## Training dataset curated from **R-SIM** database for the "miRNA" model in **RSAPred**

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1060	CCCN(CC(=O)N(CC C)CC(=O)N(CC(N)= O)CC1=CN(CCCNC( =O)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=O)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA AUUACGAAAGUAA UUGGAUCCGCAAG G	Targaprimir-96	Target_lig_62 7	Pri-miR-96 RNAC (UU loop mutant to AU basepair)	Target_159	5.3979400086 7204
1061	CCCN(CC(=O)N(CC C)CC(=O)N(CC(N)= O)CC1=CN(CCCNC( =O)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=O)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA UUUACGAAAGUAU AUGGAUCCGCAAG G	Targaprimir-96	Target_lig_62 7	Pri-miR-96 RNA1 (Drosha processing site only)	Target_160	5.9208187539 5238
1062	CCCN(CC(=0)N(CC C)CC(=0)N(CC(N)= O)CC1=CN(CCCNC( =0)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=0)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA CGAACGAAAGUUG GUGGAUCCGCAAG G	Targaprimir-96	Target_lig_62	Pri-miR-96 RNA2 (desired GG loop only)	Target_161	6.0457574905 6068
1063	CCCN(CC(=O)N(CC C)CC(=O)N(CC(N)= O)CC1=CN(CCCNC( =O)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=O)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA CGAUUUACGAAAG UAUAUGGUGGAUC CGCAAGG	Targaprimir-96	Target_lig_62 7	Pri-miR-96 RNA3 (both the Drosha target site and adjacent target site)	Target_162	4.6989700043 3602
1064	CCCN(CC(=0)N(CC C)CC(=0)N(CC(N)= O)CC1=CN(CCCNC( =0)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=0)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA CGAUUUACGAAAG UAUAUCGUGGAUC CGCAAGG	Targaprimir-96	Target_lig_62 7	Pri-miR-96 RNA4 (GG to GC loop mutation)	Target_163	5.9208187539 5238

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1065	CCCN(CC(=0)N(CC C)CC(=0)N(CC(N)= O)CC1=CN(CCCNC( =0)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=0)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	GGGAGAGGGUUUA CGAUUUACGAAAG UAAAUGGUGGAUC CGCAAGG	Targaprimir-96	Target_lig_62	Pri-miR-96 RNA5 (UU loop mutant to UA loop)	Target_164	5.8239087409 4432
1066	CCCN(CC(=0)N(CC C)CC(=0)N(CC(N)= O)CC1=CN(CCCNC( =0)CCCOC2=C(C=C (C=C2C(C) (C)C)C2=NC3=CC(= CC=C3N2)N2CC[NH +](C)CC2)C(C) (C)C)N=N1)C(=0)CC COC1=CC=CC(=C1) C1=NC2=CC(=CC=C 2N1)C1=NC2=CC(=C C=C2N1)N1CC[NH+] (C)CC1	CGCGAATTCGCGTT TTCGCGAATTCGCG	Targaprimir-96	Target_lig_62	Pri-miR-96 AT- rich DNA hairpin	Target_165	4.3010299956 6398
1190	CC(C) (C)C1=CC(=CC(=C1 OCCCC(=O)NCCCN =[N+]=[N-])C(C) (C)C)C2=NC3=C(N2) C=C(C=C3)N4CCN(C C4)C	UGGCCGAUUUUGG CACUAGCACAUUU UUGCUUGUGUCUC UCCGCUCUGAGCAA UCAUGUGCAGUGC CAAUAUGGGAAA	Benzimidazole 1 (2014)	Target_lig_67	5'UUU/3'AUA, miR-96 Precursor Processing Site	Target_202	5.8860566476 9316
1191	CC(C) (C)C1=CC(=CC1 OCCCC(=O)NCCCN =[N+]=[N-])C(C) (C)C)C2=NC3=C(N2) C=C(C=C3)N4CCN(C C4)C	UGGCCGAUUUUGG CACUAGCACAUUU UUGCUUGUGUCUC UCCGCUCUGAGCAA UCAUGUGCAGUGC CAAUAUGGGAAA	Benzimidazole 1 (2014)	Target_lig_67	5'CGAUUU/ 3'GGUAUA, Expanded miR- 96 Precursor Processing Site	Target_203	5.4685210829 5774
1271	CN1CCN(CC1)C2=C C3=C(C=C2)N=C(N3 )C4=CC5=C(C=C4)N =C(N5)C6=CC(=CC= C6)OCCCC(=O)NCC CN=[N+]=[N-]	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Targaprimir-210	Target_lig_70	Pre-miR-210	Target_227	6.1791420105 603
1331	COC1=CC=C(C=C1) NCC(CN2C3=C(C=C( C=C3)Br)C4=C2C=C C(=C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 2	Target_lig_72	miR-21 hairpin	Target_265	6.1549019599 8574
1332	COC1=CC(=C(C=C1) NCC(CN2C3=C(C=C( C=C3)Br)C4=C2C=C C(=C4)Br)O)OC	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_72	miR-21 hairpin	Target_265	5.5228787452 8034
1333	COC1=C(C=C(C=C1) NCC(CN2C3=C(C=C( C=C3)Br)C4=C2C=C C(=C4)Br)O)OC	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 4	Target_lig_72	miR-21 hairpin	Target_265	5.8538719643 2176
1334	CCOclecc(en1)NCC( Cn1c2ccc(cc2c2c1ccc( c2)Br)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_72	miR-21 hairpin	Target_265	5.3187587626 2441
1335	CCCOclecc(cc1)NCC (Cnlc2ccc(cc2c2clccc (c2)Br)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_73	miR-21 hairpin	Target_265	5.6575773191 7779
1336	OC(Cn1c2ccc(cc2c2c1 ccc(c2)Br)Br)CNc1ccc (cc1)N1CCOCC1	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 7	Target_lig_73	miR-21 hairpin	Target_265	5.6777807052 6608

