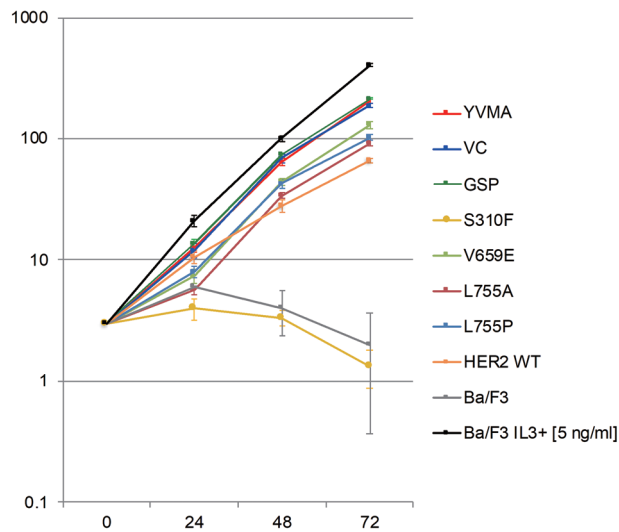
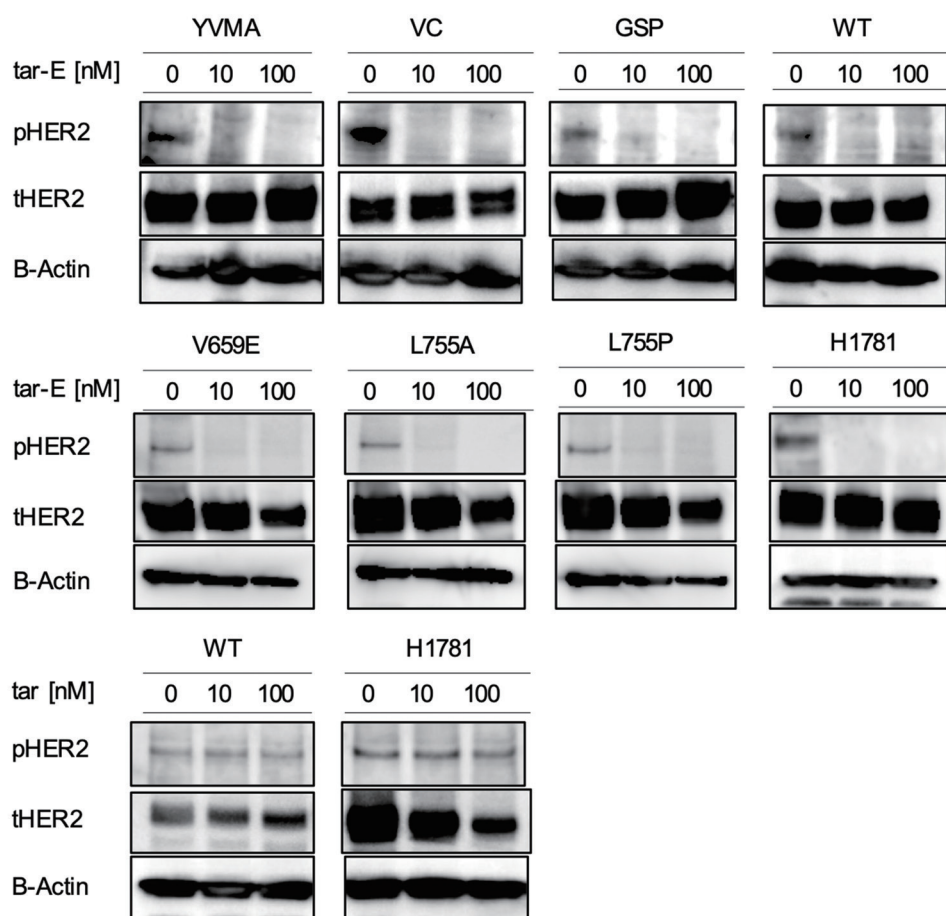


