

Timosaponin Aiii Disrupts Morphological Plasticity and Migration of Breast Adenocarcinoma Through Inhibition of Integrin Internalisation

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Abstract

Cell migration is a critical step in tumour invasion and metastasis. To acquire invasive properties, cancer cells use their surrounding environment through dynamic and bidirectional interactions and change their morphology and mode of migration. Thus, inhibition of morphological plasticity regulated by paracrine interactions may be a promising approach for anti-cancer therapy. In this study, we found that timosaponin AIII (TAIII), a steroidal saponin isolated from the roots of *Anemarrhena asphodeloides*, disrupted the morphological changes and migratory activity of breast adenocarcinoma cells promoted by paracrine interactions with mammary epithelium-derived cells. TAIII suppressed lamellipodia formation of MDA-MB-231 cells in response to exogenous stimuli from MCF10A cells, thereby inhibiting morphological changes and migration. TAIII also attenuated membrane spreading and induced contraction of HeLa cells, followed by expansion of intercellular gaps. Furthermore, we analysed the intracellular dynamics of TAIII labelled with a fluorescent dye and found that labelled TAIII was internalised in a manner dependent on dynamin. We also found that TAIII blocked internalisation of cell surface proteins including integrin b1. These results provide a novel aspect to understand how TAIII exerts pharmacological activities in suppression of cancer cell migration.

Introduction

Cell motility is closely associated with various physiological processes such as tissue formation and migration during embryonic development, wound healing, and immunological responses. The initial step in motility is the formation of lamellipodia at the leading edge, which is a thin sheet of membrane-enclosed F-actin networks [1]. In accordance with the interaction of lamellipodia with the substrate, cells generate force to start directed movement by translocation of the cell body, followed by retraction of the rear part of the cell [2].

Deregulation of cell motility is involved in pathological events, particularly cancer metastasis [3]. After dissociating from a primary lesion, cancer cells invade through the extracellular matrix (ECM) of their surrounding host tissues and enter blood vessels [4]. In this process, cancer cells communicate with their neighbouring environment, which triggers and promotes invasive properties of cancer cells. Various types of cells, including macrophages and stromal cells, have been reported to be involved in the regulation of cancer cell migration and invasion [5-8]. On the basis of the interactions with the surrounding environment, cancer cells change their morphology and mode of migration. For example, carcinoma cells acquire a mesenchymal-like cell morphology with elongated membrane protrusions, which migrate through surrounding tissues and penetrate the adjacent basement membrane. Therefore, blockade of the morphological plasticity of cancer cells may become a goal of anti-cancer therapies.

Timosaponin AIII (TAIII) is a major steroidal saponin isolated from the root of *Anemarrhena asphodeloides* Bunge [9]. TAIII has multifaceted activities and is traditionally used as an anti-pyretic, anti-inflammatory, anti-diabetic, and anti-coagulant agent in Chinese medicine [10]. It has also been reported that TAIII ameliorates learning and memory deficits in a mouse model [11]. However, the pharmacological

activity of TAIll have been mainly studied in the context of anti-cancer properties in the past decades [10,12]. A major effect of TAIll on cancer cells is cytotoxicity. TAIll causes selective cell death in various kinds of cancer cells at a certain concentration [13-15]. TAIll induces cytochrome c release and caspase activation mediated by overproduction of reactive oxygen species and mitochondrial dysfunction, which lead to apoptosis [16]. Caspase activation followed by poly (ADP)-ribose polymerase cleavage is caused by phosphorylation of JNK and p38 in response to TAIll treatment [17]. TAIll also targets machineries that regulate cell cycle progression in cancer cells, thereby leading to cell cycle arrest in G1 or G2/M phase [13].

Additionally, compelling evidences have demonstrated the inhibitory effects of TAIll on the motility of cancer cells. It has been reported that TAIll regulates proteolytic activity and mRNA expression of matrix metalloproteinase-2/9 [18-21] that play pivotal roles in cell migration and invasion [22,23]. TAIll also contributes to upregulation of microRNAs (miRNAs), which suppresses metastatic properties of cancer cells. In breast cancer cell lines, TAIll induces expression of tumour-suppressive miRNAs miR-200c/141 and reduces expression of its negative regulator, B-cell-specific Moloney murine leukaemia virus integration site 1 [24]. Furthermore, TAIll regulates the expression of miR-129-5p through inhibition of the PI3K/AKT pathway in renal carcinoma cells without altering the viability of cancer cells [25]. Taken together, the anti-metastatic activity of TAIll has been explained by transcriptional control of migration regulatory factors. However, it remains unclear whether TAIll directly controls cellular machineries involved in the generation of migratory force to promote cancer cell invasion and metastasis.

In this study, we found that TAIll disrupted morphological plasticity and migration of breast adenocarcinoma cells promoted by paracrine interactions with mammary epithelial cell lines. TAIll suppressed lamellipodia formation induced by conditioned medium and attenuated membrane ruffling and spreading regulated by exogenously expressed constitutively active forms of Rho family small GTPase Rac1. Interestingly, labelled TAIll was internalised through dynamin-dependent endocytosis pathways and TAIll significantly inhibited internalisation of integrin b1, which downregulated cell adhesion activity on ECM. These findings suggest novel targets of TAIll to inhibit cancer cell motility by the interaction with the surrounding environment.

Materials And Methods

Reagents

The natural compound library was provided by the Cooperative Research Project of the Institute of Natural Medicine, University of Toyama, Japan. TAIll, timosaponin AI (TAI), and timosaponin BIII (TBII) were purchased from Kanto Kagaku (Tokyo, Japan). Anemarrhenasaponin I (AMS) was from MedChemExpress (Monmouth Junction, NJ, USA). Sarsasapogenin was from Fujifilm-Wako Pure Chemicals (Osaka, Japan). Dynasore was from Abcam (Cambridge, UK). Laminin-332, fibronectin, and vitronectin were from ReproCELL (Yokohama, Japan), BD Biosciences (Franklin Lakes, NJ, USA), and Fujifilm-Wako Pure Chemicals, respectively.

To prepare labelled TAlII, Alexa Fluor 568 hydrazide, sodium salt was purchased from Thermo Fisher Scientific (Waltham, MA, USA). Sodium periodate, dichloromethane, chloroform, and methanol were purchased from Fujifilm-Wako Pure Chemicals. All chemicals and reagents obtained from Fujifilm-Wako were guaranteed reagent grade. Spherical silica Chromatorex Q-pack SI 30 size:60 was purchased from Fuji Silysia (Aichi, Japan). ESIMS was measured on a Shimadzu LCmS-2020 system.

