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Synthesis and biological evaluation of pyrrolopyridazine derivatives as novel HER-2 tyrosine kinase inhibitors

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ABSTRACT

A series of novel pyrrolopyridazine derivatives have been discovered to be HER-2 inhibitors. These compounds selectively inhibited HER-2 kinase activity at low nanomolar concentrations. Compound **7d** was identified as a potent HER-2 inhibitor with an IC_{50} of 4 nM.

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Receptor tyrosine kinases (RTKs) play a crucial role in signal transduction pathways that regulate cell differentiation, proliferation and angiogenesis. Inhibition of RTK activation has become a compelling approach in the development of anticancer agents.¹ The ErbB family of receptor tyrosine kinases comprises four closely related members: the epidermal growth factor receptor (EGFR, ErbB1 or HER-1), the human epidermal growth factor receptor 2 (ErbB2 or HER-2), ErbB3 (HER-3), and ErbB4 (HER-4).^{2,3} Over expression of EGFR and HER-2 has been observed in many cancers including non-small-cell lung cancer, and prostate, breast, colon and stomach cancers. Small molecules that inhibit the kinase activity of EGFR and HER-2 are considered to be new therapeutic antitumor agents.^{4,5}

Three small molecule drugs, Gefitnib (Iressa[®]),⁶ Erlotinib (Tarceva[®]),⁷ and Lapatinib (Tykerb[®]),⁸ have been approved and marketed for the treatment of cancer (Fig. 1). Several other ErbB family inhibitors are currently being investigated in various phases. These quinazoline-based small molecules have a common scaffold that mimics the adenine moiety of ATP, and therefore are potent and competitive inhibitors of ATP.⁹ During the course of our exploration of non-quinazoline scaffolds, pyrrolopyridazine derivatives (I, II, III) were found to be novel and potent HER-2 inhibitors (Fig. 2). Here, we report our progress in the synthesis and biological evaluation of these pyrrolopyridazine-based inhibitors.

As shown in Scheme 1, reaction of pyrrole diester **1** with ceric ammonium nitrate (CAN) afforded pyrrole aldehyde 2,¹⁰ which underwent a cyclization with hydrazine hydrate in acetic acid to give pyrrolo[2,3-*d*]pyridazin-4-one **3**. The pyrrolo[2,3-*d*]pyrida-

zin-4-one **3** was transformed into the 4-chloro-pyridazine derivative **4** upon treatment with POCl₃. Acid catalysed displacement of the chloride with the appropriate aniline was performed in isopropanol to give **5**. Saponification of **5** followed by coupling with different amines afforded compounds **7a–e**.¹¹

Analogs with pyrrolo[3,4-*d*]pyridazine scaffold **II** and 7-substituted pyrrolo[2,3-*d*]pyridazine scaffold **III** were synthesized by similar routes starting from different commercially available pyrroles as shown in Schemes 2 and 3. Pyrrole diester **8**¹² and **13**¹³ were transformed to aldehyde **9** and **14**, respectively, under Vilsmeier–Haack reaction conditions, which were converted to **10** and **15** via intramolecular ring-closing condensations. Chlorination of **10** and **15** followed by coupling with 3-chloro-4-(3-fluoro-benzyloxy)-phenylamine in isopropanol afforded **12** and **17**. Saponification and amidation of **17** led to the desired compounds **18a–b** as shown in Scheme 3.

Inhibition of EGFR and HER-2 tyrosine kinase activity was evaluated in enzymatic and cellular assays using A431(overexpressing EGFR) and SK-BR-3 (overexpressing HER-2).^{14,15} Table 1 summarises the in vitro potency of representative compounds. No significant EGFR tyrosine kinase inhibitory activity was observed in these compounds. Computational modeling studies provided a possible explanation for this result. A docking model of compound **7a** based on the crystal structure of EGFR complexed with Lapatinib¹⁶ (PDB code 1XKK) was developed (Fig. 3). The docked models were energy minimized in Sybyl7.3. This model showed that compound **7a** bound to EGFR in definitely different fashions when compared with Lapatinib. The pyrrolopyridazine core of **7a** deviated from the quinazoline core of Lapatinib, and was oriented in the ATP binding site such that there was a hydrogen bond between N-2 and the hinge region Met769 NH. The C-4 hydrophobic

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 $HCI = O^{2}S_{O}^{S}O = O^{2}S_{O}^{S}O = O^{2}S_{O}^{S}O = O^{2}S_{O}^{S}O_{3}H$

Gefitinib (Iressa®)

Erlotinib (Tarceva®)

Figure 1. Examples of launched ErbB family inhibitors.

