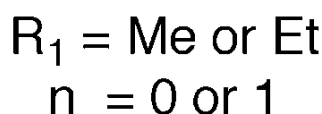
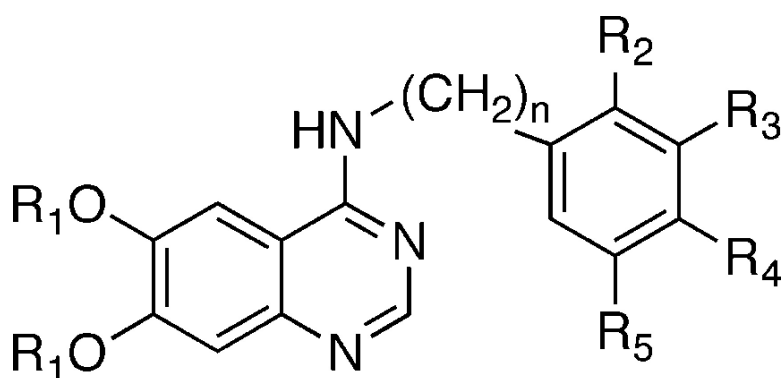


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Anilindialkoxyquinazolines: Screening Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitors for Potential Tumor Imaging Probes

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The epidermal growth factor receptor (EGFR), a long-standing drug development target, is also a desirable target for imaging. Sixteen dialkoxyquinazoline analogues, suitable for labeling with positron-emitting isotopes, have been synthesized and evaluated in a battery of in vitro assays to ascertain their chemical and biological properties. These characteristics provided the basis for the adoption of a selection schema to identify lead molecules for labeling and in vivo evaluation. A new EGFR tyrosine kinase radiometric binding assay revealed that all of the compounds possessed suitable affinity ($IC_{50} = 0.4\text{--}51\text{ nM}$) for the EGFR tyrosine kinase. All of the analogues inhibited ligand-induced EGFR tyrosine phosphorylation ($IC_{50} = 0.8\text{--}20\text{ nM}$). The HPLC-estimated octanol/water partition coefficients ranged from 2 to 5.5. Four compounds, 4-(2'-fluoroanilino)- and 4-(3'-fluoroanilino)-6,7-diethoxyquinazoline as well as 4-(3'-chloroanilino)- and 4-(3'-bromoanilino)-6,7-dimethoxyquinazoline, possess the best combination of characteristics that warrant radioisotope labeling and further evaluation in tumor-bearing mice.

Introduction

Protein tyrosine kinases (PTKs), enzymes that phosphorylate tyrosine residues on functional proteins, are common mediators of signals that regulate many cellular processes. PTKs can be divided into two subgroups on the basis of their structural characteristics, nonreceptor cytoplasmic PTKs and receptor tyrosine kinases (RTKs).¹ To date at least 20 families of receptor tyrosine kinases that share structural, most notably an intrinsic tyrosine kinase domain, and functional similarities have been classified.^{2,3} The epidermal growth factor receptor (EGFR) was one of the first oncogenes and receptor tyrosine kinases to be discovered.^{4,5} EGFR belongs to the ErbB family of receptors, which also includes ErbB2 (HER2/Neu), ErbB3, and ErbB4. These receptors are overexpressed in a variety of tumors.

EGFR tyrosine kinase phosphorylation is stimulated by epidermal growth factor (EGF) or transforming growth factor α (TGF α) binding to the extracellular ligand binding domain of the EGFR and subsequent receptor dimerization. Signal transduction initiated by these events regulates cellular proliferation, differentiation, motility, adhesion, and apoptosis. These signaling processes play an important role in normal epithelial and stromal cell morphology; moreover, overexpression or aberrant signaling from EGFR and the other ErbB

receptor family members has been shown to be a key determinant in tumor growth and proliferation.^{6,7}

Overexpression of EGFR has been found in head and neck tumors, gliomas, non-small-cell lung carcinoma and tumors of the breast, ovaries, cervix, esophagus, bladder, prostate, and kidney.⁷ There is extensive literature on the clinical significance of increased EGFR expression and signaling in tumors and the relationship to other prognostic factors.^{8,9} In a majority of tumors, poor survival rates correlate with EGFR overexpression. For example, EGFR overexpression has been detected in nearly 45% of the breast tumors studied¹⁰ and receptor overexpression is inversely correlated with patient survival.⁹ EGFR overexpression is also inversely correlated with estrogen receptor status, consistent with the failure of EGFR-positive patients to respond to hormonal therapies.^{11,12} The relevance of EGFR expression in tumors to prognostic outcome supports the years of investment toward finding an EGFR-targeted therapeutic^{13–18} and the need for a diagnostic imaging agent. Development of an imaging agent might also be a valuable tool in the search for and the characterization of EGFR-targeted therapeutics.

Two potential targets for EGFR-based probes are the extracellular domain or the intracellular tyrosine kinase domain. Radioactive probes that bind to the extracellular domain of EGFR have been generated. EGF, the 53 amino acid, 6 kDa, natural ligand, binds to the EGFR with a K_d of 0.1–1.0 nM¹⁹ and has been labeled directly with iodine-123, iodine-125, and iodine-131. In vitro studies using [¹³¹I]EGF demonstrated its cytotoxic potential.²⁰ ¹²³I-EGF has been used to image cervical cancer in humans.²¹ Despite its ability to localize in EGFR-rich tissue, radioiodinated EGF rapidly degrades in vivo, releasing radioiodine, thereby reducing the

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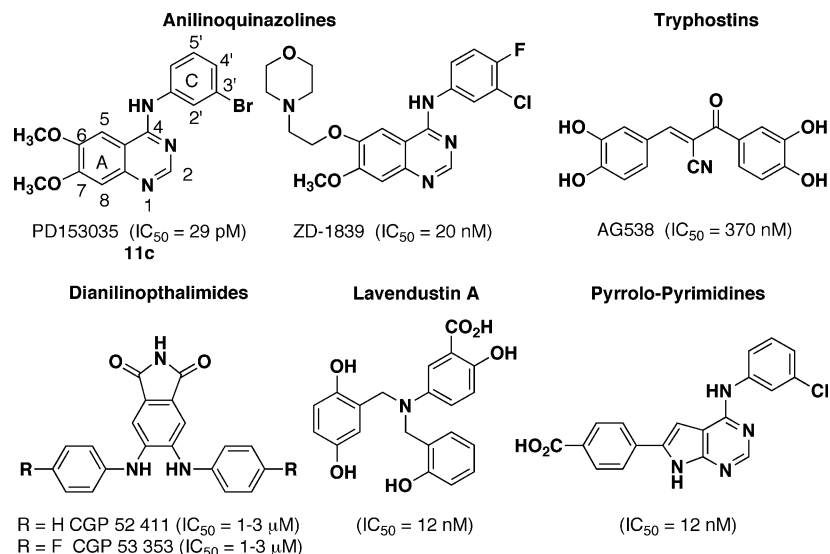


Figure 1. EGFR tyrosine kinase inhibitors^{13,16,34,55}

lifetime of the label in the tumor.²² Iodinated dextran–EGF conjugates increase retention of the iodine by tumor cells but at the cost of higher molecular weight that affects tracer distribution.²⁰ EGF has been labeled with technetium-99m using a MAG3 or 2-iminothiolane chelate and was found to accumulate in tumor xenografts with more rapid clearance than labeled monoclonal antibodies (Mabs).^{23–25} Bifunctional chelation of indium-111 to EGF using diethylenetriaminepentaacetic acid (DTPA) has been shown to have cytotoxic effects in vitro and is being investigated as a potential radiotherapeutic agent.^{26,27} The imaging potential of ¹¹¹In–EGF was found to be inferior to a labeled anti-EGFR antibody, ¹¹¹In–DTPA–Mab 528, in tumor-bearing rats.²⁸

Several Mabs against EGFR have been developed as antitumor agents. The radioiodinated Mab, EGFR1, with high affinity for the EGFR has demonstrated good uptake in MCF-7 tumors grown in nude athymic mice²⁹ and good localization in human brain gliomas.³⁰ Likewise, the indium-111-labeled C225 Mab has been tested in patients with squamous cell lung carcinoma.³¹ A direct-labeled technetium-99m anti-EGFR antibody, ior egr/r3, has demonstrated imaging sensitivity (84%), specificity (100%), and accuracy (86.5%) in human epithelial tumors.³² However, Mabs labeled with short-lived PET isotopes, in general, have demonstrated limited targeting success in large part due to pharmacokinetic constraints related mostly to their size.

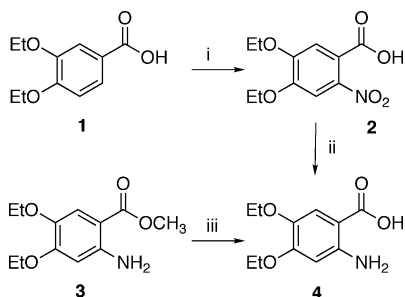
To date, no small organic molecules with affinity for the extracellular domain of the EGFR have been identified; however, a number of small molecules have been shown to be potent (nano- to picomolar) inhibitors of the intracellular EGFR tyrosine kinase at the ATP binding site. A sampling of the different compound classes that inhibit EGFR phosphorylation is shown in Figure 1. The dialkoxyanilinoquinazolines (Figure 1) were chosen as the lead compounds in the present study on the basis of the strong structure–activity data.^{33–35}

The development of imaging agents based on the small-molecule EGFR inhibitors has been a recent area of active research. A number of radiolabeled anilinoquinazoline analogues have been reported. The compounds incorporate labeled substituents on the A or C

