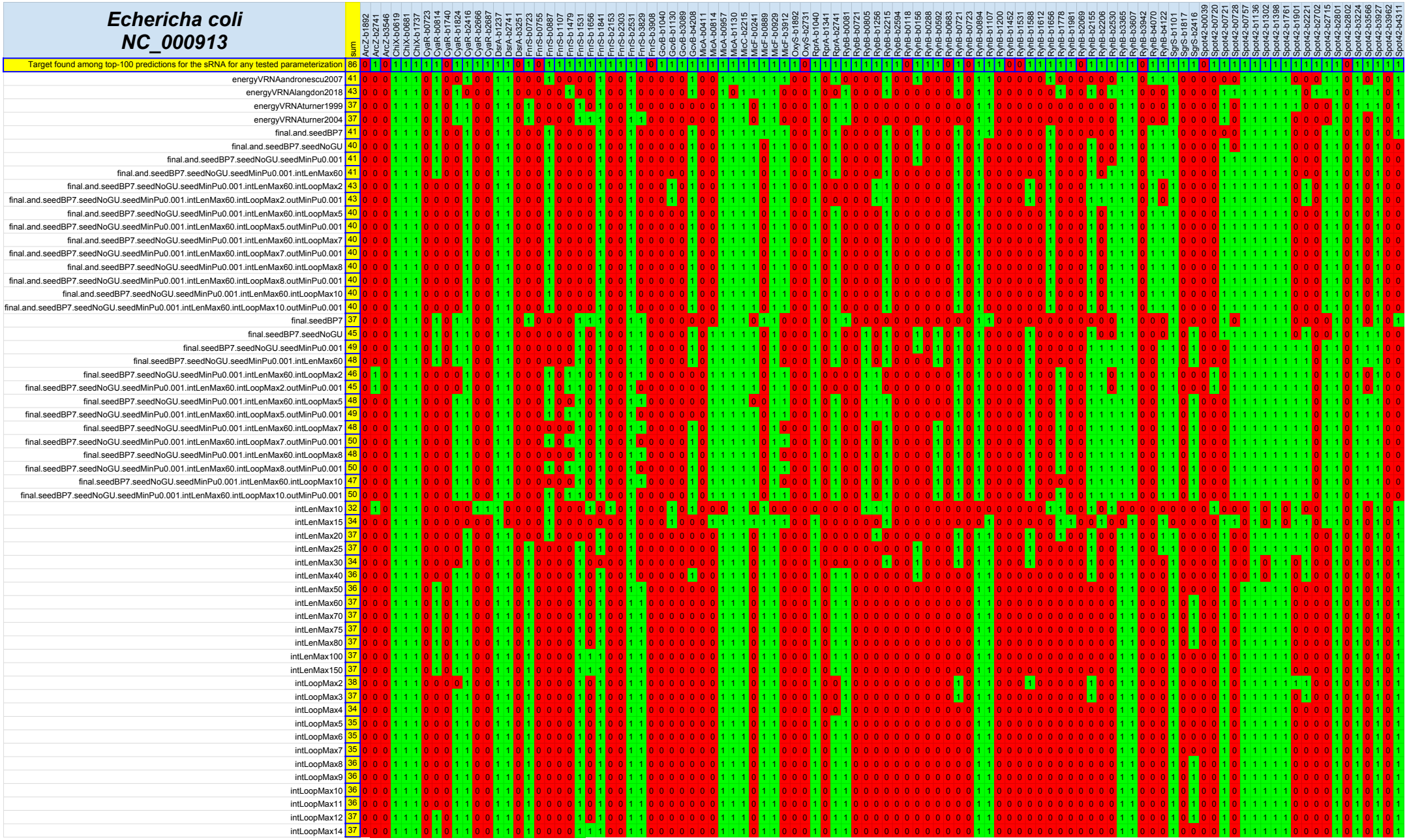
Verified sRNA-target pairs from literature

sRNA	geneID	geneName	genome	referenceDOI
ArcZ	b1892	flhD	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.08209.x
ArcZ	b2741	rpoS	NC_000913	https://doi.org/10.1038/emboj.2010.179
ArcZ	b3546	eptB	NC_000913	https://doi.org/10.1111/mmi.12257
ChiX	b0619	dpiB/citA	NC_000913	https://doi.org/10.1111/j.1365-2958.2009.06665.x
ChiX	b0681	chiP	NC_000913	https://doi.org/10.1099/mic.0.023598-0
ChiX	b1737	chbC	NC_000913	https://doi.org/10.1111/j.1365-2958.2009.06807.x
CyaR	b0723	sdhA	NC_000913	https://doi.org/10.1073/pnas.1303248110
CyaR	b0814	ompX	NC_000913	https://doi.org/10.1128/JB.01157-08
CyaR	b1740	nadE	NC_000913	https://doi.org/10.1128/JB.01157-08
CyaR	b1824	yobF	NC_000913	https://doi.org/10.1073/pnas.1303248110
CyaR	b2416	ptsI	NC_000913	https://doi.org/10.1073/pnas.1303248110
CyaR	b2666	yqaE	NC_000913	https://doi.org/10.1128/JB.01157-08
CyaR	b2687	luxS	NC_000913	https://doi.org/10.1128/JB.01157-08
DsrA	b1237	hns	NC_000913	https://doi.org/10.1073/pnas.95.21.12456
DsrA	b2741	rpoS	NC_000913	https://doi.org/10.1073/pnas.95.21.12456
DsrA	b3251	mreB	NC_000913	https://doi.org/10.1039/c4ib00102h
FnrS	b0723	sdhA	NC_000913	https://doi.org/10.1073/pnas.1303248110
FnrS	b0755	gpmA	NC_000913	https://doi.org/10.1111/j.1365-2958.2010.07044.x
FnrS	b0887	cydD	NC_000913	https://doi.org/10.1074/jbc.M109.089755
FnrS	b1107	nagZ	NC_000913	https://doi.org/10.1073/pnas.1303248110
FnrS	b1479	maeA	NC_000913	https://doi.org/10.1111/j.1365-2958.2010.07044.x
FnrS	b1531	marA	NC_000913	https://doi.org/10.1073/pnas.1303248110
FnrS	b1656	sodB	NC_000913	https://doi.org/10.1074/jbc.M109.089755
FnrS	b1841	yobA	NC_000913	https://doi.org/10.1074/jbc.M109.089755
FnrS	b2153	folE	NC_000913	https://doi.org/10.1111/j.1365-2958.2010.07044.x
FnrS	b2303	folX	NC_000913	https://doi.org/10.1111/j.1365-2958.2010.07044.x
FnrS	b2531	iscR	NC_000913	https://doi.org/10.1073/pnas.1303248110
FnrS	b3829	metE	NC_000913	https://doi.org/10.1074/jbc.M109.089755
FnrS	b3908	sodA	NC_000913	https://doi.org/10.1074/jbc.M109.089755
GcvB	b1040	csgD	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.07976.x
GcvB	b1130	phoP	NC_000913	https://doi.org/10.1371/journal.pgen.1003156
GcvB	b3089	sstT	NC_000913	https://doi.org/10.1099/mic.0.023598-0
GcvB	b4208	cycA	NC_000913	https://doi.org/10.1099/mic.0.023598-0
MicA	b0411	tsx	NC_000913	https://doi.org/10.1073/pnas.1109379108
MicA	b0814	ompX	NC_000913	https://doi.org/10.1073/pnas.1109379108
MicA	b0957	ompA	NC_000913	https://doi.org/10.1101/gad.354405
MicA	b1130	phoP	NC_000913	https://doi.org/10.1111/j.1365-2958.2010.07115.x
MicC	b2215	ompC	NC_000913	https://doi.org/10.1128/JB.186.20.6689-6697.2004
MicF	b0241	phoE	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.07994.x
MicF	b0889	lrp	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.07994.x
MicF	b0929	ompF	NC_000913	https://doi.org/10.1128/jb.178.12.3650-3653.1996
MicF	b3912	cpxR	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.07994.x
OxyS	b1892	flhD	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.08209.x
OxyS	b2731	fhlA	NC_000913	https://doi.org/10.1093/emboj/17.20.6069
RprA	b1040	csgD	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.08002.x
RprA	b1341	ydaM	NC_000913	https://doi.org/10.1111/j.1365-2958.2012.08002.x
RprA	b2741	rpoS	NC_000913	https://doi.org/10.1046/j.1365-2958.2002.03203.x
RybB	b0081	mraZ	NC_000913	https://doi.org/10.1073/pnas.1303248110
RybB	b0721	sdhC	NC_000913	https://doi.org/10.1101/gad.182493.111
RybB	b0805	fiu	NC_000913	https://doi.org/10.1073/pnas.1109379108
RybB	b1256	ompW	NC_000913	https://doi.org/10.1016/j.jmb.2006.09.004
RybB	b2215	ompC	NC_000913	https://doi.org/10.1016/j.jmb.2006.09.004
RybB	b2594	rluD	NC_000913	https://doi.org/10.1073/pnas.1109379108
RyhB	b0118	acnB	NC_000913	https://doi.org/10.1093/nar/gku649
RyhB	b0156	erpA	NC_000913	https://doi.org/10.1073/pnas.1303248110
RyhB	b0288	ykgJ	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b0592	fepB	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b0683	fur	NC_000913	https://doi.org/10.1038/sj.emboj.7601553
RyhB	b0721	sdhC	NC_000913	https://doi.org/10.1101/gad.182493.111
RyhB	b0723	sdhA	NC_000913	https://doi.org/10.1073/pnas.1303248110
RyhB	b0894	dmsA	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b1107	nagZ	NC_000913	https://doi.org/10.1073/pnas.1303248110
RyhB	b1200	dhaK	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b1452	yncE	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b1531	marA	NC_000913	https://doi.org/10.1073/pnas.1303248110
RyhB	b1588	ynfF	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b1612	fumA	NC_000913	https://doi.org/10.1101/gad.2001711
RyhB	b1656	sodB	NC_000913	https://doi.org/10.1046/j.1365-2958.2003.03727.x
RyhB	b1778	msrB	NC_000913	https://doi.org/10.1371/journal.pone.0063647
RyhB	b1981	shiA	NC_000913	https://doi.org/10.1111/j.1365-2958.2007.05733.x
RyhB	b2069	yegD	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b2155	cirA	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b2206	napA	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b2530	iscS	NC_000913	https://doi.org/10.1038/emboj.2009.116
RyhB	b3365	nirB	NC_000913	https://doi.org/10.1073/pnas.1303248110
RyhB	b3607	cysE	NC_000913	https://doi.org/10.1073/pnas.1007805107
RyhB	b3942	katG	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b4070	nrfA	NC_000913	https://doi.org/10.1093/nar/gkv1158
RyhB	b4122	fumB	NC_000913	https://doi.org/10.1093/nar/gkv1158
SgrS	b1101	ptsG	NC_000913	https://doi.org/10.1111/j.1365-2958.2006.05288.x
SgrS	b1817	manX	NC_000913	https://doi.org/10.1093/nar/gkq1219
SgrS	b2416	ptsI	NC_000913	https://doi.org/10.1073/pnas.1303248110
Spot42	b0039	caiA	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b0720	gltA	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027
Spot42	b0721	sdhC	NC_000913	https://doi.org/10.1101/gad.182493.111
Spot42	b0728	sucC	NC_000913	https://doi.org/10.1073/pnas.1303248110
Spot42	b0757	galK	NC_000913	https://doi.org/10.1101/gad.231702
Spot42	b1136	icd	NC_000913	https://doi.org/10.1073/pnas.1303248110
Spot42	b1302	puuE	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b1398	paaK	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b1761	gdhA	NC_000913	https://doi.org/10.1073/pnas.1303248110
Spot42	b1901	araF	NC_000913	https://doi.org/10.1128/JB.00691-16
Spot42	b2221	atoD	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b2702	srlA	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027
Spot42	b2715	ascF	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b2801	fucP	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b2802	fucI	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027
Spot42	b3224	nanT	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b3566	xylF	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027
Spot42	b3927	glpF	NC_000913	https://doi.org/10.1038/emboj.2012.52
Spot42	b3962	sthA	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027
Spot42	b4311	nanC	NC_000913	https://doi.org/10.1016/j.molcel.2010.12.027

sRNA	geneID	geneName	genome	referenceDOI
ArcZ	STM1682	tpx	NC_003197	https://doi.org/10.1111/j.1365-2958.2009.06857.x
ArcZ	STM2970	sdaC	NC_003197	https://doi.org/10.1111/j.1365-2958.2009.06857.x
ArcZ	STM3216	-	NC_003197	https://doi.org/10.1111/j.1365-2958.2009.06857.x
ChiX	STM0687	ybfM/chiP	NC_003197	https://doi.org/10.1101/gad.541609
ChiX	STM1313	celB	NC_003197	https://doi.org/10.1101/gad.541609
CyaR	STM0833	ompX	NC_003197	https://doi.org/10.1111/j.1365-2958.2008.06189.x
GcvB	STM0002	thrA	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM0245	metQ	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM0399	brnQ	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM0602	ybdH	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM0665	gltI	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM0959	lrp	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM1299	gdhA	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM1452	tppB	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM1746	oppA	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM2355	argT	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM2526	ndk	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3062	serA	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3064	iciA	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3225	ygjU/sstT	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3564	livK	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM3567	livJ	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM3630	dppA	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM3903	ilvE	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3909	ilvC	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
GcvB	STM3930	yifK	NC_003197	https://doi.org/10.1371/journal.pgen.1004026
GcvB	STM4351	-	NC_003197	https://doi.org/10.1101/gad.447207
GcvB	STM4398	cycA	NC_003197	https://doi.org/10.1111/j.1365-2958.2011.07751.x
MicA	STM4231	lamB	NC_003197	https://doi.org/10.1111/j.1365-2958.2007.05829.x
MicC	STM1572	ompD	NC_003197	https://doi.org/10.1038/nsmb.1631
MicF	STM0366	yahO	NC_003197	https://doi.org/10.1111/j.1365-2958.2012.08031.x
MicF	STM0959	lrp	NC_003197	https://doi.org/10.1111/j.1365-2958.2012.08031.x
MicF	STM1328	lpxR	NC_003197	https://doi.org/10.1111/j.1365-2958.2012.08031.x
RybB	STM0413	tsx	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM0687	ybfM/chiP	NC_003197	https://doi.org/10.1111/j.1365-2958.2010.07342.x
RybB	STM0999	ompF	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM1070	ompA	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM1473	ompN	NC_003197	https://doi.org/10.1016/j.molcel.2008.10.027
RybB	STM1530	-	NC_003197	http://www.bioinf.uni-freiburg.de/Publications/Wright:Thesis:2012.pdf
RybB	STM1572	ompD	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM1732	ompW	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM1995	ompS	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM2267	ompC	NC_003197	https://doi.org/10.1073/pnas.1009784107
RybB	STM2391	fadL	NC_003197	https://doi.org/10.1073/pnas.1009784107
SgrS	STM2945	sopD	NC_003197	https://doi.org/10.1073/pnas.1119414109
SgrS	STM3962	yigL	NC_003197	https://doi.org/10.1016/j.cell.2013.03.003
Spot42	STM2190	mglB	NC_003197	https://doi.org/10.15252/embj.201593360