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1338	CCOc1ccc(cc1Cl)NC C(Cn1c2ccc(cc2c2c1c cc(c2)Br)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_73	miR-21 hairpin	Target_265	5.5376020021 0104
1339	C1=CC=C(C=C1)NC C(CN2C3=C(C=C(C= C3)Br)C4=C2C=CC(= C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 10	Target_lig_73	miR-21 hairpin	Target_265	5.6777807052 6608
1342	CCOC1=CC=C(C=C1 )NCC(CN2C3=C(C=C (C=C3)Br)C4=C2C=C C(=C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 13	Target_lig_73	miR-21 hairpin	Target_265	5.9208187539 5238
1350	CCOC1=CC=C(C=C1 )NCC(CN2C3=C(C=C (C=C3)Cl)C4=C2C=C C(=C4)Cl)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_23	Target_lig_74	miR-21 hairpin	Target_265	5.4948500216 8009
1351	CCOc1ccc(cc1)NCC( Cn1c2ccc(cc2c(c1C)C )Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_24	Target_lig_74	miR-21 hairpin	Target_265	5
1352	CCOe1ccc(ce1)NCC( CN(e1ccc(ce1)Br)e1ce c(ce1)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_25	Target_lig_74	miR-21 hairpin	Target_265	5.6382721639 8241
1355	CC1=CC=CC1NC C(CN2C3=C(C=C(C= C3)Br)C4=C2C=CC(= C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_72	miR-21 hairpin	Target_265	5.6382721639 8241
1362	CC1=C(C=CC=C1Cl) NCC(CN2C3=C(C=C( C=C3)Br)C4=C2C=C C(=C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 8	Target_lig_73	miR-21 hairpin	Target_265	5.4948500216 8009
1365	C1=CC(=CC(=C1)Cl) NCC(CN2C3=C(C=C( C=C3)Br)C4=C2C=C C(=C4)Br)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_	Target_lig_73	miR-21 hairpin	Target_265	5.9208187539 5238
1366	OC(Cn1c2ccc(cc2c2c1 ccc(c2)Br)Br)CNc1ccc 2c(c1)cccc2	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 12	Target_lig_73	miR-21 hairpin	Target_265	6.0969100130 0806
1368	CC1=CC(=C(C=C1)N CC(CN2C3=C(C=C(C =C3)Br)C4=C2C=CC( =C4)Br)O)C	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Aniline_analog_ 14	Target_lig_73	miR-21 hairpin	Target_265	5.2924298239 0206
1369	C1=CC=C(C=C1)NC C(CN2C3=CC=CC=C 3C4=CC=CC=C42)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_17	Target_lig_73	miR-21 hairpin	Target_265	5.0809219076 2393
1370	CCOC1=CC=C(C=C1 )NCC(CN2C3=C(C=C (C=C3)Br)C4=CC=C C=C42)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_18	Target_lig_74	miR-21 hairpin	Target_265	5.6989700043 3602
1371	OC(Cn1c2ccc(cc2c2c1 cccc2)Br)CNc1ccc(cc 1)N1CCOCC1	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_19	Target_lig_74	miR-21 hairpin	Target_265	5.4948500216 8009
1372	CCOc1ccc(cc1)NCC( Cn1c2cc(Br)ccc2c2c1 cc(Br)cc2)O	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_20	Target_lig_74	miR-21 hairpin	Target_265	6
1373	OC(Cn1c2cc(Br)ccc2c 2c1cc(Br)cc2)CNc1cc c(cc1)N1CCOCC1	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_21	Target_lig_74	miR-21 hairpin	Target_265	5.6777807052 6608
1374	OC(Cn1c2cc(Br)ccc2c 2c1cc(Br)cc2)CNc1cc ccc1C	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_22	Target_lig_74	miR-21 hairpin	Target_265	5.5686362358 4101
1378	OC(CN(c1ccc(cc1)Br) c1ccc(cc1)Br)CNc1cc c(cc1)N1CCOCC1	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Carbazole_analo g_26	Target_lig_74	miR-21 hairpin	Target_265	5.4436974992 3271
1379	OC(CN(c1ccc(cc1)Br) c1ccc(cc1)Br)CNc1cc	GGGUUGACUGUUG AAUCUCAUGGCAA	Carbazole_analo g_27	Target_lig_74	miR-21 hairpin	Target_265	5.3565473235 1381

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	ccc1C	CCC					
1380	CNC(=0)C1=NC=CC (=C1)OC2=CC(=C(C =C2)NC(=0)NC3=CC (=C(C=C3)C1)C(F) (F)F)F	GGGUUGACUGUUG AAUCUCAUGGCAA CCC	Regorafenib	Target_lig_75	miR-21 hairpin	Target_265	6.1366771398 7954
1381	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )N)OC3C(C(C(O3)CO )OC4C(C(C(C(O4)CN )O)O)N)O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin	Target_lig_4	Pre-miR-372	Target_266	2.1307682802 6902
1382	NCC1OC(OC2C(Cn3 nnc(c3)Cn3cnc4c3ncn c4N)OC(C2O)OC2C( O)C(N)CC(C2OC2OC (CN)C(C(C2N)O)O)N )C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3a	Target_lig_75	Pre-miR-372	Target_266	3.3142582613 9774
1383	NCC1OC(OC2C(Cn3 nnc(c3)Cn3ccc(=O) [nH]c3=O)OC(C2O)O C2C(O)C(N)CC(C2O C2OC(CN)C(C(C2N) O)O)N)C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3b	Target_lig_75	Pre-miR-372	Target_266	2.9788107009 3006