Cell culture and transfection

MDA-MB-231 cells (ATCC, Manassas, VA, USA) and HeLa cells (Kyoto) were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS). MCF10A cells (ATCC) were cultured in DMEM/F-12 (Thermo Fisher Scientific) supplemented with 5% heat-inactivated horse serum (Thermo Fisher Scientific), 20 ng/mL EGF (Miltenyi Biotec, Bergisch Gladbach, Germany), 10 mg/mL insulin (Sigma-Aldrich, St. Louis, MO, USA), 0.5 mg/mL hydrocortisone (Sigma-Aldrich), and 100 ng/mL cholera toxin (Sigma-Aldrich) at 37°C with 5% CO₂. Preparation and treatment with conditioned medium from MCF10A cells were performed as described previously [26].

For transfection, Lipofectamine LTX (Thermo Fisher Scientific) was used in accordance with the manufacturer's instructions. Expression plasmids for constitutively active Val12 Rac1, Val12 CDC42, and Val14 RhoA have been described previously [26]. The expression plasmid for LifeAct-mCherry was kindly provided by Dr. Naoki Watanabe (Kyoto University) and has been described previously [27].

Scratch wound assay

MDA-MB-231 cells were seeded on 96-well Image-lock plates (Essen Bioscience) in OPTI-MEM (Thermo Fisher Scientific). After incubation overnight, the cells were pretreated with natural compounds at a concentration of 1 mM for 30 min. Then, scratches were made using a 96-pin tool (Woundmaker) in accordance with the manufacturer's instructions, followed by treatment with conditioned medium. The plate was placed in an IncuCyte device and confluence was recorded every 1 h by phase contrast scanning for 8 h at 37°C with 5% CO₂. Images were analysed using IncuCyte ZOOM software. Inhibitory activity of each compound against scratch migration was calculated as the ratio of migration confluence (0 h/8 h) and normalised to the value of DMSO.

Immunofluorescence analyses and time-lapse microscopy

Immunofluorescence analyses were performed as described previously [28] with minor modifications. Cells grown on cover slips were fixed with 4% formaldehyde in phosphate-buffered saline (PBS) containing 0.1% Triton X-100 for 10 min, blocked for 30 min with 2% dry skim milk in PBS, and then incubated for 1 h with a mouse monoclonal anti-vinculin antibody (hVIN-1; Sigma-Aldrich) diluted in blocking buffer. After three times washes with PBS for 5 min each, the cells were incubated for 30 min with Alexa Fluor 488-conjugated anti mouse IgG (Thermo Fisher Scientific) diluted in blocking buffer together with Texas Red™-X Phalloidin (Thermo Fisher Scientific). The cells were then washed with PBS

and mounted, and immunofluorescence was observed under a confocal scanning laser microscope (SP8; Leica). Images were processed using Adobe Photoshop software.

For time-lapse analysis, cells seeded on a 3.5-cm glass-bottom culture dish (Asahi Techno Glass Co., Tokyo, Japan) were observed under the confocal scanning laser microscope equipped with a temperature- and CO₂-controlled stage-top live cell incubator (Tokai Hit, Fujinomiya, Japan).

Migration assay

Cell migration assays were performed in Boyden chambers as described previously [26] with minor modifications. MDA-MB-231 cells were trypsinised, washed twice with OPTI-MEM, and then seeded at 5×10^4 cells per well in OPTI-MEM. MCF10A CM was added to the lower chamber. For TAlII treatments, the inhibitor was added to both upper and lower chambers at the indicated concentrations. After 16 h of incubation, cells that had migrated to the lower surface of the membranes were stained with 0.5% crystal violet in 20% methanol and then counted.

Cell spreading assay

HeLa cells transfected with LifeAct-mCherry-expressing plasmids were trypsinised and washed once with DMEM containing 10% FBS. After incubation for 20 min at 37°C, the cells were seeded on a glass-bottom dish or coverslip coated with fibronectin (10 µg/mL) for time-lapse imaging and immunostaining, respectively. Using the acquired phase contrast images, the cell periphery of more than 100 cells in each condition was drawn and the cell spreading area was measured by ImageJ software.

Adhesion assay

96-well culture plates were coated with 2.5 µg fibronectin, laminin-332, or vitronectin in PBS overnight at 4°C and then blocked with BSA. HeLa cells were trypsinised and washed once with DMEM containing 10% FBS. After incubation for 30 min at 37°C, 8×10^4 cells were plated in each well and incubated for 30 min. The culture plates were washed with PBS to remove unattached cells and then attached cells were fixed with 4% formaldehyde and stained with 5 mg/ml crystal violet for 10 min. After unbound dye was washed away, the cells were lysed with 2% SDS and absorbance at OD 630 was measured.

Integrin internalisation assay

An integrin internalisation assay was performed as reported previously [29] with minor modifications. Briefly, HeLa cells were plated on a fibronectin-coated glass bottom dish and cultured overnight. Surface integrin β1 was labelled with an Alexa Fluor® 488 anti-human integrin β1 antibody (clone TS2/16, Biolegend) at 4°C for 60 min. Then, the cells were incubated at 37°C to allow internalisation. Quantification was performed for one of three representative experiments. Mean pixel intensity from each cell was determined using ImageJ software.

Preparation of Alexa Fluor 568-labelled timosaponin AIII

Water (63 μL), methanol (210 μL), and chloroform (35 μL) were added to a vial that contained TAlII (0.5 mg, 0.67 μmol) and mixed well to a clear solution. Then, a 100 mM aqueous solution of sodium periodate (7 μL , 0.7 μmol) was added to the solution, followed by stirring at room temperature for 30 minutes in the dark. Alexa Fluor 568 hydrazide (0.48 mg, 0.67 μmol) was added to the reaction mixture as a methanol solution (70 μL). After stirring at room temperature for 12 hours in the dark, the solvent was removed under the flow of a nitrogen stream. To the vial containing residue, 4.5 mg silica gel (Chromatorex Q-pack SI 30) suspended in methanol (200 μL) was added and then evaporated to remove the methanol. The obtained residue was suspended in hexane/chloroform (1/1 vol/vol) and loaded onto a silica gel (Chromatorex Q-pack SI 30, 200 mg) column. Products were eluted with methanol/chloroform (1/1 vol/vol) and collected in fractions (0.2 mL each). Fractions 2–7 were combined in a recovering flask, evaporated, and dried under vacuum to produce Alexa Fluor 568-labelled TAlII (0.53 mg, 54% yield) as a dark violet film [ESIMS m/z 1428 (1427.6 calc. for $\text{C}_{72}\text{H}_{91}\text{N}_{4}\text{O}_{22}\text{S}_2^-$, M^-)].

