

*Supplementary Information*

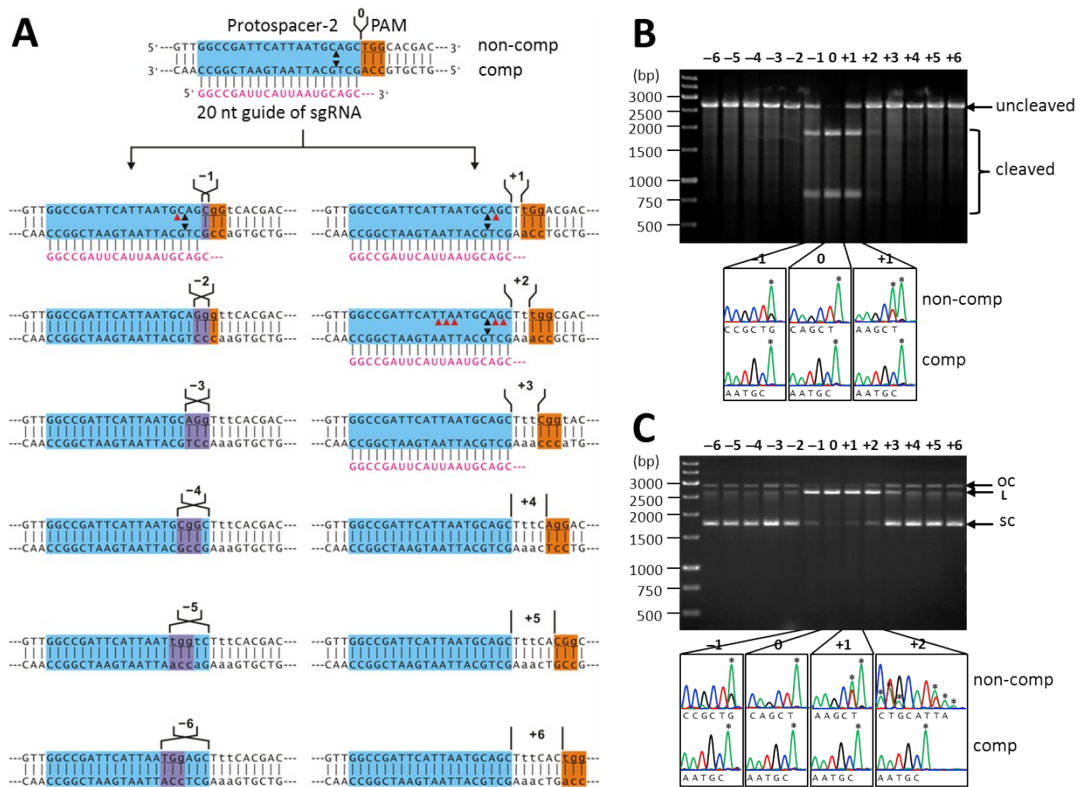
**An insight into the protospacer adjacent motif of  
*Streptococcus pyogenes* Cas9 with artificially stimulated  
RNA-guided-Cas9 DNA cleavage flexibility**

