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Role of $\beta 4$ integrin phosphorylation in human invasive squamous cell carcinoma: regulation of hemidesmosome stability modulates cell migration

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Abstract

Hemidesmosomes (HDs) are multiprotein structures that anchor epithelia to the basement membrane. During squamous cell carcinoma (SCC) invasion there is a reduction in the number of HDs, which may facilitate dissemination. Mechanisms of HD disassembly are incompletely understood. Previous work has shown that EGF-induced phosphorylation of the $\beta 4$ integrin on three of its serines, S₁₃₅₆S₁₃₆₀S₁₃₆₄, can induce HD disassembly in normal cells. Here we examine the role of $\beta 4$ integrin serine phosphorylation in SCC. We have found that around 60% of invasive cutaneous SCC show increased $\beta 4$ phosphorylation on S₁₃₅₆ when compared to carcinoma *in situ* or normal tissue. To assess mechanisms by which SCC increases $\beta 4$ phosphorylation we performed *in vitro* analyses. Compared to keratinocytes, SCC cells showed increased levels of S₁₃₅₆ phosphorylation in the absence of EGF, correlating with reduced HD-like structures. In addition, phospho-S₁₃₅₆ signal was largely segregated from other HD components. EGFR and PKC inhibitors inhibited basal levels of S₁₃₅₆ phosphorylation in SCC, suggesting that cells use intrinsic mechanisms to activate the EGF signaling pathway to induce $\beta 4$ phosphorylation. Moreover, these inhibitors stabilized HD-like structures in SCC cells and reduced their migratory ability. Mutation of S₁₃₅₆S₁₃₆₀S₁₃₆₄ in SCC cells to non-phosphorylatable alanines stabilized HD-like structures and substantially reduced migration, while mutation into phosphorylation mimicking aspartate reduced HD-like structures but had no effect on migration, suggesting that serine phosphorylation function is releasing anchorage rather than promoting migration. Altogether these results suggest that $\beta 4$ serine phosphorylation may have an important role during SCC invasion by destabilizing HDs and facilitating migration.

Keywords

Integrins; hemidesmosome; cell migration; invasion; phosphorylation

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INTRODUCTION

Squamous-cell carcinomas (SCC) are highly invasive tumors capable of metastasis (1–3). Stratified squamous epithelia, where most of the SCCs originate from, are strongly attached to the basal lamina through hemidesmosomes (HDs), multiprotein structures that provide stability (4, 5). During wound healing or SCC invasion there is an increase in HD disassembly (6, 7). In some SCC types, HD disassembly has been shown to correlate with metastatic potential (7). For this reason, there is a considerable interest to understand the mechanisms of HD disassembly.

The mechanisms of HD disassembly are not completely understood. Studies suggest that phosphorylation of the $\alpha6\beta4$ integrin, plays an important role in HD disassembly (8–12). The $\alpha6\beta4$ integrin is the main organizer of HDs (13). $\alpha6\beta4$ connects to laminin on the basal lamina and facilitates the assembly of other HD components, including plectin and BPAG1, which link $\alpha6\beta4$ to cytokeratins (4, 5, 13).

The chain of events that leads to HD disassembly in SCC is not well understood. Growth factors might trigger the initial events. Supporting this idea, EGF and MSP have been shown to induce HD disassembly *in vitro* (14–16). These factors activate signaling pathways that result in $\beta4$ phosphorylation. EGF induces phosphorylation of $\beta4$ on serine and tyrosine residues several of which have been identified and shown to be involved in HD disassembly (8, 9, 10, 11, 14). Around 95% of $\beta4$ phosphorylation induced by EGF occurs on serine (8, 15), mostly on four sites, S₁₃₅₆, S₁₃₆₀, S₁₃₆₄ and S₁₄₂₄ (8, 9, 11). In normal cells, substitution of these serines with alanine impedes phosphorylation and in a cooperative manner, can inhibit EGF-induced HD disassembly (8, 9, 11). The mechanism by which $\beta4$ serine phosphorylation induces disruption of HDs is unclear, although evidence suggests that S₁₃₅₆S₁₃₆₀S₁₃₆₄ phosphorylation controls $\alpha6\beta4$ /plectin interaction (11).

One possible scenario to explain reduced HDs in SCC, is that $\beta4$ phosphorylation may be altered, changing the balance towards disassembly. There is little information about $\beta4$ phosphorylation in SCC or its impact on HDs in cell migration. In this study, we analyzed $\beta4$ phosphorylation in primary SCC as well as in SCC cells *in vitro*. SCC frequently shows alterations in EGFR signaling (18, 19), so we analyzed phospho-S₁₃₅₆, a $\beta4$ residue whose phosphorylation is EGF-dependent. We found that S₁₃₅₆ phosphorylation in primary SCC correlates with invasiveness. *In vitro* analysis showed that SCC cells have intrinsic mechanisms to increase the basal level of $\beta4$ phosphorylation in the absence of EGF, reducing HD-like structures stability. Interestingly, SCC cells still use EGFR and PKC in the absence of exogenous EGF. Gefitinib, an EGFR kinase inhibitor, increased HD-like structures stability by reducing $\beta4$ phosphorylation, affecting cell migration as well. Mutation of $\beta4$ S₁₃₅₆S₁₃₆₀S₁₃₆₄ into alanines stabilized HD-like structures and hindered SCC migration. Our results suggest that $\beta4$ phosphorylation plays an important role in SCC progression by altering HD stability and the ability of cells to migrate. Targeting HD stability may be a method to reduce the ability of SCC to disseminate.

MATERIALS AND METHODS

Cells and reagents

Squamous cell carcinoma cell lines: A431 cells were obtained from ATCC; Colo-16 were obtained from Dr. N. Hail (University of Colorado, Denver, CO); SCC-25 were provided by Dr. A. M. Mercurio (UMass Med, Worcester, MA). HaCaT keratinocytes were obtained from Dr. S. La Flamme (Albany Medical College, Albany, NY). All cells were maintained in DMEM with 10% fetal calf serum, except SCC25 that was maintained in a 1:1 mixture of DMEM and Ham's F12 medium supplemented with 400 ng/ml hydrocortisone and 10% fetal bovine serum. Antibodies: 3E1 (β 4, Chemicon); GoH3 (α 6, Chemicon); rabbit anti- β 4 (15); anti-BPAG1 (20); anti-plectin (Santa Cruz Biotechnology); affinity-purified phospho-specific rabbit polyclonal Ab (anti-phospho S_{1356} Ab) raised against β 4 peptide DDVLR(pS)PSGSQ (custom-made, QBC, Hopkinton, MA).

