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## Design, synthesis and biological evaluation of 2-aminoquinazolin-4-(3H)-one derivatives as potential SARS-CoV-2 and MERS-CoV treatments

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### ABSTRACT

Despite the rising threat of fatal coronaviruses, there are no general proven effective antivirals to treat them. 2-Aminoquinazolin-4(3H)-one derivatives were newly designed, synthesized, and investigated to show the inhibitory effects on SARS-CoV-2 and MERS-CoV. Among the synthesized derivatives, 7-chloro-2-((3,5-dichlorophenyl)amino)quinazolin-4(3H)-one (**9g**) and 2-((3,5-dichlorophenyl)amino)-5-hydroxyquinazolin-4(3H)-one (**11e**) showed the most potent anti-SARS-CoV-2 activities ( $IC_{50} < 0.25 \mu M$ ) and anti-MERS-CoV activities ( $IC_{50} < 1.1 \mu M$ ) with no cytotoxicity ( $CC_{50} > 25 \mu M$ ). In addition, both compounds showed acceptable results in metabolic stabilities, hERG binding affinities, CYP inhibitions, and preliminary PK studies.

Coronavirus is a single positive-stranded RNA virus that was discovered in 1960 while looking for a new cold virus that infects the upper respiratory tract.<sup>1–2</sup> The clinical significance was relatively low because of its weak symptoms.<sup>3</sup> However, with the outbreak of SARS-CoV in 2003, its clinical significance has received new attention.<sup>3–4</sup> SARS-CoV spread to 4 countries, with 8,422 confirmed cases and 916 deaths, with its mortality rate of 11%, raising the public health awareness.<sup>4–5</sup> In 2012, a new outbreak of MERS-CoV occurred and spread to 27 countries by January 2020, resulting in 2519 confirmed cases and 866 deaths, with a mortality rate of 34%.<sup>6</sup> Most recently, COVID-19, which was caused by SARS-CoV-2 outbreak in Wuhan, China, in December 2019, has spread worldwide, causing 40 million confirmed cases and 1 million deaths.<sup>7</sup> Eventually, on March 11, 2020, COVID-19 was declared as third pandemic after the 1968 Hong Kong flu and 2009 influenza by the WHO.<sup>8–9</sup>

For treatments of MERS-CoV, it is generally recommended to use drugs such as interferon, immunomodulatory factor, and antiviral drugs such as ribavirin or lopinavir.<sup>10</sup> However, there are reports that interferon and ribavirin may cause side effects such as poor bone marrow function, anemia, and virus mutations.<sup>11</sup> In addition, although the

monkey model showed a therapeutic effect<sup>12</sup>, it did not show a great effect in actual clinical trials, requiring the development of a safer and more efficient MERS treatment.<sup>13</sup>

In the case of COVID-19, Remdesivir, which received urgent approval, as well as Nafamostat, and Hydroxychloroquine, are considered as promising therapeutic candidates.<sup>14</sup> However, due to side effects and low clinical effects,<sup>15</sup> safer and more effective treatments need to be developed.

Our research for novel antivirals inhibiting these fatal coronaviruses started with a biochemical high content screening (HCS) of a library containing 200,000 compounds from Korea Chemical Bank.<sup>16</sup> In the past few years, we reported inhibitors of MERS-CoV such as 2-phenylchroman-4-one derivatives,<sup>17</sup> 3-acyl-2-phenylamino-1,4-dihydroquinolin-4(1H)-one derivatives,<sup>18</sup> and 4-aniline-6-amino-quinazolin derivatives<sup>19</sup> as well as inhibitors of SARS-CoV-2 such as cyclic sulfonamide derivatives.<sup>20</sup>

In this study, the core scaffold of 3-acyl-2-phenylamino-1,4-dihydroquinolin-4(1H)-one of our lead compound (**1**)<sup>18</sup> was changed to 2-amino-quinazolin-4(3H)-one (Fig. 1). Because the aromatic rings substituted with halogen groups or electron withdrawing groups of