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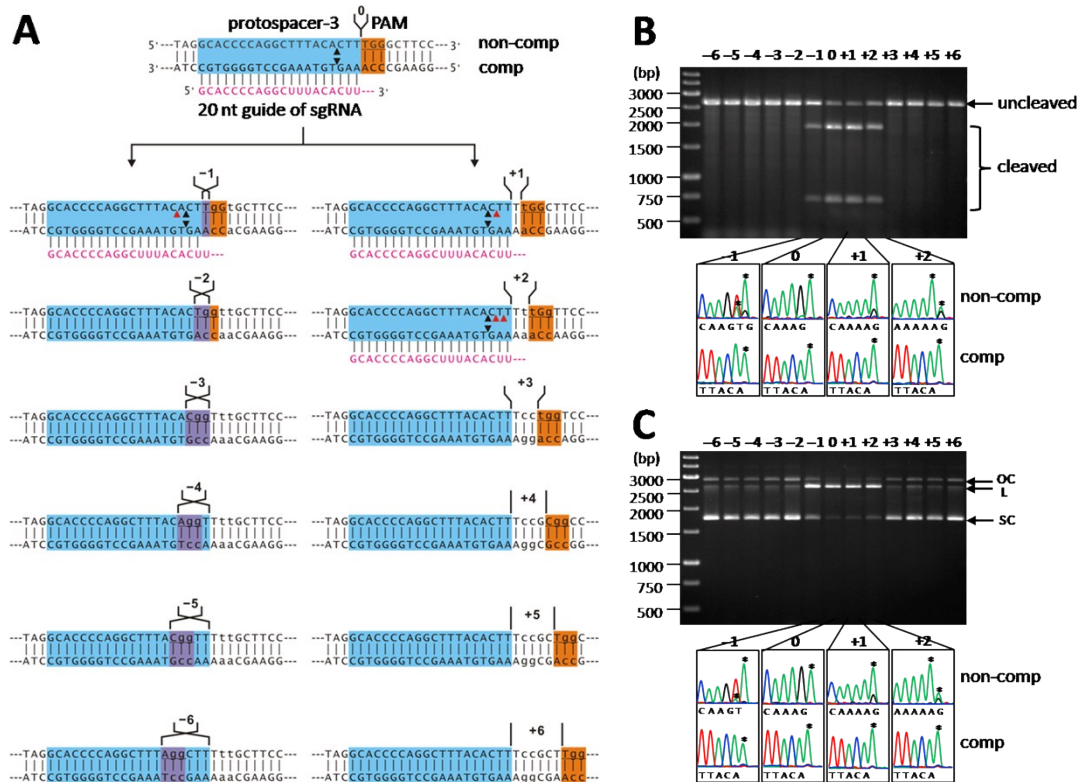
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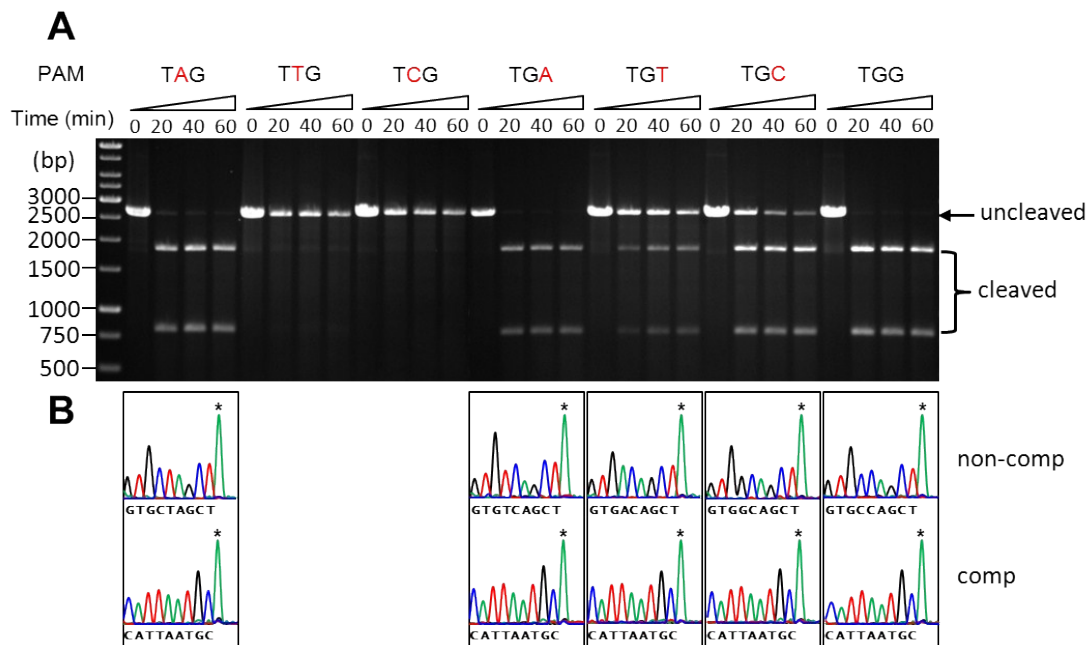
**Fig. S1. Impact of length of the interspace between protospacer-2 and NGG on SpCas9 cleavage.** (A) Schematic representation of dsDNA with interspace of variable length. Protospacer region, NGG (underlined) region and the overlapping region of NGG and protospacer are highlighted in blue, orange and purple, respectively, and sgRNA guide sequence is coloured red. The black and red triangles indicate known SpCas9 cleavage sites and newly observed cleavage sites in this study, respectively. The number with/without + and – indicate the number of base pairs of interspace or overlapping sequence between the protospacer and NGG, respectively. (B) and (C) show the cleavage results associated with dsDNA shown in (A) and end-sequencing analysis of cleaved products of *SspI*-linearized plasmid and circular plasmid, respectively. The 3' terminal A overhang, which is an artifact of the sequencing reaction, is represented by an asterisk. The following