Plasmids

β 4 shRNA-A431 cells: pLKO.1 β 4-shRNA TRCN0000057768 (Open Biosystems) against an untranslated region of β 4 was used to inhibit β 4 endogenous expression in A431. pLKO.1GFP-shRNA was used as control. Cells were puromycin selected. β 4-PCLXSN and triple mutants: β 4 integrin cDNA fused to a C-terminus myc tag was inserted in PCLXSN retroviral vector (Imgenex). A triple mutation ser \rightarrow ala or ser \rightarrow asp on $S_{1356}S_{1360}S_{1364}$ was introduced into β 4-myc using standard techniques (9). Retroviral particles were used to infect β 4 shRNA-A431 cells. As control, we used the empty vector. Cells were selected using G418 and β 4-negative cells were eliminated by FACS-sorting.

Indirect immunofluorescence

Cells were stained as described previously (15, 21). Briefly, cells grown on coverslips were extracted or not with detergent buffer containing 0.5% Triton-X-100, 100mM KCl, 200mM sucrose, 10mM EGTA, 2mM MgCl₂, and 10mM PIPES at pH6.8 for 1min., then fixed using paraformaldehyde or methanol. Cells were rinsed, blocked and stained with indicated Abs and Cy2/Cy3-conjugated secondaries. Slides were analyzed using fluorescence microscopy. Analysis of HD-like structures: collected images were background-subtracted, thresholded and fluorescence integrated density per cell was calculated using ImageJ software (NIH).

In vitro wound healing assay

Cells grown to confluency were scratched with a yellow tip and new medium containing inhibitors or not was added. Image records were collected at time 0. Wounded plates were incubated for 8–24h. Images were collected and percentage of wound closure was determined by digital analysis.

Human tissue samples

Tissue sections were obtained from the Cancer Center Tissue and Tumor Bank of UMASSMed with IRB approval. FFPE sections were stained with indicated antibodies using standard immunoperoxidase technique. Frozen sections were fixed in methanol, rinsed, blocked and stained with indicated antibodies followed by cy2/cy3-conjugated

secondaries. Slides were analyzed by fluorescence microscopy. Staining intensities were scored from 0 (absent) to 4 (very strong) by two observers (SL and IR). Colocalization analysis in frozen sections was performed using ImageJ (NIH) and Colocalization Threshold plugin (ref.22 and www.uhnresearch.ca/facilities/wcif/software/Plugins/colocalisation_threshold.html). Statistical analysis was performed using T-test.

RESULTS

Phosphorylation of the $\beta 4$ integrin in primary human SCC correlates with invasion

EGF signaling is frequently altered in SCC (18, 19). We have previously identified S₁₃₅₆ as one of the main phosphorylation sites on the $\beta 4$ integrin that is phosphorylated upon EGF stimulation in a PKC-dependent manner using peptide mapping analysis, results that have been confirmed by other groups (8, 11). We and others have also shown that S₁₃₅₆ phosphorylation contributes importantly in HD disassembly (8, 11). Therefore, we generated a phosphospecific Ab against phospho-S₁₃₅₆. To assess Ab specificity, we mutated $\beta 4$ S₁₃₅₆→A₁₃₅₆ to prevent Ab recognition of phospho-S₁₃₅₆. As shown in Fig 1A, mutation of the residue eliminated the phospho-S₁₃₅₆ Ab signal, confirming specificity. To confirm that the Ab acts in accordance to previous studies, we evaluated the kinetics of S₁₃₅₆ phosphorylation in HaCaT keratinocytes during EGF stimulation using phospho-S₁₃₅₆ Ab and Western analysis. As shown in Fig. 1B, there is little phosphorylation in non-stimulated cells, rapidly increasing to high levels after 5min and maintaining levels for 2h. Previous work suggests that phosphorylation of some of the serines in S₁₃₅₆S₁₃₆₀S₁₃₆₄ serine cluster are PKC-dependent (8, 11). Therefore we analyzed the effects of PKC stimulators and inhibitors on S₁₃₅₆ phosphorylation. As shown in Fig. 1C, EGF-dependent phosphorylation can be inhibited with conventional PKC inhibitor Go6976. Furthermore, PKC stimulator PMA induces S₁₃₅₆ phosphorylation, suggesting PKC-dependence. Altogether these results indicate that the Ab is specific and performs as expected in accordance to previous work.

To assess the prevalence of $\beta 4$ phosphorylation in human SCC, we analyzed S₁₃₅₆ phosphorylation in normal skin and primary SCC. Archival formalin fixed paraffin embedded (FFPE) or frozen sections were analyzed using immunohistochemistry (IHC) or immunofluorescence (IF) analysis, respectively.

FFPE sections from 20 cutaneous SCC were stained for phospho-S₁₃₅₆. Tissues were divided into 5 categories: normal skin (n=4); carcinoma *in situ* (n=7); well (n=4), moderately (n=5), or poorly differentiated (n=4) invasive carcinoma. Staining intensities were scored 0=absent to 4=very strong. In normal skin, $\beta 4$ was expressed in basal cells (Fig 2A) as previously described (28). Phospho-S₁₃₅₆ stain was mostly negative (Fig. 2B). In carcinoma *in situ*, $\beta 4$ was highly expressed (Fig 2C), sometimes observed in suprabasal levels, and there was little $\beta 4$ phosphorylation (Fig 2D). In invasive carcinomas, $\beta 4$ expression remained high, sometimes extending beyond the basal layer (Fig. 2E and G). In contrast to normal skin and carcinoma *in situ*, invasive SCC showed increased phospho-S₁₃₅₆ (Fig. 2F and H). The signal was specific because it was eliminated by incubating the primary antibody in the presence of competing phospho-S₁₃₅₆ peptide, or by treating the tissue with alkaline phosphatase before adding the primary antibody (results not shown). Around 60% of the invasive tumors showed high levels of phospho-S₁₃₅₆ with IHC scores

higher than normal skin (Fig. 2I). However, we found no difference between invasive carcinomas according to their degree of differentiation. $\beta 4$ phosphorylation could be found along the tumor/stroma interface (Fig. 2H), or extending deeply into the tumor (Fig. 2F) usually co-distributing with $\beta 4$. The increased phosphorylation of the $\beta 4$ integrin in invasive SCC suggests that phosphorylation may play a role during this tumor progression phase.