Statistical analysis

Results are presented as the mean \pm s.e.m. in Figs. 1d, 2b, 5b, 6c, and 6d and the mean \pm s.d. in Figs. 2d and 3d. All statistical analyses were performed using the two-tailed Student's t-test.

Results

TAlII suppresses morphological changes and migration of breast cancer cells induced by paracrine interactions with mammary epithelial cells

In our previous study [26], we demonstrated that conditioned medium from the mammary epithelial cell line MCF10A (MCF10A CM) regulates morphological plasticity and promotes migration of the breast cancer cell line MDA-MB-231. To identify natural compounds with the potential activity to disrupt the responses of breast cancer cells to paracrine interactions with mammary epithelial cells, we screened a natural compound library by a scratch wound assay. MDA-MB-231 cells were plated under a serum-deprived condition and treated with each natural compound at a concentration of 1 mM and MCF10A CM (Fig. 1a). As we reported previously [26], baicalein suppressed migration of MDA-MB-231 cells induced by MCF10A CM (Fig. 1b). Although saikosaponin a and d clearly suppressed migration of MDA-MB-231 cells, they also induced cell death during the assays (Fig. 1b). Among the compounds that inhibited migration of MDA-MB-231 cells, TAlII exhibited the strongest activity without cytotoxicity (Fig. 1b and c). The effect of TAlII in conditioned medium-induced migration was confirmed using a Boyden chamber (Fig. 1d). Therefore, we further analysed the effects of TAlII on breast cancer cell migration regulated by paracrine interactions.

First, we examined whether TAlII disrupted the morphological plasticity of breast adenocarcinoma cells regulated by conditioned medium from mammary epithelial cells. We treated MDA-MB-231 cells with MCF10A CM in the presence of TAlII for 1 h and then observed their morphology. As shown in Fig. 2a, MDA-MB-231 cells became elongated and changed their morphology in response to MCF10A CM.

However, MCF10A CM-induced morphological changes were perturbed in the presence of TAlII (Fig. 2a). These effects of TAlII were confirmed by quantification of the cell length (Fig. 2b).

Upon MCF10A CM treatment, MDA-MB-231 cells spread and formed lamellipodia at their leading edge and started migration (Fig. 2c and d, and Supplementary Video 1). Thus, we addressed whether TAlII also influenced lamellipodia formation. Time-lapse imaging analysis revealed that TAlII-treated cells did not extend lamellipodia in response to MCF10A CM. The ratio of the lamellipodia area induced by MCF10A CM in TAlII-treated cells was significantly lower than that in control cells (Fig. 2c–e and Supplementary Video 2). These results suggested that TAlII inhibited lamellipodia formation, thereby suppressing morphological plasticity and migration of breast adenocarcinoma cells induced by paracrine interactions with mammary epithelium-derived cells.

Because TAlII suppressed lamellipodia formation upon treatment with MCF10A CM, we next examined whether TAlII also affected lamellipodia formation induced by serum. MDA-MB-231 cells in normal culture medium were treated with TAlII and their morphology was observed. In this experiment, we treated cells with TAlII at a concentration of 10 mM because we did not observe any effects of TAlII at the lower concentration, which was probably due to the presence of protein components in serum, such as albumins and lipoproteins. Time-lapse analysis by phase contrast microscopy revealed that TAlII treatment disrupted lamellipodia formation (Supplementary Fig. S1a and b, and Supplementary Video 3). The number of cells with membrane ruffling was immediately and significantly decreased up to 10 min ($10.8 \pm 3.79\%$) and then temporary recovered at 20 min ($34.7 \pm 1.52\%$) but decreased again at 60 min ($7.88 \pm 1.21\%$) after treatment (Supplementary Fig. S1c). The lamellipodia area in each cell upon TAlII treatment showed similar changes (Supplementary Fig. S1d). Accordingly, the sizes of MDA-MB-231 cells became significantly smaller by approximately 50% after TAlII treatment (Supplementary Fig. S1e). Fluorescence imaging analysis revealed that actin-rich structures at the leading edge had disappeared in MDA-MB-231 cells treated with TAlII (Supplementary Fig. S1f). Together with the data from Fig. 2, TAlII negatively regulated lamellipodia formation induced by extracellular stimuli.

We also treated epidermoid carcinoma A431 cells with TAlII and observed their membrane dynamics. In the presence of serum, A431 cells tightly associated with each other and exhibited expanded lamellipodia in the cell edge and collective migration (Supplementary Fig. S2 and Video 4). However, lamellipodia formation had disappeared and migratory activity was apparently suppressed in TAlII-treated cells, while their cell-cell interactions shown by E-cadherin staining appeared intact (Supplementary Fig. S2 and Supplementary Video 5). These results corroborated the observations in breast adenocarcinoma cells that TAlII exerted inhibitory effects on cell migration by disrupting lamellipodia formation.

TAlII attenuates membrane dynamics regulated by Rac

To further investigate the effect of TAlII on membrane dynamics, we next accessed the effect of TAlII on cell spreading onto ECM. HeLa cells were transfected with the expression plasmid for LifeAct-mCherry, a widely used actin-binding peptide fused to mCherry protein [30], and replated onto a fibronectin-coated glass bottom dish. When the cells had attached to the glass-bottom dish, we treated them with TAlII and

examined their behaviours by time-lapse microscopy (Fig. 3a). After attachment to fibronectin, DMSO-treated cells showed membrane blebbing and then formed membrane ruffling and protrusions, followed by rapid spreading to the largest extension (Fig. 3b and Supplementary Video 6). We also observed dynamic remodelling of the actin cytoskeleton in response to the interaction with fibronectin (Fig. 3b and Supplementary Video 6). Moreover, vinculin-positive structures were scattered in the cell edge (Fig. 3c). Conversely, TAlII-treated cells did not form membrane blebs or ruffling. They adhered to the culture dish after replating and formed filopodia-like protrusions but could not spread on the dish (Fig. 3b and Supplementary Video 7). Vinculin-positive structures in TAlII-treated cells were obviously tiny compared with those in DMSO-treated cells (Fig. 3c). Accordingly, the cell area of TAlII-treated cells became significantly smaller, which was dependent on the dose of TAlII (Fig. 3d). These results demonstrated the specific activity of TAlII in the regulation of membrane dynamics, which is important for cell spreading.

