

Supplementary Data 1. Protein sequence alignment of Cas9 variants used in this study

Clone-1	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
Clone-2	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
Clone-3	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
WT-Cas9	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
eSpCas9	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
Cas9-HF1	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
evo-Cas9	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
Hypa-Cas9	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE
xCas9-3.7	1	MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKFKVLGNTDRHSIKKNLIGALLFDSGETAE

Clone-1	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
Clone-2	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
Clone-3	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
WT-Cas9	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
eSpCas9	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
Cas9-HF1	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
evo-Cas9	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
Hypa-Cas9	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG
xCas9-3.7	61	ATRLKRTARRRYTRRKNRICYLQEIFSNEMAKVDDSFHRLLEESFLVEEDKKHERHPIFG

Clone-1	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
Clone-2	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
Clone-3	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
WT-Cas9	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
eSpCas9	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
Cas9-HF1	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
evo-Cas9	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
Hypa-Cas9	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD
xCas9-3.7	121	NIVDEVAYHEKYPTIYHLRKKLV DSTDKADLR LIYLALAHMIKFRGHFLIEGDLNPDNSD

Clone-1	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
Clone-2	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
Clone-3	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
WT-Cas9	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
eSpCas9	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
Cas9-HF1	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
evo-Cas9	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
Hypa-Cas9	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN
xCas9-3.7	181	VDKLF IQLVQTYNQLFEENP INASGVDAKAIL SARLSKSRRENLI AQLPGEKKNGLFGN

Clone-1	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
Clone-2	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
Clone-3	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
WT-Cas9	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
eSpCas9	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
Cas9-HF1	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
evo-Cas9	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
Hypa-Cas9	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI
xCas9-3.7	241	LIALSLGLTPNFKSNFDLAEDAKLQLSKDTYDDDLNLLAQIGDQYADFLAAKNLSDAI

Clone-1	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
Clone-2	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
Clone-3	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
WT-Cas9	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
eSpCas9	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
Cas9-HF1	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
evo-Cas9	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
Hypa-Cas9	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA
xCas9-3.7	301	LLSDILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPEKYKEIFFDQSKNGYA

Clone-1 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
Clone-2 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
Clone-3 361 GYIDG[S]ASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
WT-Cas9 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
eSpCas9 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
Cas9-HF1 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
evo-Cas9 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
Hypa-Cas9 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNGSIPHQIHLGELH
xCas9-3.7 361 GYIDGGASQEEFYKFIKPILEKMDGTEELLVKNLNREDLLRKQRTFDNG[**I**]IPHQIHLGELH

Clone-1 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
Clone-2 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
Clone-3 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
WT-Cas9 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
eSpCas9 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
Cas9-HF1 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
evo-Cas9 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
Hypa-Cas9 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNFEE
xCas9-3.7 421 AILRRQEDFYFPLKDNREKIEKILTFRIPYYVGPLARGNSRFAMWTRKSEETITPWNF[E]K

Clone-1 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPA[S]L
Clone-2 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
Clone-3 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
WT-Cas9 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
eSpCas9 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
Cas9-HF1 481 VVDKGASAQSFIERMT[A]FDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
evo-Cas9 481 VVDKGASAQSFIER[V]TNFDKNLPNEKVLPKHSL[L]EYFTVYNELTK[E]KYVTEGMRKPAFL
Hypa-Cas9 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL
xCas9-3.7 481 VVDKGASAQSFIERMTNFDKNLPNEKVLPKHSLLYEYFTVYNELTKVKYVTEGMRKPAFL

Clone-1 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
Clone-2 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
Clone-3 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
WT-Cas9 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
eSpCas9 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
Cas9-HF1 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
evo-Cas9 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
Hypa-Cas9 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI
xCas9-3.7 541 SGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDSVEISGVEDRFNASLGTYHDLLKI