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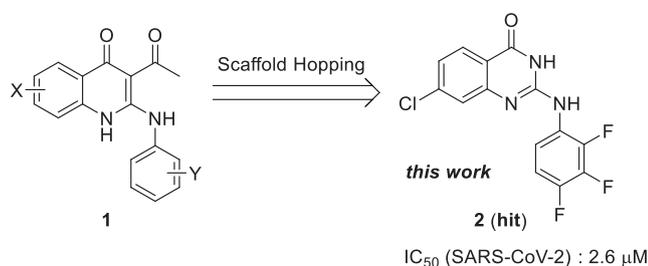


Fig. 1. Our new design of anti-coronavirus Scaffold.

dihydroquinolinones **1** were important for activity, 2-amino-quinazolin-4(3*H*)-ones with aromatic rings fixed at the similar positions were designed and evaluated to find compound **2** as a hit ( $IC_{50} = 2.6 \mu\text{M}$  for SARS-CoV-2). 2-Amino-quinazolin-4(3*H*)-one derivatives have been known to possess high activities as inhibitors of aldose reductase,<sup>21a</sup>  $K_{ATP}$  channel opener,<sup>21b</sup> anti-cancer agents,<sup>21c</sup> and anti-hyperglycemic agents,<sup>21d</sup> Here we report on the synthesis and biological effects of 2-amino-quinazolin-4(3*H*)-one derivatives.

A series of 2-amino-quinazolin-4(3*H*)-one analogues were synthesized as shown in Scheme 1<sup>22</sup>. Anthranilic acids **3** were treated with urea by heating  $150^\circ\text{C}$  for 20 h to afford quinazolinones **4**. Dichloroquinazolines **5** were prepared by reacting quinazolinones **4** with  $\text{POCl}_3$  in the presence of trimethylamine. Treatment of dichloroquinazolines **5** with 2 N NaOH led to 2-chloro-4(3*H*)-quinazolinones **6**. 2-Chloroquinazolin-4(3*H*)-ones **6** and anilines **7** with various substituents were heated in DMF at  $85^\circ\text{C}$  for 16 h to obtain 2-amino-quinazolin-4(3*H*)-ones **8**.

The anti-SARS-CoV-2 and anti-MERS-CoV activities of the synthesized compounds were evaluated by immunofluorescent assay in Vero cells.<sup>16,20</sup> In this study, Vero cells were stained using antibodies targeting spike protein for MERS-CoV and nucleocapsid protein for SARS-CoV-2, and the infection rate was measured by imaging the infected Vero cells through microscope.

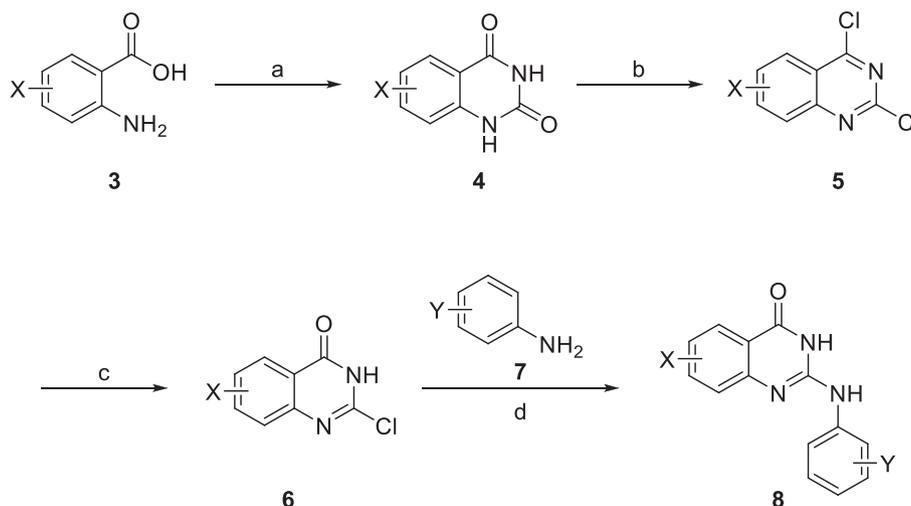
First, we began structure-activity relationship (SAR) studies for anti-SARS-CoV-2 activities by varying anilines at 2 position of quinazolinone ring of compound **2** (Table 1). For anilines with one fluoride atom, compound **9b** ( $IC_{50} = 1.4 \mu\text{M}$ ) bearing aniline substituted with fluoride at *meta*-position showed good effect, whereas compounds with *ortho*-fluoroaniline (**9a**) or *para*-fluoroaniline (**9c**) showed no inhibitory effect. In the same way, compound with anilines having two fluoride atoms at *ortho* and *para* position (**9d**) had no activity, while compound

