

Targeted Inhibitors of Epidermal Growth Factor Receptor

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Members of the epidermal growth factor receptor (EGFR) family of tyrosine kinase receptors are major regulators of cellular proliferation, differentiation, and survival. In humans, abnormal activation of EGFR is associated with the development and progression of many cancer types, which makes it an attractive target for molecular-guided therapy. Two classes of EGFR-targeted cancer therapeutics include monoclonal antibodies (mAbs), which bind to the extracellular domain of EGFR, and tyrosine kinase inhibitors (TKIs), which mostly target the intracellular part of EGFR and inhibit its activity in molecular signaling. While EGFR-specific mAbs and three generations of TKIs have demonstrated clinical efficacy in various settings, molecular evolution of tumors leads to apparent and sometimes inevitable resistance to current therapeutics, which highlights the need for deeper research in this field.

epidermal growth factor receptor (EGFR)

tyrosine kinase inhibitors (TKIs)

monoclonal antibodies (mAbs)

1. EGF Receptor Protein Family

In humans, the epidermal growth factor (EGF) receptor family (ERBB/HER) consists of four structurally related receptor tyrosine kinases (RTKs) that regulate proliferative cell signaling and play pivotal roles in both normal physiology and proliferative diseases like cancer [1]. The four family members are EGFR/ErbB1/HER1, ErbB2/Neu/HER2, ErbB3/HER3, and ErbB4/HER4 proteins [2], which are encoded, respectively, by genes EGFR, ERBB2, ERBB3, and ERBB4 [3]. These genes are located on four different chromosomes, but their products share common structural organization, including an extracellular domain, lipophilic transmembrane region, intracellular domain with tyrosine kinase activity, and a carboxy-terminal region [4].

The ERBB/HER family members are expressed in epithelial, mesenchymal, and neuronal cells and in their cellular progenitors [5]. The family members play central roles in cell proliferation, survival, differentiation, adhesion, and migration. These molecules interconnect the inner and outer compartments of the cytoplasmic membrane and trigger the cellular responses to various external stimuli by transmitting the intracellular regulatory stimuli [6]. The activated ERBB/HER receptors form regulatory complexes in which components can enter the cytoplasm and promote downstream molecular pathways (Figure 1), including well-known oncogenic pathways of RAS-RAF-MEK-ERK and AKT-PI3K-mTOR signaling axes [7]. Furthermore, apart from dimerization, EGFR molecules can also form oligomers on the cell surface, both under the action of natural ligands or in their absence [8][9]. The phenomenon of EGFR oligomerization is thought to be important for intracellular signaling because it results in a

tight organization of kinase-active molecules in a manner that is optimal for autophosphorylation in trans between adjacent dimers [10].