Clone-1 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
Clone-2 601 NKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
Clone-3 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
WT-Cas9 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
eSpCas9 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
Cas9-HF1 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
evo-Cas9 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
Hypa-Cas9 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG
xCas9-3.7 601 IKDKDFLDNEENEDILEDIVLTLTLFEDREMIEERLKYAHLFDDKVMKQLKRRRYTGWG

Clone-1 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNFQMQLIHDDSLTFKEDIQKAQVSGQGDSL
Clone-2 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNFQMQLIHDDSLTFKEDIQKAQVSGQGDSL
Clone-3 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNFQMQLIHDDSLTFKEDIQKAQVSGQGDSL
WT-Cas9 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNFQMQLIHDDSLTFKEDIQKAQVSGQGDSL
eSpCas9 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNFQMQLIHDDSLTFKEDIQKAQVSGQGDSL
Cas9-HF1 661 A[LSRKLINGIRDKQSGKTILDFLKSDFGANRNFMA]LIHDDSLTFKEDIQKAQVSGQGDSL
evo-Cas9 661 Q[LSRKLINGIRDKQSGKTILDFLKSDFGANRNF]MQLIHDDSLTFKEDIQKAQVSGQGDSL
Hypa-Cas9 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANR[A]FAA[L]IADD[SLTFKEDIQKAQVSGQGDSL
xCas9-3.7 661 RLSRKLINGIRDKQSGKTILDFLKSDFGANRNF[**I**]QLIHDDSLTFKEDIQKAQVSGQGDSL

Clone-1 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
Clone-2 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
Clone-3 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
WT-Cas9 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
eSpCas9 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
Cas9-HF1 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
evo-Cas9 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
Hypa-Cas9 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER
xCas9-3.7 721 HEHIANLAGSPAIAKKGILQTVKVVDELVKVMGRHKPENIVIEIARENQTTQKGQKNSRER

Clone-1 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
Clone-2 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
Clone-3 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
WT-Cas9 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
eSpCas9 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
Cas9-HF1 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
evo-Cas9 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
Hypa-Cas9 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH
xCas9-3.7 781 MKRIEEGIKELGSQILKEHPVENTQLQNEKLYLYYLQNGRDMYVDQELDINRLSDYDVDH

Clone-1 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
Clone-2 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
Clone-3 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
WT-Cas9 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
eSpCas9 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
Cas9-HF1 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
evo-Cas9 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
Hypa-Cas9 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL
xCas9-3.7 841 IVPQSFLKDDSIDNKVLRSDKNRGKSDNVPSEEVVKKMKNYWRQLLNAKLITQRKFDNL

Clone-1 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
Clone-2 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
Clone-3 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
WT-Cas9 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
eSpCas9 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
Cas9-HF1 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
evo-Cas9 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
Hypa-Cas9 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS
xCas9-3.7 901 TKAERGGLSELKAGFIKRQLVETRQITKHVAQILDSRMNTKYDENDKLIREVKVITLKS

Clone-1 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
Clone-2 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
Clone-3 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
WT-Cas9 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
eSpCas9 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
Cas9-HF1 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
evo-Cas9 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
Hypa-Cas9 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK
xCas9-3.7 961 KLVSDFRKDFQFYKVVREINNYHHAHDAYLNAVVGTAIIKKYPKLESEFVYGDYKVDVRK

Clone-1 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
Clone-2 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
Clone-3 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
WT-Cas9 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
eSpCas9 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
Cas9-HF1 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
evo-Cas9 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
Hypa-Cas9 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF
xCas9-3.7 1021 MIAKSEQEIGKATAKYFFYSNIMNFFKTEITLANGEIRKRPLIETNGETGEIVWDKGRDF

Clone-1 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
Clone-2 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
Clone-3 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
WT-Cas9 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
eSpCas9 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
Cas9-HF1 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
evo-Cas9 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
Hypa-Cas9 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA
xCas9-3.7 1081 ATVRKVL SMPQVNIVKKTEVQTGGFSKESILPKRNSDKLIARKKDWDPKKYGGFDSPTVA

