

Supplementary Information for:

**Structural basis of PAM recognition and target DNA unwinding by
the CRISPR-associated endonuclease Cas9**

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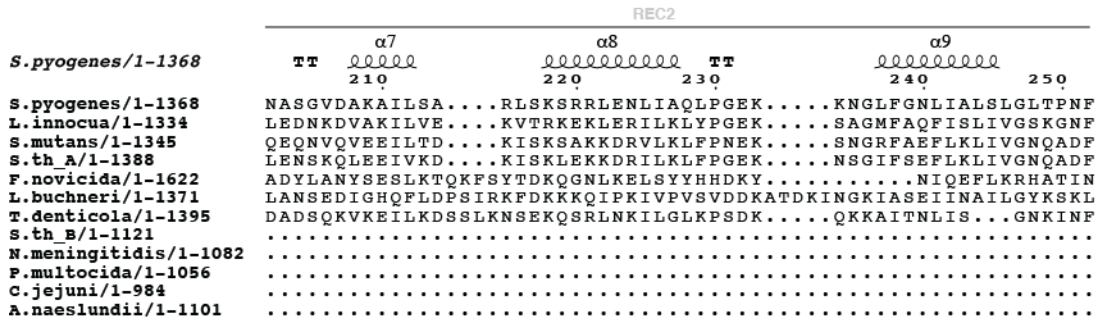
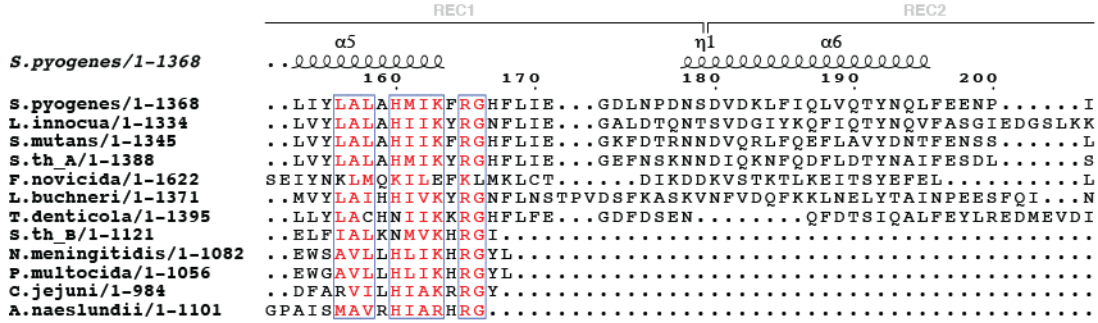
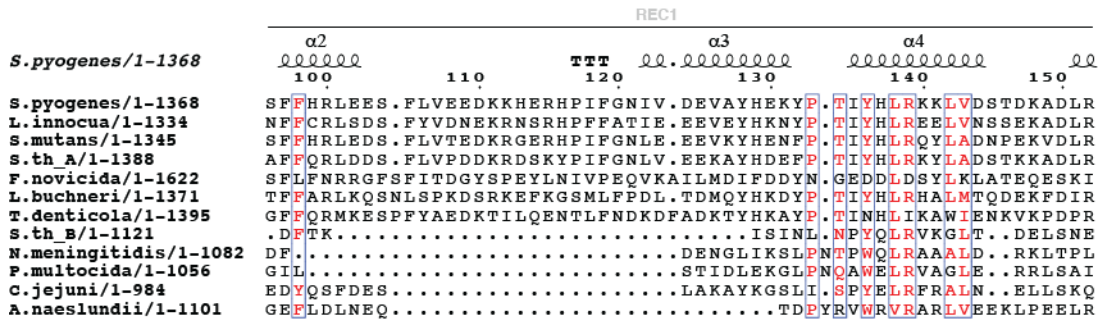
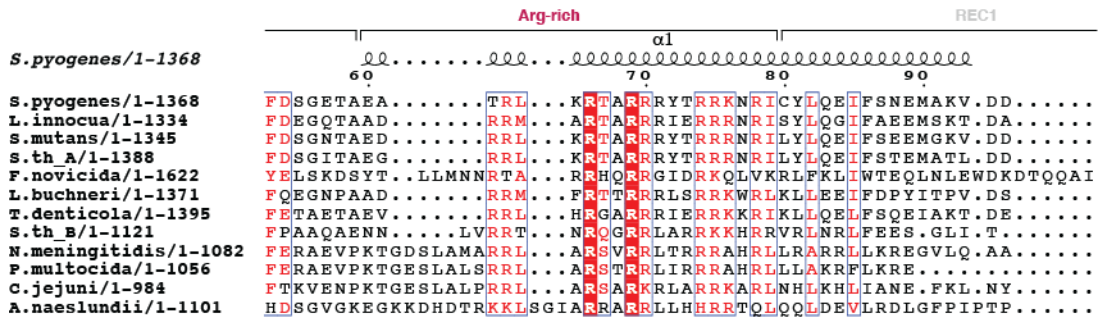
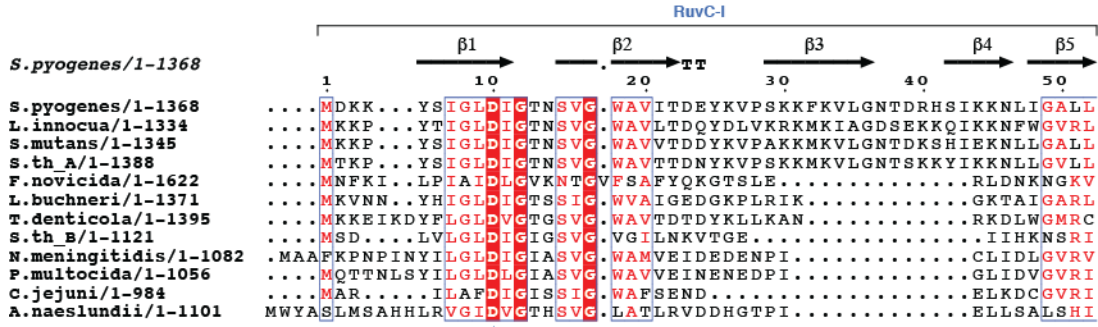
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Supplementary Figure 1 | Multiple sequence alignment of Cas9 orthologs.