Echericha coli
NC_000913
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ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
FnrS-b1479
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FnrS-b1656
FnrS-b1841
FnrS-b2153
FnrS-b2303
FnrS-b2531
FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
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MicA-b0814
MicA-b0957
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MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b0912
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OxyS-b2731
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RybB-b1656
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RybB-b1981
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RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
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Spot42-b0721
Spot42-b0728
Spot42-b0757
Spot42-b1136
Spot42-b1302
Spot42-b1398
Spot42-b1761
Spot42-b1901
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Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
energyVRNAandronescu2007
energyVRNAlangdon2018
energyVRNAturner1999
energyVRNAturner2004
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intLoopMax8
intLoopMax9
intLoopMax10
intLoopMax11
intLoopMax12
intLoopMax14


Echericha coli
NC_000913
sum
ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
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CyaR-b1740
CyaR-b1824
CyaR-b2416
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CyaR-b2687
DsrA-b1237
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GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
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MicA-b1130
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MicF-b0241
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MicF-b0912
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RprA-b2741
RybB-b0081
RybB-b0721
RybB-b0805
RybB-b1256
RybB-b2215
RybB-b2594
RybB-b0118
RybB-b0156
RybB-b0288
RybB-b0592
RybB-b0683
RybB-b0721
RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
RybB-b1588
RybB-b1612
RybB-b1656
RybB-b1778
RybB-b1981
RybB-b2069
RybB-b2155
RybB-b2206
RybB-b2530
RybB-b3365
RybB-b3607
RybB-b3942
RybB-b4070
RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
Spot42-b0721
Spot42-b0728
Spot42-b0757
Spot42-b1136
Spot42-b1302
Spot42-b1398
Spot42-b1761
Spot42-b1901
Spot42-b2221
Spot42-b2702
Spot42-b2715
Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
intLoopMax16
noSeed
outMaxE0
outMaxE-1
outMaxE-2
outMaxE-3
outMaxE-4
outMaxE-5
outMaxE-6
outMaxE-7
outMaxE-8
outMaxE-9
outMaxE-10
outMaxE-11
outMaxE-12
outMaxE-14
outMaxE-16
outMaxE-18
outMaxE-20
outMinPu0
outMinPu0.1
outMinPu0.001
outMinPu0.0001
outMinPu0.01
outMinPu0.2
outMinPu0.5
seedBP4
seedBP4.seedMinPu0
seedBP4.seedMinPu0.001
seedBP4.seedMinPu0.0001
seedBP4.seedMinPu0.1
seedBP4.seedMinPu0.01
seedBP4.seedMinPu0.1.seedNoGU
seedBP4.seedMinPu0.01.seedNoGU
seedBP4.seedMinPu0.0001.seedNoGU
seedBP4.seedMinPu0.001.seedNoGU
seedBP4.seedMinPu0.2
seedBP4.seedMinPu0.2.seedNoGU
seedBP4.seedMinPu0.5
seedBP4.seedMinPu0.5.seedNoGU
seedBP4.seedMinPu0.seedNoGU
seedBP4.seedNoGU
seedBP5
seedBP5.seedMinPu0
seedBP5.seedMinPu0.01
seedBP5.seedMinPu0.001
seedBP5.seedMinPu0.0001
seedBP5.seedMinPu0.1
seedBP5.seedMinPu0.001.seedNoGU
seedBP5.seedMinPu0.0001.seedNoGU
seedBP5.seedMinPu0.1.seedNoGU
seedBP5.seedMinPu0.01.seedNoGU
seedBP5.seedMinPu0.2
seedBP5.seedMinPu0.2.seedNoGU
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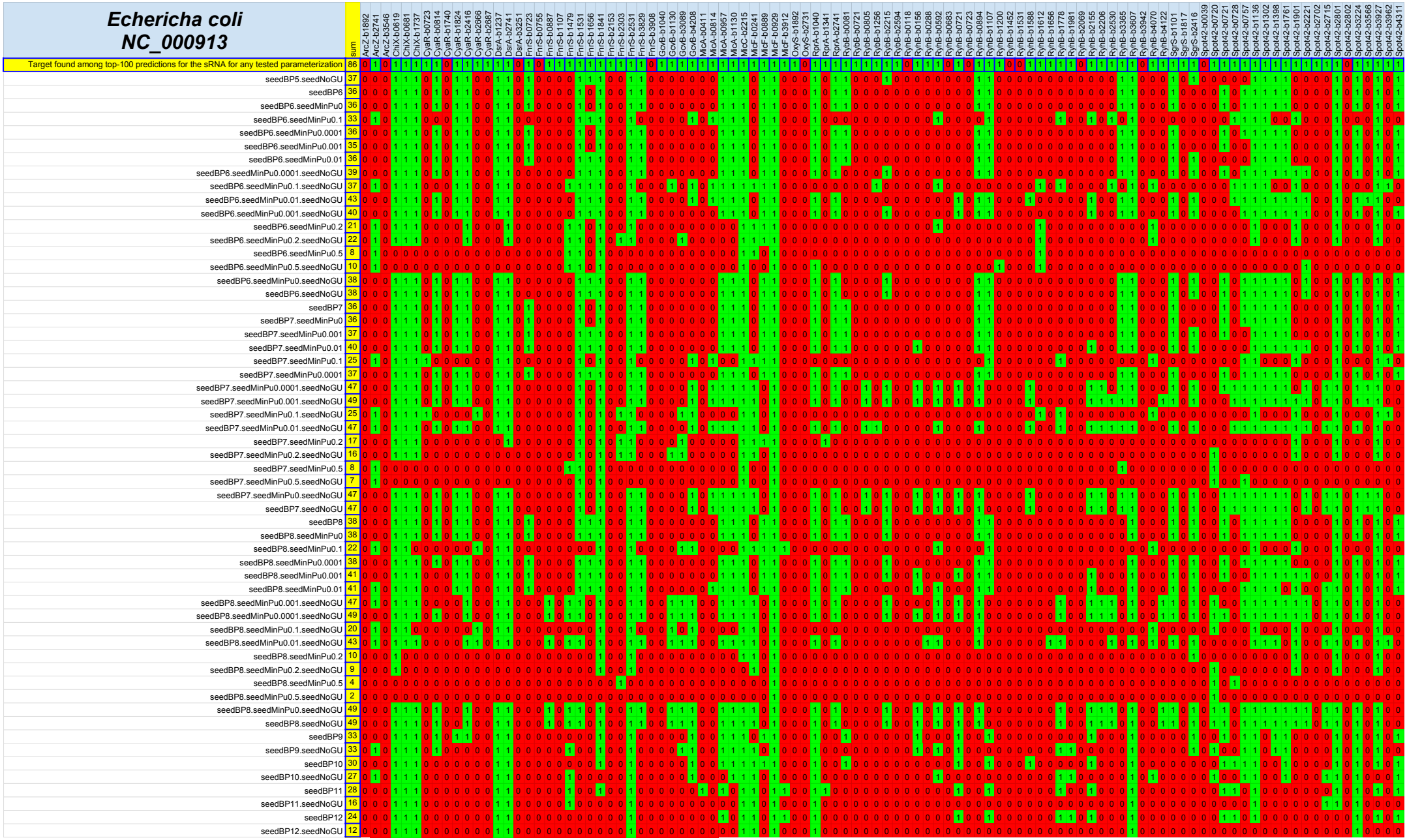
seedBP5.seedMinPu0.seedNoGU


Echericha coli
NC_000913
sum
ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
FnrS-b1479
FnrS-b1531
FnrS-b1656
FnrS-b1841
FnrS-b2153
FnrS-b2303
FnrS-b2531
FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
MicA-b1130
MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
RybB-b0721
RybB-b0805
RybB-b1256
RybB-b2215
RybB-b2594
RybB-b0118
RybB-b0156
RybB-b0288
RybB-b0592
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RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
RybB-b1588
RybB-b1612
RybB-b1656
RybB-b1778
RybB-b1981
RybB-b2069
RybB-b2155
RybB-b2206
RybB-b2530
RybB-b3365
RybB-b3607
RybB-b3942
RybB-b4070
RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
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Spot42-b0728
Spot42-b0757
Spot42-b1136
Spot42-b1302
Spot42-b1398
Spot42-b1761
Spot42-b1901
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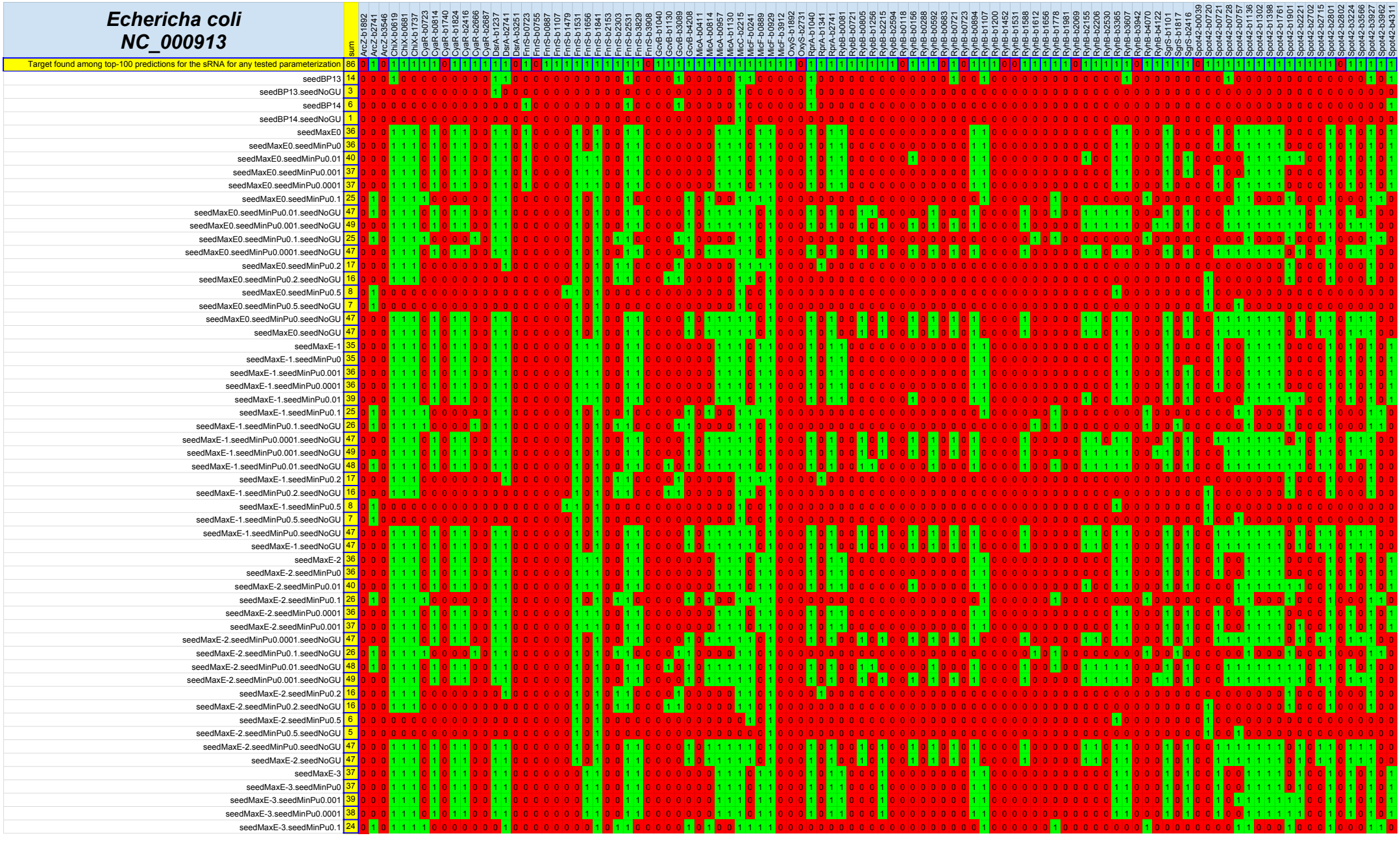
Spot42-b2715
Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedBP5.seedNoGU
seedBP6
seedBP6.seedMinPu0
seedBP6.seedMinPu0.1
seedBP6.seedMinPu0.0001
seedBP6.seedMinPu0.001
seedBP6.seedMinPu0.01
seedBP6.seedMinPu0.0001.seedNoGU
seedBP6.seedMinPu0.1.seedNoGU
seedBP6.seedMinPu0.01.seedNoGU
seedBP6.seedMinPu0.001.seedNoGU
seedBP6.seedMinPu0.2
seedBP6.seedMinPu0.2.seedNoGU
seedBP6.seedMinPu0.5
seedBP6.seedMinPu0.5.seedNoGU
seedBP6.seedMinPu0.seedNoGU
seedBP6.seedNoGU
seedBP7
seedBP7.seedMinPu0
seedBP7.seedMinPu0.001
seedBP7.seedMinPu0.01
seedBP7.seedMinPu0.1
seedBP7.seedMinPu0.0001
seedBP7.seedMinPu0.0001.seedNoGU
seedBP7.seedMinPu0.001.seedNoGU
seedBP7.seedMinPu0.1.seedNoGU
seedBP7.seedMinPu0.01.seedNoGU
seedBP7.seedMinPu0.2
seedBP7.seedMinPu0.2.seedNoGU
seedBP7.seedMinPu0.5
seedBP7.seedMinPu0.5.seedNoGU
seedBP7.seedMinPu0.seedNoGU
seedBP7.seedNoGU
seedBP8
seedBP8.seedMinPu0
seedBP8.seedMinPu0.1
seedBP8.seedMinPu0.0001
seedBP8.seedMinPu0.001
seedBP8.seedMinPu0.01
seedBP8.seedMinPu0.001.seedNoGU
seedBP8.seedMinPu0.0001.seedNoGU
seedBP8.seedMinPu0.1.seedNoGU
seedBP8.seedMinPu0.01.seedNoGU
seedBP8.seedMinPu0.2
seedBP8.seedMinPu0.2.seedNoGU
seedBP8.seedMinPu0.5
seedBP8.seedMinPu0.5.seedNoGU
seedBP8.seedMinPu0.seedNoGU
seedBP8.seedNoGU
seedBP9
seedBP9.seedNoGU
seedBP10
seedBP10.seedNoGU
seedBP11
seedBP11.seedNoGU
seedBP12
seedBP12.seedNoGU