Clone-1 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
Clone-2 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
Clone-3 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
WT-Cas9 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
eSpCas9 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
Cas9-HF1 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
evo-Cas9 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
Hypa-Cas9 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK
xCas9-3.7 1141 YSVLVVAKVEKGKSKKLKSVKELLGITIMERSSSFENPIDFLEAKGYKEVKKDLIIKLPK

Clone-1 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
Clone-2 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
Clone-3 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
WT-Cas9 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
eSpCas9 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
Cas9-HF1 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
evo-Cas9 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
Hypa-Cas9 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE
xCas9-3.7 1201 YSLFELENGRKRMLASAGELQKGNELALPSKYVNFLYLASHYEKLKGS PEDNEQKQLFVE

Clone-1 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
Clone-2 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
Clone-3 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
WT-Cas9 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
eSpCas9 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
Cas9-HF1 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
evo-Cas9 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
Hypa-Cas9 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA
xCas9-3.7 1261 QHKHYLDEIIIEQISEFSKRVILADANLDKVL SAYNKHRDKPIREQAENIIHLFTLTNLGA

Clone-1 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
Clone-2 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
Clone-3 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
WT-Cas9 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
eSpCas9 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
Cas9-HF1 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
evo-Cas9 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
Hypa-Cas9 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK
xCas9-3.7 1321 PAAFKYFDTTIDRKRYTSTKEVLDATLIHQSI TGLYETRIDLSQLGGDGGSGPPKKRKRK

Clone-1 1381 YPYDVPDYA
Clone-2 1381 YPYDVPDYA
Clone-3 1381 YPYDVPDYA
WT-Cas9 1381 YPYDVPDYA
eSpCas9 1381 YPYDVPDYA
Cas9-HF1 1381 YPYDVPDYA
evo-Cas9 1381 YPYDVPDYA
Hypa-Cas9 1381 YPYDVPDYA
xCas9-3.7 1381 YPYDVPDYA

DNA sequence alignment of Cas9 variants used in this study

Clone-1	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
Clone-2	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
Clone-3	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
WT-Cas9	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
eSpCas9	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
Cas9-HF1	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
evo-Cas9	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
Hypa-Cas9	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG
xCas9-3.7	1	ATGGACAAGAAGTACAGCATCGGCCTGGACATCGGTACCAACAGCGTGGGCTGGGCCGTG

Clone-1	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
Clone-2	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
Clone-3	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
WT-Cas9	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
eSpCas9	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
Cas9-HF1	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
evo-Cas9	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
Hypa-Cas9	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC
xCas9-3.7	61	ATCACCGACGAGTACAAGGTGCCAGCAAGAAGTTCAAGGTGCTGGGCAACACCGACCGC

Clone-1	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
Clone-2	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
Clone-3	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
WT-Cas9	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
eSpCas9	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
Cas9-HF1	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
evo-Cas9	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
Hypa-Cas9	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG
xCas9-3.7	121	CACAGCATCAAGAAGAACCTGATCGGCGCCCTGCTGTTTCGACAGCGGCGAGACCGCCGAG

Clone-1	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
Clone-2	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
Clone-3	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
WT-Cas9	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
eSpCas9	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
Cas9-HF1	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
evo-Cas9	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
Hypa-Cas9	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC
xCas9-3.7	181	GCCACCCGCCTGAAGCGCACCGCCCGCCGCGCTACACCCGCCGCAAGAACCGCATCTGC

Clone-1	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
Clone-2	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
Clone-3	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
WT-Cas9	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
eSpCas9	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
Cas9-HF1	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
evo-Cas9	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
Hypa-Cas9	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC
xCas9-3.7	241	TACCTGCAGGAGATCTTCAGCAACGAGATGGCCAAGGTGGACGACAGCTTCTTCCACCGC

Clone-1	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
Clone-2	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
Clone-3	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
WT-Cas9	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
eSpCas9	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
Cas9-HF1	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
evo-Cas9	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
Hypa-Cas9	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC
xCas9-3.7	301	CTGGAGGAGAGCTTCTTGGTGGAGGAGGACAAGAAGCAGGAGCGCCACCCCATCTTCGGC