1384	NCC1OC(OC2C(Cn3 nnc(c3)Cn3ccc(nc3=O )N)OC(C2O)OC2C(O) C(N)CC(C2OC2OC(C N)C(C(C2N)O)O)N)C (C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3c	Target_lig_75	Pre-miR-372	Target_266	3.2240256688 7063
1385	NCC1OC(OC2C(Cn3 nnc(c3)Cn3cnc4c3[nH]c(N)nc4=0)OC(C2O) OC2C(O)C(N)CC(C2 OC2OC(CN)C(C(C2N)O)O)N)C(C(C10)O)	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3d	Target_lig_75	Pre-miR-372	Target_266	3.8386319977 6503
1386	NCC1OC(OC2C(Cn3 nnc(c3)CCCC(=O)Nc 3cccc(c3)c3ncc(s3)NC (=O)C)OC(C2O)OC2 C(O)C(N)CC(C2OC2 OC(CN)C(C(C2N)O) O)N)C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3e	Target_lig_75	Pre-miR-372	Target_266	4.7958800173 4408
1387	NCC1OC(OC2C(Cn3 nnc(c3)Cc3nc4c(o3)cc c(c4)NC(=0)Nc3ccccc 3)OC(C2O)OC2C(O) C(N)CC(C2OC2OC(C N)C(C(C2N)O)O)N)C (C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3f	Target_lig_75	Pre-miR-372	Target_266	4.8569851997 4591
1388	CCCCNC(=0)Nc1ccc 2c(c1)nc(o2)Cc1nnn(c 1)CC1OC(C(C1OC1O C(CN)C(C(C1N)O)O) 0)OC1C(0)C(N)CC( C1OC1OC(CN)C(C(C 1N)O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3g	Target_lig_75	Pre-miR-372	Target_266	4.7055337738 3841
1389	NCC1OC(OC2C(Cn3 nnc(c3)Cn3cnc(c3)c3c ccc(c3)NC(=0)c3cccc c3)OC(C2O)OC2C(O) C(N)CC(C2OC2OC(C N)C(C(C2N)O)O)N)C (C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3h	Target_lig_75	Pre-miR-372	Target_266	4.7235381958 2676
1390	CCCCNC(=O)Nc1ccc c(c1)c1ncn(c1)Cc1nnn (c1)CC1OC(C(C1OC1	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU	Neomycin_nucl eobase_3i	Target_lig_75	Pre-miR-372	Target_266	4.3362990746 1035

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	OC(CN)C(C(C1N)O) O)O)OC1C(O)C(N)C C(C1OC1OC(CN)C(C (C1N)O)O)N	GGAAAGUGCUGCG ACAUUUGAGCGUC AC					
1391	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=0)NCC (=0)Nc3nc4c([nH]3)c ccc4)OC(C2O)OC2C( O)C(N)CC(C2OC2OC (CN)C(C(C2N)O)O)N )C(C(C10)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin_nucl eobase_3j	Target_lig_76	Pre-miR-372	Target_266	4.8297382846 0504
1392	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=0)Nc3c ccc(c3)c3csc(n3)NC(= O)C)OC(C2O)OC2C( O)C(N)CC(C2OC2OC (CN)C(C(C2N)O)O)N )C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-S	Target_lig_76	Pre-miR-372	Target_266	4.7958800173 4408
1393	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=0)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C10) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-3a	Target_lig_76	Pre-miR-372	Target_266	4.2798406965 9404
1394	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=0)Nc3c ccc(c3)c3css(n3)NC(= O)c3ccc(cc3)C1)OC(C 2O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)ON)C(C(C10) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-3b	Target_lig_76	Pre-miR-372	Target_266	4.3001622741 3275
1395	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)F)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-3c	Target_lig_76	Pre-miR-372	Target_266	4.6478174818 8864
1396	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)N)OC(C 2O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)ON)C(C(C1O) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-3d	Target_lig_76	Pre-miR-372	Target_266	4.5686362358 4101
1397	NCC1OC(OC2C(Cn3 nnc(c3)CCCCC(=O)N c3cccc(c3)c3csc(n3)N C(=O)C)OC(C2O)OC 2C(O)C(N)CC(C2OC 2OC(CN)C(C(C2N)O) O)N)C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-4a	Target_lig_76	Pre-miR-372	Target_266	4.5606673061 6974
1398	NCC1OC(OC2C(Cn3 nnc(c3)CCCCCC(=0) Nc3cccc(c3)c3csc(n3) NC(=0)C)OC(C2O)O C2C(O)C(N)CC(C2O C2OC(CN)C(C(2N) O)O)N)C(C(C1O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-4b	Target_lig_76	Pre-miR-372	Target_266	4.5867002359 1875
1399	NCC1OC(OC2C(Cn3 nnc(c3)c3ccc(cc3)C(= O)Nc3cccc(c3)c3csc(n 3)NC(=O)C)OC(C2O) OC2C(O)C(N)CC(C2	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC	Neomycin-4c	Target_lig_76	Pre-miR-372	Target_266	4.4710832997 2235

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	OC2OC(CN)C(C(C2N )O)O)N)C(C(C1O)O) N	AC					
1400	NCC1OC(OC2C(Cn3 nnc(c3)c3cccc(c3)C(= O)Nc3cccc(c3)c3csc(n 3)NC(=O)C)OC(C2O) OC2C(O)C(N)CC(C2 OC2OC(CN)C(C(C2N )O)O)N)C(C(C1O)O) N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-4d	Target_lig_76	Pre-miR-372	Target_266	4.3925449767 8533