Echericha coli
NC_000913
sum
ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
FnrS-b1479
FnrS-b1531
FnrS-b1656
FnrS-b1841
FnrS-b2153
FnrS-b2303
FnrS-b2531
FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
MicA-b1130
MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
RybB-b0721
RybB-b0805
RybB-b1256
RybB-b2215
RybB-b2594
RybB-b0118
RybB-b0156
RybB-b0288
RybB-b0592
RybB-b0683
RybB-b0721
RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
RybB-b1588
RybB-b1612
RybB-b1656
RybB-b1778
RybB-b1981
RybB-b2069
RybB-b2155
RybB-b2206
RybB-b2530
RybB-b3365
RybB-b3607
RybB-b3942
RybB-b4070
RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
Spot42-b0721
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Spot42-b2221
Spot42-b2702
Spot42-b2715
Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedBP13
seedBP13.seedNoGU
seedBP14
seedBP14.seedNoGU
seedMaxE0
seedMaxE0.seedMinPu0
seedMaxE0.seedMinPu0.01
seedMaxE0.seedMinPu0.001
seedMaxE0.seedMinPu0.0001
seedMaxE0.seedMinPu0.1
seedMaxE0.seedMinPu0.01.seedNoGU
seedMaxE0.seedMinPu0.001.seedNoGU
seedMaxE0.seedMinPu0.1.seedNoGU
seedMaxE0.seedMinPu0.0001.seedNoGU
seedMaxE0.seedMinPu0.2
seedMaxE0.seedMinPu0.2.seedNoGU
seedMaxE0.seedMinPu0.5
seedMaxE0.seedMinPu0.5.seedNoGU
seedMaxE0.seedMinPu0.seedNoGU
seedMaxE0.seedNoGU
seedMaxE-1
seedMaxE-1.seedMinPu0
seedMaxE-1.seedMinPu0.001
seedMaxE-1.seedMinPu0.0001
seedMaxE-1.seedMinPu0.01
seedMaxE-1.seedMinPu0.1
seedMaxE-1.seedMinPu0.1.seedNoGU
seedMaxE-1.seedMinPu0.0001.seedNoGU
seedMaxE-1.seedMinPu0.001.seedNoGU
seedMaxE-1.seedMinPu0.01.seedNoGU
seedMaxE-1.seedMinPu0.2
seedMaxE-1.seedMinPu0.2.seedNoGU
seedMaxE-1.seedMinPu0.5
seedMaxE-1.seedMinPu0.5.seedNoGU
seedMaxE-1.seedMinPu0.seedNoGU
seedMaxE-1.seedNoGU
seedMaxE-2
seedMaxE-2.seedMinPu0
seedMaxE-2.seedMinPu0.01
seedMaxE-2.seedMinPu0.1
seedMaxE-2.seedMinPu0.0001
seedMaxE-2.seedMinPu0.001
seedMaxE-2.seedMinPu0.0001.seedNoGU
seedMaxE-2.seedMinPu0.1.seedNoGU
seedMaxE-2.seedMinPu0.01.seedNoGU
seedMaxE-2.seedMinPu0.001.seedNoGU
seedMaxE-2.seedMinPu0.2
seedMaxE-2.seedMinPu0.2.seedNoGU
seedMaxE-2.seedMinPu0.5
seedMaxE-2.seedMinPu0.5.seedNoGU
seedMaxE-2.seedMinPu0.seedNoGU
seedMaxE-2.seedNoGU
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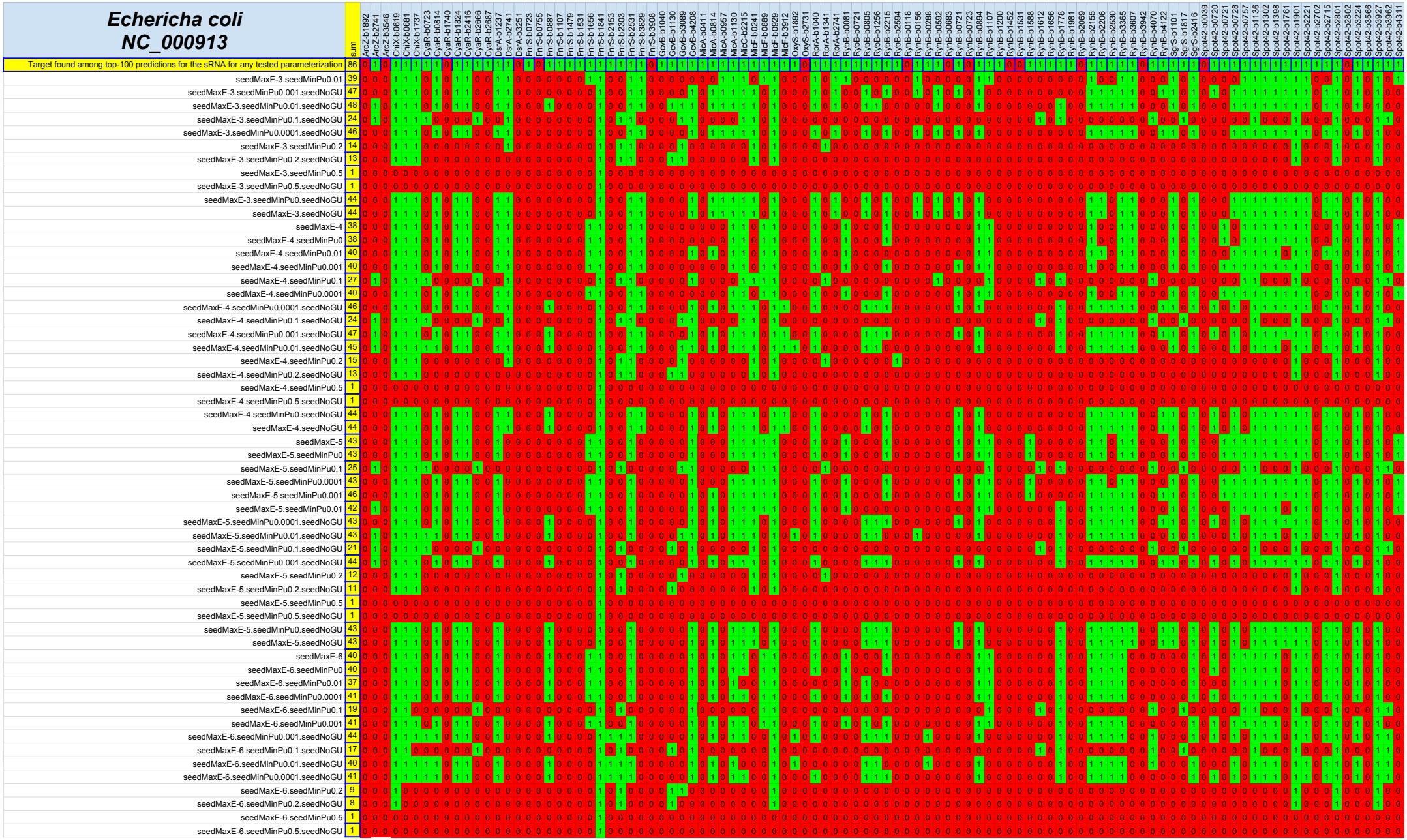
seedMaxE-3.seedMinPu0.001
seedMaxE-3.seedMinPu0.0001
seedMaxE-3.seedMinPu0.1


Echericha coli
NC_000913
sum
ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
FnrS-b1479
FnrS-b1531
FnrS-b1656
FnrS-b1841
FnrS-b2153
FnrS-b2303
FnrS-b2531
FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
MicA-b1130
MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
RybB-b0721
RybB-b0805
RybB-b1256
RybB-b2215
RybB-b2594
RybB-b0118
RybB-b0156
RybB-b0288
RybB-b0592
RybB-b0683
RybB-b0721
RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
RybB-b1588
RybB-b1612
RybB-b1656
RybB-b1778
RybB-b1981
RybB-b2069
RybB-b2155
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RybB-b2530
RybB-b3365
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RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
Spot42-b0721
Spot42-b0728
Spot42-b0757
Spot42-b1136
Spot42-b1302
Spot42-b1398
Spot42-b1761
Spot42-b1901
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Spot42-b2702
Spot42-b2715
Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
86
seedMaxE-3.seedMinPu0.01
39
seedMaxE-3.seedMinPu0.001.seedNoGU
47
seedMaxE-3.seedMinPu0.01.seedNoGU
48
seedMaxE-3.seedMinPu0.1.seedNoGU
24
seedMaxE-3.seedMinPu0.0001.seedNoGU
46
seedMaxE-3.seedMinPu0.2
14
seedMaxE-3.seedMinPu0.2.seedNoGU
13
seedMaxE-3.seedMinPu0.5
1
seedMaxE-3.seedMinPu0.5.seedNoGU
1
seedMaxE-3.seedMinPu0.seedNoGU
44
seedMaxE-3.seedNoGU
44
seedMaxE-4
38
seedMaxE-4.seedMinPu0
38
seedMaxE-4.seedMinPu0.01
40
seedMaxE-4.seedMinPu0.001
40
seedMaxE-4.seedMinPu0.1
27
seedMaxE-4.seedMinPu0.0001
40
seedMaxE-4.seedMinPu0.0001.seedNoGU
46
seedMaxE-4.seedMinPu0.1.seedNoGU
24
seedMaxE-4.seedMinPu0.001.seedNoGU
47
seedMaxE-4.seedMinPu0.01.seedNoGU
45
seedMaxE-4.seedMinPu0.2
15
seedMaxE-4.seedMinPu0.2.seedNoGU
13
seedMaxE-4.seedMinPu0.5
1
seedMaxE-4.seedMinPu0.5.seedNoGU
1
seedMaxE-4.seedMinPu0.seedNoGU
44
seedMaxE-4.seedNoGU
44
seedMaxE-5
43
seedMaxE-5.seedMinPu0
43
seedMaxE-5.seedMinPu0.1
25
seedMaxE-5.seedMinPu0.0001
43
seedMaxE-5.seedMinPu0.001
46
seedMaxE-5.seedMinPu0.01
42
seedMaxE-5.seedMinPu0.0001.seedNoGU
43
seedMaxE-5.seedMinPu0.01.seedNoGU
43
seedMaxE-5.seedMinPu0.1.seedNoGU
21
seedMaxE-5.seedMinPu0.001.seedNoGU
44
seedMaxE-5.seedMinPu0.2
12
seedMaxE-5.seedMinPu0.2.seedNoGU
11
seedMaxE-5.seedMinPu0.5
1
seedMaxE-5.seedMinPu0.5.seedNoGU
1
seedMaxE-5.seedMinPu0.seedNoGU
43
seedMaxE-5.seedNoGU
43
seedMaxE-6
40
seedMaxE-6.seedMinPu0
40
seedMaxE-6.seedMinPu0.01
37
seedMaxE-6.seedMinPu0.0001
41
seedMaxE-6.seedMinPu0.1
19
seedMaxE-6.seedMinPu0.001
41
seedMaxE-6.seedMinPu0.001.seedNoGU
44
seedMaxE-6.seedMinPu0.1.seedNoGU
17
seedMaxE-6.seedMinPu0.01.seedNoGU
40
seedMaxE-6.seedMinPu0.0001.seedNoGU
41
seedMaxE-6.seedMinPu0.2
9
seedMaxE-6.seedMinPu0.2.seedNoGU
8
seedMaxE-6.seedMinPu0.5
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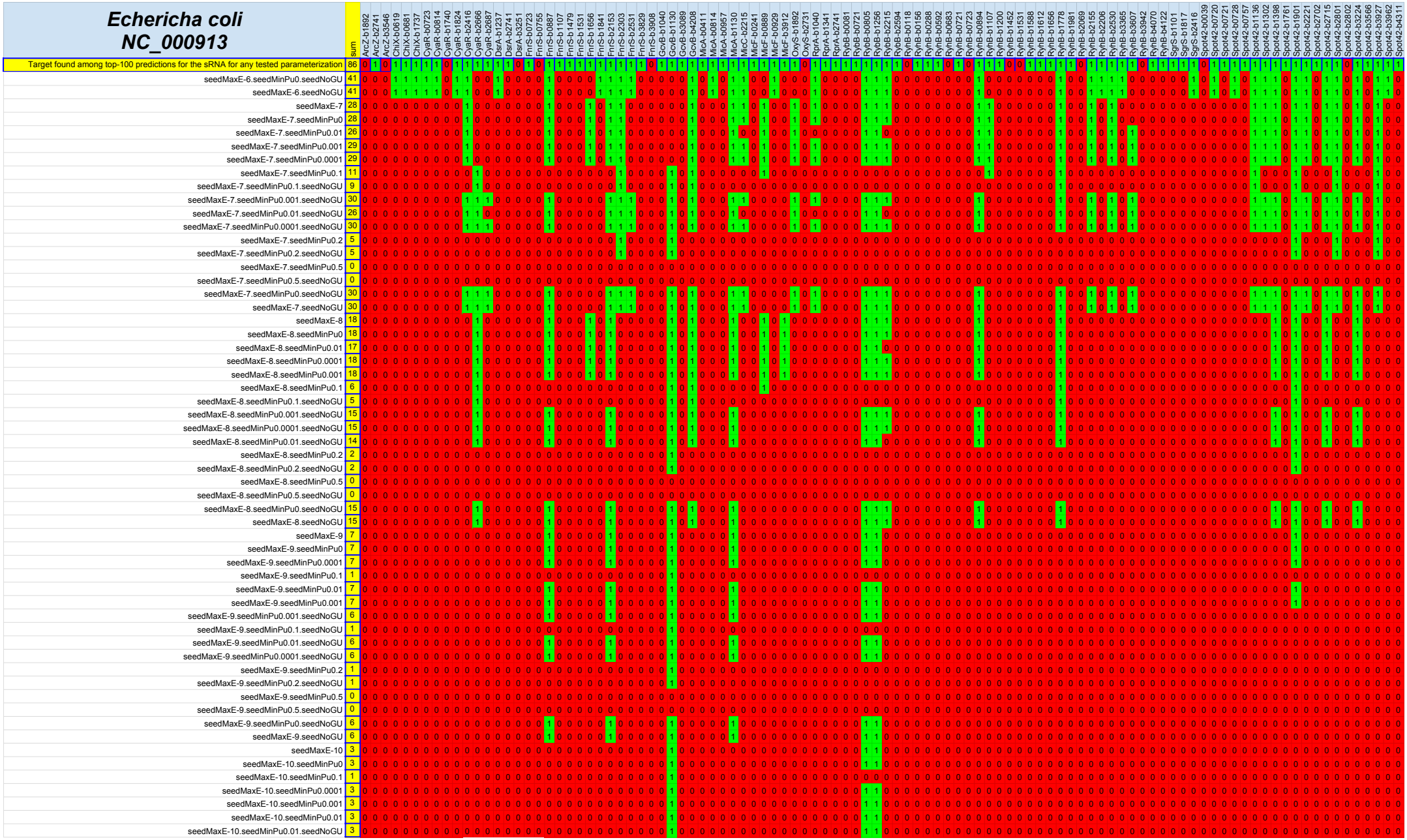
seedMaxE-6.seedMinPu0.5.seedNoGU
1