**9e** with 3,5-difluoroaniline ( $IC_{50} = 0.24 \mu\text{M}$ ), which has two fluorides at double *meta*-position, showed a highly increased activity because of the synergic effect of fluoride at *meta*-position. Changing 2,4-fluoroaniline and 3,5-difluoroaniline to 2,4-chloroaniline (**9f**) and 3,5-dichloroaniline (**9g**) showed similar activities ( $IC_{50} : >25$  and  $0.23 \mu\text{M}$ , respectively). Next, we investigated the substituent effects at *meta*-position of anilines (**9h-9m**). Electron-donating groups such as methoxy (**9h**), hydroxy (**9i**), and dihydroxy (**9j**) decreased anti-SARS-CoV-2 activities ( $IC_{50} > 11 \mu\text{M}$ ). Electron-withdrawing groups such as cyano (**9k**) and trifluoromethyl (**9l**) showed much better effects than those of electron-donating groups ( $IC_{50} = 1.7$  and  $0.68 \mu\text{M}$ , respectively), while methyl-ester (**9m**) decreased anti-SARS-CoV-2 activities ( $IC_{50} = 8.8 \mu\text{M}$ ). Then, anti-SARS-CoV-2 effects of aliphatic amine substituents such as cyclohexylmethyl amine (**9n**), isopropyl amine (**9o**), *n*-butyl amine (**9p**), and piperidinyl amine (**9q**) were evaluated to exhibit decreased potency ( $IC_{50} > 14 \mu\text{M}$ ).

In the next phase of optimization, we evaluated substituent effects of 5 to 8 positions of quinazolinone ring, having fixed with 3,5-difluoroaniline or 3,5-dichloroaniline at 2 position (Table 2). Compounds with electron-withdrawing groups, such as 7-trifluoromethyl (**10a** and **11a**) and 7-nitro (**10b** and **11b**), showed high binding affinities ( $0.20$ – $0.51 \mu\text{M}$ ). In the case of compounds with electron-donating groups, variations of inhibitory effects were shown. Compounds with 5-hydroxy (**10e** and **11e**), 8-hydroxy (**10f** and **11f**), 5,8-dichloro (**10i** and **11i**), and 7,8-dichloro (**10j** and **11j**) showed high binding affinities ( $0.15$ – $1.6 \mu\text{M}$ ). 7-Amino (**10c** and **11c**), 5-methoxy (**10d** and **11d**), 7-hydroxy (**10g** and **11g**), and 6,8-dimethyl (**10h** and **11h**) derivatives showed no inhibitory effects. The 7-*N*-substituted quinazolinone compound **11k** displayed no anti-SARS-CoV-2 activity.

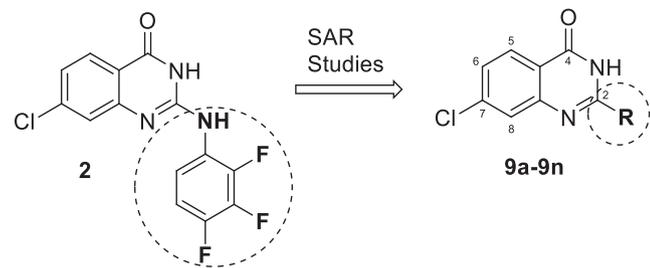
With the compounds having potent activities toward SARS-CoV-2, we tested anti-MERS-CoV activities (Table 3). All compounds except **10b** and **11b** showed good antiviral activities ( $IC_{50} = 0.39$ – $3.1 \mu\text{M}$ ). It seems that our quinazolinone compounds are potent broad spectrum coronavirus inhibitors. In particular, all the above compounds except **10a** and **11a** displayed no obvious cytotoxicity ( $CC_{50} > 25 \mu\text{M}$ ).