1401	NCC1OC(OC2C(Cn3 nnc(c3)Oc3ccc(cc3)C( =O)Nc3cccc(c3)c3csc( n3)NC(=O)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-4e	Target_lig_77	Pre-miR-372	Target_266	4.4907974776 689
1402	NCC1OC(OC2C(Cn3 nnc(c3)Oc3cccc(c3)C( =0)Nc3cccc(c3)c3csc( n3)NC(=0)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C10) O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neomycin-4f	Target_lig_77	Pre-miR-372	Target_266	4.6307841425 8986
1403	CNC1C(OC2OC(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)C)C(C(C2O)O)N)O C2C(C1O)OC(C(C2) N)OC1C(N)CC(C(1 O)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Apramycin-6a	Target_lig_77	Pre-miR-372	Target_266	3.6819366650 3724
1404	NCC1OC(OC2C(N)C C(C(C2O)O)N)C(C(C 10Cn1nnc(c1)CC(=O) Nc1cccc(c1)c1csc(n1) NC(=O)C)O)N	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	Neamine-6b	Target_lig_77	Pre-miR-372	Target_266	3.4948500216 8009
1405	O=C(Cc1nnn(c1)COC 1C(N)CC(C(C10)O)N )Nc1cccc(c1)c1csc(n1) NC(=O)C	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	2-DOS-6c	Target_lig_77	Pre-miR-372	Target_266	3.1687703061 3294
1406	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)C)OC(C2O)OC2C( O)C(N)CC(C2OC2OC (CN)C(C(C2N)O)O)N )C(C(C1O)O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-S	Target_lig_76	Pre-miR-373	Target_267	4.9507819773 2982
1407	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-3a	Target_lig_76	Pre-miR-373	Target_267	4.3665315444 2041
1408	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)Cl)OC(C 2O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)ON)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-3b	Target_lig_76	Pre-miR-373	Target_267	4.4596705252 0913

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1409	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)F)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCCCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-3c	Target_lig_76	Pre-miR-373	Target_267	4.4723700991 2866
1410	NCC1OC(OC2C(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)c3ccc(cc3)N)OC(C 2O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)ON)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-3d	Target_lig_76	Pre-miR-373	Target_267	4.3233063903 7513
1411	NCC1OC(OC2C(Cn3 nnc(c3)CCCCC(=0)N c3cccc(c3)c3csc(n3)N C(=0)C)OC(C2O)OC 2C(O)C(N)CC(C2OC 2OC(CN)C(C(C2N)O) O)N)C(C(C1O)O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4a	Target_lig_76	Pre-miR-373	Target_267	4.6363880201 0786
1412	NCC1OC(OC2C(Cn3 nnc(c3)CCCCCC(=0) Nc3cccc(c3)c3csc(n3) NC(=0)C)OC(C2O)O C2C(O)C(N)CC(C2O C2OC(CN)C(C(C2N) O)O)N)C(C(C1O)O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4b	Target_lig_76	Pre-miR-373	Target_267	4.3746875490 3833
1413	NCC1OC(OC2C(Cn3 nnc(c3)c3ccc(cc3)C(= O)Nc3cccc(c3)c3csc(n 3)NC(=O)C)OC(C2O) OC2C(O)C(N)CC(C2 OC2OC(CN)C(C(C2N )O)O)N)C(C(C1O)O) N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4c	Target_lig_76	Pre-miR-373	Target_267	4.4762535331 8844
1414	NCC1OC(OC2C(Cn3 nnc(c3)c3cccc(c3)C(= O)Nc3cccc(c3)c3csc(n 3)NC(=O)C)OC(C2O) OC2C(O)C(N)CC(C2 OC2OC(CN)C(C(C2N )O)O)N)C(C(C1O)O) N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4d	Target_lig_76	Pre-miR-373	Target_267	4.4634415574 2847
1415	NCC1OC(OC2C(Cn3 nnc(c3)Oc3ccc(cc3)C( =O)Nc3cccc(c3)c3csc( n3)NC(=O)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4e	Target_lig_77	Pre-miR-373	Target_267	4.4248121550 7234
1416	NCC1OC(OC2C(Cn3 nnc(c3)Oc3cccc(c3)C( =O)Nc3cccc(c3)c3csc( n3)NC(=O)C)OC(C2 O)OC2C(O)C(N)CC( C2OC2OC(CN)C(C(C 2N)O)O)N)C(C(C1O) O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neomycin-4f	Target_lig_77	Pre-miR-373	Target_267	4.4736607226 1016
1417	CNC1C(OC2OC(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)C)C(C(C2O)O)N)O C2C(C1O)OC(C(C2) N)OC1C(N)CC(C(C1 O)O)N	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Apramycin-6a	Target_lig_77	Pre-miR-373	Target_267	3.9956786262 1736

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1418	NCC1OC(OC2C(N)C C(C(C2O)O)N)C(C(C 1OCn1nnc(c1)CC(=O) Nc1cccc(c1)c1csc(n1) NC(=O)C)O)N	GGGAUACUCAAAA UGGGGGCCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	Neamine-6b	Target_lig_77	Pre-miR-373	Target_267	3.9746941347 3523