Primary sequences of type II-A Cas9 proteins from *Streptococcus pyogenes* (GI 15675041), *Listeria innocua* Clip 11262 (GI 16801805), *Streptococcus mutans* UA159 (GI 24379809), *Streptococcus thermophilus* LMD-9 (S.thermophilus A, GI 11662823; S.thermophilus B, GI 116627542), *Lactobacillus buchneri* NRRL B-30929 (GI 331702228), *Treponema denticola* ATCC 35405 (GI 42525843), *Francisella novicida* U112 (GI 118497352), *Campylobacter jejuni* subsp. *Jejuni* NCTC 11168 (GI 218563121), *Pasteurella multocida* subsp. *multocida* str. Pm70 (GI 218767588), *Neisseria meningitidis* Zs491 (GI 15602992) and *Actinomyces naeslundii* (GI 489880078) were aligned using MAFFT³⁶. The alignment was generated in ESPript (<http://esprict.ibcp.fr/ESPript/>) using default settings. Strictly conserved residues are shown with white letters on red background. Residues with >70% similarity are shown in red and boxed in blue. The domain organization of SpyCas9 and secondary structure are shown above the sequences. Disordered segments of the polypeptide chain are indicated with dashed lines. RuvC domain catalytic residues are denoted with blue arrowheads. HNH domain active site residues are denoted with red arrowheads. PAM-interacting residues contacting the major groove are denoted with black arrowheads. The phosphate lock loop is marked with a red box. PIM; PAM interacting domain.



