Supplementary Materials for

EGFR Signals to mTOR Through PKC and Independently of Akt in Glioma

Qi-Wen Fan, Christine Cheng, Zachary A. Knight, Daphne Haas-Kogan, David Stokoe, C. David James, Frank McCormick, Kevan M. Shokat, William A. Weiss*

*To whom correspondence should be addressed. E-mail:weiss@cgl.ucsf.edu

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Fig. S1. Inhibition or knockdown of EGFR, Akt, PI3K, and mTOR in glioma; effects of serum.

Fig. S2. PMA interferes with the ability of erlotinib to decrease phosphorylation of rpS6 in $PTEN^{wt}$ glioma cells regardless of EGFR status.

Fig. S3. Inhibition of PTEN leads to increased abundance of p-Akt in *PTEN*^{wt} LN229: *EGFR* cells.

Fig. S4. Knockdown of PKCα and PKCδ in LN229:EGFR glioma cells.

Fig. S5. Abundance of EGFR, p-PKCα, p-PKC (pan), p-rpS6, and p-Akt in normal brain and primary human glioblastoma tumors.

Fig. S6. A PKC inhibitor blocks proliferation in both *PTEN*^{wt} and *PTEN*^{mt} glioma cell lines.

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Fig S1. Inhibition or knockdown of EGFR, Akt, PI3K and mTOR in glioma; effects of serum. (A) Biochemical analysis of EGFR inhibition as a function of time and of serum concentration. Glioma cells in 1% or 10% FBS were treated with erlotinib (5 μ M) for 1 or 24h as shown. EGF (50 ng/ml) was added 15 min prior to harvest for immunoblot. Although erlotinib blocked Akt in both PTEN^{wt} LN229:EGFR and in PTEN^{mt} U373:EGFR cells irrespective of serum concentration, the ability of erlotinib to affect the abundance of p-rpS6 was limited to PTEN^{wt} cells. (B, C). Small molecule inhibition (B) or siRNA knockdown of Akt isozymes (C) failed to affect proliferation or the mTOR target p-rpS6 in PTEN^{wt} LN229:EGFR cells. (B) LN229:EGFR cells (10% FBS) treated with AktI-1/2 at doses ranging from 0.1-5 µM showed no changes in cell cycle distribution. Percentage of cells in G_0G_1 , S, and G_2M phase of the cell cycle is indicated. Data shown are means ± standard deviations for triplicate measurements. (C) Combined siRNA and small molecule inhibitor treatment in LN229: EGFR cells grown in 1% FBS for 48h led to undetectable p-Identical results were achieved using Akt3 siRNA-1 (Santa Cruz) and Akt3 siRNA-2 Akt. (Dharmacon). The abundance of p-rpS6 was either minimally affected or unaffected (note decreased levels of β -tubulin loading control in cells treated with both Akt3 siRNA-1 and AktI-1/2). whereas phosphorylation of the Akt target p-Gsk3 β was decreased. Dosages were 2 μ M (Aktl-1/2) and 5 μ M (erlotinib). (D) Small molecule inhibition of PI3K or Akt isozymes 1 and 2 failed to affect phosphorylation of the mTOR target rpS6 in PTEN^{mt} U373 MG cells. To extend results to PTEN^{mt} glioma, we analyzed both Aktl-1/2 and the PI3K inhibitor PIK-90. Inhibition of PI3K, or Akt1 and Akt2 at doses effective against p-Akt again had no affect on p-rpS6. Rapamycin inhibited rpS6 phosphorylation, a result also observed in *PTEN*^{mt} U87:MG (E).