Echericha coli
NC_000913
sum
ArcZ-b1892
ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
FnrS-b1479
FnrS-b1531
FnrS-b1656
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FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
MicA-b1130
MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
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RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
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RybB-b1588
RybB-b1612
RybB-b1656
RybB-b1778
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SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
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Spot42-b1302
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Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxE-6.seedMinPu0.seedNoGU
seedMaxE-6.seedNoGU
seedMaxE-7
seedMaxE-7.seedMinPu0
seedMaxE-7.seedMinPu0.01
seedMaxE-7.seedMinPu0.001
seedMaxE-7.seedMinPu0.0001
seedMaxE-7.seedMinPu0.1
seedMaxE-7.seedMinPu0.1.seedNoGU
seedMaxE-7.seedMinPu0.001.seedNoGU
seedMaxE-7.seedMinPu0.01.seedNoGU
seedMaxE-7.seedMinPu0.0001.seedNoGU
seedMaxE-7.seedMinPu0.2
seedMaxE-7.seedMinPu0.2.seedNoGU
seedMaxE-7.seedMinPu0.5
seedMaxE-7.seedMinPu0.5.seedNoGU
seedMaxE-7.seedMinPu0.seedNoGU
seedMaxE-7.seedNoGU
seedMaxE-8
seedMaxE-8.seedMinPu0
seedMaxE-8.seedMinPu0.01
seedMaxE-8.seedMinPu0.0001
seedMaxE-8.seedMinPu0.001
seedMaxE-8.seedMinPu0.1
seedMaxE-8.seedMinPu0.1.seedNoGU
seedMaxE-8.seedMinPu0.001.seedNoGU
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seedMaxE-8.seedMinPu0.01.seedNoGU
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seedMaxE-8.seedMinPu0.2.seedNoGU
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seedMaxE-8.seedMinPu0.seedNoGU
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seedMaxE-9
seedMaxE-9.seedMinPu0
seedMaxE-9.seedMinPu0.0001
seedMaxE-9.seedMinPu0.1
seedMaxE-9.seedMinPu0.01
seedMaxE-9.seedMinPu0.001
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seedMaxE-9.seedMinPu0.1.seedNoGU
seedMaxE-9.seedMinPu0.01.seedNoGU
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seedMaxE-9.seedNoGU
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seedMaxE-10.seedMinPu0
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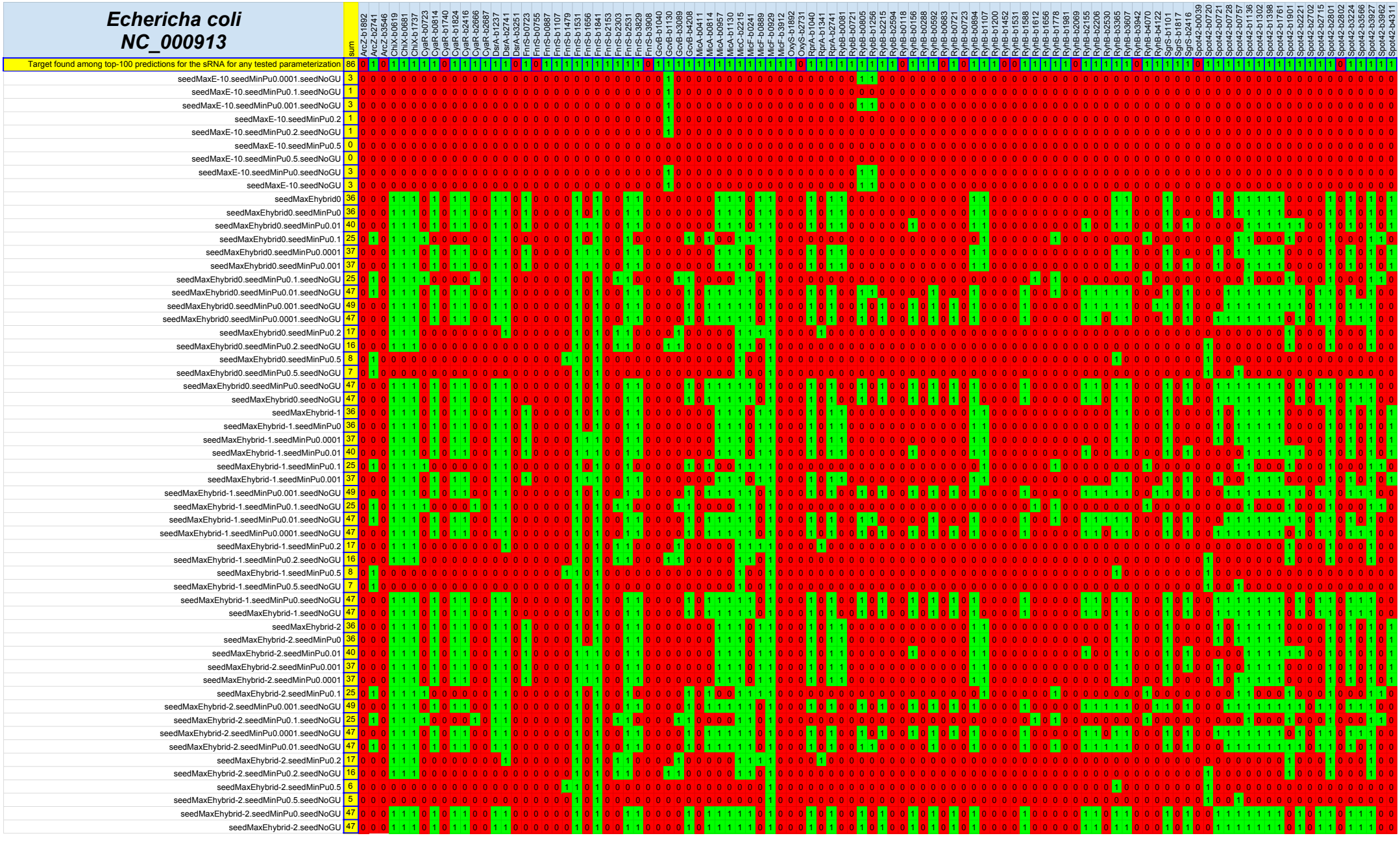
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NC_000913
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ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
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FnrS-b1656
FnrS-b1841
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FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
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MicC-b2215
MicF-b0241
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MicF-b0929
MicF-b3912
OxyS-b1892
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RprA-b2741
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RybB-b0721
RybB-b0805
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RybB-b0894
RybB-b1107
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RybB-b1452
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RybB-b1656
RybB-b1778
RybB-b1981
RybB-b2069
RybB-b2155
RybB-b2206
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RybB-b3607
RybB-b3942
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RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
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Spot42-b0728
Spot42-b0757
Spot42-b1136
Spot42-b1302
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Spot42-b2702
Spot42-b2715
Spot42-b2801
Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxE-10.seedMinPu0.0001.seedNoGU
seedMaxE-10.seedMinPu0.1.seedNoGU
seedMaxE-10.seedMinPu0.001.seedNoGU
seedMaxE-10.seedMinPu0.2
seedMaxE-10.seedMinPu0.2.seedNoGU
seedMaxE-10.seedMinPu0.5
seedMaxE-10.seedMinPu0.5.seedNoGU
seedMaxE-10.seedMinPu0.seedNoGU
seedMaxE-10.seedNoGU
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seedMaxEhybrid0.seedMinPu0
seedMaxEhybrid0.seedMinPu0.01
seedMaxEhybrid0.seedMinPu0.1
seedMaxEhybrid0.seedMinPu0.0001
seedMaxEhybrid0.seedMinPu0.001
seedMaxEhybrid0.seedMinPu0.1.seedNoGU
seedMaxEhybrid0.seedMinPu0.01.seedNoGU
seedMaxEhybrid0.seedMinPu0.001.seedNoGU
seedMaxEhybrid0.seedMinPu0.0001.seedNoGU
seedMaxEhybrid0.seedMinPu0.2
seedMaxEhybrid0.seedMinPu0.2.seedNoGU
seedMaxEhybrid0.seedMinPu0.5
seedMaxEhybrid0.seedMinPu0.5.seedNoGU
seedMaxEhybrid0.seedMinPu0.seedNoGU
seedMaxEhybrid0.seedNoGU
seedMaxEhybrid-1
seedMaxEhybrid-1.seedMinPu0
seedMaxEhybrid-1.seedMinPu0.0001
seedMaxEhybrid-1.seedMinPu0.01
seedMaxEhybrid-1.seedMinPu0.1
seedMaxEhybrid-1.seedMinPu0.001
seedMaxEhybrid-1.seedMinPu0.001.seedNoGU
seedMaxEhybrid-1.seedMinPu0.1.seedNoGU
seedMaxEhybrid-1.seedMinPu0.01.seedNoGU
seedMaxEhybrid-1.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-1.seedMinPu0.2
seedMaxEhybrid-1.seedMinPu0.2.seedNoGU
seedMaxEhybrid-1.seedMinPu0.5
seedMaxEhybrid-1.seedMinPu0.5.seedNoGU
seedMaxEhybrid-1.seedMinPu0.seedNoGU
seedMaxEhybrid-1.seedNoGU
seedMaxEhybrid-2
seedMaxEhybrid-2.seedMinPu0
seedMaxEhybrid-2.seedMinPu0.01
seedMaxEhybrid-2.seedMinPu0.001
seedMaxEhybrid-2.seedMinPu0.0001
seedMaxEhybrid-2.seedMinPu0.1
seedMaxEhybrid-2.seedMinPu0.001.seedNoGU
seedMaxEhybrid-2.seedMinPu0.1.seedNoGU
seedMaxEhybrid-2.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-2.seedMinPu0.01.seedNoGU
seedMaxEhybrid-2.seedMinPu0.2
seedMaxEhybrid-2.seedMinPu0.2.seedNoGU
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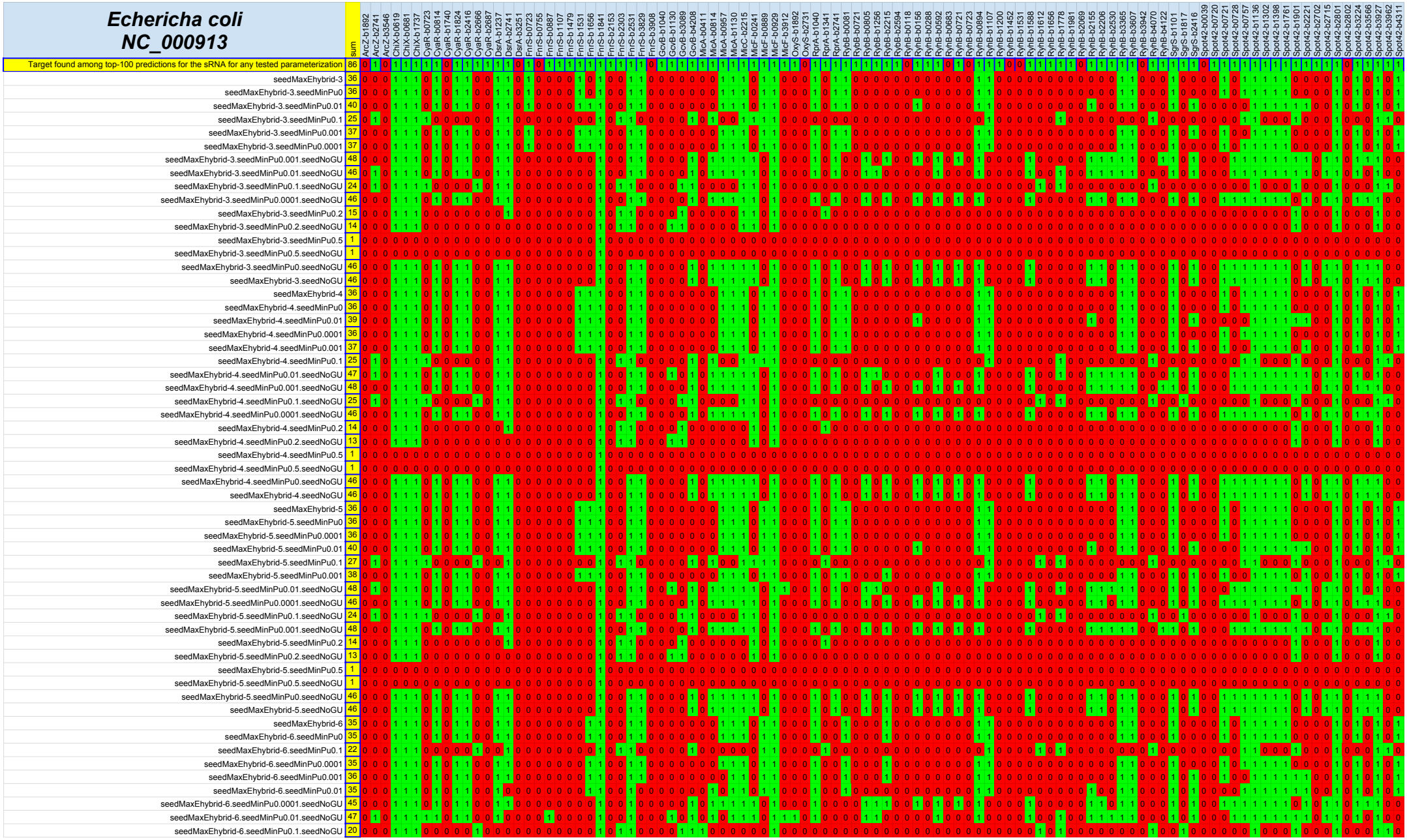
Echericha coli
NC_000913
sum
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ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
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CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
FnrS-b0723
FnrS-b0755
FnrS-b0887
FnrS-b1107
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FnrS-b1656
FnrS-b1841
FnrS-b2153
FnrS-b2303
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FnrS-b3829
FnrS-b3908
GcvB-b1040
GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
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MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b0912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
RybB-b0721
RybB-b0805
RybB-b1256
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RybB-b0118
RybB-b0156
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RybB-b0723
RybB-b0894
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RybB-b1200
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RybB-b1981
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SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
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Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxEhybrid-3
seedMaxEhybrid-3.seedMinPu0
seedMaxEhybrid-3.seedMinPu0.01
seedMaxEhybrid-3.seedMinPu0.1
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seedMaxEhybrid-3.seedMinPu0.001.seedNoGU
seedMaxEhybrid-3.seedMinPu0.01.seedNoGU
seedMaxEhybrid-3.seedMinPu0.1.seedNoGU
seedMaxEhybrid-3.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-3.seedMinPu0.2
seedMaxEhybrid-3.seedMinPu0.2.seedNoGU
seedMaxEhybrid-3.seedMinPu0.5
seedMaxEhybrid-3.seedMinPu0.5.seedNoGU
seedMaxEhybrid-3.seedMinPu0.seedNoGU
seedMaxEhybrid-3.seedNoGU
seedMaxEhybrid-4
seedMaxEhybrid-4.seedMinPu0
seedMaxEhybrid-4.seedMinPu0.01
seedMaxEhybrid-4.seedMinPu0.0001
seedMaxEhybrid-4.seedMinPu0.001
seedMaxEhybrid-4.seedMinPu0.1
seedMaxEhybrid-4.seedMinPu0.01.seedNoGU
seedMaxEhybrid-4.seedMinPu0.001.seedNoGU
seedMaxEhybrid-4.seedMinPu0.1.seedNoGU
seedMaxEhybrid-4.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-4.seedMinPu0.2
seedMaxEhybrid-4.seedMinPu0.2.seedNoGU
seedMaxEhybrid-4.seedMinPu0.5
seedMaxEhybrid-4.seedMinPu0.5.seedNoGU
seedMaxEhybrid-4.seedMinPu0.seedNoGU
seedMaxEhybrid-4.seedNoGU
seedMaxEhybrid-5
seedMaxEhybrid-5.seedMinPu0
seedMaxEhybrid-5.seedMinPu0.0001
seedMaxEhybrid-5.seedMinPu0.01
seedMaxEhybrid-5.seedMinPu0.1
seedMaxEhybrid-5.seedMinPu0.001
seedMaxEhybrid-5.seedMinPu0.01.seedNoGU
seedMaxEhybrid-5.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-5.seedMinPu0.1.seedNoGU
seedMaxEhybrid-5.seedMinPu0.001.seedNoGU
seedMaxEhybrid-5.seedMinPu0.2
seedMaxEhybrid-5.seedMinPu0.2.seedNoGU
seedMaxEhybrid-5.seedMinPu0.5
seedMaxEhybrid-5.seedMinPu0.5.seedNoGU
seedMaxEhybrid-5.seedMinPu0.seedNoGU
seedMaxEhybrid-5.seedNoGU
seedMaxEhybrid-6
seedMaxEhybrid-6.seedMinPu0
seedMaxEhybrid-6.seedMinPu0.1
seedMaxEhybrid-6.seedMinPu0.0001
seedMaxEhybrid-6.seedMinPu0.001
seedMaxEhybrid-6.seedMinPu0.01
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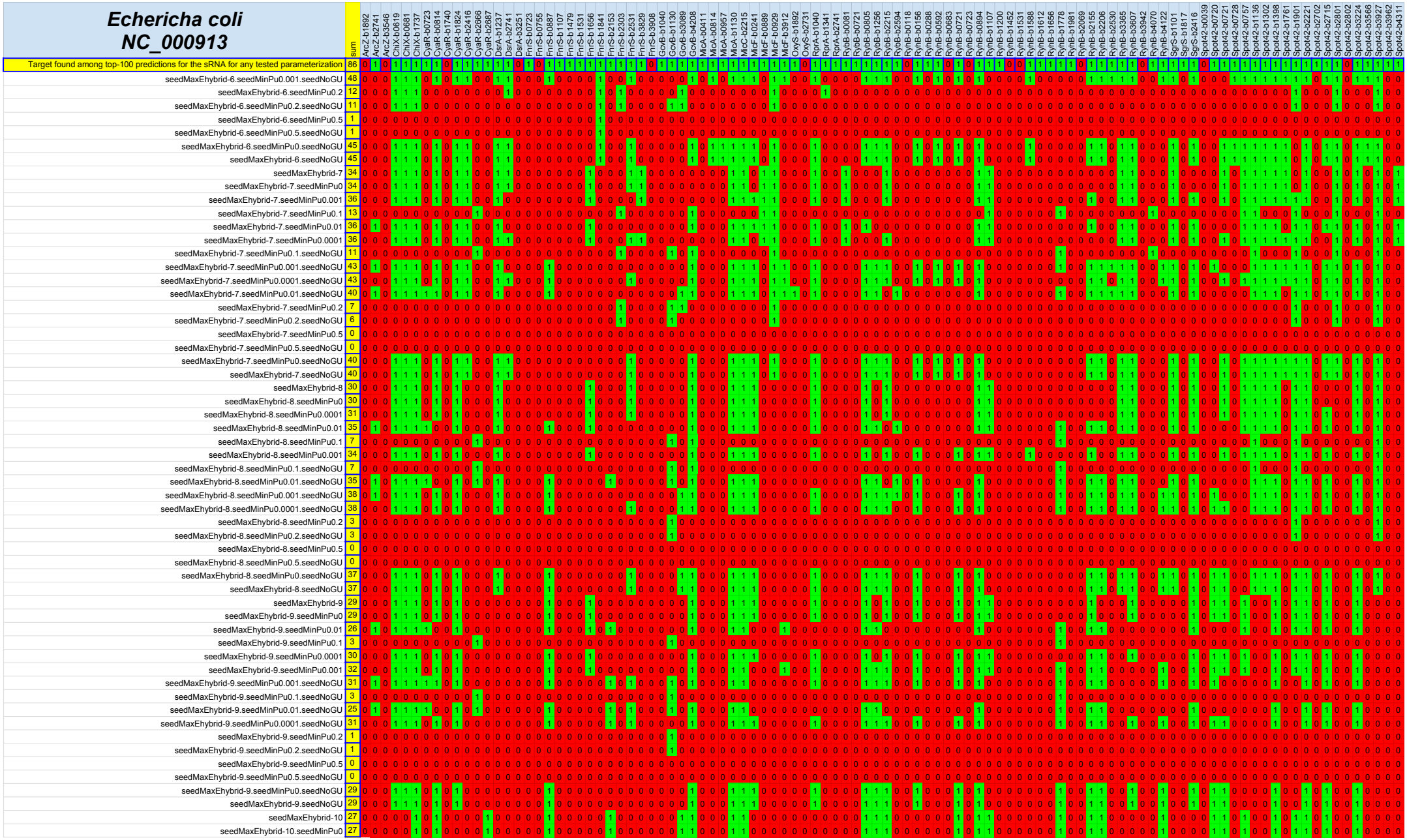
Echericha coli
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ArcZ-b2741
ArcZ-b3546
ChiX-b0619
ChiX-b0681
ChiX-b1737
CyaR-b0723
CyaR-b0814
CyaR-b1740
CyaR-b1824
CyaR-b2416
CyaR-b2666
CyaR-b2687
DsrA-b1237
DsrA-b2741
DsrA-b3251
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FnrS-b1656
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FnrS-b2153
FnrS-b2303
FnrS-b2531
FnrS-b3829
FnrS-b3908
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GcvB-b1130
GcvB-b3089
GcvB-b4208
MicA-b0411
MicA-b0814
MicA-b0957
MicA-b1130
MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
RprA-b1040
RprA-b1341
RprA-b2741
RybB-b0081
RybB-b0721
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RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
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RybB-b1656
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RybB-b1981
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RybB-b2155
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RybB-b3942
RybB-b4070
RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
Spot42-b0720
Spot42-b0721
Spot42-b0728
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Spot42-b1136
Spot42-b1302
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Spot42-b2802
Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxEhybrid-6.seedMinPu0.001.seedNoGU
seedMaxEhybrid-6.seedMinPu0.2
seedMaxEhybrid-6.seedMinPu0.2.seedNoGU
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seedMaxEhybrid-6.seedMinPu0.seedNoGU
seedMaxEhybrid-6.seedNoGU
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seedMaxEhybrid-7.seedMinPu0.001
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seedMaxEhybrid-7.seedMinPu0.01
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seedMaxEhybrid-7.seedMinPu0.001.seedNoGU
seedMaxEhybrid-7.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-7.seedMinPu0.01.seedNoGU
seedMaxEhybrid-7.seedMinPu0.2
seedMaxEhybrid-7.seedMinPu0.2.seedNoGU
seedMaxEhybrid-7.seedMinPu0.5
seedMaxEhybrid-7.seedMinPu0.5.seedNoGU
seedMaxEhybrid-7.seedMinPu0.seedNoGU
seedMaxEhybrid-7.seedNoGU
seedMaxEhybrid-8
seedMaxEhybrid-8.seedMinPu0
seedMaxEhybrid-8.seedMinPu0.0001
seedMaxEhybrid-8.seedMinPu0.01
seedMaxEhybrid-8.seedMinPu0.1
seedMaxEhybrid-8.seedMinPu0.001
seedMaxEhybrid-8.seedMinPu0.1.seedNoGU
seedMaxEhybrid-8.seedMinPu0.01.seedNoGU
seedMaxEhybrid-8.seedMinPu0.001.seedNoGU
seedMaxEhybrid-8.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-8.seedMinPu0.2
seedMaxEhybrid-8.seedMinPu0.2.seedNoGU
seedMaxEhybrid-8.seedMinPu0.5
seedMaxEhybrid-8.seedMinPu0.5.seedNoGU
seedMaxEhybrid-8.seedMinPu0.seedNoGU
seedMaxEhybrid-8.seedNoGU
seedMaxEhybrid-9
seedMaxEhybrid-9.seedMinPu0
seedMaxEhybrid-9.seedMinPu0.01
seedMaxEhybrid-9.seedMinPu0.1
seedMaxEhybrid-9.seedMinPu0.0001
seedMaxEhybrid-9.seedMinPu0.001
seedMaxEhybrid-9.seedMinPu0.001.seedNoGU
seedMaxEhybrid-9.seedMinPu0.1.seedNoGU
seedMaxEhybrid-9.seedMinPu0.01.seedNoGU
seedMaxEhybrid-9.seedMinPu0.0001.seedNoGU
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seedMaxEhybrid-9.seedMinPu0.2.seedNoGU
seedMaxEhybrid-9.seedMinPu0.5
seedMaxEhybrid-9.seedMinPu0.5.seedNoGU
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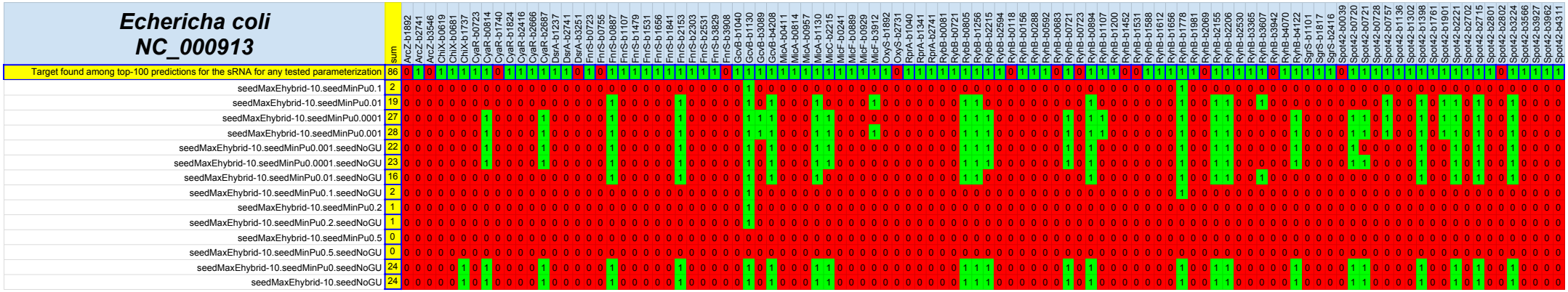
seedMaxEhybrid-9.seedNoGU
seedMaxEhybrid-10
seedMaxEhybrid-10.seedMinPu0