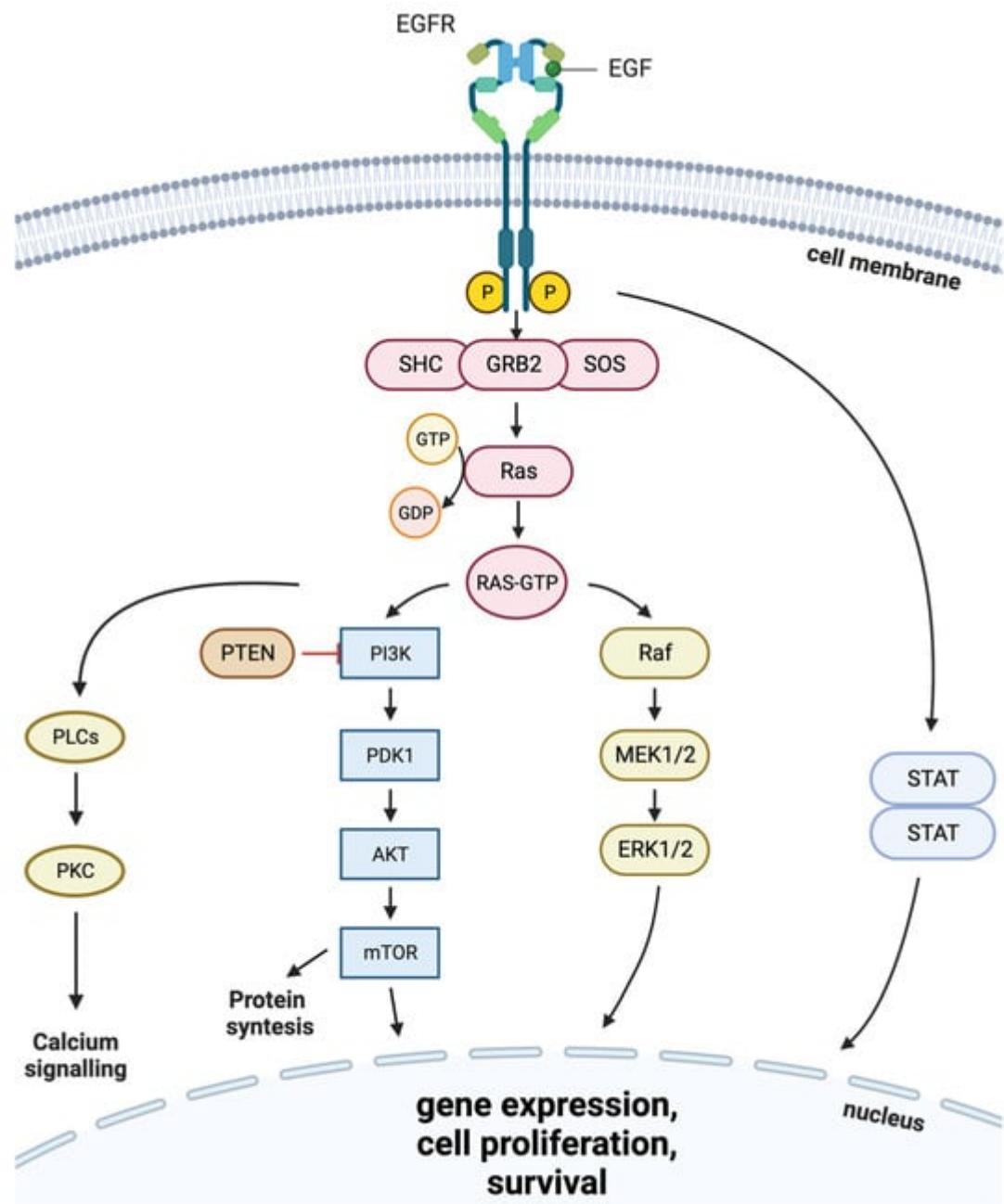


Figure 1. Intracellular signaling involving EGFR. The major regulatory pathways downstream of EGFR and other HER receptors are shown. Binding of specific ligands (e.g., EGF) leads to homo- or heterodimerization of receptors, thus resulting in conformational changes in the intracellular kinase domain, which results in phosphorylation and activation of the receptor. The signaling axes RAS-RAF-MEK-ERK and PI3K-AKT-mTOR, in turn, activate various downstream signaling pathways, thus leading to enhanced cell proliferation and survival. Created with [BioRender.com](https://biorender.com) (accessed on 1 November 2023).

Several growth factors are known to be able to bind ERBB/HER receptors and activate them. These are the members of the epidermal growth factor (EGF) family, which are generally classified into three groups. Representatives of the first one bind only to EGFR, which includes EGF [11], transforming growth factor alpha (TGF- α) [12], epigen (EPG) [13], and amphiregulin (AR) [14]. The second group has dual specificity of receptor binding and includes betacellulin (BTC) [15], heparin-binding epidermal growth factor (HB-EGF) [16], and epiregulin (EPR) [17]. The third group consists of neuregulins (NRG) and forms two subgroups depending on their ability to bind both HER3 and HER4 (NRG1 and NRG2 [18]) or only HER4 (NRG3 and NRG4 [19][20]) (Figure 2a).