Distribution pattern of $\beta 4$ phosphorylation in relation to total $\beta 4$, basement membrane and other HD components in human SCC

To assess more closely the relationship between the $\beta 4$ phosphorylation signals and total $\beta 4$, other HD components or the basement membrane in human SCC, we analyzed 9 cutaneous SCC (all invasive, 3 well and 6 moderately differentiated) and 2 normal skin frozen sections using double IF. Consistent with IHC results, normal skin showed high levels of $\beta 4$ expression and little $\beta 4$ phosphorylation (Fig. 3A–C). About 60% of the invasive SCC sections showed moderate to high levels of $\beta 4$ phosphorylation (Figs 3E,H,K and Fig. 2I). $\beta 4$ phosphorylation varied within regions, frequently showing gaps in relation to the band-like signal of $\beta 4$ (arrows Fig. 3E,F,H and I) suggesting that phosphorylation is regionally modulated. Phosphorylation was stronger at the tumor/stroma interface sometimes extending more deeply (Fig. 3E and H), similar to FFPE sections.

To assess whether the discontinuous pattern of $\beta 4$ phosphorylation corresponds to basement membrane disruptions along the tumor/stroma interface, we analyzed colocalization of phospho-S₁₃₅₆ and Laminin-332. As shown in Fig. 3G–I, Laminin-332 was continuous and did not correlate with the patchy pattern of $\beta 4$ phosphorylation, suggesting that $\beta 4$ phosphorylation is unrelated to the basement membrane. To assess if $\beta 4$ phosphorylation affects the distribution of other HD components, we evaluated colocalization of BPAG1 and phospho-S₁₃₅₆. The patchy pattern of both BPAG1 and $\beta 4$ phosphorylation frequently excluded each other (Fig. 3J–L), suggesting that $\beta 4$ phosphorylation may affect the distribution of this HD component.

Increased basal level of S₁₃₅₆ phosphorylation in SCC cells is associated with reduced HD-like structure stability

The higher levels of phosphorylation observed in invasive primary SCC and previous observations that HDs are reduced in SCC (7), prompted us to assess *in vitro* for possible mechanisms explaining these observations. Reduced HDs in SCC may be related to alterations in $\beta 4$ phosphorylation. We therefore analyzed the levels and distribution of S₁₃₅₆ phosphorylation in three SCC cell lines: A431(23), SCC-25 (24) and Colo-16 (25) in relation to HaCaT keratinocytes, which are immortalized keratinocytes capable of differentiation in organotypic cultures and formation of HD-like structures *in vitro* (29). Western analysis showed that, in comparison to HaCat keratinocytes, all SCC cells had elevated basal levels of phospho-S₁₃₅₆ in the absence of EGF (Fig. 4A and B), suggesting intrinsic mechanisms to increase phosphorylation. We then compared the phospho-S₁₃₅₆ distribution between HaCaT keratinocytes and SCC cells using IF (Fig 4C and E). The cells were dual-immunostained with $\beta 4$ and phospho-S₁₃₅₆ Abs. HaCat keratinocytes showed characteristic HD-like structures stained with the $\beta 4$ Ab that largely survived extraction with detergent buffer before fixation (Fig. 4C; Non-Ex: nonextracted; Ex: extracted). The

resistance to detergent buffer is mostly conferred by the connection of the $\beta 4$ integrin with the cytokeratins through plectin and BPAG1 (30, 31). As expected, most of the $\beta 4$ that remains after detergent extraction in HD-like structures colocalizes with plectin (Fig. 4D). Using quantitative IF, we determined that about 60% of HaCaT $\beta 4$ is resistant to detergent (Fig. 4E). The phospho-S₁₃₅₆ signal was low in HaCaT keratinocytes (Fig. 4C). In contrast, SCC cells showed a higher phospho-S₁₃₅₆ signal in all three SCC cell lines (Fig 4C), which colocalized with $\beta 4$ in HD-like structures and retraction fibers (Fig. 4E and 5A–C). However, $\beta 4$ signal was less resistant than HaCaT keratinocytes to detergent extraction (~20%, Fig 4E). Most of the phosphorylation signal in the SCC cells was not resistant to detergent (Fig 4E). These findings are consistent with the notion that $\beta 4$ phosphorylation weakens HD stability.

Previous studies suggest that phosphorylated S₁₃₅₆S₁₃₆₀S₁₃₆₄ promotes disruption of $\beta 4$ -plectin interactions (11). We therefore assessed phospho-S₁₃₅₆/plectin and plectin/total $\beta 4$ colocalization in SCC cells. A similar pattern was found for all SCC types (exemplified by SCC25 in Fig. 5), showing that while plectin usually colocalized with total $\beta 4$ in stabilized HD-like structures as previously described (Fig 5D–F; (32)), phospho-S₁₃₅₆ only partially overlapped with plectin and interestingly, within colocalized areas, an inverse gradient between the 2 signals was frequently observed (Fig. 5G–I), suggesting that plectin may be gradually removed as more $\beta 4$ becomes phosphorylated. We also assessed colocalization of phospho-S₁₃₅₆ or total $\beta 4$ with BPAG1, another linker of $\beta 4$ with cytokeratins. While BPAG1 always colocalized with parts of the total $\beta 4$ Fig 5J–L), a sharper segregation was seen between phospho-S₁₃₅₆ and BPAG1 (Fig. 5M–O) suggesting that other HD components are affected by phospho-S₁₃₅₆ as well. Altogether these results suggest that SCC cells have intrinsic mechanisms to induce $\beta 4$ phosphorylation that might disrupt interactions among HD components, decreasing HD-like structure stability.

Intrinsic activation of EGFR signaling pathway in SCC cells raises phospho-S₁₃₅₆ basal levels affecting HD-like structure stability and cell migration

To address the mechanisms that could generate an increase in the basal levels of $\beta 4$ phosphorylation in SCC, we determined whether this increase still depends on the EGF/PKC pathway. We evaluated phospho-S₁₃₅₆ in the presence or absence of EGFR inhibitor Gefitinib or PKC inhibitor Go6976. As shown in Fig. 6A, both Gefitinib and Go6976 inhibit phospho-S₁₃₅₆ basal levels in SCC cells, suggesting that activation of EGFR and PKC is still necessary in the absence of an external source of EGF and that SCC cells have intrinsic mechanisms to activate these kinases.

Since Gefitinib is used in some types of SCC chemotherapy (27), we addressed whether this drug could exert some of its anti-tumor effects through HD stabilization and modifying cell migration. As shown in Fig. 6B $\beta 4$ in HD-like structures is substantially increased using Gefitinib or Go6976. We then assessed their effect on SCC migration using *in vitro* wound healing assays. Both inhibitors efficiently reduced migration in all cells (Fig. 6C). Go6976 effect was more pronounced, suggesting that alternative signaling pathways may converge with EGF-signaling to activate PKC. These results suggest that anti-tumor activity of

Gefitinib might include inhibition of SCC cell migration through HD stabilization and inhibition of $\beta 4$ phosphorylation.