REC2

	TT	α10	η2	α11	α12
	260	270	280	290	300
<i>S. pyogenes</i> /1-1368
<i>S. pyogenes</i> /1-1368	KSNFDL	...AEDAKLQLSKDTY	DDDLNLLAQ	TGDQYAD	FLAAKNLSDAILLSDILR
<i>L. innocua</i> /1-1334	QKPFDL	...IEKSDIECAKDSY	EEDLESLLAL	TGDEYAE	LFVAAKNAYSAVVLSIIT
<i>S. mutans</i> /1-1345	KKHFEL	...EEKAPLQFSKDTY	EELLEVLLAQ	TGDNYAE	LFLSAKKLYDSILLSGFLT
<i>S. th A</i> /1-1388	RKCFNL	...DEKASLHFSKESY	DEDLETLLGY	TGDDYSV	VLKAKKLYDAILLSGFLT
<i>F. novicida</i> /1-1622	DRILD	TLLTDDLDIWNFNFEKDFD	DKNEEKLNQED	EDDKDHIA	AHLHHFVFAVNNKIKSEMA.
<i>L. buchneri</i> /1-1371	DVVVQC	TPVDSKSWALKFDEEDI	DAKLQKILPEMD	ENQQS	IIAILQNLYSQVTLNQIV.
<i>T. denticola</i> /1-1395	ADLYD	NPDLKDAEKNSISFSKDDF	DALSDDLASL	LGDSF	ELLLKAKAVYNCVLSKVI.
<i>S. th B</i> /1-1121
<i>N. meningitidis</i> /1-1082
<i>P. multocida</i> /1-1056
<i>C. jejuni</i> /1-984
<i>A. naeslundii</i> /1-1101

REC1

	α13	η3	α14	
	310	320	330	340
<i>S. pyogenes</i> /1-1368
<i>S. pyogenes</i> /1-1368	VNTEITKAPLSASM	IKRVD	DEHHQDLTL	LKALVLRQQLPEK
<i>L. innocua</i> /1-1334	VAETETNAKLSASM	IERRD	THEEDLGE	LKAFIKLHLPKH
<i>S. mutans</i> /1-1345	VTDVGTKAPLSASM	IQRV	NEHQMDLAQ	LKQFIRQLSDK
<i>S. th A</i> /1-1388	VTDNETEAPLSASM	IKRVD	NEHKEDLAL	LKKEYIRNLSLKT
<i>F. novicida</i> /1-1622	SGGRHRSQYFQ	EITNVL	DENNHQEGY	LKNFCENLHNKK
<i>L. buchneri</i> /1-1371
<i>T. denticola</i> /1-1395
<i>S. th B</i> /1-1121
<i>N. meningitidis</i> /1-1082
<i>P. multocida</i> /1-1056
<i>C. jejuni</i> /1-984
<i>A. naeslundii</i> /1-1101

REC1

	α15	α16	α17	
	370	380	390	400
<i>S. pyogenes</i> /1-1368
<i>S. pyogenes</i> /1-1368	IDGGA	SQEFYKFIKPI	L...EKMDGTEELLVKLNREDLLRKRQTFDN
<i>L. innocua</i> /1-1334	IDGKT	KQADFYKMKMTL	...ENIEGADYFIAKIEKENFLRKRQTFDN
<i>S. mutans</i> /1-1345	IDGKT	NQEFYKYLKGLL	...NKIEGSGYFLDKIEREDFLRKRQTFDN
<i>S. th A</i> /1-1388	IDGKT	NQEDFYVYLKLL	...AEFEGADYFLEKIDREDFLRKRQTFDN
<i>F. novicida</i> /1-1622
<i>L. buchneri</i> /1-1371
<i>T. denticola</i> /1-1395
<i>S. th B</i> /1-1121
<i>N. meningitidis</i> /1-1082
<i>P. multocida</i> /1-1056
<i>C. jejuni</i> /1-984
<i>A. naeslundii</i> /1-1101