Fig S2. PMA interferes with the ability of erlotinib to decrease phosphorylation of p-rpS6 in PTEN^{wt} glioma cells regardless of EGFR status. To assess signaling between EGFR, PKC α , and mTOR in the absence or presence of EGFR amplification, we analyzed (A) LN229 parental cells, (B) GBM43, a primary glioma xenograft wild-type for both *PTEN* and *EGFR*, and (C) GBM12, a primary glioma xenograft wild-type for PTEN and amplified for EGFR. PMA (100 nM) elicited appearance of a slowly migrating p-PKC α band in all cell lines and primary xenografts. In EGF-treated cells, erlotinib (5 μ M) transformed this apparent doublet to a single more rapidly migrating band. Erlotinib treatment also blocked increased p-rpS6 abundance in response to EGF. In PMA-treated cells, this doublet was still evident in the setting of erlotinib treatment and correlated with reduced activity of erlotinib against p-rpS6. (D) A dominant-active allele of PKC α [PKC α -Cat] blocks the biochemical activity of erlotinib (5 μM) against p-rpS6 in LN229:EGFR cells grown in 1% FBS. Immunoblot using antisera against C-terminus of PKCa shows levels of PKCa-Cat comparable to those observed for endogenous PKC α . Erlotinib (5 μ M) blocked induction of p-rpS6 in control cells, but was less effective in cells transduced with PKC α -Cat. The effect seen is likely an underestimate, as PKC α -Cat cells treated with both EGF and erlotinib were underloaded on blot shown (note level of β -tubulin control).



Fig S3. Inhibition of Pten leads to increased abundance of p-Akt in *PTEN^{wt}* LN229:*EGFR* cells. (**A**) *PTEN^{wt}* LN229:*EGFR* cells were treated with the Pten inhibitor bisperoxyvanadium for 24h at dosage indicated. Cells were lysed and immunoblotted as indicated. (**B**) *PTEN^{nt}* U373:*EGFR* cells were treated with erlotinib (5 μ M) in the presence of PMA (100 nM) for 24h. Immunoblot shows appearance of a slowly migrating p-PKC protein in response to PMA, which was not affected by erlotinib.



Fig S4. Knock-down of PKC α and PKC δ in LN229:*EGFR* glioma cells. (**A**) siRNAs directed against PKC α decreased the total abundance of PKC α without affecting the abundance of p-PKC α , suggesting that siRNA directed against PKC α could not be used to evaluate the role of p-PKC α . SiRNA against PKC δ blocked only the rapidly migrating phospho-PKC δ isoform, with no affect on the supershifted band, thereby excluding PKC δ as a critical intermediate between EGFR and mTOR. Erlotinib was added at 5 μ M. (**B**) Pulse chase analysis in the presence of cycloheximide (an inhibitor of protein synthesis) demonstrates that p-PKC α has $t_{1/2}$ ~48h, explaining the high levels of p-PKC α observed after efficient knock-down of total PKC. Cycloheximide (25 μ M) was added for times shown. Cultures were treated with EGF for 15 min prior to harvest. (**C**) LN229:*EGFR* cells transduced with PKC α showed increased abundance of PKC α and p-PKC α , validating the specificity of antibody reagents against these proteins. EGF dose was (50 ng/ml).



Fig S5. Abundance of EGFR, p-PKC α , p-PKC (pan), prpS6 and p-Akt in normal brain and primary human glioblastoma tumors (as compared to normal brain control and normalized to β -tubulin). Abundance of each protein in Fig 6 was quantitated using UMAX PowerLook Scanner and ImageJ software.



Fig S6. A PKC inhibitor blocks proliferation in both *PTEN*^{wt} and *PTEN*^{mt} glioma cell lines. *PTEN*^{wt} and *PTEN*^{mt} cell lines shown were treated with 5 μ M of the pan-PKC inhibitor bisindolyImaleimide I (BIM I) for 24h. Flow cytometric analysis indicates percentage of cells in G₀G₁, S, and G₂M phases of the cell cycle. That BIM treatment led to arrest at G₁ in *PTEN*^{wt} LN229:*EGFR* and at G₂ in *PTEN*^{mt} U373:*EGFR* cells argues that the proliferative arrest induced by this agent results from on-target effects, rather than from non-selective toxicity.