Prevention of $\beta 4$ phosphorylation on $S_{1356}S_{1360}S_{1364}$ restores HD-like structures in SCC cells and slows migration

Cell migration is an important component of carcinoma invasion and metastasis (33). Considering that SCC invasion correlates with $\beta 4$ phosphorylation and reduction of HD-like structures, we hypothesized that by preventing $S_{1356}S_{1360}S_{1364}$ phosphorylation in SCC, we would replenish HD-like structures and hinder migration. Previous work in normal cells has shown that triple mutation is needed to increase HD stability (8, 11). We therefore expressed in A431 a non-phosphorylatable triple mutant $\beta 4 A_{1356}A_{1360}A_{1364}$ -myc or a phosphorylation-mimicking mutant $\beta 4 D_{1356}D_{1360}D_{1364}$ -myc, using wild type (wt) $\beta 4$ -myc as control. We first silenced $\beta 4$ endogenous expression in A431 using shRNA technology (targeting $\beta 4$ non-translated region). We then expressed wt and mutant $\beta 4$ at equivalent levels (Fig. 7A). Using IF analysis and detergent extraction to assess HD-associated $\beta 4$, we found that $\beta 4$ -myc and $\beta 4$ - $A_{1356}A_{1360}A_{1364}$ -myc incorporated well into HD-like structures (Fig 7B non-extracted vs. extracted). A quantitative analysis showed that $\beta 4$ - $A_{1356}A_{1360}A_{1364}$ -myc produced twice the amount of HD-like structures as $\beta 4$ -myc (Fig 7C). In contrast, incorporation of $\beta 4 D_{1356}D_{1360}D_{1364}$ -myc into HD-like structures was reduced (Fig 7B, C). We addressed whether HD-like structure stabilization induced by $\beta 4 A_{1356}A_{1360}A_{1364}$ -myc can counter EGF-induced HD disruption. As shown in Fig 7C, $\beta 4 A_{1356}A_{1360}A_{1364}$ -myc mutant efficiently resisted EGF-induced HD-like structure disruption while $\beta 4$ -myc was largely mobilized. These data suggest that low number of HD-like structures in SCC can be reverted by preventing $\beta 4$ phosphorylation.

We then assessed the ability of $\beta 4$ phosphorylation mutants to influence cell migration using wound healing assay. As shown in Fig. 7D, $\beta 4 A_{1356}A_{1360}A_{1364}$ -myc substantially reduced A431 migration. However, the phosphorylation mimicking mutant did not affect migration. These results suggest that regulation of $\beta 4$ phosphorylation can modulate cell movement through HD destabilization, although once $\beta 4$ leaves HDs, serine phosphorylation provides no further advantage.

DISCUSSION

HDs disassemble during wound healing and carcinoma invasion. In some types of SCC, the reduction of HDs is also associated with high metastatic potential (7). In this paper we have explored the possibility that reduction of HDs in SCC is due to $\beta 4$ integrin serine phosphorylation. Consistent with this idea we have found a correlation between an increase in $\beta 4$ integrin phosphorylation and carcinoma invasion in a group of 29 primary cutaneous SCC.

This is the first report showing that $\beta 4$ phosphorylation occurs in human tissue. The analysis of the $\beta 4$ phosphorylation in FFPE and frozen tissues showed a clear increase in phospho- S_{1356} signal in ~60% of the invasive SCC. However we did not detect significant differences between the various degrees of differentiation among the invasive tumors. The phosphorylation signal was distributed unevenly, showing frequent gaps in the interface

between the SCC and stroma, suggesting that the phosphorylation may be dependent on the context of the tumor location. Considering that EGFR activation is frequently seen in SCC and that $\beta 4$ can be phosphorylated through EGF signaling, one possible scenario to explain regional variations of $\beta 4$ phosphorylation would be a concomitant activation of EGFR within the same areas. We made efforts to detect phospho-EGFR and, although we found that SCC expressed EGFR, we were not able to detect phospho-EGFR (results not shown), a problem that has also been encountered by others studying cutaneous SCC (19). Interestingly, in other types of SCC, the existence of regional variations in phospho-EGFR has been reported (34), reminiscent of $\beta 4$ phosphorylation pattern. Another possibility is that regional phosphorylation of the $\beta 4$ integrin might reflect gaps in the basement membrane, a phenomenon that frequently occurs in invasive carcinoma. However, we found no relationship between the $\beta 4$ phosphorylation pattern and the basement membrane, which mostly showed a continuous distribution of Laminin-332, suggesting that $\beta 4$ phosphorylation is not determined by the basement membrane organization.

To gain more insight into what causes an increase in $\beta 4$ phosphorylation in SCC, we used an *in vitro* model of SCC. Our *in vitro* results suggest that SCC cells acquire intrinsic mechanisms to increase $\beta 4$ phosphorylation in the absence of any added growth factor. This increase of $\beta 4$ basal phosphorylation in SCC cells is accompanied by a reduction of HD-like structures and segregation of the phosphorylation signal in relation to the HD components plectin and BPAG1, results that are consistent with the notion that $\beta 4$ phosphorylation promotes disassembly of HDs through the disruption of interactions with other HD components (8, 11). Interestingly, we also observed segregation of BPAG1 and $\beta 4$ phosphorylation signal in SCC tissues, although the complete disappearance of BPAG1 within groups of cells in SCC suggests a different mechanism of segregation whereby continuous $\beta 4$ phosphorylation might eventually affect BPAG1 turnover or expression. Clearly more studies are necessary to understand long-term effects of $\beta 4$ phosphorylation on other HD components. One shared mechanism that the three SCC cell lines use to increase the basal phosphorylation of the $\beta 4$ integrin is the maintenance of an active EGFR in the absence of added growth factor, indicated by the inhibition of $\beta 4$ phosphorylation with Gefitinib, an EGFR kinase inhibitor. An active EGFR in such conditions would suggest the possibility of autocrine secretion of EGFR ligands or sensitizing mutation/amplification of EGFR. Constitutive activation of EGFR through expression of autocrine EGFR ligands, gain-of-function mutations, or protein overexpression is found in nearly 90% of all oral cavity and head and neck SCCs (2, 27, 35). In the specific case of A431, overexpression of EGFR and autocrine stimulation with TGF α is well documented and may explain the increase in $\beta 4$ phosphorylation (36, 37). Importantly, the inhibition of $\beta 4$ phosphorylation by Gefitinib correlated with stabilized HD-like structures and reduced cell migration in all SCC cells, suggesting that EGF effect on migration may be in part due to the regulation of $\beta 4$ phosphorylation promoting HD disassembly and allowing cells to move. Downstream EGFR, PKC still seems to mediate increased S₁₃₅₆ phosphorylation in SCC as shown by the inhibitory action of Go6976, a conventional PKC inhibitor (38). Therefore SCC cells may only differ from keratinocytes in some type of dysregulation of the EGF signaling pathway rather than the activation of an alternative route.