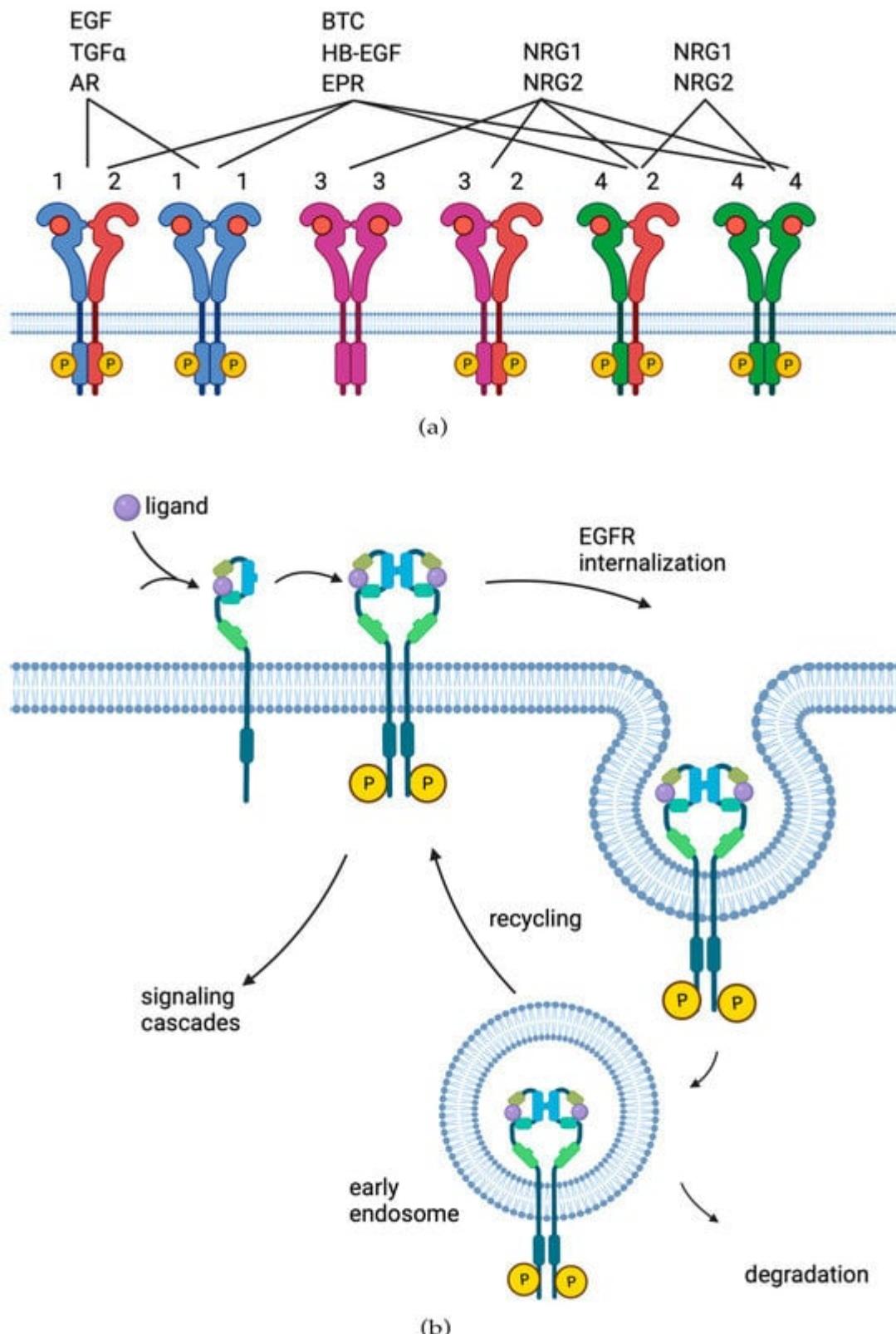


Figure 2. (a) Ligands that bind to common types of homo- and heterodimers formed by HER receptors. The following designations were used: 1—EGFR, 2—HER2, 3—HER3, and 4—HER4. (b) Dimerization, activation, and internalization of the EGFR receptor. Created with [BioRender.com](https://biorender.com) (accessed on 18 October 2023).

The inactivated forms of EGFR, HER3, and HER4 receptors exist in a pre-dimerized state. In turn, binding of the specific ligand causes rearrangement of the respective subunit of the receptor by turning the transmembrane domains. Activation leads to internalization of the receptor and trafficking to the early endosomal compartment of the cell. Next, endocytosis sorting occurs, whereby the receptor is either transported to the lysosome for further degradation or recycled to occupy a place in the cell membrane [21]. The family ligands affect receptor internalization in a different manner: upon EGF binding, the majority, but not all EGFRs, are continuously ubiquitinated and transported to lysosomes. HB-EGF and BTC also behave the same way. On the other hand, when subjected to the low pH of endosomes, TGF- α , EPR, and AR quickly separate from the receptor, which leads to de-ubiquitination of the receptor and its subsequent recycling to the plasma membrane (Figure 2b) [22].

In contrast with other HER family members, none of the ligands bind to HER2 [23]; it always exists in the dimerized state and acts as a preferred partner for heterodimerization with the other three ERBB/HER family members [24]. Also, HER2-containing heterodimers are characterized by higher affinity and broader ligand specificity than other heterodimeric ERBB/HER receptor complexes due to the slower dissociation rates of growth factors [25]. There was a controversy regarding HER3 pertaining to its kinase activity, and initially, it was posited that HER3 lacked kinase activity due to the absence of requisite residues [26].

2. EGFR Role in Cancer

Mutations and cases of overexpression of EGFR are especially frequently found in carcinomas and glioblastomas, tumors of epithelial and glial origin, respectively [27][28]. Worldwide, carcinomas are the most common type of cancer [29]. Overexpression of EGFR has been reported and implicated in the pathogenesis of many human malignancies, including head and neck [30], lung [31], breast [32], pancreatic [33], and colon cancer [34]. The EGFR-positive status of the tumor often correlates with poor prognosis and outcome, as it is beneficial for cancer cell proliferation [7][35]. EGFR overexpression was also shown to be associated with melanoma progression and promoted invasiveness and metastasis in this tumor type [36].