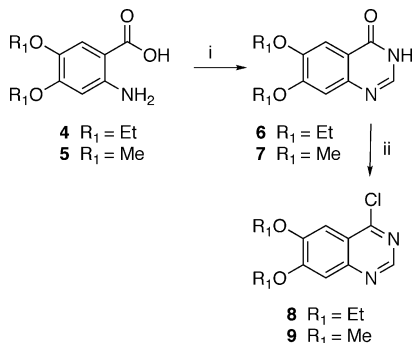
rings (see Figure 1) of the anilino- or benzylaminoquinazoline. The C ring substituents include 4-(3'-[¹²⁵I]-iodoanilino),^{36,37} 4-(3'-[¹⁸F]fluoro-5'-(trifluoromethyl)-anilino),³⁸ 4-(3',4'-dichloro-6'-[¹⁸F]fluoroanilino),³⁸ and 4-(3'-chloro-4'-[¹⁸F]fluoroanilino).³⁹ The 7-[¹⁸F]fluoroethoxy⁴⁰ and the 6- or 7-[¹¹C]methoxy^{40,41} constitute the A ring labeled substituents. Preliminary in vitro studies with the 3'-[¹²⁵I]iodo analogue demonstrated receptor-mediated uptake in cells containing high EGFR titer.³⁷ A study of the carbon-11 methoxy derivative demonstrated some uptake in human neuroblastoma xenografts in mice;^{42,43} however, the 20-min half-life of the carbon-11 may not allow adequate time for the development of good signal relative to background. Bonasera and colleagues evaluated five fluorine-18-labeled compounds.³⁸ They studied the 4-(4'-[¹⁸F]fluoroanilino)-dimethoxyquinazoline and the 4-(3',4'-dichloro-6'-[¹⁸F]fluoroanilino)-dimethoxyquinazoline in tumor-bearing mice. These tracers did not accumulate in the tumors nor was receptor-mediated uptake, based on blocking studies, observed for the latter probe. Low receptor affinity, high nonspecific binding, and probe metabolism may have contributed to the inability of these compounds to accumulate in tumor cells that overexpress EGFR.

Successful development of an imaging probe targeting a new biomarker, in this case EGFR, requires an adequate screening strategy for the selection of ligands to be carried forward for labeling and, ultimately, in vivo studies. It is neither economically nor logistically feasible to label and evaluate every compound in animal models. Likewise, navigating the structure–activity relationships in the medicinal chemistry literature can be challenging with respect to choosing an appropriate imaging lead compound. For example, the fact that a small molecule is a potent EGFR inhibitor does not necessarily guarantee that it will possess desirable EGFR imaging characteristics such as high receptor affinity, high receptor selectivity, low nonspecific binding, rapid clearance, and suitable metabolism.

In the current study, a small series of dialkoxyanilinoquinazoline EGFR inhibitors suitable for labeling with fluorine-18 (110-min half-life positron-emitter) or carbon-11 (20-min half-life positron-emitter) has been synthe-

Scheme 1^a

^a Reagents and conditions: (i) 70% HNO₃/glacial acetic acid/2 h/rt; (ii) SnCl₂/HCl/2 h/rt; (iii) NaOH/reflux/1 h, then HCl.

Scheme 2^a

^a Reagents and conditions: (i) formamidine·HCl/heat then NaOH/sonicate/1 h/rt; (ii) oxalyl chloride/DMF/1,2-dichloroethane/reflux.

sized. Appropriate assays have been developed to determine both functional and imaging characteristics, including a new radiometric binding assay to measure the affinity of the inhibitors for the enzyme. These studies provide the basis needed for the selection of ligands to be labeled and further evaluated as potential imaging agents for the noninvasive determination of EGF receptor density.

Results

Chemistry. A small library of anilino- and benzyl-dialkoxyquinazoline compounds, **10**–**17**, was prepared by coupling 4-chlorodialkoxyquinazoline **8** or **9** with the appropriate substituted aniline or benzylamine (Scheme 3). The 4-chloro-6,7-dimethoxyquinazoline was synthesized as previously reported from the 4,5-dimethoxyanthranilic acid **5**.³³ As the 4-(3'-bromoanilino)-6,7-diethoxyquinazoline was reported³³ to possess more potent biological activity than the corresponding 6,7-dimethoxy analogue, we were interested in finding a convergent synthetic route suitable for the preparation of several diethoxy analogues. The previously reported two-step conversion of the dimethoxybromoanilinoquinazoline **11c** to the diethoxy analogue **10c** (Scheme 3), using a pyridinium hydrochloride melt to give the bishydroxy intermediate followed by O-alkylation with iodoethane, proceeded in a low 5.5% yield.³³ This was inadequate for the preparation of a series of the diethoxy analogues. Thus, we produced the 4,5-diethoxyanthranilic acid **4** by two methods (Scheme 1). Initially, a small amount of methyl 2-amino-4,5-diethoxybenzoate **3** was commercially available from Aldrich Specialty Chemical. The benzoate **3** was directly converted to **4** by saponification of the methyl ester. Alternatively, when the

commercial supply was depleted, 4,5-diethoxybenzoic acid **1** was nitrated to give **2** followed by reduction of the nitro group to form the desired aminobenzoic acid **4** in moderate overall yield.

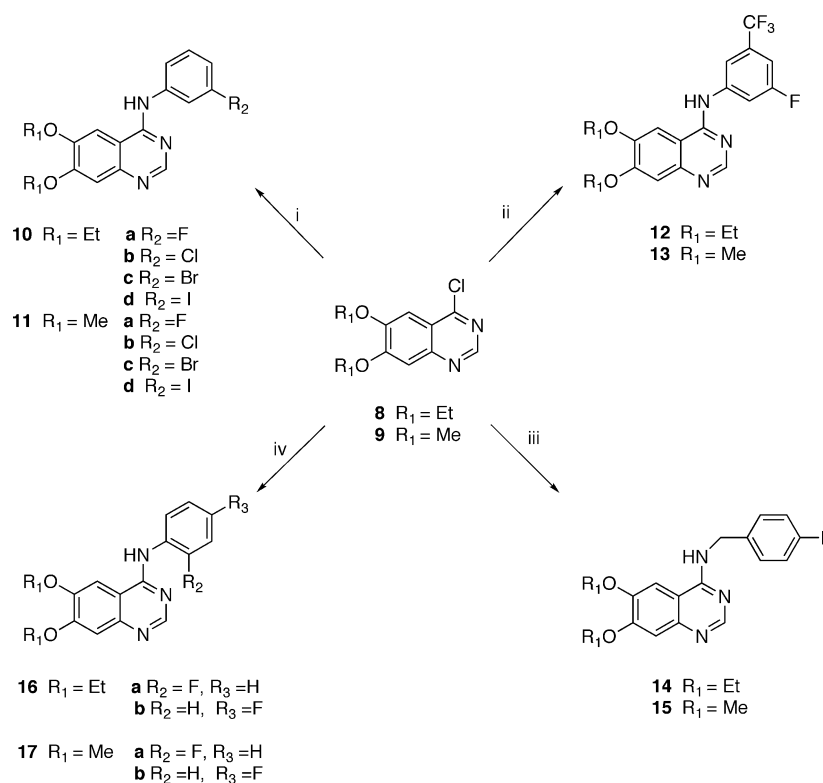
Preparation of the 4-chloroquinazolines followed a previously described procedure (Scheme 2).³³ Cyclization of the dialkoxyanthranilic acids **4** or **5** with formamidine hydrochloride at 210 °C gave the corresponding dialkoxyquinazolines, **6** or **7**, in 55–65% yield. The quinazolines were subsequently refluxed with oxalyl chloride in DMF and 1,2-dichloroethane to form the 4-chloro-dialkoxyquinazolines **8** and **9** in good yield.

A modified coupling procedure was employed for the production of the anilino- and (benzylamino)quinazolines (Scheme 3). Anhydrous DMF was used as the reaction solvent instead of the previously reported 2-propanol. The reaction was carried out at 80 °C with nearly quantitative conversion to the substituted aminoquinazoline hydrochloride (HCl) salt within 30 min to 1 h depending on the aniline substituents. The precipitated HCl salt was filtered from the DMF solution and was converted to the free base for semi-preparative normal phase HPLC purification. The pure quinazolines were reconverted to the more stable HCl salt for the biological assays.

Chemical and physical data for all of the compounds are presented in Table 1. All of the anilino- and benzylquinazoline analogues were analyzed by analytical reversed-phase HPLC and found to be greater than 99% pure. Elemental analysis of all of the dialkoxyquinazoline analogues in Table 1 agree with the calculated values to within ±0.4%

Lipophilicity Measurement. The lipophilicity of compounds can affect their tissue permeability properties, which can impact their localization in target tissues. Lipophilicity may also affect binding to low-affinity nonspecific sites that can compromise target tissue to background tissue ratios. The octanol/water partition coefficients of the quinazoline compounds were estimated by a reversed-phase HPLC method.⁴⁴ This method has been previously used by us to determine lipophilicities of steroid ligands.⁴⁵ The estimated log *P*_{ow} values are reported in Table 1. The lipophilicities generally exhibited the expected trends with a couple of noted exceptions. The lipophilicity was greater for the diethoxy series relative to the corresponding dimethoxy analogues. Within the series **10a**–**d** and **11a**–**d**, the lipophilicity increased with increasing size of the halogen from fluorine to iodine. Adding the trifluoromethyl moiety to **10a** and **11a** increased the lipophilicity by 1.7–1.8 log units. Interestingly, adding an extra methylene to produce the benzylamine did not significantly increase the compound lipophilicity (compare **14** and **16b**, **15** and **17b**). In contrast, the position of the fluorine on the aniline (C ring, Figure 1) ring did have a significant effect on the lipophilicity. The *m*- and *p*-fluoroanilino analogues, **10a** and **16b**, had similar lipophilicities, while the ortho-substituted analogue, **16a**, showed a 0.7 log unit decrease. This trend was similar for the dimethoxy analogues.

Biological Activity. The 16 quinazoline compounds were studied in a battery of in vitro assays to assess their biological properties and to develop a basis for screening these and future compounds for potential

Scheme 3^a

^a Reagents and conditions: (i) 3-haloaniline/DMF/heat; (ii) 3-fluoro-5-(trifluoromethyl)aniline/DMF/heat; (iii) 4-fluorobenzylamine/DMF/heat; (iv) 2- or 4-fluoroaniline/DMF/heat.