Our data suggest that $\beta 4$ phosphorylation might be involved in regulating the invasive ability of SCC through modulation of HD stability and cell migration. First, our results show that by inhibiting $\beta 4$ phosphorylation with Gefitinib in SCC cells, there was clear stabilization of HD-like structures, which correlated with cell migration inhibition. The effect on HD-like structure stability and cell migration was somewhat stronger using Go6976, suggesting that alternative signaling pathways may converge in the activation of PKC. Second, by preventing the phosphorylation of $\beta 4$ through mutation of the serine cluster S₁₃₅₆S₁₃₆₀S₁₃₆₄ to alanines, we were able to increase HD-like structure stability and inhibit migration even in the presence of EGF. However, a phosphorylation mimicking mutation of the serine cluster to aspartate failed to increase migration. This suggests that the role of $\beta 4$ integrin serine phosphorylation in SCC migration is to release the cell from migration-hindering attachment rather than actively promote migration. It is worth mentioning that $\beta 4$ per se was still necessary for cell migration, since silencing $\beta 4$ reduced migration, which is in agreement with previous studies suggesting that $\beta 4$ itself promotes migration and invasion through a variety of signaling pathways (39, 40). In this regard, a role of $\beta 4$ tyrosine phosphorylation in the migration-promoting function of $\beta 4$ has been previously shown (41–43, 44).

An interesting implication of our findings is that some of the antitumorigenic effects of EGFR inhibitors, such as Gefitinib, might be exerted through the stabilization of HDs by inhibiting $\beta 4$ serine phosphorylation and consequently affecting the migration capability of the tumor which ultimately impacts metastatic potential. Future studies that evaluate the restoration of HDs and reduction of $\beta 4$ phosphorylation during Gefitinib treatments could be potentially useful in assessing treatment efficiency and providing additional prognosis value.

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Abbreviations

BPAG	Bullous Pemphigoid Antigen
EGFR	epidermal growth factor receptor
FFPE	formalin fixed paraffin embedded
HD	hemidesmosome
IHC	immunohistochemistry
IF	immunofluorescence analysis
MSP	Macrophage-stimulating protein
SCC	squamous cell carcinoma
wt	wild type

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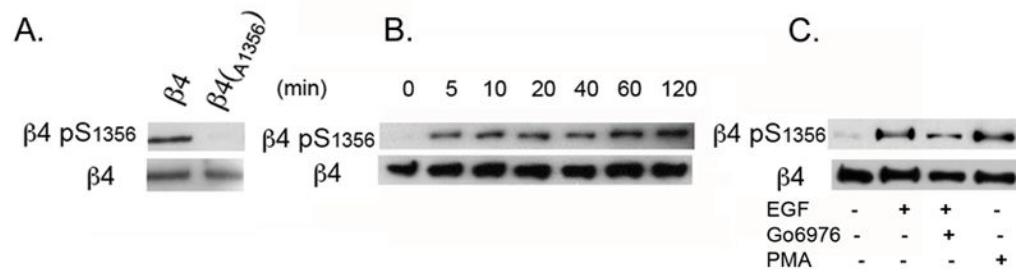


Figure 1. Regulation of $\beta 4$ phosphorylation (phospho-S₁₃₅₆) by EGF in HaCaT keratinocytes
A. Anti-phospho-S₁₃₅₆ Ab specificity. Lysates obtained from EGF-stimulated Cos 7 cells transfected with wt $\beta 4$ or mutant $\beta 4_{A1356}$ were probed with Abs against $\beta 4$ and phospho-S₁₃₅₆. **B.** Time course of S₁₃₅₆ phosphorylation in HaCaT keratinocytes. Cells were serum-starved and EGF-stimulated (50ng/ml) for different times, and analyzed by Western blotting using phospho-S₁₃₅₆ and $\beta 4$ Abs. **C.** S₁₃₅₆ phosphorylation is PKC-dependent. HaCaT keratinocytes were EGF-stimulated in the presence or absence of conventional PKC inhibitor Go6976, or with PMA, then analyzed by Western blotting using phospho-S₁₃₅₆ and $\beta 4$ Abs.

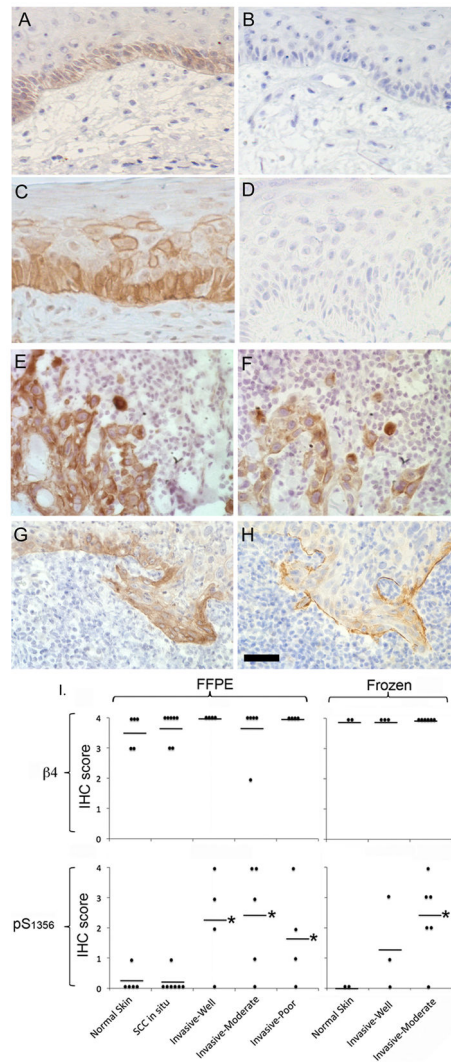


Figure 2. Phosphorylation of the $\beta 4$ integrin is increased in human SCC

A–H. FFPE sections stained with $\beta 4$ integrin (A,C,E,G) and phospho-S₁₃₅₆ (B,D,F,H). Abs were revealed by IHC in normal skin (A,B), carcinoma *in situ* (C,D), and well differentiated (E,F) or poorly differentiated (G,H) invasive cutaneous SCC. Bar=100 μ m. **I.** Frequency plots of predominant IHC score for $\beta 4$ and phospho-S₁₃₅₆ in tissue sections. FFPE (left panels) or frozen sections (right panels) were scored based on the observed staining intensity of indicated Abs, using a scale of 0 to 4. Each symbol represents a separate section and the horizontal line represents the mean score for each category. *p<0.05.