Mutations in the tyrosine kinase domain of EGFR were found in the majority of tumors that exhibited a positive response to treatment with EGFR-specific TKIs (Figure 3a in green) [37]. In some reports, the frequency of EGFR-activating mutations has strong ethnical specificity and varies by region, being as high as 46% in Asia versus only 8% in the Americas [38]. The two most common mutations of EGFR in NSCLC represent about 85–90% of all EGFR mutations [39]. The first one is a deletion of EGFR exon 19 (*del747–750*), which eliminates the leucine-arginine-glutamate-alanine motif in the tyrosine kinase domain of EGFR (*LREA deletion*), and the second one (*L858R*) is a thymine-to-guanine transversion, which results in the replacement of leucine with arginine in exon 21 codon 858 [40][41]. The third most frequent type of EGFR mutations in NSCLC is exon 20 insertions (ex20ins), which constitute 9% [42]–12% [43] of all EGFR mutations. In contrast to the other above mentioned mutations, Ex20ins is associated with poor response to treatment with TKIs [44]. It results in in-frame insertions, usually concentrated within or following the C-helix that dictates the activation status of EGFR [45]. In glioblastoma, the most frequently (~30%) occurring EGFR mutation is *EGFRΔIII* (EGFR variant III), which results from the in-frame deletion of 801 base pairs

spanning exons 2–7 of the coding sequence, resulting in ligand-independent activation of EGFR tyrosine kinase activity [46][47][48].