Because it is well-known that lamellipodia formation and membrane ruffling is regulated by activated Rac, a Rho family GTPase [31], we addressed whether TAlII influenced activation of Rac. To this end, we first transfected HeLa cells with expression plasmids for green fluorescent protein (GFP) and LifeAct-mCherry and then subjected them to time-lapse imaging analyses (Fig. 4a). As shown by fluorescence signals, HeLa cells continuously formed membrane protrusions and showed ruffling under normal culture conditions (Fig. 4b and Supplementary Videos 8 and 9). However, similar to the other cultured cells examined in this study, these membrane dynamics were suppressed by 60 min of TAlII treatment, followed by expansion of intercellular gaps (Fig. 4b and Supplementary Videos 8 and 9). We also confirmed attenuation of membrane dynamics in response to TAlII treatment by kymograph analysis (Fig. 4b). We next examined membrane and actin cytoskeleton dynamics upon TAlII treatment in HeLa cells that overexpressed GFP-tagged constitutively activated Rac1 (GFP-Val12 Rac1) together with LifeAct-mCherry. Overexpression of GFP-Val12 Rac1 induced cell spreading and frequent membrane ruffling, followed by drastic remodelling of the actin cytoskeleton at the cell edge (Fig. 4c and Supplementary Videos 10 and 11). Interestingly, upon TAlII treatment, membrane ruffling was immediately attenuated and cell contraction was observed (Fig. 4c and Supplementary Videos 10 and 11). The rapid changes of membrane dynamics in Val12 Rac1-expressing cells induced by TAlII treatment were also shown by kymograph analysis (Fig. 4c). These data demonstrated that TAlII controlled membrane ruffling independently of the regulation of Rac activity. We also examined the effects of TAlII on filopodia formation and actin stress fibres regulated by constitutively active CDC42 (Val12 CDC42) and RhoA (Val14 RhoA), respectively. Unlike Val12 Rac1-expressing cells, Val12 CDC42- and Val14 RhoA-expressing cells retained membrane and actin cytoskeleton dynamics even after TAlII treatment (Supplementary Fig. S3 and Supplementary Videos 12–15), which implied that TAlII specifically inhibited membrane dynamics regulated by Rac signalling.

Immunofluorescence analysis revealed that HeLa cells treated with TAlII also showed morphological changes with accumulation of vinculin-positive structures at the cell edge and expanded intercellular gaps (Supplementary Fig. S4). Therefore, we also assessed the structure-activity relationship among saponins isolated from the rhizome of *Anemarrhena asphodeloides* in the regulation of membrane dynamics (Supplementary Fig. S4a). TAI is a deglycosylated derivative of TAlII, which lacks glucose in its

saccharide moiety. TBII and AMS share the steroid core and disaccharide moiety that are identical to those of TAIll. TBII contains an extra sugar moiety at the end of the steroidal side chain. However, AMS does not harbour additional sugar moieties, whereas a hydroxyl group is linked to the C-15 position in its steroid core. We found that HeLa cells treated with these TAIll derivatives did not show any obvious morphological changes similar to those observed after treatment with TAIll (Supplementary Fig. S4b). We also examined sarsasapogenin, an aglycone of TAIll, and found that it had no significant effects on cell morphology or vinculin localisation (Supplementary Fig. S4b). We further tested the natural compound library to determine whether it contained inhibitory activities in the regulation of membrane dynamics. Among the compounds in the library, only shikonin and alkannin affected the morphology of HeLa cells. The cells treated with these compounds became rounded and detached from the culture dish upon treatment, probably because of their cytotoxicity (Supplementary Table S1). Triterpenoid saponins, such as astragaloside IV, ginsenosides (Rb1, Rc, Rd, Re, and Rg1), glycyrrhizic acid, and saikosaponins (a, b2, c, and d), and cardiotonic steroid bufadienolides, such as bufalin and bufotalin (Supplementary Table S1), are listed in the library. However, unlike TAIll, these compounds did not exert any obvious effects on cell morphology. Collectively, these results supported the unique activity of TAIll in the regulation of membrane ruffling.

TAIll inhibits integrin internalisation

We next clarified the mechanisms underlying how TAIll suppressed membrane ruffling. To this end, we labelled TAIll with a fluorescence dye, Alexa Fluor 568, in accordance with the results from the structure-activity relationship among TAIll derivatives and examined its intracellular behaviour. Time-lapse analysis revealed that, after attachment to the plasma membrane, labelled TAIll formed intracellular vesicle-like structures and accumulated in cytoplasm (Fig. 5a and b, and Supplementary Video 16). Intriguingly, these intracellular vesicles and accumulation of labelled TAIll were less observed when cells were pretreated with a dynamin inhibitor, dynasore [32] (Fig. 5a and b, and Supplementary Video 17), which suggested that TAIll was internalised through a dynamin-dependent pathway.

Considering the observation that TAIll was internalised through dynamin-dependent endocytic pathways (Fig. 5), we focused on endocytic trafficking of cell surface proteins. In particular, we examined behaviours of integrin $\beta 1$ in response to TAIll treatment (Fig. 6a) because TAIll-treated cells showed impaired formation of adhesion complexes positive for vinculin during spreading on fibronectin (Fig. 3c). Cell surface integrin $\beta 1$ was labelled on ice with an Alexa Fluor 488-conjugated antibody, allowed to internalise at 37°C, and then subjected to time-lapse imaging analysis. Intracellular fluorescence signals from traced integrin $\beta 1$, which was transported via endosomes on the basis of their estimated size (0.5–2 μm) [29], were observed in cells treated with DMSO up to 20 min after incubation at 37°C and its fluorescence intensity was increased over time (Fig. 6b and c, and Supplementary Videos 18 and 19). Conversely, we did not detect any intracellular signals of internalised integrin $\beta 1$ in TAIll-treated cells after incubation at 37°C, in which fluorescent signals of LifeAct-mCherry indicated cell contraction, although signals from labelled integrin traced cell shapes prior to TAIll treatment (Fig. 6b and c, and Supplementary Videos 20 and 21). Interestingly, internalisation of integrin $\beta 1$ was clearly suppressed, whereas

continuous internalisation and intracellular accumulation of labelled TAIll were observed (Supplementary Fig. S5 and Supplementary Videos 22 and 23). We next examined the intracellular dynamics of integrin b1 and TAIll. We labelled cell surface integrin b1 and incubated the cells at 37°C for 1 h to allow internalisation and then treated the cells with labelled TAIll (Supplementary Fig. S6a). Whereas any colocalisation signals were not observed on the cell surface, some internalised integrin b1 and TAIll had merged in the cytoplasm (Supplementary Fig. S6b and Supplementary Videos 24–26). We also found that TAIll suppressed internalisation of E-cadherin induced by EGTA in A431 cells (Supplementary Fig. S7). Collectively, these results suggested that TAIll inhibited machineries that regulated internalisation of specific cargo proteins such as integrin b1.