Echericha coli
NC_000913
sum
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ArcZ-b3546
ChiX-b0619
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CyaR-b0723
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CyaR-b1740
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DsrA-b1237
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GcvB-b1130
GcvB-b3089
GcvB-b4208
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MicA-b0814
MicA-b0957
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MicC-b2215
MicF-b0241
MicF-b0889
MicF-b0929
MicF-b3912
OxyS-b1892
OxyS-b2731
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RybB-b0723
RybB-b0894
RybB-b1107
RybB-b1200
RybB-b1452
RybB-b1531
RybB-b1588
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RybB-b1656
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RybB-b3942
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RybB-b4122
SgrS-b1101
SgrS-b1817
SgrS-b2416
Spot42-b0039
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Spot42-b3224
Spot42-b3566
Spot42-b3927
Spot42-b3962
Spot42-b4311
Target found among top-100 predictions for the sRNA for any tested parameterization
86
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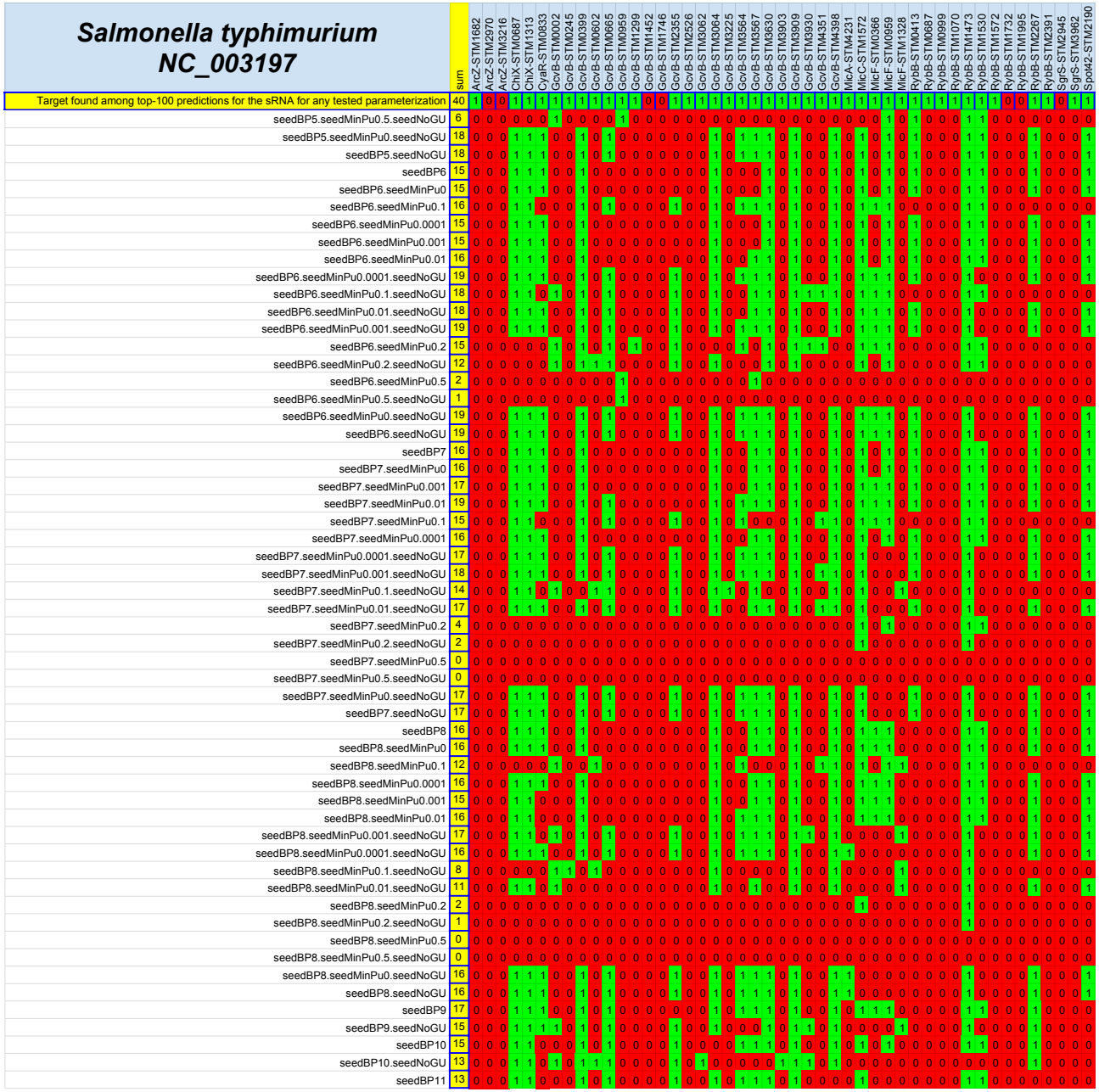
2
seedMaxEhybrid-10.seedMinPu0.01
19
seedMaxEhybrid-10.seedMinPu0.0001
27
seedMaxEhybrid-10.seedMinPu0.001
28
seedMaxEhybrid-10.seedMinPu0.001.seedNoGU
22
seedMaxEhybrid-10.seedMinPu0.0001.seedNoGU
23
seedMaxEhybrid-10.seedMinPu0.01.seedNoGU
16
seedMaxEhybrid-10.seedMinPu0.1.seedNoGU
2
seedMaxEhybrid-10.seedMinPu0.2
1
seedMaxEhybrid-10.seedMinPu0.2.seedNoGU
1
seedMaxEhybrid-10.seedMinPu0.5
0
seedMaxEhybrid-10.seedMinPu0.5.seedNoGU
0
seedMaxEhybrid-10.seedMinPu0.seedNoGU
24
seedMaxEhybrid-10.seedNoGU
24