Table 1. 4-Anilino- and 4-(Benzylamino)quinazolines Chemical and Physical Data

no.	type	R ₁	R ₂	R ₃	mp (°C)	formula	anal.	log P _{ow} ^a
10a	A	CH ₃ CH ₂	F	H	247–248	C ₁₈ H ₁₈ FN ₃ O ₂ ·HCl	C, H, N	3.71 ± 0.12
10b	A	CH ₃ CH ₂	Cl	H	260–261	C ₁₈ H ₁₈ ClN ₃ O ₂ ·HCl	C, H, N	4.31 ± 0.26
10c	A	CH ₃ CH ₂	Br	H	250–252	C ₁₈ H ₁₈ BrN ₃ O ₂	C, H, N	4.40 ± 0.25
10d	A	CH ₃ CH ₂	I	H	258–261	C ₁₈ H ₁₈ IN ₃ O ₂ ·HCl	C, H, N	4.62 ± 0.26
11a	A	CH ₃	F	H	244.5–246	C ₁₆ H ₁₄ FN ₃ O ₂ ·1.1HCl	C, H, N	2.96 ± 0.10
11b	A	CH ₃	Cl	H	230–235	C ₁₆ H ₁₄ ClN ₃ O ₂ ·HCl	C, H, N	3.51 ± 0.23
11c	A	CH ₃	Br	H	256–257.5	C ₁₆ H ₁₄ BrN ₃ O ₂ ·HCl	C, H, N	3.49 ± 0.22
11d	A	CH ₃	I	H	251–251.5	C ₁₆ H ₁₄ IN ₃ O ₂ ·HCl	C, H, N	3.65 ± 0.22
12	A	CH ₃ CH ₂	F	CF ₃	278–280	C ₁₉ H ₁₇ F ₄ N ₃ O ₂ ·HCl	C, H, N	5.49 ± 0.29
13	A	CH ₃	F	CF ₃	269–270.5	C ₁₇ H ₁₃ F ₄ N ₃ O ₂ ·HCl	C, H, N	4.66 ± 0.26
14	B	CH ₃ CH ₂	F		238.5–240	C ₁₉ H ₂₀ FN ₃ O ₂	C, H, N	3.78 ± 0.23
15	B	CH ₃	F		250–251	C ₁₇ H ₁₆ FN ₃ O ₂ ·HCl	C, H, N	3.02 ± 0.20
16a	C	CH ₃ CH ₂	F	H	252–255	C ₁₈ H ₁₈ FN ₃ O ₂ ·1.4HCl	C, H, N	3.02 ± 0.11
16b	C	CH ₃ CH ₂	H	F	220.5–222	C ₁₈ H ₁₈ FN ₃ O ₂ ·HCl	C, H, N	3.73 ± 0.23
17a	C	CH ₃	F	H	247–248	C ₁₆ H ₁₄ FN ₃ O ₂ ·HCl	C, H, N	2.20 ± 0.11
17b	C	CH ₃	H	F	231–232	C ₁₆ H ₁₄ FN ₃ O ₂ ·HCl	C, H, N	2.87 ± 0.21

^a Estimated by the reversed-phase HPLC method of Minick et al.⁴⁴

imaging agents. A radiometric binding assay was developed to determine the relative binding affinities of these compounds for the ATP binding site in the tyrosine kinase domain of the receptor. The ability of these molecules to inhibit EGFR tyrosine phosphorylation was probed. The specificity of a small subset of compounds was determined by assessing inhibition of ErbB2 and ErbB4 receptor phosphorylation. Finally, the ability of the ligand to inhibit cellular DNA synthesis, in cells dependent and not dependent on EGF for cell

proliferation, was evaluated. This assay was performed in an effort to find a test that would be amenable to high-throughput screening, the results of which would potentially correlate with receptor binding and ultimately with the pharmacokinetic distribution of the tracer in vivo.

Receptor binding is a key characteristic that these molecules must possess to be suitable imaging agents. A receptor binding assay using EGF receptors extracted from the A431 human carcinoma cell membranes was

Table 2. Biochemical Data for the EGFR Tyrosine Kinase (tk) Inhibitors

no.	receptor binding (IC ₅₀ , nM)		receptor phosphorylation (IC ₅₀ , nM)	
	EGFR tk	EGFR tk	ErbB2 tk	ErbB4 tk
10a	8.17 ± 1.57	6.5 ± 2.1		
10b	0.38 ± 0.13	1.2 ± 0.2		
10c	0.41 ± 0.09	3.2 ± 0.8	215 ± 87	50 ± 19
10d	0.64 ± 0.15	4.6 ± 2.0	69 ± 10	59 ± 29
11a	31.9 ± 7.00	6.3 ± 2.0		
11b	1.26 ± 0.00	0.8 ± 0.2		
11c	0.66 ± 0.12	2.6 ± 0.5	143 ± 52	49 ± 16
11d	1.05 ± 0.51	6.4 ± 2.7		
12	8.95 ± 3.26	>50		
13	20.0 ± 10.2	19.1 ± 2.9		
14	17.0 ± 5.0	6.6 ± 1.6	231 ± 92	>100
15	47.7 ± 14.1	10.9 ± 2.8		
16a	9.31 ± 1.19	15.8 ± 2.2		
16b	16.0 ± 3.7	23.1 ± 4.2		
17a	32.2 ± 7.4	12.8 ± 3.5		
17b	51.0 ± 11.7	19.1 ± 2.5		

developed and used to study the relative binding affinity of these compounds to the tyrosine kinase domain. Iodine-125-labeled **11d** (specific activity 583–596 Ci/mmol) was employed as the radioligand,³⁶ and nonspecific binding was determined by adding **11c**, the bromo analogue, to the assay. EGFR binding values, expressed as an IC₅₀, are shown in Table 2. All of the compounds demonstrated high affinity for the receptor with IC₅₀s ranging from 0.4 to 51 nM. In all cases the diethoxy analogues had relatively higher affinity for the receptor than the corresponding dimethoxy derivatives. **10b–10d** and **11b–11d**, the *m*-chloro-, -bromo-, and -iodoanilino analogues, exhibited the highest relative affinities (nanomolar to subnanomolar) for the tyrosine kinase domain. Three fluoroanilino analogues, **10a**, **12**, and **16a**, displayed relative affinities slightly less than 10 nM, generally considered the upper limit for imaging agents,^{46,47} while the remaining analogues had relative affinities greater than 16 nM.

While receptor binding is absolutely necessary for localization of a potential imaging agent, the ability to inhibit receptor function, in this case ligand-induced receptor tyrosine autophosphorylation, may be uncoupled from ligand binding. A series of assays were designed to examine the correlation between receptor binding and inhibition of receptor phosphorylation or inhibition of EGF-dependent DNA synthesis. The inhibition of ligand-induced EGFR tyrosine autophosphorylation is reported in Table 2. All of the compounds were potent inhibitors of EGFR tyrosine phosphorylation (kinase activity) with the exception of the 4-[3'-fluoro-5'-(trifluoromethyl)anilino]-6,7-diethoxyquinazolinone analogue **12**. There was no correlation ($r^2 = 0.02$) between binding affinity and inhibition of ligand-induced EGFR tyrosine phosphorylation. Indeed, in contrast to the results of the receptor binding assay, in the EGFR tyrosine phosphorylation assays the diethoxy analogues were not consistently better inhibitors than the corresponding dimethoxy analogues.

Specificity for the EGF receptor versus other receptor tyrosine kinases, especially ErbB2 and ErbB4, is another desirable imaging characteristic. Thus, a small set of compounds was evaluated for inhibition of ligand-induced ErbB2 and ErbB4 tyrosine phosphorylation (kinase activity). The specificity of the compounds tested

Table 3. Inhibition of DNA Synthesis (IC₅₀, nM)

no.	MCF-10A ^a	MCF-7 ^b	MCF-7/ MCF-10A
10a	99 ± 16	1923 ± 456	19.5
10b	108 ± 8	1087 ± 402	10.1
10c	78 ± 12	1695 ± 169	21.7
10d	153 ± 25	1571 ± 263	10.3
11a	59 ± 5	1956 ± 842	33.2
11b	173 ± 24	1982 ± 387	11.5
11c	211 ± 45	1613 ± 123	7.6
11d	585 ± 108	2433 ± 61	4.2
12	>3000	ND	
13	1634 ± 100	4717 ± 1014	2.9
14	188 ± 25	1433 ± 88	7.6
15	489 ± 87	>7000	
16a	141 ± 17	1101 ± 656	7.8
16b	622 ± 69	7487 ± 505	12.0
17a	179 ± 31	1950 ± 260	10.9
17b	612 ± 134	3375 ± 686	5.5

^a EGF-dependent human mammary epithelial cell line. ^b EGF-independent human mammary tumor cell line.

for EGFR appears to be high (Table 2). For the four compounds tested (**10c**, **10d**, **11c**, and **14**), the ErbB2 and ErbB4 tyrosine phosphorylation (kinase) IC₅₀ values were at least 1 order of magnitude greater than the EGFR tyrosine phosphorylation (kinase) IC₅₀ values. We do recognize that this is a comparison of phosphorylation inhibition rather than binding.

The specificity of the molecules for EGFR was also determined by examining their effect on DNA synthesis by the EGF-dependent MCF-10A human mammary epithelial cell line and the EGF-independent MCF-7 human mammary tumor cell line. These cell lines were treated with various concentrations of several of the compounds, and DNA synthesis was measured by assaying ³H-thymidine incorporation. These data were used to calculate the DNA synthesis IC₅₀ values for each compound in the two cell lines, shown in Table 3. The MCF7 DNA synthesis IC₅₀ values for the compounds were at least 1 order of magnitude higher than the corresponding MCF10A DNA synthesis IC₅₀ values. Because the DNA synthesis of MCF10A cells is EGF-dependent, these data suggest that the compounds inhibit MCF10A DNA synthesis by inhibiting the EGFR rather than some other target.