1419	O=C(Cc1nnn(c1)COC 1C(N)CC(C(C1O)O)N )Nc1ecec(c1)c1ese(n1) NC(=O)C	GGGAUACUCAAAA UGGGGGCGCUUUC CUUUUUGUCUGUA CUGGGAAGUGCUU CGAUUUUGGGGUG UCCC	2-DOS-6c	Target_lig_77	Pre-miR-373	Target_267	2.5030703519 2679
1420	CC1(C(C2=CC=CC= C2NC1=NCCCNCCC CNCCCN)N(C)C3=C C=CC=C3)C	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	PA-1	Target_lig_77 5	Pre-miR-372	Target_266	6.8239087409 4432
1421	C(CCNCCCCNCCCC N)CN	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	PA-2	Target_lig_77	Pre-miR-372	Target_266	6.2757241303 9921
1422	NCCCCN/C(=C/1\ C(=O)C=C2C(C1=O) (C)c1c(O)c(C)c(c(c1O 2)C(=O)C)O)/C	GUGGGCCUCAAAU GUGGAGCACUAUU CUGAUGUCCAAGU GGAAAGUGCUGCG ACAUUUGAGCGUC AC	PA-3	Target_lig_77	Pre-miR-372	Target_266	5.9393021596 4639
1423	CN1CCN(CC1)C2=C C3=C(C=C2)N=C(N3 )C4=CC5=C(C=C4)N =C(N5)C6=CC(=CC= C6)OCCCC(=O)NCC CN=[N+]=[N-]	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Targapremir-210	Target_lig_12	Pre-miR-210	Target_227	6.7958800173 4408
1424	CN1CCN(CC1)C2=C C3=C(C=C2)N=C(N3 )C4=CC5=C(C=C4)N =C(N5)C6=CC(=CC= C6)OCCCC(=O)NCC CN=[N+]=[N-]	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Targapremir- 210-RL	Target_lig_12	Pre-miR-210	Target_227	6.7212463990 4717
1425	N=[N+]=NCCCNC(= O)CCCOclc(cc(cc1C( C) (C)C)c1nc2c([nH]1)cc c(c2)c1nc2c([nH]1)cc( cc2)N1CCN(CC1)C)C (C)(C)C	GGGAGAGGGUUUA AUUCUUACGAAAG UAAUAAUUGGAUC CGCAAGG	Targapremir- 515/885	Target_lig_77	Pri-miR-515 drosha site (RNA 1)	Target_268	5.0457574905 6068
1426	N=[N+]=NCCCNC(= O)CCCOclc(cc(cc1C( C) (C)C)c1nc2c([nH]1)cc c(c2)c1nc2c([nH]1)cc( cc2)N1CCN(CC1)C)C (C)(C)C	GGGAGAGGGUUUA AUUUCUACGAAAG UAGCGAUUGGAUC CGCAAGG	Targapremir- 515/885	Target_lig_77 8	Pri-miR-885 drosha site (RNA 2)	Target_269	4.9208187539 5238
1427	N=[N+]=NCCCNC(= O)CCCOc1c(cc(cc1C( C) (C)C)c1nc2c([nH]1)cc c(c2)c1nc2c([nH]1)cc( cc2)N1CCN(CC1)C)C (C)(C)C	GGGAGAGGGUUUA AUUCAUACGAAAG UAUUAAUUGGAUC CGCAAGG	Targapremir- 515/885	Target_lig_77	Pri-miR-515 (RNA 3)	Target_270	5.1249387366 083
1432	N=[N+]=NCCCNC(= O)CCCOclc(cc(cc1C( C) (C)C)c1nc2c([nH]1)cc c(c2)c1nc2c([nH]1)cc( cc2)N1CCN(CC1)C)C	GGGAGAGGGUUUA AUUCAUUCUACGA AAGUAGCGUUAAU UGGAUCCGCAAGG	Targapremir-515	Target_lig_77	Pri-miR-515 and the adjacent RNA motif (RNA 4)	Target_271	7.2218487496 1636

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	(C)(C)C						
1433	CCCN(C(=0)CN(C(= O)CNCc1nnn(c1)N=[ N+]=N)CCC)CC(=0) N(CC(=0)N(CC(=0) N(CC(=0)N)CCC)CC C)CCC	GGGAGAGGUUUA AUUCAUUCUACGA AAGUAGCGUUAAU UGGAUCCGCAAGG	Compound 1a	Target_lig_78	Pri-miR-515 and the adjacent RNA motif (RNA 4)	Target_271	4.7447274948 9669
1434	CCCN(C(=0)CNCc1n nn(c1)N=[N+]=Nn1nn c(c1)CO)CC(=0)N(C C(=0)N(CC(=0)N(C C(=0)N(CC(=0)N)C CC)CCC)CCC)CCC	GGGAGAGGGUUUA AUUCAUUCUACGA AAGUAGCGUUAAU UGGAUCCGCAAGG	Compound 1b	Target_lig_78	Pri-miR-515 and the adjacent RNA motif (RNA 4)	Target_271	4.6989700043 3602
1576	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )N)OC3C(C(C(O3)CO )OC4C(C(C(C(O4)CN )O)O)N)O)O)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Neomycin	Target_lig_4	Oligonucleotide 3	Target_282	5.3979400086 7204
1577	CC1(COC(C(C1NC)O )OC2C(CC(C(C2O)O C3C(CC=C(O3)CN)N )N)N)O	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Sisomycin	Target_lig_12	Oligonucleotide 3	Target_282	5.3979400086 7204
1578	[C@H]1([C@H] (C[C@H]([C@H] ([C@H]1O)O[C@@H ]1[C@@H] (C[C@@H]([C@@H] (O1)CN)O)N)N)N)O[ C@@H]1[C@H] ([C@@H]([C@H] ([C@@H] (O1)CNc1c2cccc2nc 2c1cccc2)O)N)O	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Tobramycin	Target_lig_54	Oligonucleotide 3	Target_282	5.0969100130 0806
1579	C1C(C(C(C1NC(= O)C(CCN)O)OC2C(C (C(C(O2)CO)O)N)O) O)OC3C(C(C(C(O3)C N)O)O)O)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Amikacin	Target_lig_17	Oligonucleotide 3	Target_282	4.8239087409 4432
1580	CC(C1CCC(C(O1)OC 2C(CC(C(C2O)OC3C( C(C(CO3) (C)O)NC)O)N)N)N)N C	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Gentamycin	Target_lig_12 70	Oligonucleotide 3	Target_282	4.7958800173 4408
1581	CNC1C(OC2OC(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)C)C(C(C2O)O)N)O C2C(C1O)OC(C(C2) N)OC1C(N)CC(C(C1 O)O)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Apramycin	Target_lig_77	Oligonucleotide 3	Target_282	4.6575773191 7779