To further elucidate the effect of TAIll on integrin mediated cell functions, we performed cell adhesion assays with various ECM proteins. Trypsinised HeLa cells were treated with TAIll for 30 min and replated on culture dishes coated with each ECM protein. After incubation for another 30 min, cells associated with the ECM were measured. We examined purified fibronectin, laminin-332, and vitronectin that serve as ligands for integrins $\alpha 5\beta 1$, $\alpha 3\beta 1$, and $\alpha v\beta 1$, respectively, which are expressed in HeLa cells [33]. Adhesion assays revealed that TAIll significantly interfered with the substantial interaction of HeLa cells with purified integrin ligands in a dose-dependent manner (Fig. 6d), which demonstrated that TAIll targeted integrin heterodimers consisting of at least integrin b1. Thus, the results suggested that the effect of TAIll on the regulation of internalisation of cell surface proteins, such as integrin b1, contributed in part to its inhibitory activity against cell migration through suppression of lamellipodia formation and membrane ruffling.

Discussion

In this study, we found that TAIll exerted a strong suppressive effect on the migration of breast adenocarcinoma cells by paracrine interactions with mammary epithelium-derived cells. Lamellipodia formation and membrane ruffling induced by conditioned medium from mammary epithelial cells were clearly disrupted by TAIll. TAIll inhibited internalisation of cell surface proteins, including integrin b1, which explains its inhibitory activity against lamellipodia formation and cell migration. The overall mechanisms of TAIll in the regulation of cell motility have not been determined completely. Our study provides a novel aspect to understand how TAIll exerts anti-metastatic effects.

Moreover, we investigated intracellular behaviours of TAIll using fluorescent labelling techniques. Fluorescent labelling is a powerful strategy to perform high-resolution imaging analysis of target molecules. Indeed, fluorescent dye-labelled analogues have been used to examine intracellular localisation and trafficking of cholesterol [34]. However, labelling strategies have not been applied to saponins because the biological activities, including cytotoxicity, of saponins are closely associated with their aglycone moieties and the number and structures of monosaccharides in their sugar chains [35]. In accordance with the results from the examination of the structure-activity relationships among TAIll derivatives, we modified TAIll and evaluated intracellular behaviours of labelled TAIll. To our knowledge, this is a first report to reveal the intracellular dynamics of TAIll. The data obtained using labelled TAIll

revealed a novel function of TAlII as a negative regulator of endocytic trafficking. This would be a great benefit to address intracellular behaviours of natural compounds, which led us to elucidate their molecular functions and properly understand their pharmacological properties.

A shift in migration modes allows cancer cells to adapt and invade into their surrounding tissues. Paracrine interactions with the microenvironment play pivotal roles in regulating the changes in migration mode, such as epithelial-mesenchymal transition. To date, various factors such as the ECM and chemokines have been reported to have tumour-promoting properties [36,37]. We found that TAlII suppressed morphological plasticity and migration of breast adenocarcinoma cells induced by conditioned medium from mammary epithelial cells. The activity of conditioned medium from mammary epithelial cells is mainly mediated by secreted laminin-332 [26]. Laminins are a component in the basement membrane of mammary ducts and secreted from myoepithelial cells [7]. In particular, laminin-332 plays critical roles in regulating the motility of breast cancer cells [38,39]. Clinical investigations have demonstrated that laminin-332 is enriched in the invasion interface of breast-invasive ductal carcinoma [39]. Additionally, the expression level of laminins is significantly associated with distant metastasis-free survival of patients with breast cancers at the early stage [40]. Therefore, our observations support development of TAlII as an anti-cancer drug to suppress cancer cell metastasis and invasion dependent on the microenvironment.

Interestingly, TAlII disrupted membrane dynamics by inducing overexpression of constitutively active Rac1, but not other Rho family GTPases, in HeLa cells. These observations suggest that TAlII targets downstream of Rac, such as WAVE2 and IRSp53 [41-43], to suppress lamellipodia formation and membrane ruffling. However, our data demonstrated the inhibitory effects of TAlII on internalisation of cell surface proteins including integrin b1 and integrin-mediated cell adhesion. Integrin heterodimers play a crucial role in formation of adhesion complexes that serve as platforms to induce lamellipodia formation and promote cell migration [44]. Integrins are also required to stabilise lamellipodia formation [45]. The spatiotemporal regulation of cell surface integrins through trafficking systems gives impact to their functions in the context of cell migration [46-50]. Furthermore, a ligand of integrin heterodimers, laminin-332, plays important roles in the regulation of morphological plasticity and cell migration of MDA-MB-231 cells [26]. Collectively, our results lead us to propose that the activity of TAlII to block internalisation of integrin b1 should contribute, in part, to its inhibitory effect on cell migration through negative regulation of lamellipodia formation. This conclusion is consistent with a previous observation that demonstrated the importance of endocytosis to regulate lamellipodia formation and cancer cell invasion [51].

A recent study has demonstrated that TAlII inhibits platelet aggregation induced by U46619 *in vitro* and prevents thrombus formation *in vivo* [52]. Anti-platelet and anti-thrombotic activities of TAlII are reported to be mediated by decreased adenosine diphosphate (ADP) secretion caused by suppression of thromboxane A2 receptor activity and the Gq signalling pathway. Apart from the suppression of ADP secretion, blockade of integrin activation is an important strategy to interfere with platelet crosslinking and platelet-derived thrombus formation. In fact, inhibitors of glycoprotein IIb/IIIa, an integrin complex on

platelets, such as abciximab, eptifibatide, and tirofiban, are used clinically [53]. Therefore, we speculate that the anti-platelet and anti-thrombotic activity of TAlII might, in part, be also mediated by the negative regulation of integrin internalisation. TAlII also exerts anti-angiogenesis effects through inactivation of VEGF signalling [15]. Receptor internalisation and subsequent cytoplasmic trafficking are required for activation of signalling cascades mediated by proangiogenic growth factors in endothelial cells *in vitro* and *in vivo* [54]. Our observations might also explain the pharmacological activity of TAlII in the context of endocytic regulation of signal transduction.