We selected the compounds for the purpose of further pharmacological investigations on SARS-CoV-2. In the case of electron-withdrawing substituents in the aromatic ring of quinazolinone, compound **9g** with chloro substituent at position 7 was selected because it has the highest SI (110). And we chose compound **11e** with electron-donating substituent (hydroxyl) at 5 position because it is the most active on SARS-CoV-2 and has the highest SI (168). Compound **9g** and **11e** were further evaluated for their microsomal stabilities, cytotoxicities, human ether a-go-go (hERG) bindings, plasma protein bindings



Scheme 1. Synthesis of 2-aminoquinazolin-4(3*H*)-ones derivatives. Reagents and conditions: (a) Urea,  $150^\circ\text{C}$ , 20 h; (b)  $\text{POCl}_3$ , TEA,  $115^\circ\text{C}$ , 17 h; (c) 2 N NaOH, rt, 20 h; (d) DMF,  $85^\circ\text{C}$ , 16 h.

**Table 1**  
Lead optimization of 2-anilino groups.



Compound	R	SARS-CoV-2		
		IC <sub>50</sub> <sup>a</sup> (μM)	CC <sub>50</sub> <sup>b</sup> (μM)	SI <sup>c</sup>
<b>2</b>	2,3,4-F <sub>3</sub> -Ph-NH-	2.6	>25	9.4
<b>9a</b>	2-F-Ph-NH-	>25	>25	1
<b>9b</b>	3-F-Ph-NH-	1.4	>25	18
<b>9c</b>	4-F-Ph-NH-	>25	>25	1
<b>9d</b>	2,4-F <sub>2</sub> -Ph-NH-	>25	>25	1
<b>9e</b>	3,5-F <sub>2</sub> -Ph-NH-	0.24	18	74
<b>9f</b>	2,4-Cl <sub>2</sub> -Ph-NH-	>25	>25	1
<b>9g</b>	3,5-Cl <sub>2</sub> -Ph-NH-	0.23	>25	110
<b>9h</b>	3-CH <sub>3</sub> O-Ph-NH-	11	>25	2.3
<b>9i</b>	3-HO-Ph-NH-	>25	>25	1
<b>9j</b>	3,5-(HO) <sub>2</sub> -Ph-NH-	>25	>25	1
<b>9k</b>	3-CN-Ph-NH-	1.7	>25	15
<b>9l</b>	3-CF <sub>3</sub> -Ph-NH-	0.68	>25	37
<b>9m</b>	3-CH <sub>3</sub> O(O = )C-Ph-NH-	8.8	>25	2.8
<b>9n</b>	Cyclohexylmethyl-NH-	14	>25	1.7
<b>9o</b>	<i>i</i> -Pr-NH-	>25	>25	1
<b>9p</b>	<i>n</i> -Bu-NH-	>25	>25	1
<b>9q</b>	piperidinyl-N-	>25	>25	1

<sup>a,b</sup> IC<sub>50</sub> and CC<sub>50</sub> were derived from the results of at least two dependent experiment in Vero cells infected with SARS-CoV-2

<sup>c</sup> SI(selective index) = CC<sub>50</sub>/IC<sub>50</sub> for inhibiting SARS-CoV-2 infection.

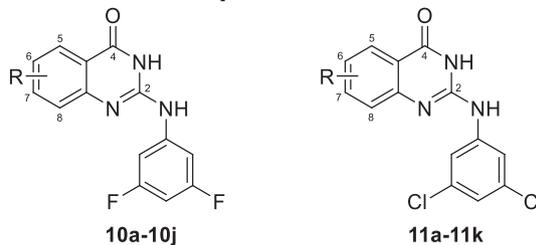
(PPB) and CYP inhibitions (Table 4 and 5). First, the results of microsomal stability show that **9g** is stable in mouse, rat, and human and **11e** is microsomally more unstable than **9g**, which seems to be due to the presence of -OH in the structure. In the cytotoxicity assay, **9g** and **11e** showed a little toxicity in HFL-1 (CC<sub>50</sub> = 7.5 and 9.6 μM), but did not show toxicity in the rest of the cells (Table 4).