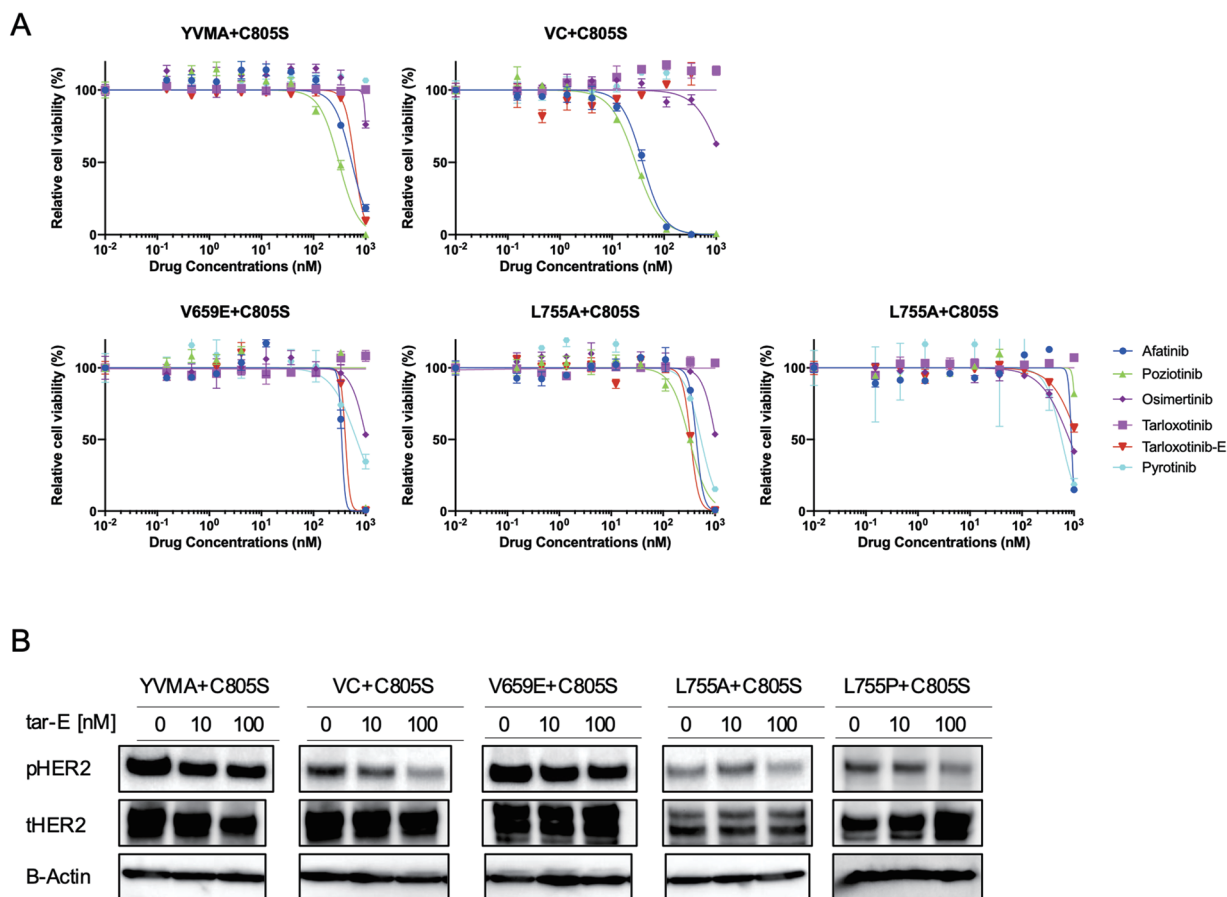
**Figure S1** The structures of tarloxotinib (prodrug) and tarloxotinib-E (active drug). Tarloxotinib conversions to active form tarloxotinib-E, an irreversible EGFR/HER2 inhibitor, under hypoxic condition in tumor microenvironment.



**Figure S2** Growth of transfected Ba/F3 cells with *HER2* point mutations. Ba/F3 cells transfected with *HER2* V659E, L755A or L755P proliferated in the absence of IL-3. On the other hand, Ba/F3 cells with the S310F mutation were not viable under IL-3-independent growth conditions. YVMA, A775\_G776insYVMA; VC, G776\_delinsVC; GSP, P780\_Y781insGSP.



**Figure S3** Western blotting of pHER2, tHER2 and  $\beta$ -actin extracted from Ba/F3 cells transfected with WT *HER2*, *HER2* with exon 20 insertions, and *HER2* with point mutations that were treated with the indicated concentrations of tarloxotinib-E or tarloxotinib. YVMA, A775\_G776insYVMA; VC, G776\_delinsVC; GSP, P780\_Y781insGSP; tar-E, tarloxotinib-E; pHER2, phospho-HER2; tHER2, total-HER2.



**Figure S4** Growth inhibition curves and western blotting of tarloxotinib-E resistant clones. (A) Growth inhibition curves of Ba/F3 cells with a secondary C805S mutation generated through ENU mutagenesis show the resistance of these cells to all *HER2* TKIs. (B) Western blotting of p-*HER2*, total *HER2* and  $\beta$ -actin extracted from the TR clones treated with each *HER2* TKI. Phosphorylation of *HER2* in C805S clones was not decreased by tarloxotinib-E. YVMA, A775\_G776insYVMA; VC, G776\_delinsVC; GSP, P780\_Y781insGSP; tar-E, tarloxotinib-E; pHER2, phospho-*HER2*; tHER2, total-*HER2*.

**Table S1** Designed primers for HER2 mutation mutagenesis

HER2 mutation		Sequence
S310F	Forward	5'- TGGGATTCTGCACCCTCGTCTGCCCC -3'
	Reverse	5'- GGGTGCAGAATCCCACGTCCGTAGAAAGGTAG -3'
V659E	Forward	5'- CGGTGGAAGGCATTCTGCTGGTCGTGGTCTTG -3'
	Reverse	5'- GAATGCCTTCCACCGCAGAGATGATGGACGTC -3'
L755A	Forward	5'- TCAAAGTGGCGAGGGAAAACACATCCCCC -3'
	Reverse	5'- CCCTCGCCACTTTGATGGCCACTGGAA -3'
L755P	Forward	5'- AGTGCCGAGGGAAAACACATCCCCCAAAG -3'
	Reverse	5'- TTTTCCCTCGGCACTTTGATGGCCACTG -3'

**Table S2** Designed primers for HER3 copy number analysis

	Sequence
Forward	5'- TGGGTTATGAGTACATGGATGTG -3'
Reverse	5'- CTTCATCTGGAGTTGTGCCTG -3'