Discussion

There are four well-established criteria for the development of disease-specific radioprobes that would be sensitive to changes in binding site concentration.^{46,47} They are (i) to identify a binding site whose concentration changes as a function of a specific disease, (ii) to design and to produce a radioprobe that selectively targets the binding site, (iii) to evaluate sensitivity as a function of altered binding site concentration, and (iv) to evaluate sensitivity relative to the selected disease. The EGF receptor overexpression in various tumors satisfies the first criterion. Identifying lead molecules to test as imaging probes and developing an underlying selection process to identify future candidate molecules, the subject of the present effort, begins to address the second criterion.

Designing and producing an enzyme- or receptor-binding radioprobe involves several steps, often an iterative process, intended to obtain a thorough understanding of the biochemical and physiological behavior

of the probe to match against a set of desirable imaging characteristics.^{46,47} A receptor-binding radiotracer should meet the following criteria: (i) high affinity for the desired enzyme or receptor, typically <10 nM; (ii) appropriate lipophilicity (coupled to cell membrane or blood-brain barrier penetration, typically 1.5–3.0); (iii) high selectivity for the enzyme or receptor (e.g. low affinity for receptors within the same family or similar proteins, typically >10:1); (iv) suitable metabolic properties (labeled metabolites can alter the distribution profile of the probe); and (v) rapid clearance from nontarget tissues and the body (necessary for good target-to-background ratio, typically >3:1, and lower radiation dose to the subject). In vitro data for a series of dialkoxyquinazoline EGFR-targeted compounds have been gathered and used to select probes for labeling and in vivo evaluation. These data were used to establish a ligand selection process. The process is detailed in the context of the following discussion.

The dialkoxyquinazolines (Figure 1) were chosen as the lead compounds in the present study on the basis of the reported structure inhibition relationships.^{33–35} Fry and colleagues demonstrated that the quinazoline backbone and the 6,7-dimethoxy moieties were necessary for enhanced EGFR tyrosine kinase inhibition. The 6,7-diethoxy analogue **10c** exhibited a 4-fold lower inhibition IC₅₀ value (6 pM vs 25 pM), and halogen substitution at the 2', 3', 4', and 5' positions, even 3',4'-dibromination, of the anilino ring ("C" ring Figure 1) was well-tolerated. The fluorobenzylamino analogue was not previously studied, but on the basis of the radiochemical availability of this analogue, it was included here.⁴⁸ The unsubstituted benzyl compound was evaluated as an inhibitor and found to have a 3-fold lower inhibition IC₅₀ value (10 nM) compared to the corresponding unsubstituted anilino compound (29 nM).^{33,35} The 3-fluoro-5-(trifluoromethyl)aniline as well as 2-, 3-, and 4-fluoroaniline can be synthesized with fluorine-18, so the corresponding nonradioactive analogues were added to the study.⁴⁹ Inhibition data from the 4-[3'-(trifluoromethyl)anilino]-6,7-dimethoxyquinazoline (inhibition IC₅₀ = 0.24 nM) and the 3'-fluoroanilino analogue (inhibition IC₅₀ = 3.8 nM) supported their inclusion.

The preparation of the analogues for this study was straightforward. The 4-chloro-dimethoxy- and 4-chloro-diethoxy-quinazoline intermediates were synthesized and coupled with the appropriate aniline or benzylamine to yield the desired anilino- or (benzylamino)-quinazolines in good yield. The approach outlined herein improved upon the chemistry previously described by Bridges et al.³³ and provides a suitable starting point to generate libraries of anilino- or (benzylamino)-quinazolines with any dialkoxy or mixed alkoxy substituents in the 6 and 7 positions.

A receptor binding assay was developed to study the affinity of the compounds for the tyrosine kinase binding site and determine the correlation, if any, between binding and inhibition values. In vitro equilibrium binding assays provide a good means of segregating potential receptor-based imaging agents from those whose unfavorable kinetics (high binding potential, B_{\max}/K_d) might lead to flow-mediated distribution.⁴⁷ As all of the literature for the quinazolines and other classes of compounds targeting the EGFR tk reported

inhibition constants, the following question was posed: Could these values be used as a primary determinant for potential imaging probes? All of the compounds possessed suitable affinity for the EGFR tk (IC₅₀ range = 0.4–51 nM). Likewise, all of the compounds were nanomolar inhibitors of ligand-induced EGFR tyrosine phosphorylation (IC₅₀ range = 0.8–23.1 nM) in whole cells. Yet, there was a complete lack of correlation between the matched set of binding and phosphorylation data. On the basis of this small set of compounds, the phosphorylation data may not be used to predict receptor affinity. This is likely due to the fact that inhibition of ligand-induced receptor phosphorylation is not only a function of receptor binding affinity but is also a function of the ability to penetrate a cell and gain access to the receptor. Thus, the receptor binding assay along with a measure of cell penetration (inhibition of EGFR dependent DNA synthesis, see below) will be integral as an initial screen for tracers. Compounds **10a–d**, **11b–d**, **12**, and **16a** meet the generally accepted minimum affinity for receptor-based imaging agents, 10 nM.

A subsequent screen of selected lead compounds for the inhibition of ligand-induced tyrosine phosphorylation will identify those molecules that readily access the receptor. For example, compound **12** has a suitable affinity for EGFR in vitro but displays minimal inhibition of ligand-induced receptor tyrosine phosphorylation. Thus, compound **12** probably possesses limited penetration of cells and/or limited EGFR access. Therefore, compound **12** is likely to be of limited value as a potential EGFR-specific tumor imaging agent. Compounds **10a–d**, **11b–d**, and **16a** are potent inhibitors of EGFR phosphorylation.

In cases where facilitated transport is nonexistent, cell membrane permeability is a direct function of the diffusion coefficient and compound lipophilicity. If the partition coefficient value is too low, the compound will not cross a cell membrane; if the value is too high, hydrophobic interactions with lipids and proteins will dominate, leading to high nonspecific binding, compromising image contrast. It has been demonstrated for radiotracers crossing the blood-brain barrier that a log $P_{o/w}$ range of 1.5–3 is optimal.⁵⁶ While this range works equally well for nonbrain receptor-targeted tracers, estrogen radioprobes with higher lipophilicity (up to log $P = 4.5$) have been successful tumor imaging agents.⁵⁷ Therefore, the log $P_{o/w}$ range for the selection of the EGFR probes in the current study has been set from 1.5 to 4. The HPLC-derived lipophilicities (log $P_{o/w}$) ranged from 2.2 to 5.5. Of those compounds previously chosen, **10a**, **11b–d**, and **16a** fall within this desired range.

Selectivity for the chosen receptor is the final in vitro test. One preliminary measure of receptor selectivity that was employed in this study was to evaluate ErbB2 and ErbB4 inhibition. All of the compounds were much less potent inhibitors of ligand-induced ErbB2 and ErbB4 phosphorylation than of ligand-induced EGFR phosphorylation. Indeed, the compounds tested here exhibit between 1 and 2 orders of magnitude of selectivity for the EGFR over ErbB2 or ErbB4. All of the compounds tested may be reasonable EGFR-selective tumor-imaging agents. The optimal agent would exhibit

the least amount of avidity for other RTKs. In many types of tumor cells ErbB2 is frequently overexpressed at levels comparable to the levels of EGFR overexpression.⁷ Similarly, ErbB4 is expressed in the normal mammary epithelium⁵⁹ and infrequently overexpressed in tumor tissue.^{58, 59} These observations highlight the need for EGFR-selective imaging agents that exhibit minimal binding to ErbB2 and ErbB4, among other RTKs, to determine and understand EGFR expression using molecular imaging techniques.

The quinazoline compounds were also assayed for inhibition of cellular DNA synthesis in EGF-dependent (MCF10A) and EGF-independent (MCF7) cell lines. This assay, like the inhibition of ligand-induced receptor tyrosine phosphorylation assays, not only assesses the specificity of a compound for EGFR but also provides a measure of a compounds' ability to penetrate cells and specifically target the EGFR. Thus, it is not surprising that compound **12**, which we hypothesized failed to inhibit ligand-induced EGFR tyrosine phosphorylation because it exhibits limited cell penetration, also failed to inhibit EGF-dependent and -independent DNA synthesis.

In general, the MCF10A DNA synthesis IC₅₀ values are much lower than the MCF7 DNA synthesis IC₅₀ values. This suggests that the compounds inhibit DNA synthesis in the MCF10A cells by primarily targeting EGFR. Indeed, this assay has been used to connote that lavendustin A analogues, which inhibit EGFR kinase activity, inhibit cell proliferation by inhibiting tubulin polymerization.⁵⁰ This indicates that at least some of the compounds inhibit DNA synthesis in the MCF7 cells by targeting a protein other than the EGFR. Compounds that may target ATPases other than the EGFR would be expected to exhibit reduced MCF7/MCF10A DNA synthesis IC₅₀ ratios. Compounds with reduced ratios include **11d**, **13**, and **17b**. Such molecules may not be specific for the EGFR and may not be appropriate for further investigation as potential EGFR-specific tumor imaging agents. Compounds **10a–d**, **11b,c**, **14**, **16a,b**, and **17a** exhibit elevated MCF7/MCF10A DNA synthesis IC₅₀ ratios and by this criterion may be suitable for further investigation as potential EGFR-specific tumor-imaging agents.

These criteria taken together support the further investigation of compounds **10a**, **11b**, **11c**, and **16a**. Compounds **10a** and **16a**, 4-(3'-fluoroanilino)- and 4-(2'-fluoroanilino)-6,7-diethoxyquinazoline, respectively, may both be labeled with the 110-min half-life fluorine-18. Compound **11b**, 4-(3'-chloroanilino)-6,7-dimethoxyquinazoline, may be labeled with the 20-min half-life carbon-11; however, as pointed out earlier, the short half-life may not afford the time necessary to achieve the desired imaging pharmacokinetics. Compound **11c**, 4-(3'-bromoanilino)-6,7-dimethoxyquinaxoline, may be a candidate for bromine-76 (16.2h positron emitter) labeling.