We found suppression of integrin $\beta 1$ internalisation induced by TAlII regardless of continuous intracellular accumulation of TAlII itself, which demonstrated that TAlII regulated endocytosis of a specific subclass of cargoes. Additionally, this result also raised a question how TAlII regulated internalisation of cell surface proteins. TAlII is considered to direct membrane-associating proteins in accordance with computational docking approaches demonstrating that TAlII would associate with the receptor for advanced glycation end products (RAGE) to modulate RAGE/MAPK signalling pathways [55]. Furthermore, cardiac glycosides, digoxin, and ouabain bind to cell surface Na^+, K^+ -ATPase to exert its enzymatic activity [56]. Therefore, TAlII might interact with its membrane targets to regulate internalisation of cargo proteins. However, in this study, we did not observe clear colocalised signals of labelled TAlII and integrin $\beta 1$ at the plasma membrane. It has been reported that abrogation of the recycling pathway impairs integrin internalisation and downregulation of migratory activity [29]. Taken together with our observation that labelled TAlII colocalised with integrin $\beta 1$ in intracellular vesicles of cells treated with labelled TAlII after internalisation of integrin $\beta 1$ was allowed, TAlII might disrupt the regulatory machineries for trafficking vesicles, thereby suppressing internalisation of cell surface cargoes. More than two decades, numerous efforts were made to reveal the overall network of integrin trafficking and various membrane-associating proteins were found to be involved in integrin trafficking [29,57]. It would also be interesting to explore the molecular targets of TAlII, which might expand our knowledge of the regulatory machineries for cargo transport.

Herbal medicines have been used as therapies against various diseases. While numerous compounds have been isolated, little is known about how these compounds exert their fascinating pharmacological effects. In this study, we uncovered the activity of TAlII in suppression of lamellipodia formation and migration of breast cancer cells, which would be mediated, in part, through blocking internalisation of cell surface proteins including integrin. However, it remains unresolved how TAlII blocks internalisation of cell surface proteins. Further studies are needed to clarify the mechanisms underlying the regulation of internalisation by TAlII, which would contribute to re-evaluating traditional medicines for drug development.

Abbreviations

AMS: Anemarrhenasaponin I

CM: Conditioned medium

ECM: Extracellular matrix

GFP: Green fluorescent protein

miRNAs: microRNAs

RAGE: Receptor for advanced glycation end products

TAllI: Timosaponin AllI

TBI: Timosaponin BI

Declarations

Ethics declarations

Funding

JSPS KAKENHI (17K08339) for Toshimasa Ishizaki

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Conflicts of interest/Competing interests

We have no conflicts of interest or competing interests.

Availability of data and material

It can be supplied if necessary.

Code availability

IncuCyte ZOOM software and Adobe Photoshop software are licensed. ImageJ is opensource software.

Ethics approval and consent to participate

Not applicable

Consent for publication

Not applicable

Authors' contributions

T.T. and T.I. conceived the study design. T.T., K.H., and T.I. performed the experiments and analyzed the data, with contributions from T.A., S.M., T.S., and A.K. produced materials for the experiments. T.T wrote

the manuscript, with contributions from K.H., T.S. and T.I.. All the authors have read and confirmed the final version of the manuscript.

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Figures

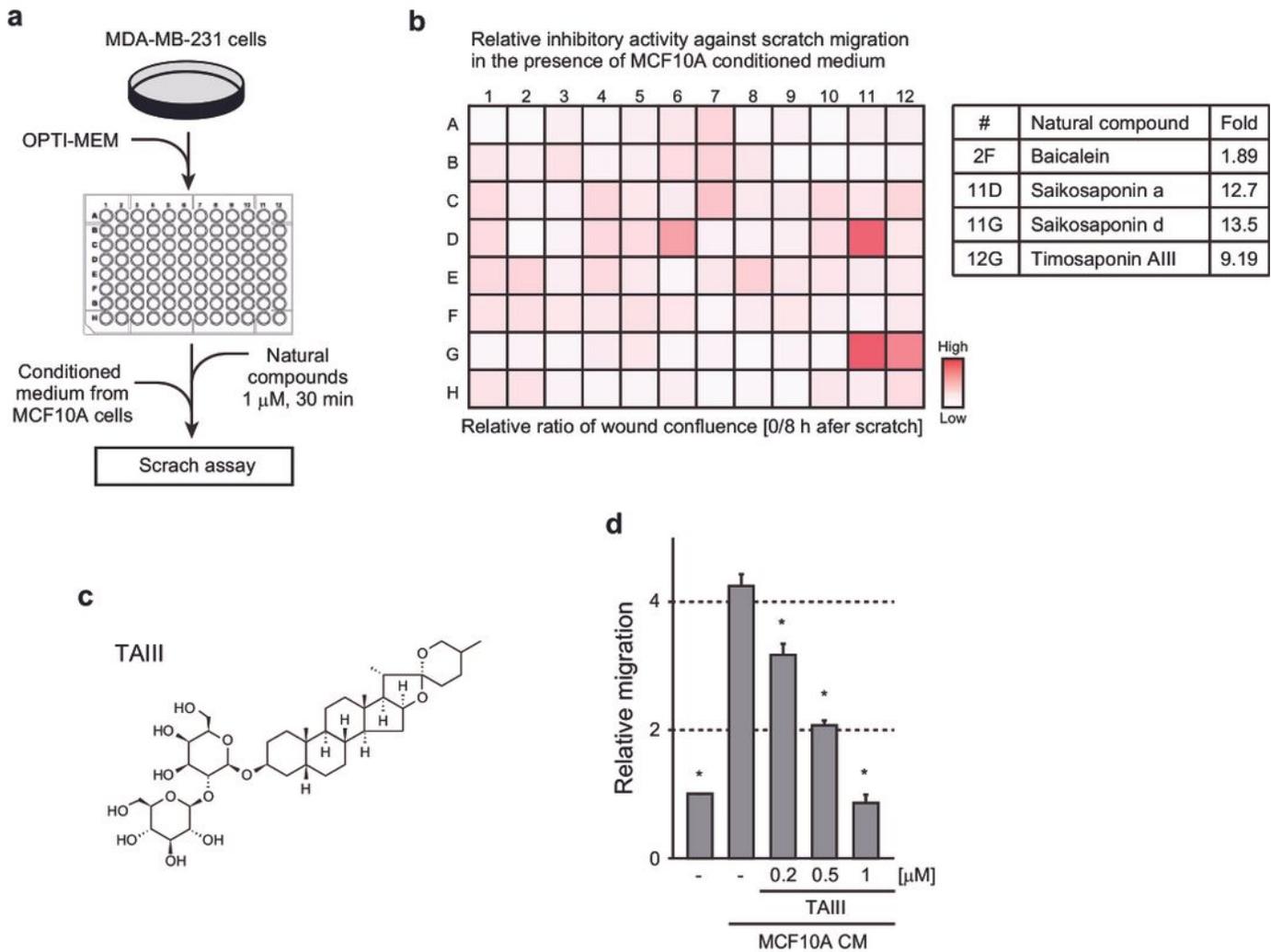


Figure 1 TAIll disturbed morphological changes and migration of breast adenocarcinoma cells induced by paracrine interactions with mammary epithelial cells.

