019.1	60	CGG		CVG	CAGCFG	CFNCWNWJ	II
YP_002367408.1	60	CGG		CVG	CAGCFG	CFNCWNWJ	II
YP_001375144.1	46			CFG	CFGCFG	CFNCWNWJ	FFI
YP_001645091.1	62	CGG	GRCGGF	RFCFGCF	CFSCFG	CGGCSNCSN	CFDGFNGT
ZP_03233379.1	54	CGG	AGRCGGF	RFCFGCF	CFGCFG	CGGCGSCGS	CSNCFNGT
NP_978937.1	54	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFNGT
NP_832311.1	58	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFDGF
YP_002367295.1	56	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFDGF
ZP_00744092.1	56	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFDGF
ZP_00239435.1	55	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFDGF
YP_002445982.1	56	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGGCGG	CSNCFDGF
YP_002338620.1	49	CGG	FRCGGF	RFCFGCF	CFGCFG	CGGCGG	CSNCFNGT
NT03BL1107.1	48	C	WSCAGH	SCWSCMH	CSWSCAG	HC	CWSCWHGG