abbreviations were used: OC, open circular DNA; L, linear DNA; SC, supercoiled DNA; comp, complementary strand; non-comp, non-complementary strand.



**Fig. S2. Impact of length of the interspace between protospacer-3 and NGG on SpCas9 cleavage.** (A) Schematic representation of dsDNA with interspace of variable length. The protospacer region, NGG (underlined) region and the overlapping region of NGG and protospacer are highlighted in blue, orange and purple, respectively, and sgRNA guide sequence is coloured red. The black and red indicate known cleavage sites of SpCas9 and cleavage sites that were observed for the first time as part of this study, respectively. The number with/without + and – indicate the number of base pairs associated with interspace or overlapping sequence between the

protospacer and NGG, respectively. (B) and (C) represent the cleavage results from the dsDNA shown in (A) and end-sequencing analysis of the cleavage products of *SspI*-linearized plasmid and circular plasmid, respectively. The 3' terminal A overhang which is an artifact of the sequencing reaction is represented by an asterisk. The following abbreviations were used: OC, open circular DNA; L, linear DNA; SC, supercoiled DNA; comp, complementary strand; non-comp, non-complementary strand.



**Fig. S3. The influence of single guanosine mutations in the NGG sequence of PAM on SpCas9-mediated cleavage using *SspI*-linearized plasmid on protospacer-2. Cleavage results (A) and end-sequencing results of cleavage products generated by treatment with SpCas9 for 60 min (B).**

**Table S1. Oligonucleotides used in this study**

Oligonucleotides (5'→3') used for <i>in vitro</i> transcription templates		
protospacer-1	N1P1	GGAATTGTGAGCGGAGAAGAGTTTTAGAGCTAGAAATAGC
	N1P3	GATCACTAATACGACTCACTATAGGAATTGTGAGCGGAGAAG A
	N1-2P1	GGAATTGTGAGCGGAGAAAGTTTTAGAGCTAGAAATAGC
	N1-2P3	GATCACTAATACGACTCACTATAGGAATTGTGAGCGGAGAAA G
	N1-3P1	GGAATTGTGAGCGGAGAAGGGTTTTAGAGCTAGAAATAGC
	N1-3P3	GATCACTAATACGACTCACTATAGGAATTGTGAGCGGAGAAG G
protospacer-2	N2P1	GGCCGATTCATTAATGCAGCGTTTTAGAGCTAGAAATAGC
	N2P3	GATCACTAATACGACTCACTATAGGCCGATTCATTAATGCAGC
	N2-2P1	GGCCGATTCATTAATGCAGGGTTTTAGAGCTAGAAATAGC
	N2-2P3	GATCACTAATACGACTCACTATAGGCCGATTCATTAATGCAGG
	N2-4P1	GGCCGATTCATTAATGCGGCGTTTTAGAGCTAGAAATAGC
	N2-4P3	GATCACTAATACGACTCACTATAGGCCGATTCATTAATGCGGC
	N2-5P1	GGCCGATTCATTAATTGGTCGTTTTAGAGCTAGAAATAGC
	N2-5P3	GATCACTAATACGACTCACTATAGGCCGATTCATTAATTGGTC
	N2-6P1	GGCCGATTCATTAATGGAGCGTTTTAGAGCTAGAAATAGC
	N2-6P3	GATCACTAATACGACTCACTATAGGCCGATTCATTAATGGAGC
protospacer-3	N3P1	GCACCCAGGCTTTACACTTGTTTTAGAGCTAGAAATAGC
	N3P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTACACTT
	N3-2P1	GCACCCAGGCTTTACACTGGTTTTAGAGCTAGAAATAGC
	N3-2P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTACACTG
	N3-3P1	GCACCCAGGCTTTACACGGTTTTAGAGCTAGAAATAGC
	N3-3P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTACACG G
	N3-4P1	GCACCCAGGCTTTACAGGTGTTTTAGAGCTAGAAATAGC
	N3-4P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTACAGG T
	N3-5P1	GCACCCAGGCTTTACGGTTGTTTTAGAGCTAGAAATAGC
	N3-5P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTACGGTT
	N3-6P1	GCACCCAGGCTTTAGGCTTGTGTTTTAGAGCTAGAAATAGC
	N3-6P3	GATCACTAATACGACTCACTATAGCACCCAGGCTTTAGGCTT
	P2-4	AGCACCGACTCGGTGCCACTTTTCCAAGTTGATAACGGACTA GCCTTATTTAACT
Oligonucleotides (5'→3') used for site-directed mutagenesis		
protospacer <sup>-1</sup>	N1TGG-S	TGTGAGCGGAGAAGATGGTCACACAGGAAACAGCTATGAC
	N1TGG-A	TCCTGTGTGACCATCTTCTCCGCTCACAATTCCACACAAC
	N1+1-S	TGTGAGCGGAGAAGATTGGCACACAGGAAACAGCTATGAC
	N1+1-A	TTCTGTGTGCCAATCTTCTCCGCTCACAATTCCACACAA