In hERG channel inhibition assay, it was found that both compounds **9g** and **11e** did not interact with hERG channel. In the PPB assay, both compounds **9g** and **11e** showed high binding rates (97–100%). The results of CYP inhibition assay show that **9g** showed little inhibition of the CYP enzyme and **11e** displayed some inhibition (>65% CYP inhibition at 10 μM) in CYP1A2 and CYP3A4.

The preliminary pharmacokinetic properties of **9g** and **11e** were investigated by intravenous (*i.v.*) and oral (*p.o.*) routes in rats, with 2 mg/kg and 10 mg/kg, respectively (Table 6). The oral bioavailability of **9g** and **11e** were 15.6% and 7.8%, respectively, partly because the clearance of compound **11e** is higher than that of **9g**. The PK profile of **9g** seemed to be acceptable for the discovery of anti-coronavirus drugs.

In conclusion, we designed and developed 2-aminoquinazolin-4(3H)-one derivatives as potent inhibitors against both SARS-CoV-2 and MERS-CoV. Among them, 7-chloro-2-((3,5-dichlorophenyl)amino)quinazolin-4(3H)-one (**9g**) and 2-((3,5-dichlorophenyl)amino)-5-hydroxyquinazolin-4(3H)-one (**11e**) were considered as new drug candidates because both have high anti-SARS-CoV-2 activities [(**9g**, IC<sub>50</sub> = 0.23 μM), (**11e**, IC<sub>50</sub> = 0.15 μM)] and anti-MERS-CoV activities [(**9g**, IC<sub>50</sub> = 0.93 μM), (**11e**, IC<sub>50</sub> = 1.02 μM)] with no cytotoxicity (CC<sub>50</sub> > 25 μM). Our two lead compounds also showed good microsomal stabilities, relatively low cytotoxicities, low hERG binding affinities and CYP inhibitions. The PK profile of **9g** seemed to be acceptable for the discovery of antivirals. 2-Aminoquinazolinone derivatives were found to be a promising new scaffold against coronaviruses and further optimizations to increase pharmacokinetic profiles are currently underway.

**Table 2**  
SAR studies of 2-anilinoquinazolin-4(3H)-one derivatives



Compound	R	SARS-CoV-2		
		IC <sub>50</sub> <sup>a</sup> (μM)	CC <sub>50</sub> <sup>b</sup> (μM)	SI <sup>c</sup>
<b>10a</b>	7-CF <sub>3</sub>	0.20	7.6	38
<b>10b</b>	7-NO <sub>2</sub>	0.41	>25	61
<b>10c</b>	7-NH <sub>2</sub>	>25	>25	1
<b>10d</b>	5-CH <sub>3</sub> O	>25	>25	1
<b>10e</b>	5-OH	0.47	>25	54
<b>10f</b>	8-OH	1.6	>25	15
<b>10g</b>	7-OH	>25	>25	1
<b>10h</b>	6,8-(CH <sub>3</sub> ) <sub>2</sub>	>25	>25	1
<b>10i</b>	5,8-Cl <sub>2</sub>	0.33	>25	76
<b>10j</b>	7,8-Cl <sub>2</sub>	0.25	>25	98
<b>11a</b>	7-CF <sub>3</sub>	0.21	7.1	34
<b>11b</b>	7-NO <sub>2</sub>	0.51	>25	49
<b>11c</b>	7-NH <sub>2</sub>	>25	>25	1
<b>11d</b>	5-CH <sub>3</sub> O	>25	>25	1
<b>11e</b>	5-OH	0.15	>25	168
<b>11f</b>	8-OH	0.37	>25	65
<b>11g</b>	7-OH	>25	>25	1
<b>11h</b>	6,8-(CH <sub>3</sub> ) <sub>2</sub>	>25	>25	1
<b>11i</b>	5,8-Cl <sub>2</sub>	0.35	>25	71
<b>11j</b>	7,8-Cl <sub>2</sub>	0.24	>25	103
<b>11k</b>	7-Morpholinyl	>25	>25	1

<sup>a,b</sup> IC<sub>50</sub> and CC<sub>50</sub> were derived from the results of at least two dependent experiment in Vero cells infected with SARS-CoV-2

<sup>c</sup> SI(selective index) = CC<sub>50</sub>/IC<sub>50</sub> for inhibiting SARS-CoV-2 infection.