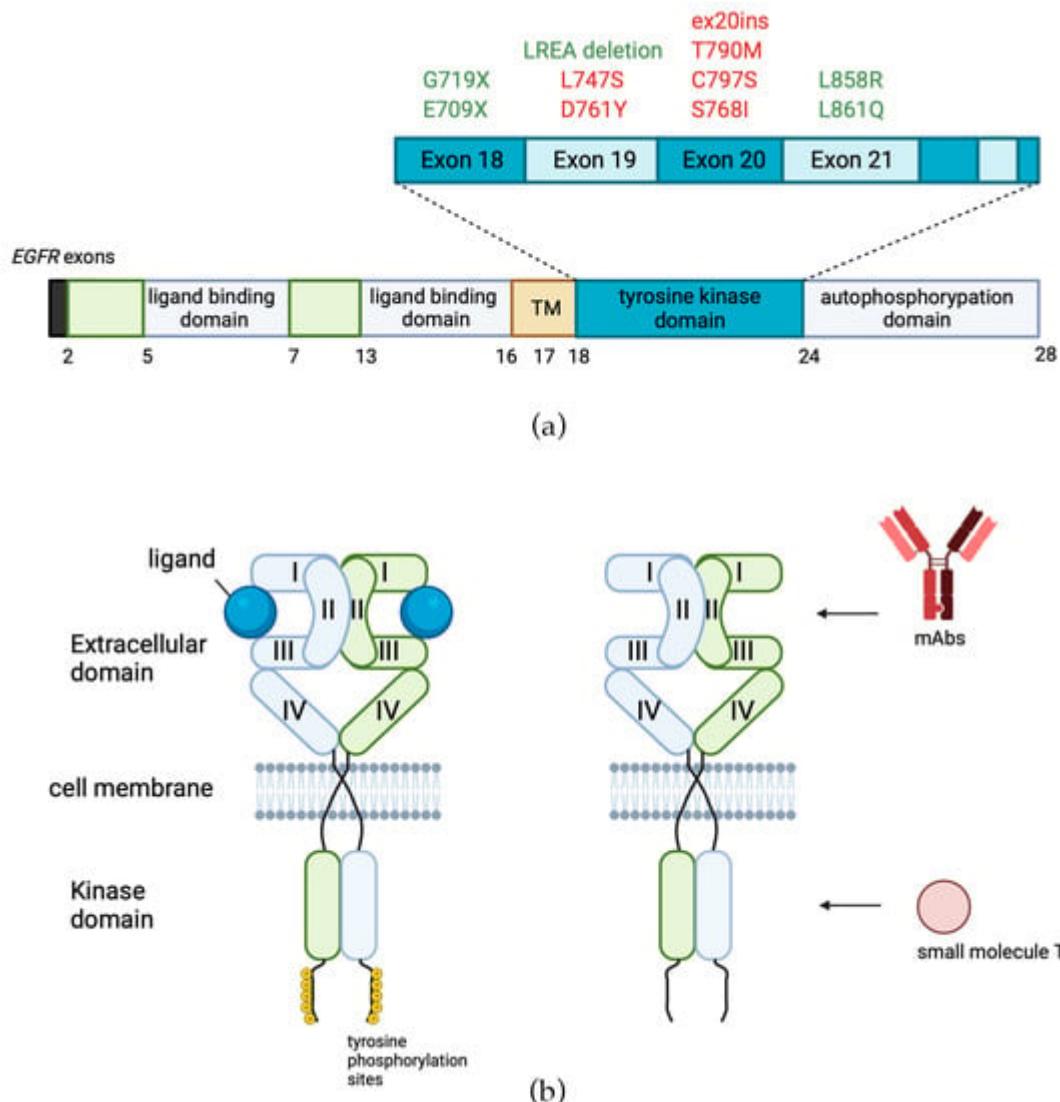


Figure 3. (a) Structure of EGFR gene. EGFR exons 18–21 encode the tyrosine kinase domain and may contain mutations, playing a crucial role in the development and progression of different cancers with a strong proven relationship to resistance (red) and sensitivity (green) to specific TKIs. **(b)** Domain view of EGFR protein. Left, a schematic diagram of ligand-bound dimerized EGFR. Right, sites of inhibition of EGFR activity by different targeted drugs (mAb: monoclonal antibodies; TKIs: tyrosine kinase inhibitors). Created with [BioRender.com](https://biorender.com) (accessed on 18 October 2023).

The *LREA* deletion of exon 19 of EGFR is shown to increase EGFR autophosphorylation and to activate downstream pathways AKT and STAT, thus promoting survival and cell growth [49]. With this mutation, the EGFR dimer exhibits increased stability as it tightens the molecular contacts of arginine ARG744 and asparagines ASP974 and ASP976 from the reciprocal monomers [50].

The 21st exon point mutation *L858R* is also a common activation mutation of *EGFR*, accounting for nearly 40% of all *EGFR* mutations. The *L858R* mutation locks the kinase in a constitutively active state by preventing the activation loop segment (residue 858 and flanking residues) from adopting the inactive, helical conformation, which leads to about 50-fold greater activity of the mutant *EGFR* [51].

EGFRvIII (EGFR with 2–7 exon deletion) lacks a ligand-binding domain and is constitutively active. It is the most common *EGFR* mutation occurring in glioblastoma [46]. *EGFRvIII* pathologic isoform does not contain amino acids 6-273 of wild-type *EGFR*, and it results in the formation of a new glycine residue at the junction site. This alteration imitates the effects of ligand binding and triggers changes in the receptor conformation by ultimately activating downstream signaling pathways [52].

In addition, the patients may have uncommon *EGFR* mutations. The use of next-generation sequencing (NGS) has become a novel diagnostic method for detection, which led to the identification of increasingly rare or atypical *EGFR* mutations. For example, *EGFR* fusion mutation *EGFR–SEPT14* was found in a patient with colorectal adenocarcinoma. The exon 24 of *EGFR* was fused to the exon 10 on *SEPT14* while retaining the *EGFR* tyrosine kinase domain. This tumor appeared to be sensitive to erlotinib treatment, and the patient developed a partial response following therapy [53].

In addition, activating mutations of downstream genes of regulatory kinases involved in the Ras/MAPK signaling pathway, such as KRAS, NRAS, and BRAF, are exceptionally frequent and appear in more than 90% of pancreatic, ~32% of lung, and ~52% of colon cancers [54].

3. EGFR-Targeted Therapies

Since *EGFR* is frequently overexpressed and/or mutated in multiple cancer types, it has prompted the development of a number of specific targeted therapeutics. Currently, there are two classes of *EGFR*-specific cancer drugs: monoclonal antibodies (mAbs), which bind to the extracellular domain of the transmembrane receptor and block its dimerization, and small-molecule tyrosine kinase inhibitors (TKIs), which bind to the adenosine triphosphate (ATP) binding site [55] (Figure 3b, Table 1). In turn, TKIs can be classified according to the mechanism of binding with the receptor tyrosine kinase domain: type I (binding with ATP site in mainly active conformation), type II (binding with ATP site plus back pocket, DFG(Asp855-Gly857)-out, in inactive conformation), type I½ (binding to a DFG-in, in inactive conformation), type III inhibitors binding to allosteric sites, and type IV inhibitors which generally form covalent adducts with their target protein [56][57]. *EGFR*-targeted drugs are currently widespread, globally approved, and are used worldwide for hundreds of thousands of patients per year.

Table 1. Characterization of *EGFR*-targeting inhibitors.