Interestingly, Bonasera et al.³⁸ have recently labeled three of the analogues studied in this paper, **11a**, **13**, and **17b**, and injected one of them, **17b**, into A431 tumor-bearing mice. Compound **13** was not studied in vivo due to low measured EGFR inhibition values, and compound **11a** was not readily labeled with fluorine-18. Tumor uptake of [¹⁸F]**17b** at 30 min was greater than 1% of the injected dose per gram (%ID/g) of tissue,

but the tumor-to-blood ratio was only 0.6. At 60 min the tumor uptake rose slightly to 1.34% ID/g with a tumor-to-blood ratio of 1.62. This uptake may be associated with EGFR targeting, but receptor-blocking studies were not performed to demonstrate receptor-mediated uptake. Bone uptake, an indicator of metabolic defluorination, was also not assessed. A second fluorine-labeled analogue, 4-(3',4'-dichloro-6'-fluoroanilino)-6,7-dimethoxyquinazoline, was tested in tumor-bearing mice. Tumor accumulation of the probe never exceeded blood levels and the uptake was not receptor-mediated. Several factors may have contributed to the limited tumor accumulation in vivo, including high nonspecific binding or diminished availability due to untoward metabolism.

It is clear from the preliminary studies that development of an effective EGFR imaging probe may present some challenges. Thus, effective screening of candidate probes and establishment of baseline data are essential for a radiopharmaceutical development program targeting EGFR.

Conclusion

Anilindialkoxyquinazolines suitable for labeling with radioisotopes were readily prepared using an approach amenable to a synthetic library of aminoquinazoline analogues. All of the dialkoxy analogues possessed suitable affinity for the EGF receptor and all analogues were potent inhibitors of ligand-induced EGFR phosphorylation. They exhibited a range of lipophilicities based on the A ring and C ring substituents. Selectivity, as determined by comparison of ErbB2 and ErbB4 receptor inhibition to EGFR inhibition, favored the EGFR by 1–2 orders of magnitude. On the basis of measures of affinity, lipophilicity, and selectivity, analogues **10a**, **11b**, **11c**, and **16a**, were selected for further evaluation as tumor imaging probes.

Experimental Section

Chemistry. Unless otherwise noted, all solvents and reagents were obtained from commercial suppliers (Aldrich Chemical, Co., Lancaster Synthesis, Inc., VWR, etc.) and were used without further purification. Melting points were determined using a Mel-Temp melting point apparatus and are reported uncorrected. NMR spectra were recorded on either a Bruker VBAMX 300 300 MHz or AMX 400 400 MHz. Chemical shifts are reported in ppm (δ) relative to an internal standard. Elemental analyses were performed by the Microanalytical Laboratory in the College of Chemistry, University of California, Berkeley. Mass spectral data was obtained on a Perkin-Elmer SCIEX mass spectrometer at the SynPep Corp. facility (Dublin, CA) or on a VG ProSpec mass spectrometer at the Mass Spectrometry Facility in the College of Chemistry, University of California, Berkeley.

2-Nitro-4,5-diethoxybenzoic Acid (2). A flask immersed in a room-temperature water bath was charged with 3,4-diethoxybenzoic acid **1** (12.9 g, 61.4 mmol) and acetic acid (glacial, 52 mL). Over a 15-min period, HNO₃ (70%, 54 mL) was added dropwise with stirring. The deep orange solution was stirred for an additional 125 min at room temperature. The reaction was quenched upon addition of 110 g of ice. A yellow precipitate formed that was filtered and washed with H₂O (3 × 50 mL). The resulting yellow-white solid was dissolved in ether (150 mL). The ether was washed with 1 N NaOH (3 × 60 mL). The aqueous washings were combined and acidified with concentrated HCl, resulting in the production of a pale yellow precipitate. The precipitate was filtered, washed with H₂O (3 × 100 mL), and dissolved in ether. The

ether was dried over MgSO_4 and concentrated in vacuo to yield 10.55 g (67%) of **2**: mp 142–145 °C; $^1\text{H NMR}$ (CDCl_3) δ 7.36 (s, 1H, *H*-6), 7.21 (s, 1H, *H*-3), 4.20 (q, 2H, CH_2CH_3 , $J = 6.9$ Hz), 4.17 (q, 2H, CH_2CH_3 , $J = 6.9$ Hz), 1.50 (t, 6H, CH_2CH_3 , $J = 7.1$ Hz); EI MS [M^+] 255 (100).

2-Amino-4,5-diethoxybenzoic Acid (4). Method a. A round-bottom flask containing 2-amino-4,5-diethoxy-methyl benzoate **3** (1.00 g, 4.19 mmol, Sigma-Aldrich Rare Chemicals) was mixed with 6.25 N NaOH (2.67 mL, 16.7 mmol) and water (5.0 mL). The solution was refluxed for 1 h to give a clear brown solution. After cooling at ambient temperature for 10 min, water (15 mL) was added to the flask and the solution was titrated to pH 6 with 1 N HCl. The solution was further cooled in an ice bath for 30 min. The precipitate was filtered, washed with water (100 mL) and pentanes (50 mL), and dried in vacuo over P_2O_5 overnight to give 0.83 g (76%) of 2-amino-4,5-diethoxybenzoic acid as a pale yellow solid.

Method b. An oven-dried round-bottomed flask (250 mL) equipped with a stirring bar was immersed in a room-temperature water bath and charged with $\text{SnCl}_2 \cdot 2\text{H}_2\text{O}$ (24.9 g, 110 mmol) and HCl (concentrated, 100 mL). 2-Nitro-4,5-diethoxybenzoic acid **2** (1.64 g, 6.43 mmol) was added and the mixture was stirred for 120 min at room temperature. The mixture was diluted with HCl (concentrated, 40 mL), filtered, and washed with concentrated HCl. The resulting white solid was taken up in H_2O (500 mL) and filtered to remove the remaining undissolved material. The pH of the filtrate was adjusted to 4.5 using NH_4OH . The deep purple solution was extracted with CH_2Cl_2 (3 \times 300 mL). The organic washings were combined, dried over MgSO_4 , and concentrated in vacuo, to yield 0.64 g (44%) of a purple-white powder

: mp 120–126 °C; $^1\text{H NMR}$ (CDCl_3) δ 7.37 (s, 1H, *H*-6), 6.11 (s, 1H, *H*-3), 4.06 (q, 2H, CH_2CH_3 , $J = 6.8$ Hz), 4.01 (q, 2H, CH_2CH_3 , $J = 6.8$ Hz), 1.46 (t, 3H, CH_2CH_3 , $J = 6.8$ Hz), 1.41 (t, 3H, CH_2CH_3 , $J = 6.8$ Hz); EI MS [M^+] 225 (100).

6,7-Diethoxyquinazolin-4-one (6). A modified literature procedure³³ was developed to produce **6**. A 500 mL round-bottom flask equipped with an air condenser was charged with 2-amino-4,5-diethoxybenzoic acid **4** (1.36 g, 6.04 mmol) and formamidine hydrochloride (0.70 g, 8.76 mmol). The solids were thoroughly mixed and then heated to 200 °C under an argon atmosphere for 15 min. The heating block temperature was adjusted to 80 °C and the solution cooled to 80 °C over 40 min. Dilute NaOH (0.33 N, 20 mL) was added to the flask. The mixture was sonicated at room temperature for 1 h, producing in a dark gray-purple suspension. The solid was filtered and washed with water (200 mL), pentanes (200 mL), and ethyl acetate (200 mL) to give 6,7-diethoxyquinazolin-4-one as an off-white solid. Drying in vacuo overnight over P_2O_5 gave 0.80 g (57%) of **6**: mp 248–251.5 °C; $^1\text{H NMR}$ (CD_3OD) δ 7.98 (s, 1H, *ArH*), 7.55 (s, 1H, *ArH*), 7.11 (s, 1H, *ArH*), 4.20 (q, 2H, CH_2CH_3 , $J = 6.8$ Hz), 4.16 (q, 2H, CH_2CH_3 , $J = 6.8$ Hz), 1.48 (t, 3H, CH_2CH_3 , $J = 6.8$ Hz), 1.46 (t, 3H, CH_2CH_3 , $J = 6.8$ Hz); EI MS [M^+] 234 (100).

6,7-Dimethoxyquinazolin-4-one (7). 2-Amino-4,5-dimethoxybenzoic acid **5** (5.0 g, 25.3 mmol) was converted to **7** in 65% yield (3.64 g) following the procedure for compound **6**: mp 278–278.5 °C (lit.³³ mp 295–298 °C); $^1\text{H NMR}$ [$(\text{CD}_3)_2\text{SO}$] δ 12.07 (br s, 1H, *NH*), 7.98 (s, 1H, *ArH*), 7.43 (s, 1H, *ArH*), 7.11 (s, 1H, *ArH*), 3.90 (s, 3H, OCH_3), 3.87 (s, 3H, OCH_3).

4-Chloro-6,7-diethoxyquinazoline (8). Following a modified literature procedure,³³ DMF (0.94 mL, 12.1 mmol) was added dropwise to a solution of 1,2-dichloroethane (8.1 mL) and oxalyl chloride (1.1 mL, 12.6 mmol) stirring under argon, vigorously releasing gas. Following cessation of the gas production, 6,7-diethoxyquinazolin-4-one **6** (1.89 g, 8.1 mmol) was added to the thick white slurry and then refluxed for 2.5 h, resulting in a yellow-brown suspension. The reaction was quenched by addition of Na_2HPO_4 (0.5 M, 16.8 mL) followed by stirring in an ice bath for 1 h. The suspension was filtered and washed with water (200 mL) to isolate 4-chloro-6,7-diethoxyquinazoline **8** as a pale gray solid (1.48 g, 73%): mp 139–140 °C; $^1\text{H NMR}$ (CDCl_3) δ 8.82 (s, 1H, *ArH*), 7.35 (s, 1H, *ArH*), 7.28 (s, 1H, *ArH*), 4.26 (m, 4H, OCH_2CH_3), 1.55 (m, 6H,

OCH_2CH_3); APCI MS [$M + 1$] 253.1 (100), 255.1 (33). Anal. ($\text{C}_{12}\text{H}_{13}\text{ClN}_2\text{O}_2$) C, H, N.