Clone-1 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
Clone-2 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
Clone-3 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
WT-Cas9 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
eSpCas9 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
Cas9-HF1 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
evo-Cas9 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
Hypa-Cas9 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG
xCas9-3.7 361 AACATCGTGGACGAGGTGGCCTACCACGAGAAGTACCCCACCATCTACCACCTGCGCAAG

Clone-1 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
Clone-2 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
Clone-3 421 AAGCTGGTGGACAGCACCTGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
WT-Cas9 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
eSpCas9 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
Cas9-HF1 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
evo-Cas9 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
Hypa-Cas9 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC
xCas9-3.7 421 AAGCTGGTGGACAGCACCGACAAGGCCGACCTGCGCCTGATCTACCTGGCCCTGGCCAC

Clone-1 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
Clone-2 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
Clone-3 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
WT-Cas9 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
eSpCas9 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
Cas9-HF1 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
evo-Cas9 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
Hypa-Cas9 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC
xCas9-3.7 481 ATGATCAAGTTCCGCGGCCACTTCCTGATCGAGGGCGACCTGAACCCCGACAACAGCGAC

Clone-1 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
Clone-2 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
Clone-3 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
WT-Cas9 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
eSpCas9 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
Cas9-HF1 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
evo-Cas9 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
Hypa-Cas9 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC
xCas9-3.7 541 GTGGACAAGCTGTTTCATCCAGCTGGTGCAGACCTACAACCAGCTGTTTCGAGGAGAACCCC

Clone-1 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
Clone-2 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
Clone-3 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
WT-Cas9 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
eSpCas9 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
Cas9-HF1 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
evo-Cas9 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
Hypa-Cas9 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC
xCas9-3.7 601 ATCAACGCCAGCGGCGTGGACGCCAAGGCCATCCTGAGCGCCCGCCTGAGCAAGAGCCGC

Clone-1 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
Clone-2 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
Clone-3 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
WT-Cas9 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
eSpCas9 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
Cas9-HF1 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
evo-Cas9 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
Hypa-Cas9 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC
xCas9-3.7 661 CGCCTGGAGAACCTGATCGCCCAGCTGCCC GGCGAGAAGAAGAACGGCCTGTTTCGGCAAC

Clone-1 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
Clone-2 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
Clone-3 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
WT-Cas9 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
eSpCas9 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
Cas9-HF1 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
evo-Cas9 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
Hypa-Cas9 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG
xCas9-3.7 721 CTGATCGCCCTGAGCCTGGGCCTGACCCCCAACTTCAAGAGCAACTTCGACCTGGCCGAG

Clone-1 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
Clone-2 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
Clone-3 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
WT-Cas9 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
eSpCas9 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
Cas9-HF1 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
evo-Cas9 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
Hypa-Cas9 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC
xCas9-3.7 781 GACGCCAAGCTGCAGCTGAGCAAGGACACCTACGACGACGACCTGGACAACCTGCTGGCC

Clone-1 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
Clone-2 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
Clone-3 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
WT-Cas9 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
eSpCas9 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
Cas9-HF1 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
evo-Cas9 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
Hypa-Cas9 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC
xCas9-3.7 841 CAGATCGGCGACCAGTACGCCGACCTGTTCTTGGCCGCAAGAACCTGAGCGACGCCATC

Clone-1 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
Clone-2 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
Clone-3 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
WT-Cas9 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
eSpCas9 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
Cas9-HF1 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
evo-Cas9 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
Hypa-Cas9 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC
xCas9-3.7 901 CTGCTGAGCGACATCCTGCGCGTGAACACCGAGATCACCAAGGCCCCCTGAGCGCCAGC

Clone-1 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
Clone-2 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
Clone-3 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
WT-Cas9 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
eSpCas9 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
Cas9-HF1 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
evo-Cas9 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
Hypa-Cas9 961 ATGATCAAGCGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC
xCas9-3.7 961 ATGATCAAGCTGCTACGACGAGCACCACCAGGACCTGACCCTGCTGAAGGCCCTGGTGCGC