	N1+2-S	TGTGAGCGGAGAAGATTTGGACACAGGAAACAGCTATGAC
	N1+2-A	TTTCCTGTGTCCAAATCTTCTCCGCTCACAATTCCACACA
	N1+3-S	TGTGAGCGGAGAAGATTTTGGCACAGGAAACAGCTATGAC
	N1+3-A	GTTTCCTGTGCCAAAATCTTCTCCGCTCACAATTCCACAC
	N1+4-S	TGTGAGCGGAGAAGAATTTTGGACAGGAAACAGCTATGAC
	N1+4-A	TGTTTCCTGTCCAAAATCTTCTCCGCTCACAATTCCACA
protospacer-1	N1+5-S	TGTGAGCGGAGAAGAATTTTTGGCAGGAAACAGCTATGAC
	N1+5-A	CTGTTTCCTGCCAAAATCTTCTCCGCTCACAATTCCAC
	N1-1-S	TGTGAGCGGAGAAGAGGTTTACACAGGAAACAGCTATGAC
	N1-1-A	CCTGTGTGAACCTCTTCTCCGCTCACAATTCCACACAACA
	N1-2-S	TGTGAGCGGAGAAAGTTTTACACAGGAAACAGCTATGAC
	N1-2-A	CTGTGTGAAACCTTTCTCCGCTCACAATTCCACACAACAT
	N1-3-S	TGTGAGCGGAGAAGGTTTTACACAGGAAACAGCTATGAC
	N1-3-A	CTGTGTGAAAACCTTCTCCGCTCACAATTCCACACAACAT
protospacer-2	N2+1-S	TTAATGCAGCTTGGACGACAGGTTTCCCGACTGGAAAGCG
	N2+1-A	AACCTGTCGTCCAAGCTGCATTAATGAATCGGCCAACGCG
	N2+2-S	TAATGCAGCTTTGGCGACAGGTTTCCCGACTGGAAAGCGG
	N2+2-A	AAACCTGTCGCCAAAGCTGCATTAATGAATCGGCCAACGCG
	N2+3-S	TAATGCAGCTTTCCGGTACAGGTTTCCCGACTGGAAAGCGG
	N2+3-A	GGAAACCTGTACCGAAAGCTGCATTAATGAATCGGCCAAC
	N2+4-S	TAATGCAGCTTTCAGGACAGGTTTCCCGACTGGAAAGCGG
	N2+4-A	GAAACCTGTCCTGAAAGCTGCATTAATGAATCGGCCAACG
	N2+5-S	TAATGCAGCTTTCACGGCAGGTTTCCCGACTGGAAAGCGG
	N2+5-A	GGGAAACCTGCCGTGAAAGCTGCATTAATGAATCGGCCAA
	N2+6-S	TAATGCAGCTTTCAGTGGAGGTTTCCCGACTGGAAAGCGG
	N2+6-A	CGGAAACCTCCAGTAAAGCTGCATTAATGAATCGGCCA
	N2-1-S	ATTAATGCAGCGGTCACGACAGGTT
	N2-1-A	AACCTGTCGTGACCGCTGCATTAAT
	N2-2-S	CGATTCATTAATGCAGGGTTCACGACAGGTTTCCC
	N2-2-A	GGGAAACCTGTCGTGAACCCTGCATTAATGAATCG
	N2-3-S	ATTAATGCAGGTTTACGACAGGTTTCCCGACTGGAAAGC
	N2-3-A	ACCTGTCGTGAAACCTGCATTAATGAATCGGCCAACGCGC
	N2-4-S	TCATTAATGCGGCTTTCACGACAGGTTTCCCGACTGGAAA
	N2-4-A	ACCTGTCGTGAAAGCCGCATTAATGAATCGGCCAACGCGC
	N2-5-S	ATTCATTAATTGGTCTTTCACGACAGGTTTCCCGACTGGA
	N2-5-A	ACCTGTCGTGAAAGACCAATTAATGAATCGGCCAACGCGC
	N2-6-S	TTCATTAATGGAGCTTTCACGACAGGTTTCCCGACTGGAA
	N2-6-A	ACCTGTCGTGAAAGCTCCATTAATGAATCGGCCAACGCGC
	N2TAG-S	ATTAATGCAGCTAGCACGACAGGTT
	N2TAG-A	AACCTGTCGTGCTAGCTGCATTAAT
	N2TTG-S	ATTAATGCAGCTTGCACGACAGGTT
	N2TTG-A	AACCTGTCGTGCAAGCTGCATTAAT
	N2TCG-S	TAATGCAGCTCGCACGACAGGTTTCCCGACTGGAAAGCGG