Salmonella typhimurium NC_003197

	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
energyVRNAandronescu2007	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	1	0	1	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
energyVRNAlangdon2018	14	0	0	0	1	1	1	1	0	1	0	0	0	0	0	1	0	0	0	0	0	1	1	0	1	0	0	1	0	0	0	1	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	1
energyVRNAturner1999	19	0	0	0	1	1	1	0	0	1	0	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	1	0	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	1	0	1
energyVRNAturner2004	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	1	0	1	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU	20	0	0	0	1	1	1	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	1	0	1	0	0	1	1	0	0	1	1	0	0	1	1	0	0	0	1	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001	20	0	0	0	1	1	1	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	1	0	1	0	0	1	1	0	0	1	1	0	0	1	1	0	0	0	1	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60	20	0	0	0	1	1	1	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	1	0	1	0	0	1	1	0	0	1	1	0	0	1	1	0	0	0	1	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax2	16	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	0	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax2.outMinPu0.001	16	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	0	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax5	17	0	0	0	1	1	0	1	0	1	0	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax5.outMinPu0.001	17	0	0	0	1	1	0	1	0	1	0	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax7	19	0	0	0	1	1	0	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	1	0	1	0	0	1	1	0	0	1	1	0	0	1	1	0	0	0	1	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax7.outMinPu0.001	18	0	0	0	1	1	0	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax8	19	0	0	0	1	1	0	1	0	1	0	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax8.outMinPu0.001	18	0	0	0	1	1	0	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax10	19	0	0	0	1	1	0	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.and.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax10.outMinPu0.001	18	0	0	0	1	1	0	1	0	1	0	1	0	0	0	1	0	0	0	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	1	0	1	0	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60	18	0	0	0	1	1	1	0	0	1	0	1	0	1	0	0	0	0	1	0	0	1	1	0	1	0	1	1	1	0	1	0	1	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax2	15	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	1	1	0	1	0	1	1	0	1	0	0	0	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax2.outMinPu0.001	15	0	0	0	1	1	0	0	0	1	0	0	0	0	0	1	1	0	1	0	1	1	1	0	1	0	0	0	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax5	16	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	0	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax5.outMinPu0.001	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	0	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax7	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax7.outMinPu0.001	18	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	1	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax8	18	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	1	0	1	0	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax8.outMinPu0.001	19	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax10	18	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	1	0	1	0	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
final.seedBP7.seedNoGU.seedMinPu0.001.intLenMax60.intLoopMax10.outMinPu0.001	19	0	0	0	1	1	0	0	0	1	0	1	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax10	14	0	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	1	1	1	1	0	1	0	1	0	0	0	1	1	0	0	1	0	0	1
intLenMax15	12	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	1	1	0	0	0	0	1	1	0	0	0	1	0	0	1
intLenMax20	14	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	0	0	1	1	1	1	0	0	0	0	1	1	0	0	0	1	0	0	1	0	1
intLenMax25	15	0	0	0	1	1	0	1	0	0	0	1	0	0	0	0	1	1	0	0	0	1	0	0	0	1	0	0	1	0	0	1	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	0	1
intLenMax30	14	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	1	0	1	0	0	1	0	1	1	1	1	0	0	0	0	1	1	0	0	0	1	0	0	1	0	1
intLenMax40	15	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	1	0	0	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax50	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	1	0	0	1	1	1	0	1	0	0	1	0	1	0	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
intLenMax60	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax70	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax75	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax80	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax100	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLenMax150	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLoopMax2	14	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	1	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	0	0	0	1	1	0	0	0	1	0	0	0	1	0	0	1
intLoopMax3	14	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	1	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	0	0	0	1	1	0	0	0	1	0	0	0	1	0	0	1
intLoopMax4	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	0	1	1	0	0	0	1	0	0	0	1
intLoopMax5	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	0	1	0	1	0	0	1	0	1	0	1	0	1	1	0	0	0	0	1	1	0	0	0	1	0	0	1
intLoopMax6	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	0	0	1	0	1	0	1	0	1	0	0	1	0	1	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
intLoopMax7	17	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	0	0	1	0	1	0	1	0	1	0	0	1	0	1	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
intLoopMax8	17	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
intLoopMax9	16	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLoopMax10	15	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLoopMax11	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
intLoopMax12	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1

Salmonella typhimurium NC_003197	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
intLoopMax14	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
intLoopMax16	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
noSeed	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE0	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-1	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-2	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-3	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-4	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-5	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-6	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-7	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-8	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-9	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-10	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMaxE-11	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	0	0
outMaxE-12	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	0	0
outMaxE-14	14	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	0	0
outMaxE-16	11	0	0	0	0	1	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	0	1	0	1	0	0	0	1	1	0	0	0	0	0	0	0	0	0
outMaxE-18	7	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
outMaxE-20	4	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
outMinPu0	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMinPu0.1	17	0	0	0	1	1	0	1	0	0	1	0	1	0	0	0	1	1	0	0	0	0	0	1	0	1	0	0	1	0	1	1	1	1	0	0	0	0	1	1	0	0	0	1	0	0	0	0
outMinPu0.001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMinPu0.0001	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMinPu0.01	13	0	0	0	1	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0	1	0	0	1	0	1	1	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	1	1
outMinPu0.2	11	0	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	1	1	1	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0
outMinPu0.5	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
seedBP4	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.001	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.0001	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.1	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.01	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.1.seedNoGU	20	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.01.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.0001.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.001.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedMinPu0.2	15	0	0	0	0	0	1	0	0	1	0	1	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP4.seedMinPu0.2.seedNoGU	15	0	0	0	0	0	1	0	0	1	0	1	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP4.seedMinPu0.5	9	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	0	1	0	0	0	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0	0
seedBP4.seedMinPu0.5.seedNoGU	11	1	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0
seedBP4.seedMinPu0.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP4.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.01	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.001	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.0001	15	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.1	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP5.seedMinPu0.001.seedNoGU	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.0001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.1.seedNoGU	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	1	1	1	0	1	0	0	0	0	1	0	0	0	0	0	0	0
seedBP5.seedMinPu0.01.seedNoGU	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1
seedBP5.seedMinPu0.2	14	0	0	0	0	0	0	0	0	1	0	1	0	0	0	0	1	0	0	0	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP5.seedMinPu0.2.seedNoGU	16	0	0	0	0	0	0	1	0	1	0	1	0	0	0	0	1	0	0	0	0	1	1	0	1	1	1	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	0	0	0	0	0
seedBP5.seedMinPu0.5	7	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	0	1	1	0	0	0	0	0	0	0

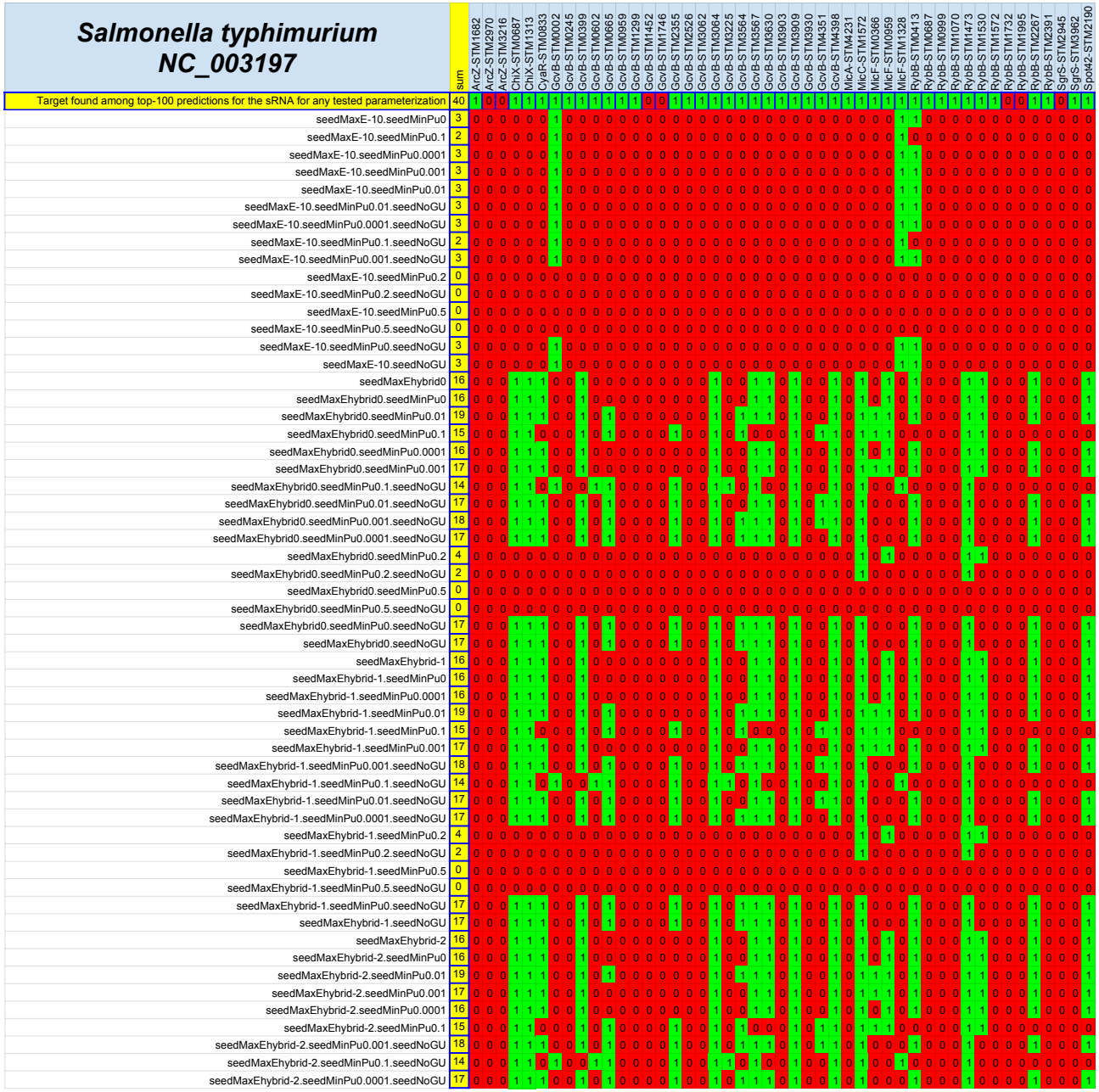

Salmonella typhimurium
NC_003197
sum
ArcZ-STM1682
ArcZ-STM2970
ArcZ-STM3216
ChiX-STM0687
ChiX-STM1313
CyaR-STM0833
GcvB-STM0002
GcvB-STM0245
GcvB-STM0399
GcvB-STM0602
GcvB-STM0665
GcvB-STM0959
GcvB-STM1299
GcvB-STM1452
GcvB-STM1746
GcvB-STM2355
GcvB-STM2526
GcvB-STM3062
GcvB-STM3064
GcvB-STM3225
GcvB-STM3564
GcvB-STM3567
GcvB-STM3630
GcvB-STM3903
GcvB-STM3909
GcvB-STM3930
GcvB-STM4351
GcvB-STM4398
MicA-STM4231
MicC-STM1572
MicF-STM0366
MicF-STM0959
MicF-STM1328
RybB-STM0413
RybB-STM0687
RybB-STM0999
RybB-STM1070
RybB-STM1473
RybB-STM1530
RybB-STM1572
RybB-STM1732
RybB-STM1995
RybB-STM2267
RybB-STM2391
SgrS-STM2945
SgrS-STM3962
Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization
seedBP5.seedMinPu0.5.seedNoGU
seedBP5.seedMinPu0.seedNoGU
seedBP5.seedNoGU
seedBP6
seedBP6.seedMinPu0
seedBP6.seedMinPu0.1
seedBP6.seedMinPu0.0001
seedBP6.seedMinPu0.001
seedBP6.seedMinPu0.01
seedBP6.seedMinPu0.0001.seedNoGU
seedBP6.seedMinPu0.1.seedNoGU
seedBP6.seedMinPu0.01.seedNoGU
seedBP6.seedMinPu0.001.seedNoGU
seedBP6.seedMinPu0.2
seedBP6.seedMinPu0.2.seedNoGU
seedBP6.seedMinPu0.5
seedBP6.seedMinPu0.5.seedNoGU
seedBP6.seedMinPu0.seedNoGU
seedBP6.seedNoGU
seedBP7
seedBP7.seedMinPu0
seedBP7.seedMinPu0.001
seedBP7.seedMinPu0.01
seedBP7.seedMinPu0.1
seedBP7.seedMinPu0.0001
seedBP7.seedMinPu0.0001.seedNoGU
seedBP7.seedMinPu0.001.seedNoGU
seedBP7.seedMinPu0.1.seedNoGU
seedBP7.seedMinPu0.01.seedNoGU
seedBP7.seedMinPu0.2
seedBP7.seedMinPu0.2.seedNoGU
seedBP7.seedMinPu0.5
seedBP7.seedMinPu0.5.seedNoGU
seedBP7.seedMinPu0.seedNoGU
seedBP7.seedNoGU
seedBP8
seedBP8.seedMinPu0
seedBP8.seedMinPu0.1
seedBP8.seedMinPu0.0001
seedBP8.seedMinPu0.001
seedBP8.seedMinPu0.01
seedBP8.seedMinPu0.001.seedNoGU
seedBP8.seedMinPu0.0001.seedNoGU
seedBP8.seedMinPu0.1.seedNoGU
seedBP8.seedMinPu0.01.seedNoGU
seedBP8.seedMinPu0.2
seedBP8.seedMinPu0.2.seedNoGU
seedBP8.seedMinPu0.5
seedBP8.seedMinPu0.5.seedNoGU
seedBP8.seedMinPu0.seedNoGU
seedBP8.seedNoGU
seedBP9
seedBP9.seedNoGU
seedBP10
seedBP10.seedNoGU
seedBP11