Clone-1 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
Clone-2 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
Clone-3 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
WT-Cas9 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
eSpCas9 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
Cas9-HF1 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
evo-Cas9 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
Hypa-Cas9 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC
xCas9-3.7 1021 CAGCAGCTGCCCCGAGAAGTACAAGGAGATCTTCTTCGACCAGAGCAAGAACGGCTACGCC

Clone-1 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
Clone-2 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
Clone-3 1081 GGCTACATCGACGGCAAGCGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
WT-Cas9 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
eSpCas9 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
Cas9-HF1 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
evo-Cas9 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
Hypa-Cas9 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG
xCas9-3.7 1081 GGCTACATCGACGGCGGCCAGCCAGGAGGAGTTCTACAAGTTCATCAAGCCCATCCTG

Clone-1 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
Clone-2 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
Clone-3 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
WT-Cas9 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
eSpCas9 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
Cas9-HF1 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
evo-Cas9 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
Hypa-Cas9 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC
xCas9-3.7 1141 GAGAAGATGGACGGCACCAGGAGGAGCTGCTGGTGAAGCTGAACCGCGAGGACCTGCTGCGC

Clone-1 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
Clone-2 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
Clone-3 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
WT-Cas9 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
eSpCas9 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
Cas9-HF1 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
evo-Cas9 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
Hypa-Cas9 1201 AAGCAGCGCACCTTCGACAACGGCAGCATCCCCACCAGATCCACCTGGGCGAGCTGCAC
xCas9-3.7 1201 AAGCAGCGCACCTTCGACAACGGCATTCATCCCCACCAGATCCACCTGGGCGAGCTGCAC

Clone-1 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
Clone-2 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
Clone-3 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
WT-Cas9 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
eSpCas9 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
Cas9-HF1 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
evo-Cas9 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
Hypa-Cas9 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC
xCas9-3.7 1261 GCCATCCTGCGCCGCCAGGAGGACTTCTACCCCTTCTGAAGGACAACCGCGAGAAGATC

Clone-1 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
Clone-2 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
Clone-3 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
WT-Cas9 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
eSpCas9 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
Cas9-HF1 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
evo-Cas9 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
Hypa-Cas9 1321 GAGAAGATCCTGACCTTCCGCATCCCCTACTACGTGGGCCCCCTGGCCCGCGGCAACAGC
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Clone-1 1381 CGCTTCGCCTGGATGACCCGCAAGAGCGAGGAGACCATCACCCCTGGAACCTTCGAGGAG
Clone-2 1381 CGCTTCGCCTGGATGACCCGCAAGAGCGAGGAGACCATCACCCCTGGAACCTTCGAGGAG
Clone-3 1381 CGCTTCGCCTGGATGACCCGCAAGAGCGAGGAGACCATCACCCCTGGAACCTTCGAGGAG
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eSpCas9 1381 CGCTTCGCCTGGATGACCCGCAAGAGCGAGGAGACCATCACCCCTGGAACCTTCGAGGAG
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Clone-1 1441 GTGGTGGACAAGGGCGCCAGCGCCCAGAGCTTCATCGAGCGCATGACCAACTTCGACAAG
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Clone-3 1441 GTGGTGGACAAGGGCGCCAGCGCCCAGAGCTTCATCGAGCGCATGACCAACTTCGACAAG
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eSpCas9 1441 GTGGTGGACAAGGGCGCCAGCGCCCAGAGCTTCATCGAGCGCATGACCAACTTCGACAAG
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evo-Cas9 1441 GTGGTGGACAAGGGCGCCAGCGCCCAGAGCTTCATCGAGCGCATGACCAACTTCGACAAG
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xCas9-3.7 1441 GTGGTGGACAAGGGCGCCAGCGCCCAGAGCTTCATCGAGCGCATGACCAACTTCGACAAG