Lapatinib (Tykerb®)



Figure 2. General structure of the pyrrolopyridazine-based compounds.

group extended out into the aqueous phase instead of extending into the deep hydrophobic pocket. This might explain why compound **7a** was almost devoid of EGFR activity.

Moderate growth inhibitions toward A431 cells in compounds **7b**, **7c** and **7e** might be caused by the inhibition of other kinases. It was noticeable that compounds **7a** and **7d** exhibited similar potency as Lapatinib (0.003 μ M) against HER-2 kinase, but was found to be >100 times more selective than the closely related EGFR ki-



Scheme 1. Synthesis of type I derivatives. Reagents and conditions: (a) 4.0 equiv CAN, THF/HOAc/H₂O, rt, 1.5 h, 65%; (b) 1.1 equiv NH₂NH₂·H₂O, HOAc, 100 °C, 2 h, 79%; (c) 2.0 equiv POCl₃, CH₃CN, 80 °C, 1.5 h, 95%; (d) 1.5 equiv R¹NH₂, 2.5 equiv Et₃N, *i*-PrOH, aq HCl, 80 °C, 4 h, 50%; (e) 1 N NaOH, EtOH, reflux, then 1 N HCl, pH 1, 91%; (f) 1.5 equiv R⁵NH₂, Et₃N, EDCl, HOBt, DMF, rt, 17–58%.



Scheme 2. Synthesis of type II derivatives. Reagents and conditions: (a) 2.0 equiv POCl₃, DMF, 100 °C, 2 h, 77%; (b) 1.5 equiv NH₂NH₂·H₂O, EtOH, 90 °C, 3 h, 72%; (c) 2.0 equiv POCl₃, CH₃CN, 80 °C, 5 h; (d) 1.1 equiv 3-chloro-4-(3-fluoro-benzyloxy) -phenylamine, *i*-PrOH, 90 °C, 7 h, 51%.



Scheme 3. Synthesis of type **III** derivatives. Reagents and conditions: (a) 2.0 equiv POCl₃, DMF, 100 °C, 3 h, 21%; (b) 1.5 equiv NH₂NH₂·H₂O, EtOH 90 °C, 1.5 h, 44%; (c) 2.0 equiv POCl₃, CH₃CN, 80 °C, 3 h; (d) 1.1 equiv 3-chloro-4-(3-fluoro-benzyloxy)-phenylamine, *i*-PrOH, 90 °C, 3 h, 34%; (e) 1 N NaOH, EtOH, reflux, then 1 N HCl, pH 1, 85%; (f) 1.5 equiv R⁵NH₂, Et₃N, EDCl, HOBt, DMF, rt, 62–85%.

Table 1

Enzyme and cellular inhibitory activity assay results

No.	R ¹	R ⁵	EGFR IC_{50}^{a} (μM)	A431 IC_{50}^{a} (μM)	HER-2 IC_{50}^{a} (μM)	SK-BR-3 IC_{50}^{a} (µM)
7a	CI F	N_N_	1.28	1.16	0.007	0.96
7b	Cl F		>10	0.98	ND	2.24
7c	CI F		>10	0.257	ND	0.843
7d	CI F		5–10	12.65	0.004	>5
7e		N N	>10	0.867	ND	>5
12	-	-	>10	1.21	ND	0.934
18a	-	N_N_	>10	77.22	ND	3.84
18b	-		>10	15.14	ND	0.62
Tarceva®b	_	_	0.015	0.42	1.52	1.89
Tykerb ^{®b}	_	_	0.019	0.14	0.003	0.124

ND: not determined.

^a Average values (at least two experiments). ^b Tarceva® and Tykerb® were prepared according to the literatures.^{7,8}



Figure 3. Predicted binding mode of compound **7a** (yellow) modeled in the X-ray structure of the Lapatinib/EGFR kinase complex. The Lapatinib molecule is shown in magenta.

nase. The HER-2 selectivities of the compounds **7a** and **7d** cannot be explained adequately due to lack of an HER-2 X-ray structure. Further studies are planned to investigate these selectivities. Compounds **7a**, **7c**, **12** and **18** were also found to be low-micromolar inhibitors toward SK-BR-3 cells.

In conclusion, pyrrolopyridazine derivatives were prepared and found to be low nanomolar HER-2 selective inhibitors over EGFR. Further studies of these compounds are in progress.

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