a. Schematic experimental procedure for screening of natural compound library. MDA-MB-231 cells were plated in a 96-well culture plate under serum deprivation condition (OPTI-MEM). Cells were pretreated with each natural compound for 30 min and then treated with MCF10A CM and subjected to scratch assays. b. Heat map of the inhibitory activities of each compound. The relative inhibitory activity was calculated as the relative ratio of wound confluency at 0 h to 8 h and normalised by the value of DMSO. c. Structure of TAIll. d. MDA-MB-231 cells were subjected to Boyden chamber migration assays in the presence of MCF10A CM and indicated concentrations of TAIll. Each column shows the mean \pm s.e.m. of three independent experiments. * $p < 0.05$ compared with cells treated with MCF10A CM alone.

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Figure 1

TAIll disrupts morphological changes and migration of breast adenocarcinoma cells induced by paracrine interactions with mammary epithelial cells. a. Schematic experimental procedure for screening of natural compound library. MDA-MB-231 cells were plated in a 96-well culture plate under serum deprivation condition (OPTI-MEM). Cells were pretreated with each natural compound for 30 min and then treated with MCF10A CM and subjected to scratch assays. b. Heat map of the inhibitory activities of each

compound. The relative inhibitory activity was calculated as the relative ratio of wound confluency at 0 h to 8 h and normalised by the value of DMSO. c. Structure of TAIII. d. MDA-MB-231 cells were subjected to Boyden chamber migration assays in the presence of MCF10A CM and indicated concentrations of TAIII. Each column shows the mean \pm s.e.m. of three independent experiments. * $p < 0.05$ compared with cells treated with MCF10A CM alone.

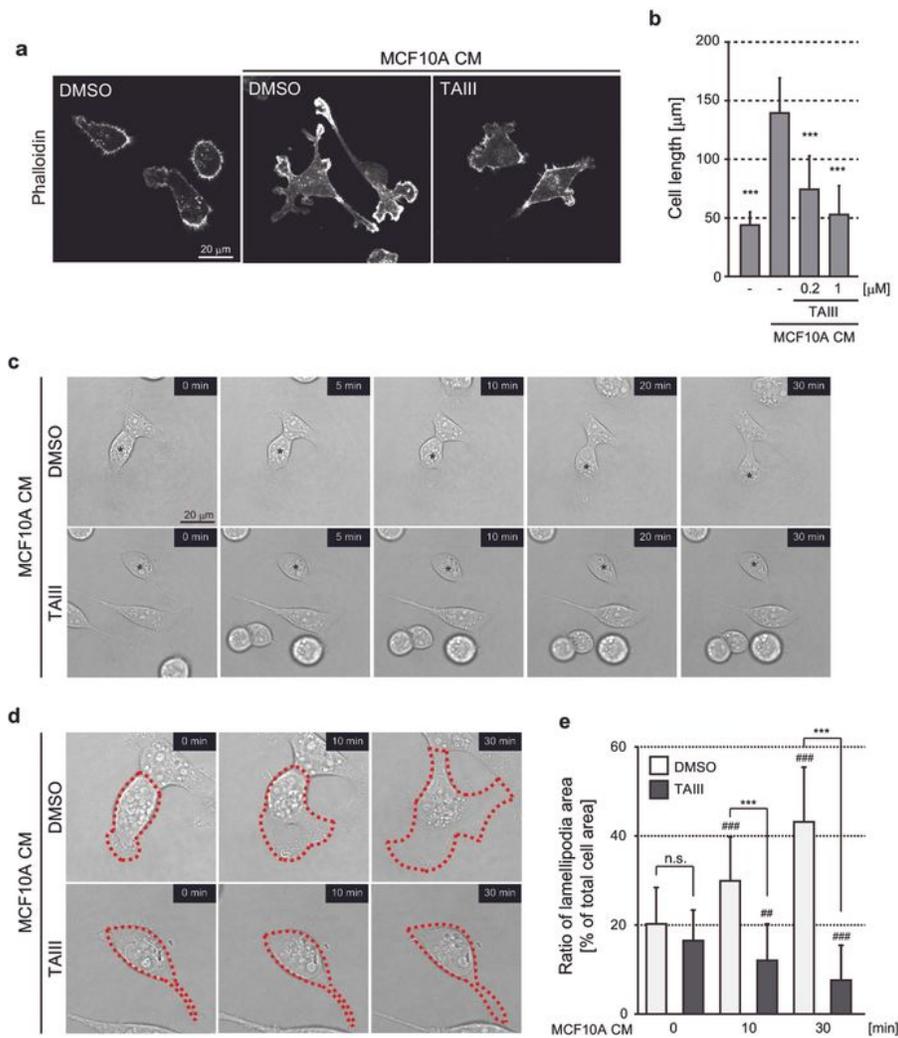


Figure 2 TAIII suppressed lamellipodia formation induced by MCF10A conditioned medium.

a. MDA-MB-231 cells treated with MCF10A CM in the presence of 1 μ M TAIII for 1 h were fixed and stained with phalloidin. Scale bar: 20 μ m. b. Quantification of the length of MDA-MB-231 cells treated with MCF10A CM in the presence of TAIII at the indicated concentrations for 1 h. Each column shows the mean \pm s.e.m. of three independent experiments. *** $p < 0.005$ compared with cells treated with MCF10A CM alone. c. MDA-MB-231 cells were pretreated with DMSO or 1 μ M TAIII and then further treated with MCF10A CM and observed under time-lapse phase-contrast microscopy. A series of images over the indicated time course is shown. Asterisks indicate identical cells among frames in each treatment. Scale bar: 20 μ m. d. Magnified images of cells labelled with asterisks in (c) at the indicated time after MCF10A CM treatment. Total cell areas are outlined with dotted lines. e. The ratio of lamellipodia area after MCF10A CM treatment ($n > 30$). Each column shows the mean \pm s.d. ### $p < 0.01$ and #### $p < 0.005$ compared with cells at 0 min in DMSO or TAIII treatment. *** $p < 0.005$ compared TAIII with DMSO treated cells at each time points.