1582	CCNC1CC(C(C(C10 C2C(C(C(CO2) (C)O)NC)O)O)OC3C( CC=C(O3)CN)N)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Netilmicin	Target_lig_12 71	Oligonucleotide 3	Target_282	4.4559319556 4972
1583	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )O)O)OC3C(C(C(O 3)CO)O)N)O)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Kanamycin-A	Target_lig_12 72	Oligonucleotide 3	Target_282	4.3565473235 1381
1584	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )N)OC3C(C(C(O3)CO )O)O)O)N	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Ribostamycin	Target_lig_12 73	Oligonucleotide 3	Target_282	4.0457574905 6068
1585	CC(C1C(C(C(O1)O C2C(CC(C(C2O)OC3 C(C(C(CO3) (C)O)NC)O)N)N)N)O )O)O	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Geneticin	Target_lig_14	Oligonucleotide 3	Target_282	3.9956786262 1736

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
1586	CC1C(C(C(O1)OC2C( C(C(C(C2O)O)N=C( N)N)O)N=C(N)N)OC 3C(C(C(C(O3)CO)O) O)NC)(C=O)O	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Streptomycin	Target_lig_63	Oligonucleotide 3	Target_282	3.8728952016 3519
1587	CNC1CC(C(C(C10)O C2C3C(C(C(O2)CO) O)OC4(O3)C(C(C(C O4)C(CO)N)O)O)O)O	CCGACUGAUGUUG ACUGUUGAAUCUC AUGGCAACACCAG UCGG	Hygromycin	Target_lig_3	Oligonucleotide 3	Target_282	3.5833594926 6172
1588	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )N)OC3C(C(C(O3)CO )OC4C(C(C(C(O4)CN )O)O)N)O)O)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Neomycin	Target_lig_4	Oligonucleotide 4	Target_283	5.5228787452 8034
1589	CC1(COC(C(C1NC)O )OC2C(CC(C(C2O)O C3C(CC=C(O3)CN)N )N)N)O	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Sisomycin	Target_lig_12	Oligonucleotide 4	Target_283	5.3979400086 7204
1590	[C@H]1([C@H] (C[C@H]([C@H] ([C@H]1O)O[C@@H ]1[C@@H] (C[C@@H]([C@@H] (O1)CN)O)N)N)O[ C@@H]1[C@H] ([C@@H]([C@H] ([C@@H] (O1)CNc1c2cccc2nc 2c1cccc2)O)N)O	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Tobramycin	Target_lig_54	Oligonucleotide 4	Target_283	5.0969100130 0806
1591	C1C(C(C(C1NC(= O)C(CCN)O)OC2C(C (C(C(O2)CO)O)N)O) O)OC3C(C(C(C(O3)C N)O)O)O)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Amikacin	Target_lig_17	Oligonucleotide 4	Target_283	4.8538719643 2176
1592	CC(C1CCC(C(O1)OC 2C(CC(C(C2O)OC3C( C(C(CO3) (C)O)NC)O)N)N)N)N C	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Gentamycin	Target_lig_12 70	Oligonucleotide 4	Target_283	4.8860566476 9316
1593	CNC1C(OC2OC(Cn3 nnc(c3)CCC(=O)Nc3c ccc(c3)c3csc(n3)NC(= O)C)C(C(C2O)O)N)O C2C(C1O)OC(C(C2) N)OC1C(N)CC(C(C1 O)O)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Apramycin	Target_lig_77	Oligonucleotide 4	Target_283	4.6382721639 8241
1594	CCNC1CC(C(C(C10 C2C(C(C(C02) (C)0)NC)0)0)OC3C( CC=C(03)CN)N)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Netilmicin	Target_lig_12	Oligonucleotide 4	Target_283	4.4685210829 5774
1595	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )O)O)OC3C(C(C(C(O 3)CO)O)N)O)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Kanamycin-A	Target_lig_12 72	Oligonucleotide 4	Target_283	4.3467874862 2466
1596	C1C(C(C(C(C1N)OC2 C(C(C(C(O2)CN)O)O )N)OC3C(C(C(O3)CO )O)O)O)N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Ribostamycin	Target_lig_12 73	Oligonucleotide 4	Target_283	4.1079053973 0952
1597	CC(C1C(C(C(O1)O C2C(CC(C(C2O)OC3 C(C(C(CO3) (C)O)NC)O)N)N)N)O )O)O	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Geneticin	Target_lig_14	Oligonucleotide 4	Target_283	4.0268721464 003
1598	CC1C(C(C(O1)OC2C( C(C(C(C2O)O)N=C( N)N)O)N=C(N)N)OC 3C(C(C(C(O3)CO)O)	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Streptomycin	Target_lig_63	Oligonucleotide 4	Target_283	3.9393021596 4639

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	O)NC)(C=O)O						
1599	CNC1CC(C(C(C10)O C2C3C(C(C(O2)CO) O)OC4(O3)C(C(C(C O4)C(CO)N)O)O)O)O )N	CCGACUGAUGUUG AXUGUUGAAUYUC AUGGCAACACCAG UCGG	Hygromycin	Target_lig_3	Oligonucleotide 4	Target_283	3.5361070110 1409
1729	N=[N+]=NCCCNC(= O)CCCOc1ccc(cc1)c1 nc2c([nH]1)cc(cc2)c1c cc(cc1)N1CCN(CC1) C	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Compound1	Target_lig_96	Pre-miR-21	Target_227	4.7447274948 9669
1730	CN1CCN(CC1)e1ece( ec1)e1ece2e(e1) [nH]e(n2)e1ece(ec1)O CCCC(=0)N1CCN(C C1)e1nnec2e1ecs2	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Compound9	Target_lig_96	Pre-miR-21	Target_227	6