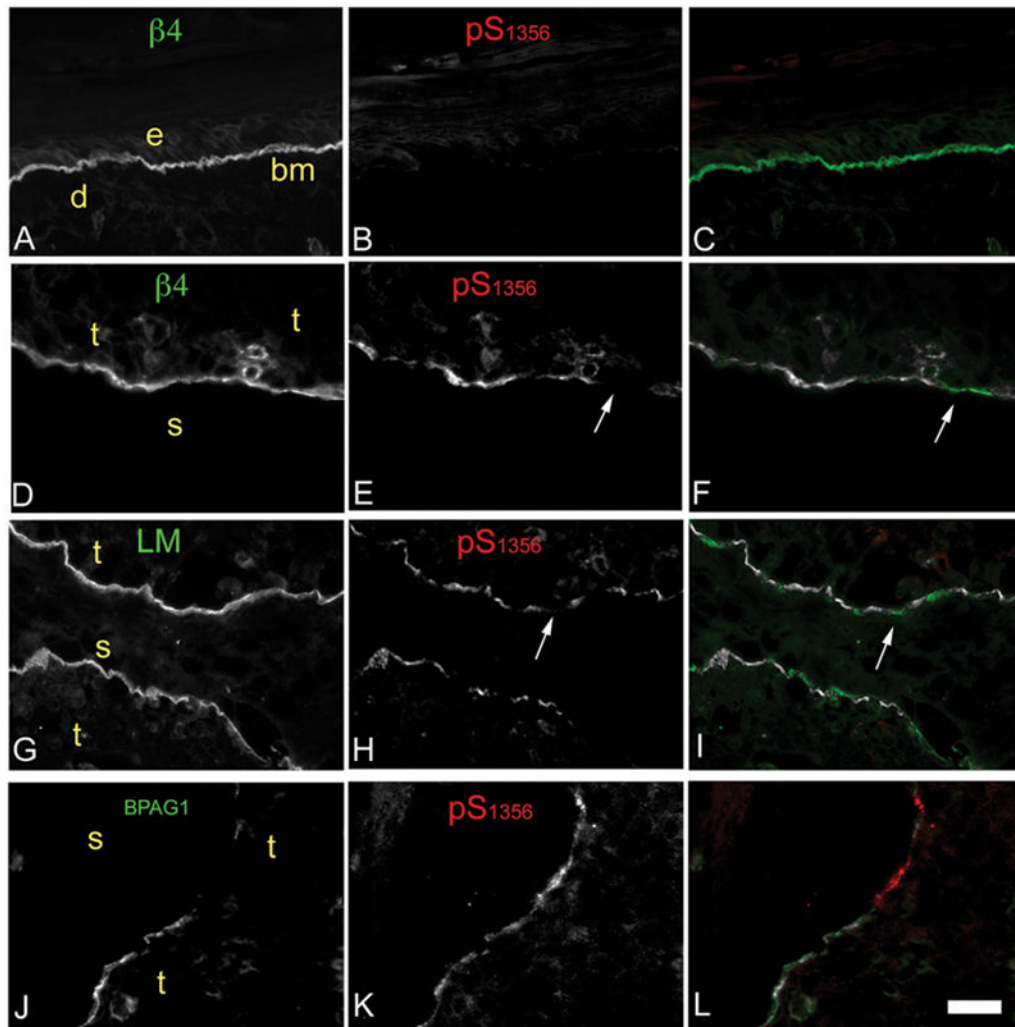


Figure 3. Distribution pattern of $\beta 4$ phosphorylation in human SCC frozen sections: discontinuous patches that can exclude other HD components

Frozen sections from skin (A–C) or invasive SCC (D–L) were dual-immunostained using Abs against (indicated within image): phospho- S_{1356} (red); and $\beta 4$, Laminin-332 or BPAG1 (green). Colocalization in the third column was determined by threshold correlation analysis (22), which shows correlating areas above threshold in white (colocalizing) or below threshold in red or green (non-colocalizing). e, epidermis; d, dermis; bm, basement membrane; t, tumor; s, stroma. Bar=100 μ m. Notice that $\beta 4$ phosphorylation may range from mildly discontinuous (E) to highly discontinuous (H). Laminin-332 is mostly continuous and does not follow phospho- S_{1356} gaps (G–I). While both BPAG1 and phospho- S_{1356} phosphorylation are discontinuous they show segregation more frequently (J–L).

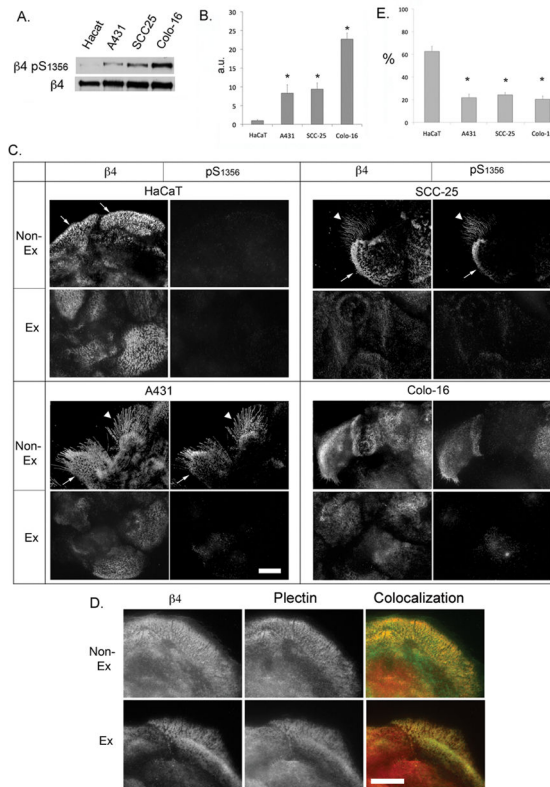


Figure 4. Increased basal levels of $\beta 4$ phosphorylation on S₁₃₅₆ in SCC cells correlates with a reduction of HD-like structures

A. Cell lysates from serum-starved HaCaT keratinocytes or SCC cells (A431, SCC-25, Colo-16) were analyzed by Western blotting using phospho-S₁₃₅₆ and $\beta 4$ Abs. **B.** Bands were quantified by densitometry. **C.** IF analysis of total $\beta 4$ and phospho-S₁₃₅₆ spatial distribution in HaCaT keratinocytes and SCC cells. Cells grown on coverslips were serum-starved and detergent-extracted to identify HD-associated $\beta 4$ (“extracted”), or not to assess total $\beta 4$ (“non-extracted”), then fixed and stained using phospho-S₁₃₅₆ and $\beta 4$ Abs. HaCaT keratinocytes show characteristic HD-like structures identified by $\beta 4$ staining on the basal aspect of the cell (thin arrows). Phospho-S₁₃₅₆ signal can be observed in SCC cells (A431, SCC-25 and Colo-16) in HD-like structures (thin arrows) and retraction fibers (arrowheads). Bar=10um. **D.** Dual immunostaining analysis of detergent-extracted or non-extracted HaCaT cells using anti- $\beta 4$ (green) and anti-plectin antibodies (red). After detergent extraction most of the detergent-resistant $\beta 4$ colocalizes with plectin in HD-like structures. Bar=10um. **E.** Percentage of detergent-resistant $\beta 4$ in HD-like structures: Using IF analysis, the integrated fluorescence density for $\beta 4$ was calculated for cells extracted or not with detergent buffer before fixation as described in Methods, and expressed as % of detergent-resistant $\beta 4$ / total $\beta 4$. Data shown are means \pm SE of >200 cells. *p<0.05.