4-Chloro-6,7-dimethoxyquinazoline (9). 6,7-Dimethoxyquinazolin-4-one **7** (3.40 g, 16.5 mmol) was converted to **9** in 52% yield (1.91 g) following the procedure for compound **8**: mp 184–186 °C; $^1\text{H NMR}$ (CDCl_3) δ 8.86 (s, 1H, *ArH*), 7.38 (s, 1H, *ArH*), 7.32 (s, 1H, *ArH*), 4.05 (s, 6H, OCH_3); APCI MS [$M + 1$] 225.1 (100), 227.1 (30). Anal. ($\text{C}_{10}\text{H}_9\text{ClN}_2\text{O}_2$) C, H, N.

4-(3'-Chloroanilino)-6,7-diethoxyquinazoline Hydrochloride (10b). General Coupling Procedure. A solution of 4-chloro-6,7-diethoxyquinazoline **8** (71 mg, 0.32 mmol) and 3-chloroaniline (40 μL , 0.38 mmol) in 3 mL of DMF was heated at 80 °C under argon for 40 min. The reaction was cooled at room temperature for 1 h. Ethyl acetate (2 mL) was added. The resulting precipitate was filtered and further washed with 20 mL of ethyl acetate to give the HCl salt **10a** (100 mg, 90%). The salt was converted to the free base by dissolving **10b** in a mixture of 3 mL of ethyl acetate and 3 mL of 1 N NaOH. The biphasic mixture was stirred vigorously for several minutes. The ethyl acetate layer was filtered, washed with water (3 \times 1 mL), and dried over MgSO_4 . The ethyl acetate was filtered and the volume was reduced to less than 1 mL. The solution was applied to a normal phase semipreparative HPLC (Whatman M9/50 partisol 10 column, 70:30 EtOAc: hexane, 6 mL/min., UV 254 nm) for purification. The fraction containing the free base was concentrated to dryness and the residue was dissolved in MeOH (8 mL) with gentle heating. HCl (1 N, 3 mL) was added and the solution was placed in an ice bath. The precipitate was filtered and washed with ethyl acetate to give 72 mg (65%) of the 4-(3-chloroanilino)-6,7-diethoxyquinazoline hydrochloride salt **10b**: mp 260–261 °C; $^1\text{H NMR}$ (CD_3OD) δ 8.72 (s, 1H, *H*-2), 7.97 (s, 1H, *Ar-H*), 7.87 (s, 1H, *H*-2'), 7.65 (d, 1H, *H*-4', $J = 8.0$ Hz), 7.47 (t, 1H, *H*-5' $J = 8.1$ Hz), 7.35 (d, 1H, *H*-6', $J = 8.0$ Hz), 7.20 (s, 1H, *Ar-H*), 4.31 (q, 4H, OCH_2CH_3 , $J = 6.9$ Hz), 1.55 (t, 6H, OCH_2CH_3 , $J = 7.0$ Hz); EI MS [M^+] 343 (100), 345 (34). Anal. ($\text{C}_{18}\text{H}_{18}\text{ClN}_3\text{O}_2$) C, H, N.

4-(3'-Fluoroanilino)-6,7-diethoxyquinazoline Hydrochloride (10a). Similar treatment of **8** (0.021 g, 0.083 mmol) with 3-fluoroaniline yielded **10a** (77%): mp 247.0–248.0 °C; $^1\text{H NMR}$ (CD_3OD) δ 8.73 (s, 1H, *H*-2), 7.99 (s, 1H, *H*-5), 7.66 (d, $J = 10.6$ Hz, 1H, *H*-2'), 7.53 (m, 1H, *H*-5'), 7.53 (m, 1H, *H*-6'), 7.21 (s, 1H, *H*-8), 7.08 (dd, $J = 9.1$, 7.1 Hz, 1H, *H*-4'), 4.33 (q, $J = 6.9$ Hz, 4H, $\text{CH}_3\text{CH}_2\text{O}$), 1.55 (t, $J = 6.9$ Hz, 3H, $\text{CH}_3\text{CH}_2\text{O}$), 1.54 (t, $J = 6.9$ Hz, 3H, $\text{CH}_3\text{CH}_2\text{O}$); EI MS [M] 327. Anal. ($\text{C}_{18}\text{H}_{18}\text{FN}_3\text{O}_2 \cdot \text{HCl}$) C, H, N.

4-(3'-Bromoanilino)-6,7-diethoxyquinazoline Hydrochloride (10c). Similar treatment of **8** (59.5 mg, 0.24 mmol) with 3-bromoaniline yielded **10c** (50%): mp 250–252 °C (lit.³³ mp 155–167 °C, free base); $^1\text{H NMR}$ [$(\text{CD}_3)_2\text{SO}$] δ 11.17 (br s, 1H, *NH*), 8.86 (s, 1H, *H*-2), 8.19 (s, 1H, *Ar-H*), 7.99 (s, 1H, *H*-2'), 7.73 (d, 1H, *H*-4', $J = 7.6$ Hz), 7.51 (d, 1H, *H*-6', $J = 7.5$ Hz), 7.45 (t, 1H, *H*-5' $J = 8.0$ Hz), 7.23 (s, 1H, *Ar-H*), 4.27 (m, 4H, OCH_2CH_3 , $J = 7.2$ Hz), 1.45 (t, 6H, OCH_2CH_3 , $J = 6.8$ Hz); HREIMS calcd for $\text{C}_{18}\text{H}_{18}\text{BrN}_3\text{O}_2$ m/z (M^+) 387.05824, 389.05619, found 387.05776, 389.05532. Anal. ($\text{C}_{18}\text{H}_{18}\text{BrN}_3\text{O}_2$) C, H, N.

4-(3'-Iodoanilino)-6,7-diethoxyquinazoline Hydrochloride (10d). Similar treatment of **8** (116.1 mg, 0.46 mmol) with 3-iodoaniline yielded **10d** (95%): mp 258.5–261 °C; $^1\text{H NMR}$ [$(\text{CD}_3)_2\text{SO}$] δ 11.19 (br s, 1H, *NH*), 8.86 (s, 1H, *H*-2), 8.19 (s, 1H, *Ar-H*), 8.10 (s, 1H, *H*-2'), 7.75 (d, 1H, *H*-4', $J = 8.0$ Hz), 7.67 (d, 1H, *H*-6', $J = 8.0$ Hz), 7.30 (s, 1H, *Ar-H*), 7.29 (t, 1H, *H*-5' $J = 8.0$ Hz), 4.27 (m, 4H, OCH_2CH_3), 1.44 (t, 6H, OCH_2CH_3 , $J = 6.8$ Hz). APCI MS [M^+] 436.0 (100). Anal. ($\text{C}_{18}\text{H}_{18}\text{IN}_3\text{O}_2 \cdot \text{HCl}$) C, H, N.

4-(3'-Fluoroanilino)-6,7-dimethoxyquinazoline Hydrochloride (11a). Similar treatment of **9** (0.032 g, 0.14 mmol) with 3-fluoroaniline yielded **11a** (27%): mp 244.5–246.0 °C (lit.³³ mp 253–254 °C); $^1\text{H NMR}$ (CD_3OD) δ 8.72 (s, 1H, *H*-2), 7.82 (s, 1H, *H*-5), 7.47 (d, $J = 10.6$ Hz, 1H, *H*-2'), 7.33 (m, 1H, *H*-5'), 7.33 (m, 1H, *H*-6'), 7.03 (s, 1H, *H*-8), 6.89 (dd, $J = 8.6$, 7.3 Hz, 1H, *H*-4'), 3.88 (s, 6H, CH_3O); EI MS [M] 299. Anal. ($\text{C}_{16}\text{H}_{14}\text{FN}_3\text{O}_2 \cdot 1.1\text{HCl}$) C, H, N.

4-(3'-Chloroanilino)-6,7-dimethoxyquinazoline Hydrochloride (11b). Similar treatment of **9** (71.1 mg, 0.32 mmol) with 3-chloroaniline yielded **11b** (90%): mp 230–235 °C (lit.³³ mp 261–262 °C); ¹H NMR (CD₃OD) δ 8.66 (s, 1H, *H*-2), 7.93 (s, 1H, *Ar*-*H*), 7.90 (s, 1H, *H*-2'), 7.66 (d, 1H, *H*-4', *J* = 8.0 Hz), 7.44 (t, 1H, *H*-5' *J* = 8.1 Hz), 7.29 (d, 1H, *H*-6', *J* = 8.0 Hz), 7.21 (s, 1H, *Ar*-*H*), 4.02 (s, 6H, OCH₃); HREIMS calcd for C₁₆H₁₄ClN₃O₂ (*M*+*z*) 315.07745, 317.07450, found 315.07682, 317.07303. Anal. (C₁₆H₁₄ClN₃O₂·HCl) C, H, N.

4-(3'-Bromoanilino)-6,7-dimethoxyquinazoline Hydrochloride (11c). Similar treatment of **9** (312 mg, 1.39 mmol) with 3-bromoaniline yielded **11c** (87%): mp 256–257.5 °C; ¹H NMR [(CD₃)₂SO] δ 11.35 (br s, 1H, *NH*), 8.88 (s, 1H, *H*-2), 8.29 (s, 1H, *Ar*-*H*), 8.01 (s, 1H, *H*-2'), 7.77 (d, 1H, *H*-4', *J* = 8.0 Hz), 7.50 (d, 1H, *H*-6', *J* = 8.0 Hz), 7.45 (t, 1H, *H*-5' *J* = 8.0 Hz), 7.32 (s, 1H, *Ar*-*H*), 4.00 (s, 3H, OCH₃), 3.99 (s, 3H, OCH₃); EI MS [*M*+*z*] 359 (100), 361 (97). Anal. (C₁₆H₁₄BrN₃O₂·HCl) C, H, N.