1731	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=0)N(Cc1ccnc c1)CCNC(=0)c1ccc(c (c1)C)F	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Compound7	Target_lig_96	Pre-miR-21	Target_227	6.8927900303 5213
1732	Coclccc(cc1OCC1=N C2N(C1)C=CC=C2)N C(=0)CCCOclccc(cc 1)c1[nH]c2c(n1)ccc(c 2)c1ccc(cc1)N1CCN( CC1)C	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	Compound8	Target_lig_96	Pre-miR-21	Target_227	6.5702477199 9759
1733	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=O)N1CCN(C C1)c1nncc2c1ccs2	GUUGACUGUUGAA UCUCAUGGCAAC	Compound9	Target_lig_96	miR-21 WT	Target_301	6.4736607226 1016
1734	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=O)N1CCN(C C1)c1nncc2c1ccs2	GUUGACUGUUGAA UCUCAAUGGUCAA C	Compound9	Target_lig_96	miR-21 BP	Target_302	5
1735	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=O)N1CCN(C C1)c1nncc2c1ccs2	GUUGACUGUUGAA UCUCAAUGGUCAA C	Compound9	Target_lig_96	miR-21 A bulge	Target_303	6.6382721639 8241
1736	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=O)N1CCN(C C1)c1nncc2c1ccs2	GUUGACUGUUGAA UCUCAUGGUCAAC	Compound9	Target_lig_96	miR-21 U bulge	Target_304	5
1737	CN1CCN(CC1)c1ccc( cc1)c1ccc2c(c1) [nH]c(n2)c1ccc(cc1)O CCCC(=O)N1CCN(C C1)c1nncc2c1ccs2	GUUGACUGUUGAA UCUCAUAGCAAC	Compound9	Target_lig_96	miR-21 G-A	Target_305	5.4814860601 2211
1775	C1=CC(=CC(=C1)C2 =NC3=C(N2)C=C(C= C3)N)C4=NC5=C(N4 )C=C(C=C5)N	GCAGAGGUUGCCC UUGGUGAAUUCGC UUUAUUUAUGUUG AAUCACACAAAGG CAACUUUUGU	Compound C1	Target_lig_99	Pre-miR-377 dicer processing site	Target_312	4.5376020021 0104
1776	CN1CCN(CC1)e1ec( Ne2nece(e2)Ne2ece3e (n2)ecce3)ne(n1)C	GCAGAGGUUGCCC UUGGUGAAUUCGC UUUAUUUAUGUUG AAUCACACAAAGG CAACUUUUGU	Compound C2	Target_lig_99 5	Pre-miR-377 5'-G_G/3'-CAC bulge	Target_313	5.5228787452 8034
1777	CCOC(=0)CCCOC1= CC(=CC(=C1)C2=NC 3=C(N2)C=C(C=C3) N)C4=NC5=C(N4)C=	GCAGAGGUUGCCC UUGGUGAAUUCGC UUUAUUUAUGUUG AAUCACACAAAGG	Compound C1- COOEt	Target_lig_99	Pre-miR-377 dicer processing site	Target_312	6.0969100130 0806

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	C(C=C5)N	CAACUUUUGU					
1778	C#CCc1nc(Nc2nccc(c 2)Nc2ccc3c(n2)cccc3) cc(n1)N1CCN(CC1)C	GCAGAGGUUGCCC UUGGUGAAUUCGC UUUAUUUAUGUUG AAUCACACAAAGG CAACUUUUGU	Compound C2- Ak	Target_lig_99	Pre-miR-377 5'-G_G/3'-CAC bulge	Target_313	5.7958800173 4408
1779	CCCN(C(=0)CN(C(= O)CN(C(=O)CN(C(= O)CCCOc1cc(cc(1)c 1[nH]c2c(n1)ccc(c2)N )c1[nH]c2c(n1)ccc(c2) N)CCC)CCC)CCC)C C(=0)N(CC(=0)N)C CCn1cnc(c1)CN1CCN (CC1)c1nc(CC)nc(c1) Nc1nccc(c1)Nc1ccc2c (n1)cccc2	GCAGAGGUUGCCC UUGGUGAAUUCGC UUUAUUUAUGUUG AAUCACACAAAGG CAACUUUUGU	Targapremir-377	Target_lig_99	Pre-miR-377	Target_312	6.7212463990 4717
1794	N=[N+]=NCCCNC(= O)CCCOclecc(cc1)c1 nc2c([nH]1)ccc(c2)c1c cc(cc1)N1CCN(CC1) C	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	miR-21 binder compound 1	Target_lig_10 08	miR-21	Target_227	4.6989700043 3602
1795	CCCN(C(=0)CN(C(= 0)NCc1nnn(c1)CCCN C(=0)CCCOc1ccc(cc 1)c1nc2c([nH]1)ccc(c 2)c1ccc(cc1)N1CCN( CC1)C)CCC(=0) N(CC(=0)N(Cc1nnn( c1)CCCNC(=0)CCC Oc1ccc(cc1)c1nc2c([n H]1)ccc(c2)c1ccc(cc1) N1CCN(CC1)C)CC(= 0)N)CCC	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	miR-21 binder compound 2	Target_lig_10 09	miR-21	Target_227	6.0087739243 0751
1797	NC(=[NH2+])C1=CC =C(NC2=CC=C(C=C 2)C2=CC3=CC=C(C= C3N2)C(N)=[NH2+]) C=C1	GUGCAGGUAGUGA UAUGUGCAUCUAC UGCAC	Compound 2	Target_lig_67	miR-17 Dicer site	Target_316	6.9208187539 5238
1798	NC(=[NH2+])C1=CC =C(NC2=CC=C(C=C 2)C2=CC3=CC=C(C= C3N2)C(N)=[NH2+]) C=C1	GUGCAGGUAGUGA UAUGUGCAUCUAC UGCAC	Compound 2	Target_lig_67	miR-17 G- bulge	Target_317	4.8239087409 4432
1799	NC(=[NH2+])C1=CC =C(NC2=CC=C(C=C 2)C2=CC3=CC=C(C= C3N2)C(N)=[NH2+]) C=C1	GUGCAGGUAGUGA UAUGUGCAUCUAC CUGCAC	Compound 2	Target_lig_67	miR-17 U- bulge	Target_318	4.7212463990 4717
1800	NC(=[NH2+])C1=CC =C(NC2=CC=C(C=C 2)C2=CC3=CC=C(C= C3N2)C(N)=[NH2+]) C=C1	GUGCAGGUAGAUG AUAUGUGCAUCUA CCUGCAC	Compound 2	Target_lig_67	miR-17 Base pairing control	Target_319	4
1801	NC(=[NH2+])C1=CC =C(NC2=CC=C(C=C 2)C2=CC3=CC=C(C= C3N2)C(N)=[NH2+]) C=C1	UGCAGAUAGUGAA GUAGAUAGGCAUC UACUGCA	Compound 2	Target_lig_67	miR-18a Dicer site	Target_320	6.9065783148 3777