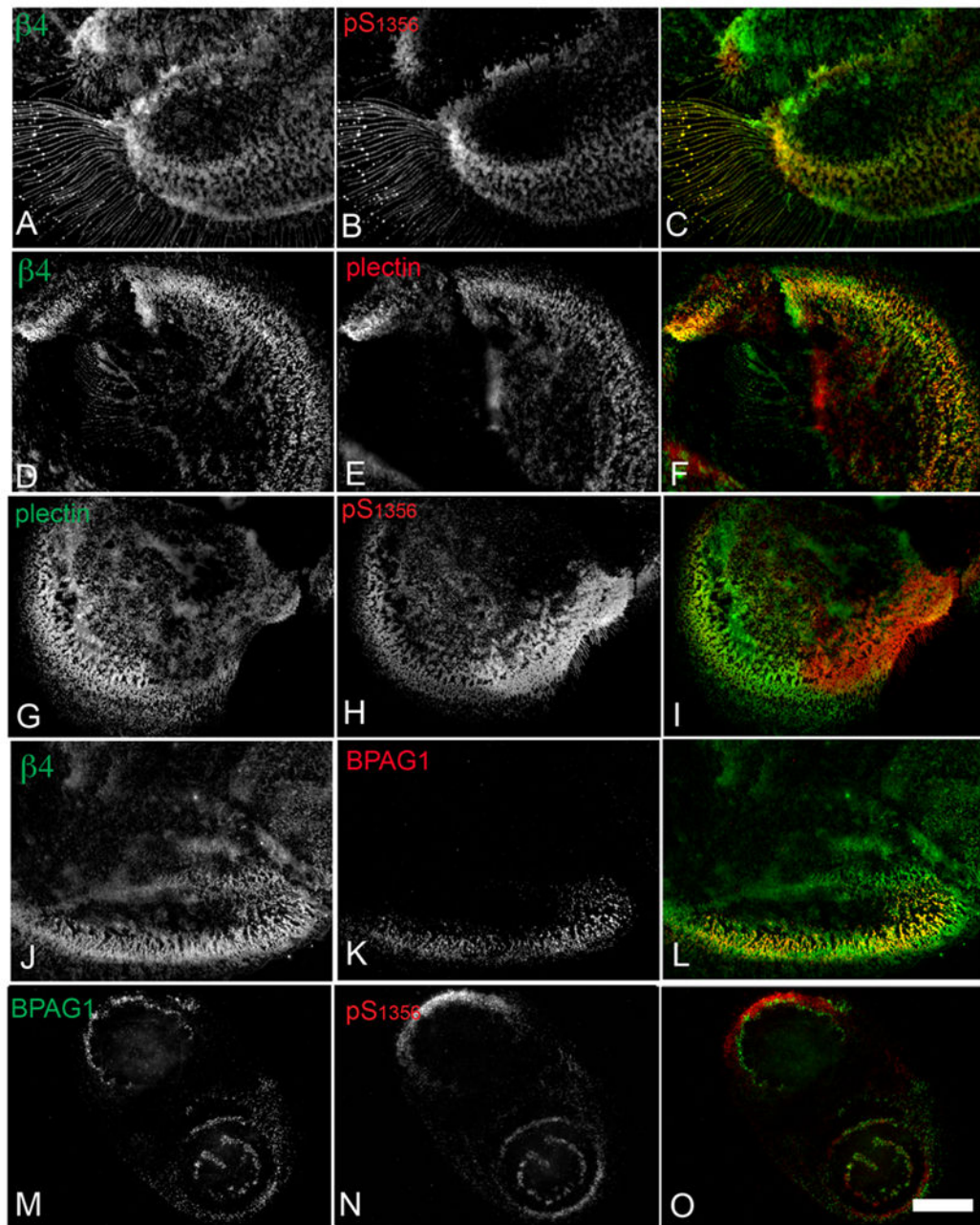


Figure 5. Phospho-S₁₃₅₆ is partly segregated from other HD components in SCC cells
 SCC-25 cells were grown in coverslips, fixed and stained for IF analysis using the indicated Ab. The third column shows colocalization in yellow. β 4 and phospho-S₁₃₅₆ colocalize in retraction fibers and a portion of the HD-like structures (A–C). While HD component plectin colocalizes well with total β 4 in HD-like structures (D–F), it is partially segregated from the phospho-S₁₃₅₆ signal (G–I). Notice inverse gradients between phospho-S₁₃₅₆ and plectin signals in colocalized areas. HD component BPAG1 colocalizes well with β 4 (J–L) whereas it is highly segregated from phospho-S₁₃₅₆ signal (M–O). Bar= 10 μ m.

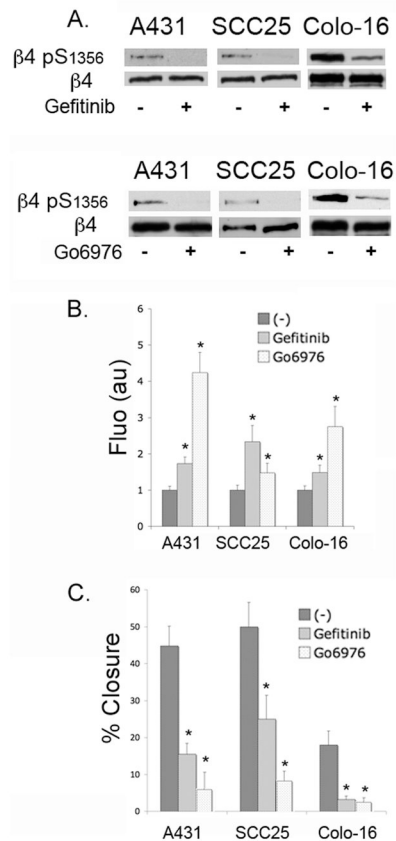


Figure 6. EGFR and PKC inhibitors reduce S₁₃₅₆ basal phosphorylation in SCC cells, stabilizing HD-like structures and inhibiting cell migration