	N2TCG-A	CCTGTCGTGCGAGCTGCATTAATGAATCGGCCAACGCGCG
	N2TGA-S	ATTAATGCAGCTGACACGACAGGTT
	N2TGA-A	AACCTGTCGTGTCAGCTGCATTAAT
	N2TGT-S	ATTAATGCAGCTGTCACGACAGGTT
	N2TGT-A	AACCTGTCGTGACAGCTGCATTAAT
	N2TGC-S	ATTAATGCAGCTGCCACGACAGGTT
protospacer-2	N2TGC-A	AACCTGTCGTGGCAGCTGCATTAAT
	N2TAA-S	TAATGCAGCTAACACGACAGGTTTCCCGACTGGAAAGCGG
	N2TAA-A	ACCTGTCGTGTTAGCTGCATTAATGAATCGGCCAACGCGC
	N2TAT-S	TAATGCAGCTATCACGACAGGTTTCCCGACTGGAAAGCGG
	N2TAT-A	ACCTGTCGTGATAGCTGCATTAATGAATCGGCCAACGCGC
	N2TAC-S	TAATGCAGCTACCACGACAGGTTTCCCGACTGGAAAGCGG
	N2TAC-A	ACCTGTCGTGGTAGCTGCATTAATGAATCGGCCAACGCGC
	N2TTA-S	TAATGCAGCTTACACGACAGGTTTCCCGACTGGAAAGCGG
	N2TTA-A	ACCTGTCGTGTAAGCTGCATTAATGAATCGGCCAACGCGC
	N2TTT-S	ATTAATGCAGCTTTCACGACAGGTT
	N2TTT-A	AACCTGTCGTGAAAGCTGCATTAAT
	N2TTC-S	TAATGCAGCTTCCACGACAGGTTTCCCGACTGGAAAGCGG
	N2TTC-A	ACCTGTCGTGGAAGCTGCATTAATGAATCGGCCAACGCGC
	N2TCA-S	TAATGCAGCTCACACGACAGGTTTCCCGACTGGAAAGCGG
	N2TCA-A	ACCTGTCGTGTGAGCTGCATTAATGAATCGGCCAACGCGC
	N2TCT-S	TAATGCAGCTCTCACGACAGGTTTCCCGACTGGAAAGCGG
	N2TCT-A	ACCTGTCGTGAGAGCTGCATTAATGAATCGGCCAACGCGC
	N2TCC-S	ATTAATGCAGCTCCCACGACAGGTT
	N2TCC-A	AACCTGTCGTGGGAGCTGCATTAAT
	protospacer-3	N3TGG-S
N3TGG-A		GAGCCGGAAGACCAAAGTGTAAGCCTGGGGTGCCTAATG
N3+1-S		TTTACACTTTTGGCTTCCGGCTCGTATGTTGTGTGGAATT
N3+1-A		AGCCGGAAGCCAAAAGTGTAAGCCTGGGGTGCCTAATGA
N3+2-S		TTTACACTTTTTGGTTCCGGCTCGTATGTTGTGTGGAATT
N3+2-A		CGAGCCGGAACCAAAAAGTGTAAGCCTGGGGTGCCTAAT
N3+3-S		TTTACACTTTCCTGGTCCGGCTCGTATGTTGTGTGGAATT
N3+3-A		ACGAGCCGGACCAGGAAAGTGTAAGCCTGGGGTGCCTAA
N3+4-S		TTTACACTTTCCGCGGCCGGCTCGTATGTTGTGTGGAATT
N3+4-A		TACGAGCCGGCCGCGGAAAGTGTAAGCCTGGGGTGCCTA
N3+5-S		TTTACACTTTCCGCTGGCGGCTCGTATGTTGTGTGGAATT
N3+5-A		ATACGAGCCGCCAGCGGAAAGTGTAAGCCTGGGGTGCCT
N3+6-S		TTTACACTTTCCGCTTGGGGCTCGTATGTTGTGTGGAATT
N3+6-A		CATACGAGCCCAAGCGGAAAGTGTAAGCCTGGGGTGCC
N3-1-S		CTTTACACTTGGTGCTTCCGGCTCGTATGTTGTGTGGAAT
N3-1-A		GCCGGAAGCACCAAGTGTAAGCCTGGGGTGCCTAATGAG
N3-2-S		GCTTTACACTGGTTGCTTCCGGCTCGTATGTTGTGTGGAA
N3-2-A		GCCGGAAGCAACCAGTGTAAGCCTGGGGTGCCTAATGAG