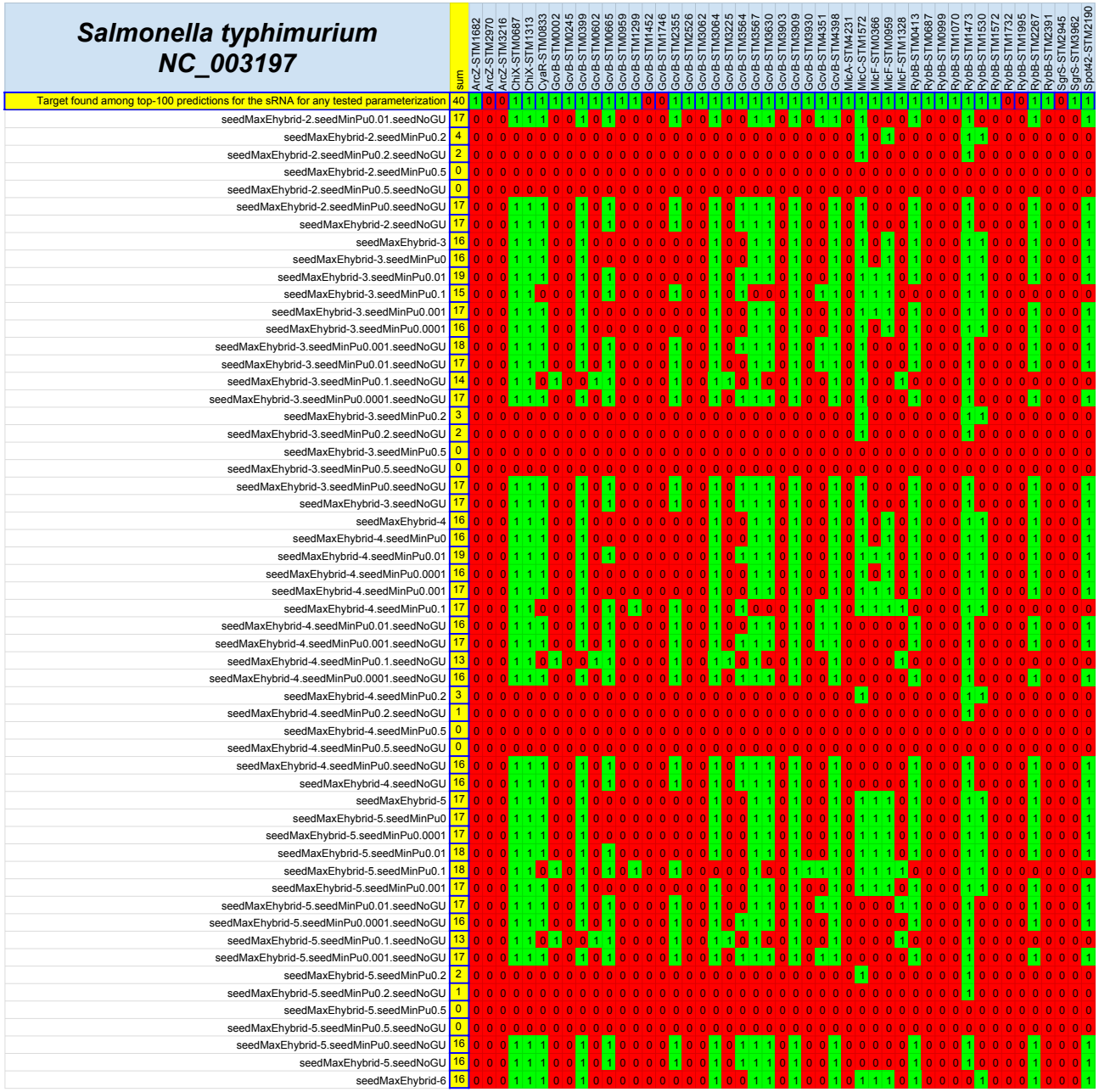
Salmonella typhimurium NC_003197	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
seedBP11.seedNoGU	6	0	0	0	1	1	0	0	0	0	1	1	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP12	10	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	1	0	1	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0
seedBP12.seedNoGU	5	0	0	0	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP13	5	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
seedBP13.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedBP14	5	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
seedBP14.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE0	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
seedMaxE0.seedMinPu0	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
seedMaxE0.seedMinPu0.01	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE0.seedMinPu0.001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE0.seedMinPu0.0001	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE0.seedMinPu0.1	15	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	0	0	0	1	0	1	1	0	1	1	1	1	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.01.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	0	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE0.seedMinPu0.001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE0.seedMinPu0.1.seedNoGU	14	0	0	0	1	1	0	1	0	0	1	1	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	1	0	1	0	1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.0001.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE0.seedMinPu0.2	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.2.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE0.seedMinPu0.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE0.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-1	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-1.seedMinPu0	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	0	1	0	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-1.seedMinPu0.001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-1.seedMinPu0.0001	16	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	0	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-1.seedMinPu0.01	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-1.seedMinPu0.1	15	0	0	0	1	1	0	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	0	0	0	1	0	1	1	0	1	1	1	1	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.1.seedNoGU	14	0	0	0	1	1	0	1	0	0	1	1	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	1	0	1	0	1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.0001.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-1.seedMinPu0.001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-1.seedMinPu0.01.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	0	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-1.seedMinPu0.2	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.2.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-1.seedMinPu0.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-1.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-2	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-2.seedMinPu0	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-2.seedMinPu0.01	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-2.seedMinPu0.1	16	0	0	0	1	1	0	0	0	1	0	1	0	1	0	0	1	0	0	1	0	1	0	0	0	1	0	1	1	0	1	1	1	1	1	1	0	0	0	0	1	1	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.0001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-2.seedMinPu0.001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	1	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	1
seedMaxE-2.seedMinPu0.0001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-2.seedMinPu0.1.seedNoGU	14	0	0	0	1	1	0	1	0	0	1	1	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	0	1	0	1	0	1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.01.seedNoGU	17	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	0	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-2.seedMinPu0.001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-2.seedMinPu0.2	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.2.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-2.seedMinPu0.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-2.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-3	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1

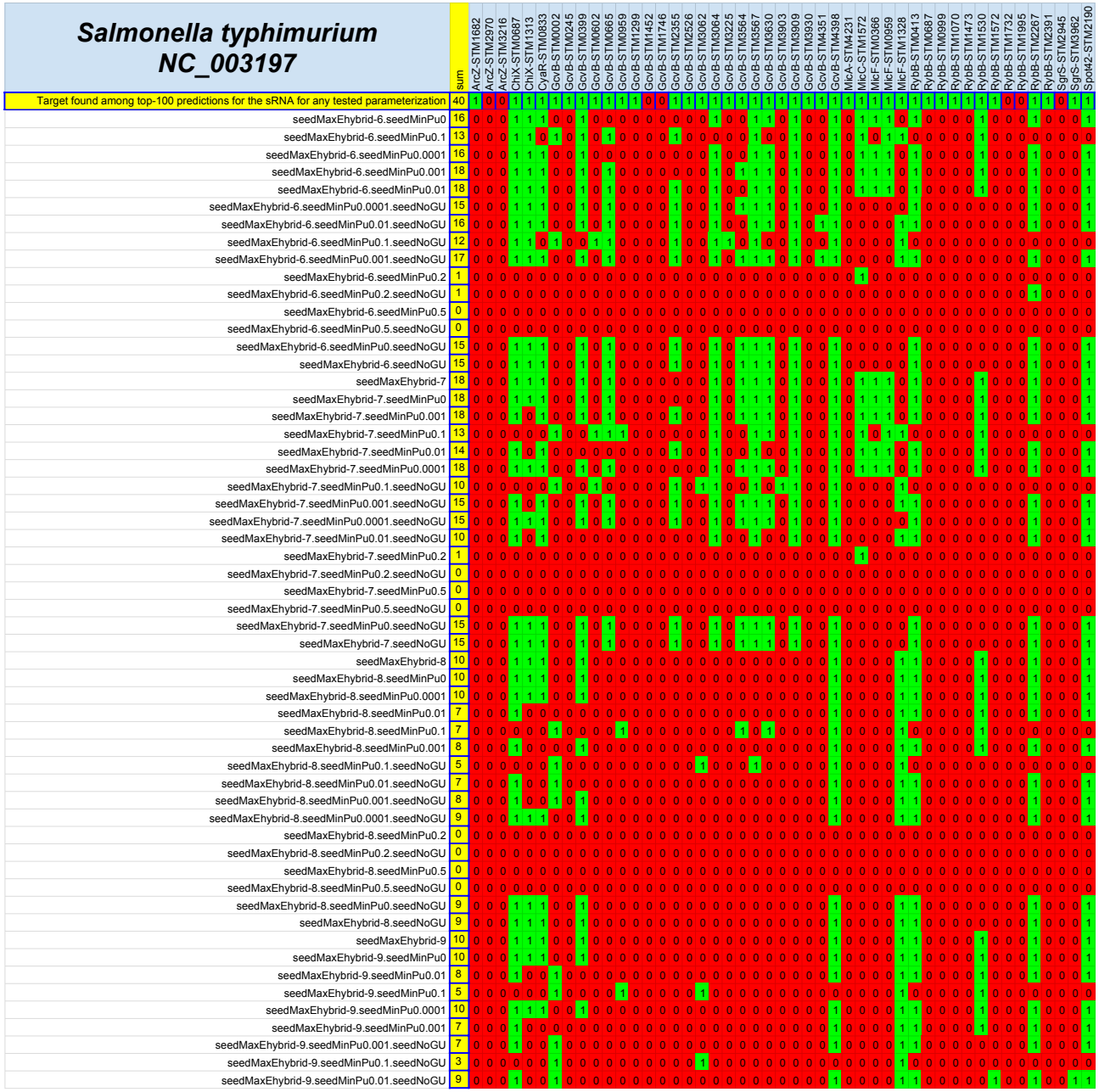
Salmonella typhimurium NC_003197

	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
seedMaxE-3.seedMinPu0	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.001	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.0001	17	0	0	0	1	1	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	1	1	0	1	0	0	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.1	17	0	0	0	1	1	0	0	0	1	0	1	1	0	1	0	0	1	1	0	0	0	0	1	1	0	0	1	1	1	1	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.01	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.01.seedNoGU	17	0	0	0	1	1	1	0	0	0	0	1	0	0	0	0	1	0	0	0	1	0	0	1	1	0	1	0	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.1.seedNoGU	14	0	0	0	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1	1	0	1	0	0	1	0	1	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.0001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	1	0	0	0	1	0	0	0	1	0	0	0	0	0	1	0	0	0	1
seedMaxE-3.seedMinPu0.2	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.2.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-3.seedMinPu0.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-3.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	1	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-4	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.01	17	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.001	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.1	15	0	0	0	1	1	0	1	0	0	0	1	0	1	0	0	0	0	0	0	0	1	0	0	1	0	1	1	0	1	1	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.0001	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.0001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	1	0	0	0	1	0	0	0	1	0	0	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.1.seedNoGU	14	0	0	0	1	1	0	1	0	0	0	1	0	0	0	0	0	0	0	1	1	0	1	0	1	1	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.001.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.01.seedNoGU	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-4.seedMinPu0.2	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.2.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-4.seedMinPu0.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-4.seedNoGU	18	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-5	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.1	13	0	0	0	0	1	0	1	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1	0	0	1	0	0	1	0	1	1	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.0001	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.001	19	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	1	1	1	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.01	17	0	0	0	1	1	1	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	1	0	0	1	0	1	1	0	1	0	0	0	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.0001.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.01.seedNoGU	13	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.1.seedNoGU	12	0	0	0	0	1	0	1	0	0	1	0	0	0	0	0	1	0	1	1	0	0	1	0	1	1	0	0	1	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.001.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-5.seedMinPu0.2	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.2.seedNoGU	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-5.seedMinPu0.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-5.seedNoGU	16	0	0	0	1	1	1	0	0	1	0	1	0	0	0	0	0	0	1	0	1	1	1	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	1	0	0	0	0	1	0	0	0	1
seedMaxE-6	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-6.seedMinPu0	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-6.seedMinPu0.01	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-6.seedMinPu0.0001	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-6.seedMinPu0.1	12	0	0	0	0	1	0	1	0	0	1	0	1	0	0	0	0	0	1	0	0	0	1	0	1	0	0	1	0	0	0	1	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.001	14	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	1	1	1	1	0	0	0	1	1	0	0	0	1	0	0	0	1
seedMaxE-6.seedMinPu0.001.seedNoGU	11	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0	1	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1
seedMaxE-6.seedMinPu0.1.seedNoGU	10	0	0	0	0	1	0	1	0	0	1	0	0	0	0	0	1	0	1	1	0	0	0	0	1	1	0	0	1	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.01.seedNoGU	11	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0	1	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1

Salmonella typhimurium NC_003197	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
seedMaxE-6.seedMinPu0.0001.seedNoGU	11	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-6.seedMinPu0.2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-6.seedMinPu0.seedNoGU	11	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-6.seedNoGU	11	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-7	9	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0	9	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0.01	8	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	0	0	0	1
seedMaxE-7.seedMinPu0.001	9	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0.0001	9	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0.1	9	0	0	0	0	0	0	1	0	0	1	0	1	0	0	0	0	0	0	0	0	0	1	0	1	0	0	1	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.1.seedNoGU	7	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	0	0	1	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.001.seedNoGU	8	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0.01.seedNoGU	7	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1
seedMaxE-7.seedMinPu0.0001.seedNoGU	10	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	1	0	0	0	0	0	1	0	0	1
seedMaxE-7.seedMinPu0.2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-7.seedMinPu0.seedNoGU	10	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	1	0	0	0	0	0	1	0	0	1
seedMaxE-7.seedNoGU	10	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	1	0	0	0	0	0	1	1	0	1	0	0	0	0	0	1	0	0	1
seedMaxE-8	10	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-8.seedMinPu0	10	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-8.seedMinPu0.01	8	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	0	0	0	1
seedMaxE-8.seedMinPu0.0001	10	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-8.seedMinPu0.001	9	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	0	0	1
seedMaxE-8.seedMinPu0.1	7	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.1.seedNoGU	4	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.001.seedNoGU	7	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	1	0	0
seedMaxE-8.seedMinPu0.0001.seedNoGU	8	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	1	0	0
seedMaxE-8.seedMinPu0.01.seedNoGU	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1
seedMaxE-8.seedMinPu0.2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-8.seedMinPu0.seedNoGU	8	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	1	0	0
seedMaxE-8.seedNoGU	8	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	1	0	0
seedMaxE-9	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedMinPu0	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedMinPu0.0001	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedMinPu0.1	3	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.01	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1
seedMaxE-9.seedMinPu0.001	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedMinPu0.001.seedNoGU	6	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedMinPu0.1.seedNoGU	3	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.01.seedNoGU	5	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1
seedMaxE-9.seedMinPu0.0001.seedNoGU	6	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	0	0
seedMaxE-9.seedMinPu0.2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxE-9.seedMinPu0.seedNoGU	6	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-9.seedNoGU	6	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1	0	0	1
seedMaxE-10	3	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0