A. Serum-starved SCC cells were treated or not with EGFR inhibitor Gefitinib (10uM, upper panel) or conventional PKC inhibitor Go6976 (2uM, lower panel) for one hour, then lysed and analyzed by Western blotting using phospho-S₁₃₅₆ and β4 Abs. **B.** Effects of Gefitinib and Go6976 on HD-like structures stability. SCC cells were grown on coverslips and treated or not with inhibitor for 1h, then detergent-extracted, fixed and processed for IF using anti-β4 Ab. The integrated fluorescence density for the HD-like structures was quantified as described in Methods, and expressed in arbitrary units per cell. Data shown are means±SE of >200 cells. *p<0.05. **C.** Effects of Gefitinib and Go6976 on SCC cell migration using *in vitro* wound healing assay. Confluent SCC cells were scratched with a yellow tip. Inhibitors were added or not and the wound was allowed to close. Images captured at the start and end of the experiment were used to quantify wound closure (%). Data shown are means±SE of three independent experiments *p<0.05.

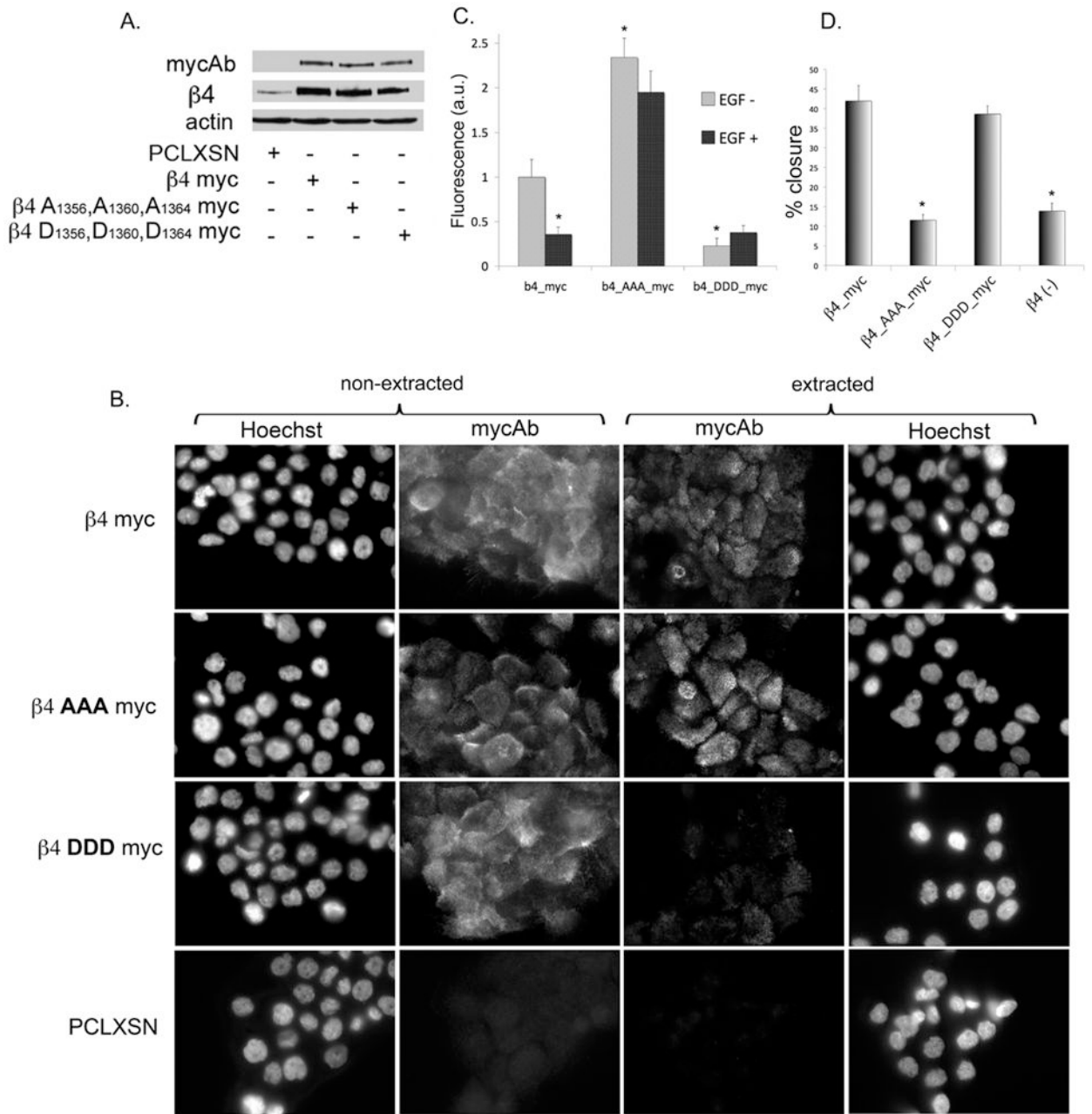


Figure 7. Abrogation of $\beta 4$ phosphorylation sites S₁₃₅₆S₁₃₆₀S₁₃₆₄ in SCC cells increases HD-like structure stability and hinders cell migration

Wild type $\beta 4$ myc or triple mutants containing either a Ser→Ala ($\beta 4$ -AAA-myc) or Ser→Asp ($\beta 4$ -DDD-myc) substitutions on S₁₃₅₆S₁₃₆₀S₁₃₆₄, were stably expressed in $\beta 4$ shRNA-silenced A431 cells. **A.** Cells expressing the $\beta 4$ constructs were analyzed by Western blotting using anti-myc Ab, showing similar level of expression. **B.** Analysis of HD-like structures stability in $\beta 4$ phosphorylation mutants. Cells grown on coverslips were detergent-extracted or not before fixation, and processed for IF using anti-myc Ab. Notice that only wt $\beta 4$ -myc and $\beta 4$ -AAA-myc mutant were substantially retained in HD-like structures after extraction. **C.** Resistance of $\beta 4$ phosphorylation mutants to EGF-induced

HD-like structures disassembly. Cells on coverslips were treated or not with EGF for 30 minutes before being detergent-extracted, fixed and processed for IF using anti-myc Ab to analyze incorporation into HD-like structures. Quantitation of HD-like structures by digital image analysis is expressed in arbitrary units of fluorescence integrated density per cell. * $p < 0.05$. **D.** Effect of $\beta 4$ mutants on cell migration using *in vitro* wound healing assay. Cells were assayed as in previous figure.