REC1

	η4	α18	α19	α20
	410	420	430	440
<i>S. pyogenes</i> /1-1368
<i>S. pyogenes</i> /1-1368	GSIPH	QIHLG	ELHAILRR	QEDF.Y...PFLKD...NRE...KIEKILT
<i>L. innocua</i> /1-1334	GAIPH	QLHLE	LEAILHQ	QAKY.Y...PFLKE...NYD...KIKSLVT
<i>S. mutans</i> /1-1345	GSIPH	QIHLQ	EMRAILRR	QAEF.Y...PFLAD...NQD...RIEKLIT
<i>S. th A</i> /1-1388	GSIPY	QIHLQ	EMRAILDR	QAKF.Y...PFLAK...NKE...RIEKLIT
<i>F. novicida</i> /1-1622	AKYSY	KDLCN	ELKQKVT	KAGLVDFLLELDP CRTIP
<i>L. buchneri</i> /1-1371	GVIPH	QLHQR	ELDEIEH	QSKY.Y...PWLAEINPNKHLHLAKYKIEELVA
<i>T. denticola</i> /1-1395	AEPY	QLRKM	LEKILSNA	EKKH.F...SFLKQKDEKGLSHSE...KIIIMLLT
<i>S. th B</i> /1-1121	NVPPT	SAYRS	ELALRIL	QQTQ.F.N...PQITD...EFINRYLELIT
<i>N. meningitidis</i> /1-1082
<i>P. multocida</i> /1-1056
<i>C. jejuni</i> /1-984
<i>A. naeslundii</i> /1-1101

REC1

	α21
	450
<i>S. pyogenes</i> /1-1368
<i>S. pyogenes</i> /1-1368	FRIPY
<i>L. innocua</i> /1-1334	FRIPY
<i>S. mutans</i> /1-1345	FRIPY
<i>S. th A</i> /1-1388	FRIPY
<i>F. novicida</i> /1-1622	LDNQYPNWQ
<i>L. buchneri</i> /1-1371	FRVPY
<i>T. denticola</i> /1-1395	FKIPY
<i>S. th B</i> /1-1121	GKRKY
<i>N. meningitidis</i> /1-1082	TQRP
<i>P. multocida</i> /1-1056	WQKPA
<i>C. jejuni</i> /1-984	YKRAL
<i>A. naeslundii</i> /1-1101	FKADS

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L. buchneri/1-1371
T. denticola/1-1395
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C. jejuni/1-984
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RuvC-II

S. pyogenes/1-1368

α36 α37 β9

720 730 740 750 760

S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th_A/1-1388
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L. buchneri/1-1371
T. denticola/1-1395
S. th_B/1-1121
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A. naeslundii/1-1101

RuvC-II HNH HNH

α38 η7 η8

770 780 790 800

S. pyogenes/1-1368

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L. innocua/1-1334
S. mutans/1-1345
S. th_A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th_B/1-1121
N. meningitidis/1-1082
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A. naeslundii/1-1101

HNH

α39 β10 η9 β11

810 820 830 840 850 860

S. pyogenes/1-1368

S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th_A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th_B/1-1121
N. meningitidis/1-1082
F. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

HNH

η10 α40 α41

870 880 890

S. pyogenes/1-1368

S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th_A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th_B/1-1121
N. meningitidis/1-1082
F. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

HNH RuvC-III

η11 α42 α43 β12

900 910 920 930 940

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L. innocua/1-1334
S. mutans/1-1345
S. th_A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th_B/1-1121
N. meningitidis/1-1082
F. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

PIM (Topo) PIM (CTD)

α51 β20 η14 β21 β22 TT

.....00 → 1190 1200 → 1210

S. pyogenes/1-1368
S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th B/1-1121
N. meningitidis/1-1082
P. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

PIM (CTD)

β23 β24 α52 α53

→ 1220 → 1230 0000000000 1240 1250 00000000

S. pyogenes/1-1368
S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th B/1-1121
N. meningitidis/1-1082
P. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

PIM (CTD)

α54 α55

.....00 0000000000.....00 000.....0000000000 1300 000000

1270 1280 1290 1300

S. pyogenes/1-1368
S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th B/1-1121
N. meningitidis/1-1082
P. multocida/1-1056
C. jejuni/1-984
A. naeslundii/1-1101

PIM (CTD)

α56 η15 β25 β26 major groove η16

000.....000000 1310 1320 → 1330 → 1340 00.0

S. pyogenes/1-1368
S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th A/1-1388
F. novicida/1-1622
L. buchneri/1-1371
T. denticola/1-1395
S. th B/1-1121
N. meningitidis/1-1082
P. multocida/1-1056
C. jejuni/1-984
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PIM (CTD)

β27 β28

→ 1350 → 1360

S. pyogenes/1-1368
S. pyogenes/1-1368
L. innocua/1-1334
S. mutans/1-1345
S. th A/1-1388
F. novicida/1-1622
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