Salmonella typhimurium
NC_003197
sum
ArcZ-STM1682
ArcZ-STM2970
ArcZ-STM3216
ChiX-STM0687
ChiX-STM1313
CyaR-STM0833
GcvB-STM0002
GcvB-STM0245
GcvB-STM0399
GcvB-STM0602
GcvB-STM0665
GcvB-STM0959
GcvB-STM1299
GcvB-STM1452
GcvB-STM1746
GcvB-STM2355
GcvB-STM2526
GcvB-STM3062
GcvB-STM3064
GcvB-STM3225
GcvB-STM3564
GcvB-STM3567
GcvB-STM3630
GcvB-STM3903
GcvB-STM3909
GcvB-STM3930
GcvB-STM4351
GcvB-STM4398
MicA-STM4231
MicC-STM1572
MicF-STM0366
MicF-STM0959
MicF-STM1328
RybB-STM0413
RybB-STM0687
RybB-STM0999
RybB-STM1070
RybB-STM1473
RybB-STM1530
RybB-STM1572
RybB-STM1732
RybB-STM1995
RybB-STM2267
RybB-STM2391
SgrS-STM2945
SgrS-STM3962
Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxE-10.seedMinPu0
seedMaxE-10.seedMinPu0.1
seedMaxE-10.seedMinPu0.0001
seedMaxE-10.seedMinPu0.001
seedMaxE-10.seedMinPu0.01
seedMaxE-10.seedMinPu0.01.seedNoGU
seedMaxE-10.seedMinPu0.0001.seedNoGU
seedMaxE-10.seedMinPu0.1.seedNoGU
seedMaxE-10.seedMinPu0.001.seedNoGU
seedMaxE-10.seedMinPu0.2
seedMaxE-10.seedMinPu0.2.seedNoGU
seedMaxE-10.seedMinPu0.5
seedMaxE-10.seedMinPu0.5.seedNoGU
seedMaxE-10.seedMinPu0.seedNoGU
seedMaxE-10.seedNoGU
seedMaxEhybrid0
seedMaxEhybrid0.seedMinPu0
seedMaxEhybrid0.seedMinPu0.01
seedMaxEhybrid0.seedMinPu0.1
seedMaxEhybrid0.seedMinPu0.0001
seedMaxEhybrid0.seedMinPu0.001
seedMaxEhybrid0.seedMinPu0.1.seedNoGU
seedMaxEhybrid0.seedMinPu0.01.seedNoGU
seedMaxEhybrid0.seedMinPu0.001.seedNoGU
seedMaxEhybrid0.seedMinPu0.0001.seedNoGU
seedMaxEhybrid0.seedMinPu0.2
seedMaxEhybrid0.seedMinPu0.2.seedNoGU
seedMaxEhybrid0.seedMinPu0.5
seedMaxEhybrid0.seedMinPu0.5.seedNoGU
seedMaxEhybrid0.seedMinPu0.seedNoGU
seedMaxEhybrid0.seedNoGU
seedMaxEhybrid-1
seedMaxEhybrid-1.seedMinPu0
seedMaxEhybrid-1.seedMinPu0.0001
seedMaxEhybrid-1.seedMinPu0.01
seedMaxEhybrid-1.seedMinPu0.1
seedMaxEhybrid-1.seedMinPu0.001
seedMaxEhybrid-1.seedMinPu0.001.seedNoGU
seedMaxEhybrid-1.seedMinPu0.1.seedNoGU
seedMaxEhybrid-1.seedMinPu0.01.seedNoGU
seedMaxEhybrid-1.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-1.seedMinPu0.2
seedMaxEhybrid-1.seedMinPu0.2.seedNoGU
seedMaxEhybrid-1.seedMinPu0.5
seedMaxEhybrid-1.seedMinPu0.5.seedNoGU
seedMaxEhybrid-1.seedMinPu0.seedNoGU
seedMaxEhybrid-1.seedNoGU
seedMaxEhybrid-2
seedMaxEhybrid-2.seedMinPu0
seedMaxEhybrid-2.seedMinPu0.01
seedMaxEhybrid-2.seedMinPu0.001
seedMaxEhybrid-2.seedMinPu0.0001
seedMaxEhybrid-2.seedMinPu0.1
seedMaxEhybrid-2.seedMinPu0.001.seedNoGU
seedMaxEhybrid-2.seedMinPu0.1.seedNoGU
seedMaxEhybrid-2.seedMinPu0.0001.seedNoGU


Salmonella typhimurium
NC_003197
sum
ArcZ-STM1682
ArcZ-STM2970
ArcZ-STM3216
ChiX-STM0687
ChiX-STM1313
CyaR-STM0833
GcvB-STM0002
GcvB-STM0245
GcvB-STM0399
GcvB-STM0602
GcvB-STM0665
GcvB-STM0959
GcvB-STM1299
GcvB-STM1452
GcvB-STM1746
GcvB-STM2355
GcvB-STM2526
GcvB-STM3062
GcvB-STM3064
GcvB-STM3225
GcvB-STM3564
GcvB-STM3567
GcvB-STM3630
GcvB-STM3903
GcvB-STM3909
GcvB-STM3930
GcvB-STM4351
GcvB-STM4398
MicA-STM4231
MicC-STM1572
MicF-STM0366
MicF-STM0959
MicF-STM1328
RybB-STM0413
RybB-STM0687
RybB-STM0999
RybB-STM1070
RybB-STM1473
RybB-STM1530
RybB-STM1572
RybB-STM1732
RybB-STM1995
RybB-STM2267
RybB-STM2391
SgrS-STM2945
SgrS-STM3962
Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxEhybrid-2.seedMinPu0.01.seedNoGU
seedMaxEhybrid-2.seedMinPu0.2
seedMaxEhybrid-2.seedMinPu0.2.seedNoGU
seedMaxEhybrid-2.seedMinPu0.5
seedMaxEhybrid-2.seedMinPu0.5.seedNoGU
seedMaxEhybrid-2.seedMinPu0.seedNoGU
seedMaxEhybrid-2.seedNoGU
seedMaxEhybrid-3
seedMaxEhybrid-3.seedMinPu0
seedMaxEhybrid-3.seedMinPu0.01
seedMaxEhybrid-3.seedMinPu0.1
seedMaxEhybrid-3.seedMinPu0.001
seedMaxEhybrid-3.seedMinPu0.0001
seedMaxEhybrid-3.seedMinPu0.001.seedNoGU
seedMaxEhybrid-3.seedMinPu0.01.seedNoGU
seedMaxEhybrid-3.seedMinPu0.1.seedNoGU
seedMaxEhybrid-3.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-3.seedMinPu0.2
seedMaxEhybrid-3.seedMinPu0.2.seedNoGU
seedMaxEhybrid-3.seedMinPu0.5
seedMaxEhybrid-3.seedMinPu0.5.seedNoGU
seedMaxEhybrid-3.seedMinPu0.seedNoGU
seedMaxEhybrid-3.seedNoGU
seedMaxEhybrid-4
seedMaxEhybrid-4.seedMinPu0
seedMaxEhybrid-4.seedMinPu0.01
seedMaxEhybrid-4.seedMinPu0.0001
seedMaxEhybrid-4.seedMinPu0.001
seedMaxEhybrid-4.seedMinPu0.1
seedMaxEhybrid-4.seedMinPu0.01.seedNoGU
seedMaxEhybrid-4.seedMinPu0.001.seedNoGU
seedMaxEhybrid-4.seedMinPu0.1.seedNoGU
seedMaxEhybrid-4.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-4.seedMinPu0.2
seedMaxEhybrid-4.seedMinPu0.2.seedNoGU
seedMaxEhybrid-4.seedMinPu0.5
seedMaxEhybrid-4.seedMinPu0.5.seedNoGU
seedMaxEhybrid-4.seedMinPu0.seedNoGU
seedMaxEhybrid-4.seedNoGU
seedMaxEhybrid-5
seedMaxEhybrid-5.seedMinPu0
seedMaxEhybrid-5.seedMinPu0.0001
seedMaxEhybrid-5.seedMinPu0.01
seedMaxEhybrid-5.seedMinPu0.1
seedMaxEhybrid-5.seedMinPu0.001
seedMaxEhybrid-5.seedMinPu0.01.seedNoGU
seedMaxEhybrid-5.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-5.seedMinPu0.1.seedNoGU
seedMaxEhybrid-5.seedMinPu0.001.seedNoGU
seedMaxEhybrid-5.seedMinPu0.2
seedMaxEhybrid-5.seedMinPu0.2.seedNoGU
seedMaxEhybrid-5.seedMinPu0.5
seedMaxEhybrid-5.seedMinPu0.5.seedNoGU
seedMaxEhybrid-5.seedMinPu0.seedNoGU
seedMaxEhybrid-5.seedNoGU
seedMaxEhybrid-6


Salmonella typhimurium
NC_003197
sum
ArcZ-STM1682
ArcZ-STM2970
ArcZ-STM3216
ChiX-STM0687
ChiX-STM1313
CyaR-STM0833
GcvB-STM0002
GcvB-STM0245
GcvB-STM0399
GcvB-STM0602
GcvB-STM0665
GcvB-STM0959
GcvB-STM1299
GcvB-STM1452
GcvB-STM1746
GcvB-STM2355
GcvB-STM2526
GcvB-STM3062
GcvB-STM3064
GcvB-STM3225
GcvB-STM3564
GcvB-STM3567
GcvB-STM3630
GcvB-STM3903
GcvB-STM3909
GcvB-STM3930
GcvB-STM4351
GcvB-STM4398
MicA-STM4231
MicC-STM1572
MicF-STM0366
MicF-STM0959
MicF-STM1328
RybB-STM0413
RybB-STM0687
RybB-STM0999
RybB-STM1070
RybB-STM1473
RybB-STM1530
RybB-STM1572
RybB-STM1732
RybB-STM1995
RybB-STM2267
RybB-STM2391
SgrS-STM2945
SgrS-STM3962
Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization
seedMaxEhybrid-6.seedMinPu0
seedMaxEhybrid-6.seedMinPu0.1
seedMaxEhybrid-6.seedMinPu0.0001
seedMaxEhybrid-6.seedMinPu0.001
seedMaxEhybrid-6.seedMinPu0.01
seedMaxEhybrid-6.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-6.seedMinPu0.01.seedNoGU
seedMaxEhybrid-6.seedMinPu0.1.seedNoGU
seedMaxEhybrid-6.seedMinPu0.001.seedNoGU
seedMaxEhybrid-6.seedMinPu0.2
seedMaxEhybrid-6.seedMinPu0.2.seedNoGU
seedMaxEhybrid-6.seedMinPu0.5
seedMaxEhybrid-6.seedMinPu0.5.seedNoGU
seedMaxEhybrid-6.seedMinPu0.seedNoGU
seedMaxEhybrid-6.seedNoGU
seedMaxEhybrid-7
seedMaxEhybrid-7.seedMinPu0
seedMaxEhybrid-7.seedMinPu0.001
seedMaxEhybrid-7.seedMinPu0.1
seedMaxEhybrid-7.seedMinPu0.01
seedMaxEhybrid-7.seedMinPu0.0001
seedMaxEhybrid-7.seedMinPu0.1.seedNoGU
seedMaxEhybrid-7.seedMinPu0.001.seedNoGU
seedMaxEhybrid-7.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-7.seedMinPu0.01.seedNoGU
seedMaxEhybrid-7.seedMinPu0.2
seedMaxEhybrid-7.seedMinPu0.2.seedNoGU
seedMaxEhybrid-7.seedMinPu0.5
seedMaxEhybrid-7.seedMinPu0.5.seedNoGU
seedMaxEhybrid-7.seedMinPu0.seedNoGU
seedMaxEhybrid-7.seedNoGU
seedMaxEhybrid-8
seedMaxEhybrid-8.seedMinPu0
seedMaxEhybrid-8.seedMinPu0.0001
seedMaxEhybrid-8.seedMinPu0.01
seedMaxEhybrid-8.seedMinPu0.1
seedMaxEhybrid-8.seedMinPu0.001
seedMaxEhybrid-8.seedMinPu0.1.seedNoGU
seedMaxEhybrid-8.seedMinPu0.01.seedNoGU
seedMaxEhybrid-8.seedMinPu0.001.seedNoGU
seedMaxEhybrid-8.seedMinPu0.0001.seedNoGU
seedMaxEhybrid-8.seedMinPu0.2
seedMaxEhybrid-8.seedMinPu0.2.seedNoGU
seedMaxEhybrid-8.seedMinPu0.5
seedMaxEhybrid-8.seedMinPu0.5.seedNoGU
seedMaxEhybrid-8.seedMinPu0.seedNoGU
seedMaxEhybrid-8.seedNoGU
seedMaxEhybrid-9
seedMaxEhybrid-9.seedMinPu0
seedMaxEhybrid-9.seedMinPu0.01
seedMaxEhybrid-9.seedMinPu0.1
seedMaxEhybrid-9.seedMinPu0.0001
seedMaxEhybrid-9.seedMinPu0.001
seedMaxEhybrid-9.seedMinPu0.001.seedNoGU
seedMaxEhybrid-9.seedMinPu0.1.seedNoGU
seedMaxEhybrid-9.seedMinPu0.01.seedNoGU

Salmonella typhimurium NC_003197	sum	ArcZ-STM1682	ArcZ-STM2970	ArcZ-STM3216	ChiX-STM0687	ChiX-STM1313	CyaR-STM0833	GcvB-STM0002	GcvB-STM0245	GcvB-STM0399	GcvB-STM0602	GcvB-STM0665	GcvB-STM0959	GcvB-STM1299	GcvB-STM1452	GcvB-STM1746	GcvB-STM2355	GcvB-STM2526	GcvB-STM3062	GcvB-STM3064	GcvB-STM3225	GcvB-STM3564	GcvB-STM3567	GcvB-STM3630	GcvB-STM3903	GcvB-STM3909	GcvB-STM3930	GcvB-STM4351	GcvB-STM4398	MicA-STM4231	MicC-STM1572	MicF-STM0366	MicF-STM0959	MicF-STM1328	RybB-STM0413	RybB-STM0687	RybB-STM0999	RybB-STM1070	RybB-STM1473	RybB-STM1530	RybB-STM1572	RybB-STM1732	RybB-STM1995	RybB-STM2267	RybB-STM2391	SgrS-STM2945	SgrS-STM3962	Spot42-STM2190
Target found among top-100 predictions for the sRNA for any tested parameterization	40	1	0	0	1	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1	1	0	1	1
seedMaxEhybrid-9.seedMinPu0.0001.seedNoGU	11	0	0	0	1	1	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-9.seedMinPu0.2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-9.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-9.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-9.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-9.seedMinPu0.seedNoGU	11	0	0	0	1	1	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-9.seedNoGU	11	0	0	0	1	1	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-10	8	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-10.seedMinPu0	8	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-10.seedMinPu0.1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.01	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1
seedMaxEhybrid-10.seedMinPu0.0001	7	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-10.seedMinPu0.001	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	0	0	0	1
seedMaxEhybrid-10.seedMinPu0.001.seedNoGU	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	1	1	0	0	1
seedMaxEhybrid-10.seedMinPu0.0001.seedNoGU	11	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	1	0	0	0	0	0	1	1	0	0	1
seedMaxEhybrid-10.seedMinPu0.01.seedNoGU	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	0	0	0	0	1	0	0	0	0	0	0	1
seedMaxEhybrid-10.seedMinPu0.1.seedNoGU	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.2.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.5.seedNoGU	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
seedMaxEhybrid-10.seedMinPu0.seedNoGU	10	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	0	0	0	0	0	0	1	1	0	0	1
seedMaxEhybrid-10.seedNoGU	10	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	1	0	1	0	0	0	0	0	0	1	1	0	0	1