4-(3'-Iodoanilino)-6,7-dimethoxyquinazoline Hydrochloride (11d). Similar treatment of **9** (276 mg, 1.23 mmol) with 3-iodoaniline yielded **11d** (92%): mp 251–251.5 °C (lit.³³ mp 273 °C); ¹H NMR [(CD₃)₂SO] δ 11.33 (br s, 1H, *NH*), 8.90 (s, 1H, *H*-2), 8.30 (s, 1H, *Ar*-*H*), 8.28 (s, 1H, *H*-2'), 7.79 (d, 1H, *H*-4', *J* = 8.0 Hz), 7.69 (d, 1H, *H*-6', *J* = 8.0 Hz), 7.35 (s, 1H, *Ar*-*H*), 7.31 (t, 1H, *H*-5' *J* = 8.0 Hz), 4.04 (s, 3H, OCH₃), 4.02 (s, 3H, OCH₃); APCI MS [*M*+*z*] 408.0 (100). Anal. (C₁₆H₁₄I-N₃O₂·HCl) C, H, N.

4-[3'-Fluoro-5'-(trifluoromethyl)anilino]-6,7-diethoxyquinazoline Hydrochloride (12). Similar treatment of **8** (58.5 mg, 0.23 mmol) with 3-fluoro-5-(trifluoromethyl)aniline yielded **12** (92%): mp 278–280 °C; ¹H NMR (CDCl₃) δ 8.70 (s, 1H, *H*-2), 8.09 (d, 1H, *H*-4', *J*_{H,F} = 10.9), 7.62 (s, 1H, *Ar*-*H*), 7.26 (s, 1H, *H*-6'), 7.22 (br s, 1H, *NH*), 7.06 (d, 1H, *H*-2', *J*_{H,F} = 8.0 Hz), 7.00 (s, 1H, *Ar*-*H*), 4.24 (m, 4H, OCH₂CH₃), 1.55 (m, 6H, OCH₂CH₃); HREIMS calcd for C₁₉H₁₇F₄N₃O₂ (*M*+*z*) 395.12569, found 395.12596. Anal. (C₁₉H₁₇F₄N₃O₂·HCl) C, H, N.

4-[3'-Fluoro-5'-(trifluoromethyl)anilino]-6,7-dimethoxyquinazoline Hydrochloride (13). Similar treatment of **9** (150 mg, 0.67 mmol) with 3-fluoro-5-(trifluoromethyl)aniline yielded **13** (19%): mp 269–270.5 °C; ¹H NMR (CDCl₃) δ 8.72 (s, 1H, *H*-2), 8.08 (d, 1H, *H*-4', *J*_{H,F} = 10.8), 7.63 (s, 1H, *Ar*-*H*), 7.38 (br s, 1H, *NH*), 7.27 (s, 1H, *H*-6'), 7.06 (d, 1H, *H*-2', *J*_{H,F} = 8.4 Hz), 7.02 (s, 1H, *Ar*-*H*), 4.03 (s, 3H, OCH₃), 4.01 (s, 3H, OCH₃); APCI MS [*M*+*z*] 368.1 (100). Anal. (C₁₇H₁₃F₄N₃O₂·HCl) C, H, N.

4-[(4'-Fluorobenzyl)amino]-6,7-diethoxyquinazoline Hydrochloride (14). Similar treatment of **8** (66.9 mg, 0.26 mmol) with 4-fluorobenzylamine yielded **14** (84%): mp 238.5–240 °C; ¹H NMR (CDCl₃) δ 8.56 (s, 1H, *H*-2), 7.37 (dd, 2H, *H*-2', *H*-6', *J*_{H,H} = 8.2, *J*_{H,F} = 5.5), 7.18 (s, 1H, *Ar*-*H*), 7.03 (t, 2H, *H*-3', *H*-5', *J*_{H,F} = *J*_{H,F} = 8.6 Hz), 6.84 (s, 1H, *Ar*-*H*), 5.53 (br s, 1H, *NH*), 4.81 (d, 2H, -CH₂-, *J* = 5.4), 4.20 (q, 2H, OCH₂CH₃, *J* = 7.0 Hz), 4.12 (q, 2H, OCH₂CH₃, *J* = 7.0 Hz), 1.51 (t, 3H, OCH₂CH₃, *J* = 7.0 Hz), 1.49 (t, 3H, OCH₂CH₃, *J* = 7.0 Hz); EI MS [*M*+*z*] 341 (100). Anal. (C₁₉H₂₀FN₃O₂) C, H, N.

4-[(4'-Fluorobenzyl)amino]-6,7-dimethoxyquinazoline Hydrochloride (15). Similar treatment of **9** (320 mg, 0.143 mmol) with 4-fluorobenzylamine yielded **15** (22%): mp 250–251 °C; ¹H NMR [(CD₃)₂SO] δ 10.37 (br s, 1H, *NH*), 8.82 (s, 1H, *H*-2), 8.00 (s, 1H, *Ar*-*H*), 7.45 (dd, 2H, *H*-2', *H*-6', *J*_{H,H} = 8.4, *J*_{H,F} = 5.6), 7.22 (s, 1H, *Ar*-*H*), 7.19 (t, 2H, *H*-3', *H*-5', *J*_{H,F} = *J*_{H,F} = 8.8 Hz), 4.91 (d, 2H, -CH₂-, *J* = 5.2), 3.95 (s, 3H, OCH₃), 3.94 (s, 3H, OCH₃); APCI MS [*M*+*z*] 314.2 (100). Anal. (C₁₇H₁₆FN₃O₂·HCl) C, H, N.

4-(2'-Fluoroanilino)-6,7-diethoxyquinazoline Hydrochloride (16a). Similar treatment of **8** (0.024 g, 0.095 mmol) with 2-fluoroaniline yielded **16a** (61%): mp 220.5–222.0 °C; ¹H NMR (CD₃OD) δ 8.62 (s, 1H, *H*-2), 7.91 (s, 1H, *H*-5), 7.57 (dd, *J* = 7.7, 7.5 Hz, 1H, *H*-5'), 7.44 (m, 1H, *H*-4'), 7.32 (d, *J* = 7.7 Hz, 1H, *H*-6'), 7.28 (m, 1H, *H*-3'), 7.20 (s, 1H, *H*-8), 4.32 (q, *J* = 6.6 Hz, 2H, CH₃CH₂O), 4.29 (q, *J* = 6.6 Hz, 2H, CH₃CH₂O), 1.55 (t, *J* = 6.6 Hz, 3H, CH₃CH₂O), 1.54 (t, *J* = 6.6 Hz,

3H, CH₃CH₂O); HREIMS calcd for C₁₈H₁₈FN₃O₂ (*M*+*z*) 327.13831, found 327.13779. Anal. (C₁₈H₁₈FN₃O₂·1.4HCl) C, H, N.

4-(4'-Fluoroanilino)-6,7-diethoxyquinazoline Hydrochloride (16b). Similar treatment of **8** (0.10 g, 0.40 mmol) with 4-fluoroaniline yielded **16b** (97%): mp 252.0–255.0 °C; ¹H NMR ((CD₃)₂SO) δ 10.06 (s, 1H, *NH*), 8.77 (d, *J* = 1.8 Hz, 1H, *H*-2), 8.27 (s, 1H, *H*-5), 7.68 (m, 2H, *H*-2', *H*-6'), 7.32 (s, 1H, *H*-8), 7.29 (dd, *J* = 6.7, 6.6 Hz, 2H, *H*-3', *H*-5'), 4.25 (q, *J* = 5.1 Hz, 2H, CH₃CH₂O), 4.20 (q, *J* = 5.1 Hz, 2H, CH₃CH₂O), 1.41 (t, *J* = 5.1 Hz, 3H, CH₃CH₂O), 1.40 (t, *J* = 5.1 Hz, 3H, CH₃CH₂O); APCI MS [*M*+*z*] 328.3. Anal. (C₁₈H₁₈FN₃O₂·HCl) C, H, N.

4-(2'-Fluoroanilino)-6,7-dimethoxyquinazoline Hydrochloride (17a). Similar treatment of **9** (0.033 g, 0.15 mmol) with 2-fluoroaniline yielded **17a** (50%): mp 231.0–232.0 °C; ¹H NMR (CD₃OD) δ 8.65 (s, 1H, *H*-2), 7.97 (s, 1H, *H*-5), 7.58 (ddd, *J* = 7.7, 7.5 Hz, 1.10 Hz, 1H, *H*-5'), 7.44 (m, 1H, *H*-4'), 7.33 (d, *J* = 7.7 Hz, 1H, *H*-6'), 7.29 (m, 1H, *H*-3'), 7.25 (s, 1H, *H*-8), 4.09 (s, 3H, CH₃O), 4.07 (s, 3H, CH₃O); HREIMS calcd for C₁₆H₁₄FN₃O₂ (*M*+*z*) 299.10701, found 299.10648. Anal. (C₁₆H₁₄FN₃O₂·HCl) C, H, N.

4-(4'-Fluoroanilino)-6,7-dimethoxyquinazoline Hydrochloride (17b). Similar treatment of **9** (0.10 g, 0.45 mmol) with 4-fluoroaniline yielded **17b** (94%): mp 247.0–248.0 °C; ¹H NMR ((CD₃)₂SO) δ 10.12 (br s, 1H, *NH*), 8.79 (s, 1H, *H*-2), 8.25 (s, 1H, *H*-5), 7.68 (m, 2H, *H*-2', *H*-6'), 7.30 (dd, *J* = 7.0, 6.2 Hz, 2H, *H*-3', *H*-5'), 7.30 (s, 1H, *H*-8), 3.98 (s, 3H, CH₃O), 3.96 (s, 3H, CH₃O); APCI MS [*M*+*z*] 300.1. Anal. (C₁₆H₁₄FN₃O₂·HCl) C, H, N.

log *P* Determinations. log *P* values were estimated from the log *k'*_w values determined by HPLC chromatography following the procedure of Minick.⁴⁴ The solvents were HPLC-grade methanol, 1-octanol, *n*-decylamine, and distilled, deionized water. The standards [*p*-anisidine (*t*_R = 0.95), acetophenone (*t*_R = 1.58), *p*-bromoaniline (*t*_R = 2.26), naphthalene (*t*_R = 3.30), pyrene (*t*_R = 4.88)] were obtained from Aldrich and used without further purification. A ThermoQuest HPLC system equipped with an autoinjector, pump, diode-array UV/vis detector, and an Es Industries MC8 column (4.6 × 150 mm, 5μ, 60Å) was used for these measurements. The organic mobile phase was methanol containing 0.25% v/v 1-octanol. The aqueous mobile phase was octanol-saturated water containing 0.02 M 3-morpholinopropanesulfonic acid (MOPS) buffer, 0.15% v/v *n*-decylamine, adjusted to pH 7.4. The flow rate was set at 1 mL/min. The quinazolines and standards were dissolved in methanol to a final concentration of approximately 0.1 mg/mL and 10 μL was injected onto the column. The column void volume was estimated from the retention time of uracil, which was included as a nonretained internal reference standard with each injection. The log *k'*_w was determined by linear extrapolation of the compound residence time (retention volume less void volume) versus the methanol concentration over a range of 60 to 85% methanol mobile phase.