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Clone-3 1501 AACCTGCCCCAACGAGAAGGTGCTGCCCAAGCACAGCCTGCTGTACGAGTACTTCACCGTG
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Clone-3 1561 TACAACGAGCTGACCAAGGTGAAGTACGTGACCGAGGGCATGCGCAAGCCCGCCTTCCTG
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Clone-3 1621 AGCGGCGAGCAGAAGAAGGCCATCGTGGACCTGCTGTTCAAGACCAACCGCAAGGTGACC
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evo-Cas9 1621 AGCGGCGAGCAGAAGAAGGCCATCGTGGACCTGCTGTTCAAGACCAACCGCAAGGTGACC
Hypa-Cas9 1621 AGCGGCGAGCAGAAGAAGGCCATCGTGGACCTGCTGTTCAAGACCAACCGCAAGGTGACC
xCas9-3.7 1621 AGCGGCGAGCAGAAGAAGGCCATCGTGGACCTGCTGTTCAAGACCAACCGCAAGGTGACC

Clone-1 1681 GTGAAGCAGCTGAAGGAGGACTACTTCAAGAAGATCGAGTGCTTCGACAGCGTGGAGATC
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Clone-3 1681 GTGAAGCAGCTGAAGGAGGACTACTTCAAGAAGATCGAGTGCTTCGACAGCGTGGAGATC
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eSpCas9 1681 GTGAAGCAGCTGAAGGAGGACTACTTCAAGAAGATCGAGTGCTTCGACAGCGTGGAGATC
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evo-Cas9 1681 GTGAAGCAGCTGAAGGAGGACTACTTCAAGAAGATCGAGTGCTTCGACAGCGTGGAGATC
Hypa-Cas9 1681 GTGAAGCAGCTGAAGGAGGACTACTTCAAGAAGATCGAGTGCTTCGACAGCGTGGAGATC
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Clone-3 1741 AGCGGCGTGGAGGACCGCTTCAACGCCAGCCTGGGCACCTACCACGACCTGCTGAAGATC
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eSpCas9 1741 AGCGGCGTGGAGGACCGCTTCAACGCCAGCCTGGGCACCTACCACGACCTGCTGAAGATC
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evo-Cas9 1741 AGCGGCGTGGAGGACCGCTTCAACGCCAGCCTGGGCACCTACCACGACCTGCTGAAGATC
Hypa-Cas9 1741 AGCGGCGTGGAGGACCGCTTCAACGCCAGCCTGGGCACCTACCACGACCTGCTGAAGATC
xCas9-3.7 1741 AGCGGCGTGGAGGACCGCTTCAACGCCAGCCTGGGCACCTACCACGACCTGCTGAAGATC

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Clone-3 1861 TTGACCCTGACCCTGTTTCGAGGACCGCGAGATGATCGAGGAGCGCCTGAAGACCTACGCC
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evo-Cas9 1861 CTGACCCTGACCCTGTTTCGAGGACCGCGAGATGATCGAGGAGCGCCTGAAGACCTACGCC
Hypa-Cas9 1861 CTGACCCTGACCCTGTTTCGAGGACCGCGAGATGATCGAGGAGCGCCTGAAGACCTACGCC
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Clone-1 1921 CACCTGTTTCGACGACAAGGTGATGAAGCAGCTGAAGCGCCGCCGCTACACCGGCTGGGGC
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Clone-3 1921 CACCTGTTTCGACGACAAGGTGATGAAGCAGCTGAAGCGCCGCCGCTACACCGGCTGGGGC
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evo-Cas9 1921 CACCTGTTTCGACGACAAGGTGATGAAGCAGCTGAAGCGCCGCCGCTACACCGGCTGGGGC
Hypa-Cas9 1921 CACCTGTTTCGACGACAAGGTGATGAAGCAGCTGAAGCGCCGCCGCTACACCGGCTGGGGC
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Clone-3 1981 CGCCTGAGCCGCAAGCTTATCAACGGCATCCGCGACAAGCAGAGCGGCAAGACCATCCTG
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evo-Cas9 1981 CAGCTGAGCCGCAAGCTTATCAACGGCATCCGCGACAAGCAGAGCGGCAAGACCATCCTG
Hypa-Cas9 1981 CGCCTGAGCCGCAAGCTTATCAACGGCATCCGCGACAAGCAGAGCGGCAAGACCATCCTG
xCas9-3.7 1981 CGCCTGAGCCGCAAGCTTATCAACGGCATCCGCGACAAGCAGAGCGGCAAGACCATCCTG