Tyrosine Kinase Inhibitors					
Drug	Tumor Type	Therapeutic Indication	Molecular Target	Inhibitor Type	Molecular Markers of Efficiency
<i>First Generation</i>					
<i>Gefitinib</i>	Advanced or metastatic NSCLC	First-line therapy for NSCLC carrying EGFR-activating mutations	EGFR: ATP-binding site	I	Activating mutations of EGFR: Exon 19 deletions; L858R
<i>Erlotinib</i>	Advanced or metastatic NSCLC, pancreatic cancer	First-line therapy for NSCLC carrying EGFR-activating mutations With gemcitabine: first-line treatment option for patients with locally advanced and metastatic pancreatic carcinoma	EGFR: ATP-binding site	I	Activating mutations of EGFR: Exon 19 deletions; L858R
<i>Lapatinib</i>	Metastatic breast cancer	With capecitabine: the treatment of HER2-positive MBC in patients who have previously received therapy (anthracycline, a taxane, trastuzumab) With letrozole: the treatment of postmenopausal women with hormone receptor positive MBC that overexpresses the HER2 receptor for whom hormonal therapy is indicated	ATP-binding site of EGFR and HER2	1½	HER2-positive status of tumor
<i>Second Generation</i>					
<i>Afatinib</i>	Metastatic NSCLC	First-line therapy for metastatic NSCLC carrying EGFR-activating mutations	ATP-binding site of EGFR, HER2, and HER4	IV	Activating mutations of EGFR: Exon 19 deletions; L858R
<i>Neratinib</i>	Breast cancer	Extended adjuvant treatment of patients with early stage HER2-positive breast cancer, to follow adjuvant trastuzumab based therapy With capecitabine: the treatment of patients with	ATP-binding site of EGFR, HER2, and HER4	IV	HER2-positive status of tumor

Tyrosine Kinase Inhibitors					
Drug	Tumor Type	Therapeutic Indication	Molecular Target	Inhibitor Type	Molecular Markers of Efficiency
<i>Dacomitinib</i>	Metastatic NSCLC	advanced or metastatic HER2-positive BC who have received two or more prior anti-HER2 based regimens in the metastatic setting First-line therapy for metastatic NSCLC carrying EGFR-activating mutations	ATP-binding site of EGFR, HER2, and HER4	IV	Activating mutations of <i>EGFR</i> : Exon 19 deletions; <i>L858R</i>
• <i>Third Generation</i>					
<i>Osimertinib</i>	Advanced or metastatic NSCLC	Adjuvant and first-line therapy for metastatic NSCLC carrying EGFR-activating mutations The treatment of adult patients with metastatic EGFR <i>T790M</i> mutation-positive NSCLC, whose disease has progressed on or after EGFR TKI therapy	ATP-binding site of the EGFR	IV	Activating mutations of <i>EGFR</i> : Exon 19 deletions; <i>L858R</i> The secondary <i>T790M</i> resistance mutation
<i>Almonertinib</i>	Advanced NSCLC	Adjuvant therapy for advanced NSCLC patients with <i>T790M</i> mutant EGFR who had developed resistance to first- and second-generation EGFR TKIs like gefitinib and afatinib	ATP-binding site of the EGFR	IV	Activating mutations of <i>EGFR</i> : Exon 19 deletions; <i>L858R</i> The secondary <i>T790M</i> resistance mutation
<i>Lazertinib</i>	Advanced NSCLC	Treatment of locally advanced or metastatic NSCLC carrying EGFR <i>T790M</i> mutation	ATP-binding site of the EGFR	IV	Activating mutations of <i>EGFR</i> : Exon 19 deletions; <i>L858R</i> The secondary <i>T790M</i> resistance mutation
<i>Furmonertinib</i>	Locally advanced or metastatic NSCLC	Treatment of locally advanced or metastatic EGFR <i>T790M</i> + NSCLC that developed after	ATP-binding site of the EGFR		The secondary <i>T790M</i> resistance mutation

Tyrosine Kinase Inhibitors				
Drug	Tumor Type	Therapeutic Indication	Molecular Target	Molecular Markers of Efficiency
progression on treatment with first-generation EGFR TKIs				
Monoclonal Antibodies				
Drug	Tumor Type	Therapeutic Indication	Molecular Target	Molecular Markers of Efficiency
<i>Cetuximab</i>	Advanced or metastatic SCCHN, metastatic CRC	With radiation therapy: treatment of locally or regionally advanced SCCHN	The binding site in domain III of EGFR	KRAS wild-type status of EGFR-overexpressing tumor
		With platinum-based therapy with fluorouracil: metastatic SCCHN		
		Metastatic SCCHN progressing after platinum-based therapy		
		With FOLFIRI: first-line treatment of KRASwt EGFR-overexpressing mCRC		
		With irinotecan in patients who are refractory to irinotecan-based chemotherapy: treatment of KRASwt EGFR-overexpressing mCRC; as a single-agent in patients who have failed oxaliplatin-and irinotecan-based chemotherapy or who are intolerant to irinotecan		
		Single agent treatment of metastatic CRC with disease progression on or following fluoropyrimidine, oxaliplatin, and irinotecan chemotherapy regimens		
<i>Panitumumab</i>	Metastatic CRC	With gemcitabine and cisplatin: first-line treatment of patients with metastatic NSCLC	The binding site in domain III of EGFR	RAS wild-type status of EGFR-overexpressing tumor
		Distinct from the Active State. <i>J. Biol. Chem.</i> 2020, 295, 13353–13362.		
<i>Necitumumab</i>	Metastatic NSCLC	Distinct from the Active State. <i>J. Biol. Chem.</i> 2020, 295, 13353–13362.	The binding site in domain III of EGFR	EGFR-overexpressing status of tumor