**Table 3**  
Anti-MERS-CoV activity of 2-Aminoquinazolin-4(3H)-ones Derivatives

Compound	MERS-CoV			SARS-CoV-2 IC <sub>50</sub> <sup>d</sup> (μM)
	IC <sub>50</sub> <sup>a</sup> (μM)	CC <sub>50</sub> <sup>b</sup> (μM)	SI <sup>c</sup>	
<b>9e</b>	0.39	>25	65	0.24
<b>9g</b>	0.93	>25	27	0.23
<b>10a</b>	0.73	>25	35	0.20
<b>11a</b>	0.96	7.3	7.6	0.21
<b>10b</b>	8.5	>25	2.9	0.41
<b>11b</b>	>25	>25	1.0	0.51
<b>10e</b>	1.1	>25	22	0.47
<b>11e</b>	1.02	>25	25	0.15
<b>10f</b>	3.1	>25	8.0	1.64
<b>11f</b>	1.7	>25	14	0.37
<b>10i</b>	0.79	>25	31	0.33
<b>11i</b>	0.62	>25	42	0.35
<b>10j</b>	0.83	>25	30	0.25
<b>11j</b>	0.88	>25	31	0.24

<sup>a,b</sup> IC<sub>50</sub> and CC<sub>50</sub> were derived from the results of at least two dependent experiment in Vero cells infected with MERS-CoV

<sup>c</sup> SI(selective index) = CC<sub>50</sub>/IC<sub>50</sub> for inhibiting MERS-CoV infection.

<sup>d</sup> IC<sub>50</sub> were derived from the results of at least two dependent experiment in Vero cells infected with SARS-CoV-2

### Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

**Table 4**  
Result of Microsomal stability and cytotoxicity of **9g** and **11e**

Compound	MS <sup>a</sup>			Cytotoxicity (μM) <sup>b</sup>			
	m <sup>a</sup>	r <sup>a</sup>	h <sup>a</sup>	HFL-1	L929	NIH 3 T3	CHO-K1
<b>9g</b>	112	105	99	7.5	31	>100	11
<b>11e</b>	42	54	55	9.6	27	60	28

<sup>a</sup> % of remaining after 30 min<sup>b</sup> Cell information. HFL-1: human embryonic lung cell line, L929: NCTC clone 929, mouse fibroblast cell line, NIH 3 T3 : mouse embryonic fibroblast cell line, CHO-K1 : Chinese hamster ovary cell line.**Table 5**  
Result of hERG, PPB, CYP inhibition of **9g** and **11e**

Compound	hERG(10 μM)	PPB (5 μM)		CYP inhibition (10 μM)				
		r	h	1A2	2C9	2C19	2D6	3A4
<b>9g</b>	38	100	100	15	18	<1	20	32
<b>11e</b>	28	97	100	65	34	12	46	65

**Table 6**  
Rat pharmacokinetic study of **9g** and **11e**

Compounds	<b>9g</b>		<b>11e</b>	
	I.V., 2 mg/kg	P.O., 10 mg/kg	I.V., 5 mg/kg	P.O., 10 mg/kg
Tmax (h)	NA <sup>b</sup>	2.5	NA	1.0
Cmax (μg/h)	NA	0.9	NA	0.06
T <sub>1/2</sub> (h)	2.2	5.7	9.2	42
AUC (μg-h/mL)	8.5	6.6	6.5	1.01
CL (L/h/kg)	0.25	NA	0.67	NA
V <sub>ss</sub> (L/Kg)	1.6	NA	6.8	NA
F <sub>t</sub> (%)	NA	15.6	NA	7.8

<sup>a</sup> All results are the mean of experiments using three rats. <sup>b</sup> NA: not applicable

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## Appendix A. Supplementary data

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