Clone-1 2041 GACTTCCTGAAGAGCGACGGCTTCGCCAACCGCAACTTCATGCAGCTGATCCACGACGAC
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Clone-3 2041 GACTTCCTGAAGAGCGACGGCTTCGCCAACCGCAACTTCATGCAGCTGATCCACGACGAC
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eSpCas9 2041 GACTTCCTGAAGAGCGACGGCTTCGCCAACCGCAACTTCATGCAGCTGATCCACGACGAC
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evo-Cas9 2041 GACTTCCTGAAGAGCGACGGCTTCGCCAACCGCAACTTCATGCAGCTGATCCACGACGAC
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xCas9-3.7 2041 GACTTCCTGAAGAGCGACGGCTTCGCCAACCGCAACTTCATCCAGCTGATCCACGACGAC

Clone-1 2101 AGCCTGACCTTCAAGGAGGACATCCAGAAGGCCAGGTGAGCGGCCAGGGCGACAGCCTG
Clone-2 2101 AGCCTGACCTTCAAGGAGGACATCCAGAAGGCCAGGTGAGCGGCCAGGGCGACAGCCTG
Clone-3 2101 AGCCTGACCTTCAAGGAGGACATCCAGAAGGCCAGGTGAGCGGCCAGGGCGACAGCCTG
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Clone-3 2401 GTGGAGAACACCCAGCTGCAGAACGAGAAGCTGTACCTGTACTACCTGCAGAACGGCCGC
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Clone-3 2461 GACATGTACGTGGACCAGGAGCTGGACATCAACCGCCTGAGCGACTACGACGTGGACCAC
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Hypa-Cas9 2461 GACATGTACGTGGACCAGGAGCTGGACATCAACCGCCTGAGCGACTACGACGTGGACCAC
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eSpCas9 2521 ATCGTGCCCCAGAGCTTCTG**GC**GGACGACAGCATCGACAACAAGGTGCTGACCCGCAGC
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eSpCas9 2581 GACAAGAACCGCGGCAAGAGCGACAACGTGCCAGCGAGGAGGTGGTGAAGAAGATGAAG
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Clone-3 2641 AACTACTGGCGCCAGCTGCTGAACGCCAAGCTGATCACCCAGCGCAAGTTCGACAACCTG
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eSpCas9 2641 AACTACTGGCGCCAGCTGCTGAACGCCAAGCTGATCACCCAGCGCAAGTTCGACAACCTG
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Hypa-Cas9 2641 AACTACTGGCGCCAGCTGCTGAACGCCAAGCTGATCACCCAGCGCAAGTTCGACAACCTG
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Clone-3 2701 ACCAAGGCCGAGCGCGGCGGCCTGAGCGAGCTGGACAAGGCCGGCTTCATCAAGCGCCAG
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Cas9-HF1 2701 ACCAAGGCCGAGCGCGGCGGCCTGAGCGAGCTGGACAAGGCCGGCTTCATCAAGCGCCAG
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eSpCas9 2761 CTGGTGGAGACCCGCCAGATCACCAAGCACGTGGCCCAGATCCTGGACAGCCGCATGAAC
Cas9-HF1 2761 CTGGTGGAGACCCGC**GC**GATCACCAAGCACGTGGCCCAGATCCTGGACAGCCGCATGAAC
evo-Cas9 2761 CTGGTGGAGACCCGCCAGATCACCAAGCACGTGGCCCAGATCCTGGACAGCCGCATGAAC
Hypa-Cas9 2761 CTGGTGGAGACCCGCCAGATCACCAAGCACGTGGCCCAGATCCTGGACAGCCGCATGAAC
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Clone-3 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC
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eSpCas9 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC
Cas9-HF1 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC
evo-Cas9 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC
Hypa-Cas9 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC
xCas9-3.7 2821 ACCAAGTACGACGAGAACGACAAGCTGATCCGCGAGGTGAAGGTGATCACCCCTGAAGAGC