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4. First Generation of EGFR-Targeted Drugs

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Gefitinib

(a)

Afatinib

(d)

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Erlotinib

(b)

Neratinib

(e)

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Lapatinib

(c)

Dacomitinib

(f)

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5 patients with HER2-positive metastatic breast cancer. **Figure 4b** shows the analysis of EGFR with a Meyer–von Staudinger hydroxyl structure superimposed with the EGFR structure. **Figure 4c** shows the EGFR mutants and their complexes: Mechanism of Activation and Inhibition.

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The anti-proliferative effects of neratinib were examined *in vitro* across a panel of 115 cancer cell lines by ATPlite 1-step luminescence assay system for analysis of cell viability. In this panel, there were 22 cell lines harboring point mutations or amplifications of the *HER2* ($n = 9$), *HER3* ($n = 10$), or *EGFR* ($n = 10$) genes, and neratinib was proven to be the effective drug with IC_{50} s comparable to other TKIs in this study [81].

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In xenograft models overexpressing *HER2* (BT474) and *EGFR* (SKOV-3 and A431), neratinib dose-dependently inhibited tumor growth almost by 70–90% in xenografts of BT474, 30–60% in xenografts of SK-OV-3, and ~32–44% in xenografts of A431 [79].

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Dacomitinib (Vizimpro, PF-00299804, **Figure 4f**) [82], another second-generation EGFR inhibitor, was approved by the FDA in 2018 as the first-line treatment of patients with metastatic NSCLC with EGFR-activating mutations (exon 19 deletion or exon 21 substitution L858R) (**Table 1**) [83]. This drug also has activity against *EGFR*, *HER2*, and *HER4* receptors, which are inhibited through irreversible covalent binding of the drug at the edge of the ATP-binding cleft of tyrosine kinase domain [84]. For *EGFR*, irreversible inhibition is achieved by interacting with *EGFR* C797, similar to afatinib [85].

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NSCLC cell lines harboring endogenous *EGFR* T790M mutation, dacomitinib proved itself as an effective agent in *vitro*. In cell lines with L858R mutation or wild-type *EGFR*, dacomitinib had 10 times lower IC_{50} (μ M) than reversible *EGFR* inhibitor gefitinib. For example, the IC_{50} of dacomitinib in the H3255 cell line carrying L858R mutation was 0.007 μ M versus 0.075 μ M for gefitinib. In wild-type *EGFR* cell lines H1819 and Calu-3, IC_{50} values of dacomitinib were 0.029 and 0.063 μ M versus 0.42 and 1.4 μ M for gefitinib, respectively [86].

Suppression of E2F-1 Expression and Telomerase Activity. *Anticancer Res.* 2006, 26, 3387–3391.

16. Third Generation of EGFR-Targeted Drugs

Gefitinib (ZD1839) (Iressa) Tablets. *Oncologist* 2003, 8, 303–306.

First-generation *EGFR*-targeted low molecular mass therapeutics erlotinib and gefitinib have the disadvantage of being reversible inhibitors, and they are proven to be ineffective against the secondary *EGFR* mutations, such as Erlotinib (Tarceva) Tablets. *Oncologist* 2005, 10, 461–466.

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the T790M substitution, which has been found in over 50% of *EGFR*-mutant NSCLC cases with acquired resistance to *EGFR* inhibitors [87].

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BLU-945 (Figure 6e) was obtained by Blueprint Medicines by optimizing the molecules from ~25,000 compound library of designed small molecule kinase inhibitors and showed *in vitro* sub-nanomolar activities against the EGFR T790M and EGFR-T790M/C797S mutants. In addition, it reduced EGFR phosphorylation in Ba/F3 cells in *L858R/T790M/C797S* mutants with $IC_{50} = 3.2$ nM and in *ex19del/T790M/C797S* mutants with $IC_{50} = 4.0$ nM. In

vivo Inhibition and Enhanced Detection of the T790M Mutation Using a 72S EGFR Nucleated Acid-Based Assay. *Cancer Res.* 2011, 17, 1169–1180.