1809	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GCUGAUUUCUUUU GGUGUUCAGAGGG GUCGGUGGUGGUG GUGGUGGUGGUUC UAGCACCAUCUGA AAUCGGUUA	CMBL-3aL	Target_lig_10	R-Seq1	Target_321	6.6615435063 954
1810	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/	GCUGAUUUCUUUU GGUGUUCAGAGUG GUCGGUGGUUGGU	CMBL-3aL	Target_lig_10 18	R-Seq2	Target_322	5.8827287043 4424

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GGUGGGUGGU UCUAGCACCAUCUG AAAUCGGUUA					
1811	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GCUGAUUUCUUUU GGUGUUCAGAGAG GUCGGUGGUUGGU GGUGGGUGGUGGU UCUAGCACCAUCUG AAAUCGGUUA	CMBL-3aL	Target_lig_10	R-Seq3	Target_323	6.1662156253 4352
1812	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GCUGAUUUCUUUU GGUGUUCAGAGUG GGGUGAGCUUUCU AGCACCAUCUGAA AUCGGUUA	CMBL-3aL	Target_lig_10	R-Seq4	Target_324	6.8696662315 0499
1813	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GUGCAUUGUAGUU GCAUUGCAUGUUC UGGUGGUACCCAU CGAAUGUUUCCAC AGUGCAUCAC	CMBL-3aL	Target_lig_10	Pre-miR-33a	Target_325	8.8601209135 9876
1814	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GUGCAUUGUAGUU GCAUUGCAUGUUC UGGUAGUACCCAU CGAAUGUUUCCAC AGUGCAUCAC	CMBL-3aL	Target_lig_10	Pre-miR-33a- G31A	Target_326	8.6497519816 6584
1815	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	GUGCAUUGUAGUU GCAUUGCAUGUUC UGGUGAUACCCAU CGAAUGUUUCCAC AGUGCAUCAC	CMBL-3aL	Target_lig_10	Pre-miR-33a- G32A	Target_327	8.5934598195 6604
1816	NCCCC(=O)NCCCN 1Cc2ccc3c(n2)nc(cc3) NC(=O)OCC/C=C/ CCOC(=O)Nc2nc3nc( C1)ccc3cc2	UGCCUACUGAGCU GAAACACAGUUGG UUUGUGUACACUG GCUCAGUUCAGCA GGAACAG	CMBL-3aL	Target_lig_10	Pre-miR-24-2	Target_328	8.6903698325 741
1820	N=[N+]=NCCCNC(= O)CCCOclecc(c1)c1 nc2c([nH]1)ccc(c2)c1 nc2c([nH]1)cc(cc2)N1 CCN(CC1)C	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	SMB1	Target_lig_12 78	Pre-miR-210	Target_227	6.7958800173 4408
1821	CCCN(C(=0)NCc1nn n(c1)CCCCC(=0)CC COc1c(cc(cc1C(C) (C)C)c1[nH]c2c(n1)cc (cc2)N1CCN(CC1)C) C(C) (C)C)CC(=0)N(C(=0) )CCCOc1ccc(cc1)c1[n H]c2c(n1)cc(cc2)N1C CN(CC1)CCC	GGGAGAGGGUUUA AUUACGAAAGUAA UUGGAUCCGCAAG G	SMB2	Target_lig_10 22	Pri-miR-96	Target_159	7.6989700043 3602
1822	CCCN(C(=0)CN(C(= O)NCc1nnn(c1)CCCC C(=0)CCCOc1ccc(cc 1)c1nc2c([nH]])ccc(c 2)c1ccc(cc1)N1CCN( CC1)C)CCC)CC(=0) N(CC(=0)N(Cc1nnn( c1)CCCCC(=0)CCC Oc1ccc(cc1)c1nc2c([n H]1)ccc(c2)c1ccc(cc1) N1CCN(CC1)C)CC(= O)N)CCC	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	SMB3	Target_lig_10 23	Pre-miR-21	Target_227	6.0087739243 0751
1823	CCCN(C(=0)CN(C(= 0)NCc1nnn(c1)CCCC C(=0)CCCOc1c(cc(cc 1C(C) (C)C)c1nc2c([nH]1)cc c(c2)c1ccc(cc1)N1CC N(CC1)C)C(C) (C)C)CCC)CC(=0)N(	UAGCUUAUCAGAC UGAUGUUGACUGU UGAAUCUCAUGGC AACACCAGUCGAU GGGCUGU	SMB4	Target_lig_12 79	Pre-miR- 17/18a/20a	Target_331	6.9208187539 5238

Entry_ID	SMILES	Target_RNA_sequence	Molecule_name	Molecule_ID	Target_RNA_ name	Target_RNA_ ID	pKd
	CC(=0)N(Cc1nnn(c1) CCCCC(=0)CCCOc1 c(cc(c1C(C) (C)C)c1nc2c([nH]1)cc c(c2)c1ccc(cc1)N1CC N(CC1)C)C(C) (C)C)CC(=0)N)CCC						
1824	OC(COP(=O) (OC1C(O)C(OC1n1cn c2c1ncnc2N)COP(=O) (OC1C(O)C(OC1n1cn c2c1ncnc2N)COP(=O) (OC1C(O)C(OC1n1cn c2c1ncnc2N)COP(=O) (OC1C(O)C(OC1n1cn c2c1ncnc2N)COP(=O) (O(O)O(OOC1n1cn c2c1ncnc2N)COP(=O) (O(O)O(OOC)CCCCCCCCCCCCCCCCCCCCCCCCCC	AGCCCCUGCCCACC GCACACUGCGCUGC CCCAGACCCACUGU GCGUGUGACAGCG GCUGA	SMC1	Target_lig_10 24	Pre-miR-210	Target_227	6.7212463990 4717
1825	CCCN(C(=0)N(Cc1n nn(c1)CCCCC(=0)CC COc1c(cc(cc1C(C) (C)C)c1nc2c([nH]1)cc c(c2)N1CCN(CC1)C) C(C) (C)CCCCCCCCCCOP (=0) (OC1C(O)C(OC1n1cn c2c1ncnc2N)COP(=0) (OC1C(O)C(OC1n1cn c2c1ncnc2C([nH]1)C cc(c2)N1CCN(CC1)C)	GGGAGAGGGUUUA AUUACGAAAGUAA UUGGAUCCGCAAG G	SMC2	Target_lig_10 25	Pri-miR-96	Target_159	7.0757207139 3812
1826	CCCN(C(=0)N(Cc1n nn(c1)CCCCC(=0)CC COc1c(cc(cc1C(C) (C)C)c1nc2c([nH]1)cc c(c2)N1CCN(CC1)C) C(C) (C)CCC(=0)NCCCC NCCCNC(=0)c1csc(n 1)c1csc(n1)CCNC(=0)C(C(NC(=0)C(C(C(C(C(C(C(C(0)C(=0)N)C(C(C(C(C(C(0)C(=0)N)C(C(C(C(C(C(0)C(=0)N)N)C(C(=0)N)N)C(C(=0)N)N)C(C(C(=0)N)N)C(C(=0)N)N(C(C(=0)N)C(C(C(=0)N)N)C(C(=0)N)N(C(C(=0)N)C(C(=0)C(C(=0)N)C((=0)C(C(=0)N)C((=0)C(C(=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)C((=0)N)C((=0)N)C((=0)C((=0)	GGGAGAGGGUUUA AUUACGAAAGUAA UUGGAUCCGCAAG G	SMC2-2	Target_lig_10 26	Pri-miR-96	Target_159	7.1938200260 1611