Clone-1 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
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Clone-3 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
WT-Cas9 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
eSpCas9 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
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evo-Cas9 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
Hypa-Cas9 2881 AAGCTGGTGAGCGACTTCCGCAAGGACTTCCAGTTCTACAAGGTGCGCGAGATCAACAAC
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Clone-1 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG
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WT-Cas9 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG
eSpCas9 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG
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evo-Cas9 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG
Hypa-Cas9 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG
xCas9-3.7 2941 TACCACCACGCCCACGACGCCTACCTGAACGCCGTGGTGGGCACCGCCCTGATCAAGAAG

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Clone-3 3001 TACCCCAAGCTGGAGAGCGAGTTCGTGTACGGCGACTACAAGGTGTACGACGTGCGCAAG
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Hypa-Cas9 3001 TACCCCAAGCTGGAGAGCGAGTTCGTGTACGGCGACTACAAGGTGTACGACGTGCGCAAG
xCas9-3.7 3001 TACCCCAAGCTGGAGAGCGAGTTCGTGTACGGCGACTACAAGGTGTACGACGTGCGCAAG

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Clone-3 3061 ATGATCGCCAAGAGCGAGCAGGAGATCGGCAAGGCCACCGCCAAGTACTTCTTCTACAGC
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evo-Cas9 3061 ATGATCGCCAAGAGCGAGCAGGAGATCGGCAAGGCCACCGCCAAGTACTTCTTCTACAGC
Hypa-Cas9 3061 ATGATCGCCAAGAGCGAGCAGGAGATCGGCAAGGCCACCGCCAAGTACTTCTTCTACAGC
xCas9-3.7 3061 ATGATCGCCAAGAGCGAGCAGGAGATCGGCAAGGCCACCGCCAAGTACTTCTTCTACAGC

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Hypa-Cas9 3121 AACATCATGAACTTCTTCAAGACCGAGATCACCCCTGGCCAACGGCGAGATCCGCAAGCGC
xCas9-3.7 3121 AACATCATGAACTTCTTCAAGACCGAGATCACCCCTGGCCAACGGCGAGATCCGCAAGCGC

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Hypa-Cas9 3181 CCCCTGATCGAGACCAACGGCGAGACCGGCGAGATCGTGTGGGACAAGGGCCGCGACTTC
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eSpCas9 3241 GCCACCGTGC GCAAGGTGCTGAGCATGCCCCAGGTGAACATCGTGAAGAAGACCGAGGTG
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Hypa-Cas9 3661 CAGAAGGGCAACGAGCTGGCCCTGCCAGCAAGTACGTGAACTTCCTGTACCTGGCCAGC
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evo-Cas9 3721 CACTACGAGAAGCTGAAGGGCAGCCCCGAGGACAACGAGCAGAAGCAGCTGTTTCGTGGAG
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Hypa-Cas9 4021 GAGGTGCTGGACGCCACCCTGATCCACCAGAGCATCACCGGTCTGTACGAGACCCGCATC
xCas9-3.7 4021 GAGGTGCTGGACGCCACCCTGATCCACCAGAGCATCACCGGTCTGTACGAGACCCGCATC

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Clone-3 4141 TACCCCTACGACGTGCCCGACTACGCC
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eSpCas9 4141 TACCCCTACGACGTGCCCGACTACGCC
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Hypa-Cas9 4141 TACCCCTACGACGTGCCCGACTACGCC
xCas9-3.7 4141 TACCCCTACGACGTGCCCGACTACGCC