	N3-3-S	GGCTTTACACGGTTTGCTTCCGGCTCGTATGTTGTGTGGA
	N3-3-A	GCCGGAAGCAAACCGTGTAAGCCTGGGGTGCCTAATGAG
	N3-4-S	AGGCTTTACAGGTTTTGCTTCCGGCTCGTATGTTGTGTGG
	N3-4-A	GCCGGAAGCAAACCGTGTAAGCCTGGGGTGCCTAATGAG
	N3-5-S	CAGGCTTTACGGTTTTTGCTTCCGGCTCGTATGTTGTGTG
	N3-5-A	GCCGGAAGCAAAAACCGTGTAAGCCTGGGGTGCCTAATGAG
protospacer-3	N3-6-S	CCAGGCTTTAGGCTTTTTGCTTCCGGCTCGTATGTTGTGT
	N3-6-A	GCCGGAAGCAAAAAGCCTAAGCCTGGGGTGCCTAATGAG
Oligonucleotides (5' → 3') used as PAMmers at protospacer-2		
N2PL	TGGCACGACAGGTTTCCCG	
N2PTTC	TTCCACGACAGGTTTCCCG	
N2PM	AATGCAGCTGGCACGACAG	
N2PR	GATTCATTAATGCAGCTGG	
N2PE	AATGCAGCTGGCACGACAGGTTTCCCG	
N2PA	CGGGAAACCTGTCGTGGGA	