Biological Methods. Cell Lines and Cell Culture. The CEM human T lymphocyte cell line engineered to express ErbB4 (CEM4) and its culture conditions have been described previously.^{51,52} Briefly, these cells were propagated in RPMI supplemented with 10% heat-inactivated fetal bovine serum and 300 μg/mL G418. The BaF3 mouse lymphoid cell lines engineered to express either EGFR (BaF3/EGFR) or ErbB2 and ErbB3 together (BaF3/2+3) and the culture conditions for these cell lines have been described earlier.⁵³ Briefly, these cells were propagated in RPMI supplemented with 10% fetal bovine serum, 300 μg/mL G418, and 10% medium conditioned by WeHI cells. This conditioned medium serves as a source for Interleukin 3.

MCF-10A human mammary epithelial cells and MCF-7, MDA-MB-231, and MDA-MB-453 human mammary tumor cell lines were obtained from the American Type Culture Collection (ATCC). These lines were propagated according to ATCC recommendations.

Inhibition of Receptor Tyrosine Phosphorylation. The assay for inhibition of ErbB family receptor tyrosine phospho-

rylation was adapted from a previously described protocol.^{52,53} Briefly, 200 mL cultures of CEM/4, BaF3/EGFR, or BaF3/2+3 cells were grown to saturation density ($\sim 10^6$ cells/mL) and were incubated for 24 h at 37 °C in serum-free medium to reduce basal levels of receptor tyrosine phosphorylation. The cells were collected by centrifugation and resuspended in serum-free medium at a final concentration of $\sim 10^7$ cells/mL (~ 20 mL of cells). Cells were transferred to microcentrifuge tubes in 1-mL aliquots and putative kinase inhibitors were added to the cells. Each tyrosine kinase inhibitor was tested at three to five different concentrations. The inhibitors were dissolved in 5 μ L DMSO; hence, cells treated with 5 μ L DMSO were used as a solvent control. Cells were incubated in the presence of inhibitor for 2 h at 37 °C and then incubated on ice for 20 min. Chilling the cells reduces the amount of ligand-induced receptor downregulation.⁵³

Ligand was then added to the appropriate samples at a final concentration of 100 ng/mL, and the samples were mixed and incubated on ice for 7 min. Recombinant human epidermal growth factor (EGF; Sigma) was used as the ligand for EGFR, whereas neuregulin1 β (NRG1 β ; R&D Systems) was used as the ligand for ErbB3 and ErbB4. Note that because ErbB3 lacks kinase activity, ligand-induced ErbB2 and ErbB3 phosphorylation in the BaF3/2+3 cells is the result of ligand-induced ErbB2–ErbB3 heterodimerization and ErbB2 kinase activity.⁵³ Following incubation with ligand, the cells were collected by centrifugation, the supernatant was removed by aspiration, and the cells were resuspended in an isotonic lysis buffer containing 0.5% NP40/Igepal CA-630 (nonionic detergent; Sigma).

The cells were incubated for 20 min on ice to permit lysis. The samples were centrifuged for 10 min at 4 °C to collect the nuclei and cellular debris. The supernatants (cell lysates) were transferred to fresh tubes. Concanavalin A Sepharose (Amersham/Pharmacia) beads were added to each sample (35 μ L of a 50% v/v slurry) and the samples were incubated at 4 °C for 30 min. Concanavalin A Sepharose precipitates the cellular glycoproteins, which include ErbB family receptors. The precipitated glycoproteins were washed three times with 500 μ L of ice-cold lysis buffer and then were eluted by boiling the beads for 5 min in 80 μ L of reducing SDS protein sample buffer. The beads were collected by centrifugation and half of the eluted glycoproteins (40 μ L) were recovered and resolved by SDS/PAGE on a 7.5% acrylamide gel.

The resolved glycoproteins were electroblotted onto nitrocellulose (BiotraceNT, Gelman Sciences). The resulting blot was blocked by incubation for 45 min at room temperature in a solution consisting of 5% bovine serum albumin (Sigma) dissolved in Tris-buffered normal saline (TBS) supplemented with 0.05% Tween-20 (TBS-T). The blot was then probed with a mouse monoclonal antiphosphotyrosine antibody (4G10, Upstate Biotechnology). The blot was washed with TBS-T five times for 6 min each, and primary antibody binding was detected by probing the blot with a goat anti-mouse antibody conjugated to horseradish peroxidase (HRP; Pierce). The blot was washed with TBS-T 12 times for 10 min each, after which HRP activity was visualized by enhanced chemiluminescence (ECL; Amersham Pharmacia Biotech). The resulting chemilumigrams were digitized using a Linotype-Hell Jade flatbed scanner and the amount of receptor tyrosine phosphorylation was quantified using NIH Image software. The amount of receptor tyrosine phosphorylation in samples from cells treated with a putative receptor tyrosine kinase inhibitor was compared to a standard curve generated using samples from cells treated with DMSO solvent control. This enabled us to determine the concentration of a given tyrosine kinase inhibitor that was necessary to cause a 50% reduction in receptor tyrosine phosphorylation. This value is reported as the receptor tyrosine phosphorylation IC₅₀ value.

Inhibition of Cellular DNA Synthesis. The assay for inhibition of cellular DNA synthesis was adapted from a previously described protocol.⁵⁴ Briefly, human mammary (tumor) cells were seeded in 1-mL aliquots into 24-well culture dishes at a density of 10^5 cells/well. Cells were incubated for

24 h at 37 °C, and a tyrosine kinase inhibitor dissolved in DMSO was added to each well in a volume of 10 μ L. Each tyrosine kinase inhibitor was assayed at three to five different concentrations and each concentration was assayed using three to four wells of cells. Cells treated with 10 μ L of DMSO served as the solvent control. Cells were then incubated for 48 h at 37 °C. ³H-Thymidine (1.5 μ Ci, Amersham Pharmacia Biotech) dissolved in a 1.5 μ L of an aqueous solution was added to each well, and the cells were incubated for an additional 2 h at 37 °C. The culture medium was aspirated from the wells, and the cells were rinsed once with 1 mL of ice-cold phosphate-buffered saline (PBS) and once with 1 mL of ice-cold 10% trichloroacetic acid (TCA). Incorporated ³H-thymidine was precipitated by incubating the cells for at least 30 min at 4 °C in 1 mL of 10% TCA. Following incubation, the TCA solution was aspirated from each well and the precipitated (incorporated) ³H-thymidine was solubilized by incubating the cells for 30 min at 95 °C in 500 μ L of 3% perchloric acid. The perchloric acid extracts were transferred to scintillation vials containing 10 mL of Cytosint scintillation cocktail (ICN). The incorporated ³H-thymidine was assayed by scintillation counting on a Packard Tricarb scintillation counter. The amount of ³H-thymidine incorporation observed in the cells treated with the solvent control was divided by two to determine the amount of half-maximal ³H-thymidine incorporation. Dose–response curves for each combination of putative tyrosine kinase inhibitor and cell line were then constructed using the ³H-thymidine incorporation data. The dose–response curves and the half-maximal ³H-thymidine values were used to calculate the concentration of each inhibitor required to inhibit ³H-thymidine incorporation by 50% in a given cell line. This value is reported as the DNA synthesis IC₅₀ value.

In Vitro EGFR Binding Assay. EGFR tyrosine kinase receptor binding was determined by a competitive radiometric assay using [¹²⁵I]-4-(3'-iodoanilino)-6,7-dimethoxyquinazoline³⁶ as the radiotracer (specific activity = 590 Ci/mmol). Various concentrations (10^{-11} M– 10^{-6} M) of the quinazoline compounds were prepared in binding buffer (10 mM HEPES, 1 mM EDTA, 5 mM MgCl₂, 0.1% BSA, 10 μ g/mL leupeptin, 10 μ g/mL pepstatin, 0.5 μ g/mL aprotin, and 200 μ g/mL bacitracin (pH 7.4)). Commercially available (Receptor Biology, Beltsville, MD) A431 human carcinoma cell membrane were diluted in ice-cold binding buffer (50 μ L of 0.06 μ g/ μ L stock solution) and homogenized with a hand-held homogenizer. This preparation was added to the buffer solution followed by the addition of 1 μ Ci of the radiotracer to initiate the binding assay. The mixture was incubated at room temperature in the dark for 60 min. The incubation was terminated by the addition of 5 mL of ice-cold buffer (10 mM HEPES, 1 mM EDTA, 5 mM MgCl₂, and 0.1% BSA (pH 7.4)), and the solutions were filtered through polyethylenamine-soaked (0.5% solution, 30 min) GF/B filter paper (Brandel, Gaithersburg, MD) using a Brandel cell harvester, followed by two washes (5 mL each) with wash buffer. The filter paper was dried and counted for 10 min using a TM Analytic γ well counter. Nonspecific binding was determined by adding 1 μ M 4-(3'-bromoanilino)dimethoxyquinazoline to the assay. Inhibition constants at 50% of specific binding (IC₅₀) were derived from specific binding versus concentration curves. Triplicate assays were performed for each compound.

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Supporting Information Available: Results from elemental analysis. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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