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For two other fourth-generation EGFR inhibitors, JIN-A02 and BBT-176, the successful application in both in vitro 90. Yang, J.C.-H.; Camidge, D.R.; Yang, C.-T.; Zhou, J.; Guo, R.; Chiu, C.-H.; Chang, G.-C.; Shiah, and in vivo studies were published: JIN-A02 inhibited *ex19del/T790M/C797S* and *L858R/T790M/C797S* EGFR-H.-S.; Chen, Y.; Wang, C.-C.; et al. Safety, Efficacy, and Pharmacokinetics of Almonertinib (HS-mutant Ba/F3 cells (IC_{50} = 51.0 and 49.2 nM, respectively) and resulted in tumor regression in 10296) in Pretreated Patients with EGFR-Mutated Advanced NSCLC: A Multicenter, Open-Label, *ex19del/T790M/C797S* Ba/F3 xenograft mouse models [105]. The IC_{50} values of BBT-176 for Ba/F3 cells Phase 1 Trial. *J. Thorac. Oncol.* 2020, 15, 1907–1918, engineered to express EGFR *19Del/C797S*, EGFR *19Del/T790M/C797S*, and EGFR *L858R/C797S* and 91. 1858R/T790M/C797S were [106]. Park, C.-H.; Kim, S.; Yun, M.R.; Kang, H.N.; Pyo, K.-H.; Lee, S.S.; Koh, J.S.; et al. YH25448, an Irreversible EGFR-TKI with Potent Intracranial Activity in EGFR-Mutant Non-Small Cell Lung Cancer. *Clin. Cancer Res.* 2019, 25, 3575–3587.

8. EGFR-Specific Therapeutic Monoclonal Antibodies

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large amounts of EGFR. The idea was that it was possible to stop EGFR-overexpressing tumor proliferation 93. Vasconcelos, P.E.N.S.; Kobayashi, I.S.; Kobayashi, S.S.; Costa, D.B. Preclinical Characterization through Interference with the EGFR Signaling [107]. Because EGFR permeates the cell membrane, the idea arose that monoclonal antibodies could be an effective therapeutic against tumors with increased expression of this receptor. EGFR-specific mAbs function similarly by disrupting pro-tumor growth and survival signaling through 94. Arnold, A.; Ganti, A.K. Clinical Utility of Mohocertinib in the Treatment of NSCLC—Patient binding to growth factor receptors, thus altering their activation state or preventing ligand binding [108].

Selection and Reported Outcomes. *OncoTargets Ther.* 2023, 16, 559–569. -Cetuximab (Erbitux, Merck Serono) was the first monoclonal antibody targeting the EGFR receptor, a human-mouse chimeric anti-EGFR mAb with the human IgG1 constant region [109]. It exhibits a strong affinity for Human 95. Wu, L.; Ke, L.; Zhang, Z.; Yu, J.; Meng, X. Development of EGFR TKIs and Options to Manage EGFR and effectively hinders ligand binding, ultimately resulting in the suppression of receptor phosphorylation and downstream signaling pathways [110]. The primary effect of cetuximab binding to EGFR is steric blockage of 96. Papadimitrakopoulou, V.A.; Wu, Y.L.; Han, J.P.; Alim, M.; Rajanayagam, S.S.; John, T.; Okamoto, I.; Yang, J.C.-H.; Bulusu, K.C.; Laus, G.; et al. Analysis of Resistance Mechanisms to It was found that the inhibition of cell growth induced by blocking EGFR activation of cetuximab deals with the Osimertinib in Patients with EGFR T790M Advanced NSCLC from the AURAS Study. *Ann. Oncol.* induction of cell cycle arrest and apoptosis. Cetuximab induced cell accumulation in the G1 phase and increased 2018, 29, viii741.

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Chicklas, S.; Hsieh, J.; et al. **Abstract 1262: BLU-701 Is a Highly Potent, Brain-Penetrant and WT-Sparing next-Generation EGFR TKI for the Treatment of Sensitizing (Ex19del, L858R) and C797S Resistance Mutations in Metastatic NSCLC**. *Cancer Res.* **2021**, *81*, 1262.

The drug effectively inhibited the growth of tumor cell lines of epidermal, pancreatic, and colorectal origins with EGFR overexpression *in vitro* [\[19\]](#). Additionally, necitumumab has significant antitumor activity in various human 104. Paveri, L.; Schram, S.; Campbell, J.; Guo, J.; Medelsohn, J.; Chen, M.; Albayya, F.; Dineen, T.; Zhang, Z.; Iliou, M.; et al. **Abstract 3328: Antitumor Activity of BLU-945 and BLU-701 as Single Agents and in Combination in EGFR L858R-Driven Models of NSCLC**. *Cancer Res.* **2022**, *82*, 3328.

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