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TAllI suppresses lamellipodia formation induced by MCF10A CM. a. MDA-MB-231 cells were pretreated with DMSO or 1 μ M TAllI and then further treated with MCF10A CM and observed under time-lapse phase-contrast microscopy. A series of images over the indicated time course is shown. Asterisks indicate identical cells among frames in each treatment. Scale bar: 20 μ m. b. Magnified images of cells labelled with asterisks in (a) at the indicated time after MCF10A CM treatment. Total cell areas are outlined with dotted lines. c. The ratio of lamellipodia area after MCF10A CM treatment ($n > 30$). Each column shows the mean \pm s.d. ## $p < 0.01$ and ### $p < 0.005$ compared with cells at 0 min in DMSO or TAllI treatment. *** $p < 0.005$ compared TAllI with DMSO treated cells at each time points.

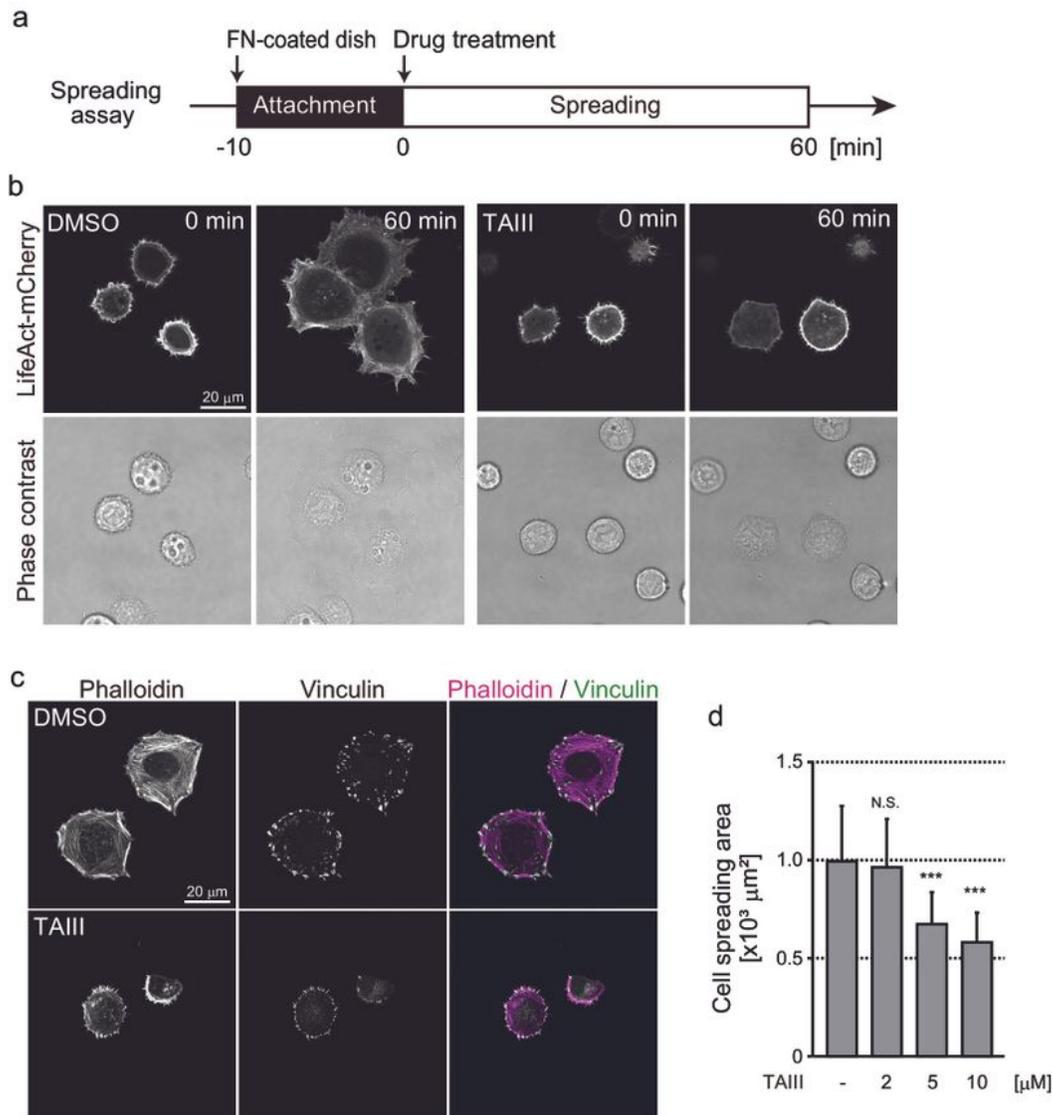


Figure 3 TAIII inhibited cell spreading on ECM.

a. Schematic experimental procedure for the cell-spreading assays in HeLa cells treated with TAIII. b. HeLa cells transfected with LifeAct-mCherry expression plasmid were plated onto fibronectin-coated dish. After attached, cells were treated with 10 μM TAIII and subjected to time-lapse imaging analysis. Fluorescence and phase contrast images were taken until 60 min after TAIII treatment. c. HeLa cells subjected to cell-spreading assay were fixed at 60 min after TAIII treatment and stained for vinculin and phalloidin. Bar; 20 μm . d. Quantification of cell spreading area at 60 min after TAIII treatment at the indicated concentrations. $n > 30$. Each column shows the mean \pm s.d. *** $p < 0.005$ compared with cells treated with DMSO.

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Figure 3

TAIII inhibits cell spreading on fibronectin. a. Schematic experimental procedure for the cell-spreading assays in HeLa cells treated with TAIII. b. HeLa cells transfected with LifeAct-mCherry expression plasmid were plated onto fibronectin-coated dish. After attached, cells were treated with 10 μM TAIII and subjected to time-lapse imaging analysis. Fluorescence and phase contrast images were taken until 60 min after TAIII treatment. c. HeLa cells subjected to cell-spreading assay were fixed at 60 min after TAIII treatment

and stained for vinculin and phalloidin. Bar; 20 μ m. d. Quantification of cell spreading area at 60 min after TAlII treatment at the indicated concentrations. $n > 30$. Each column shows the mean \pm s.d. *** $p < 0.005$ compared with cells treated with DMSO.

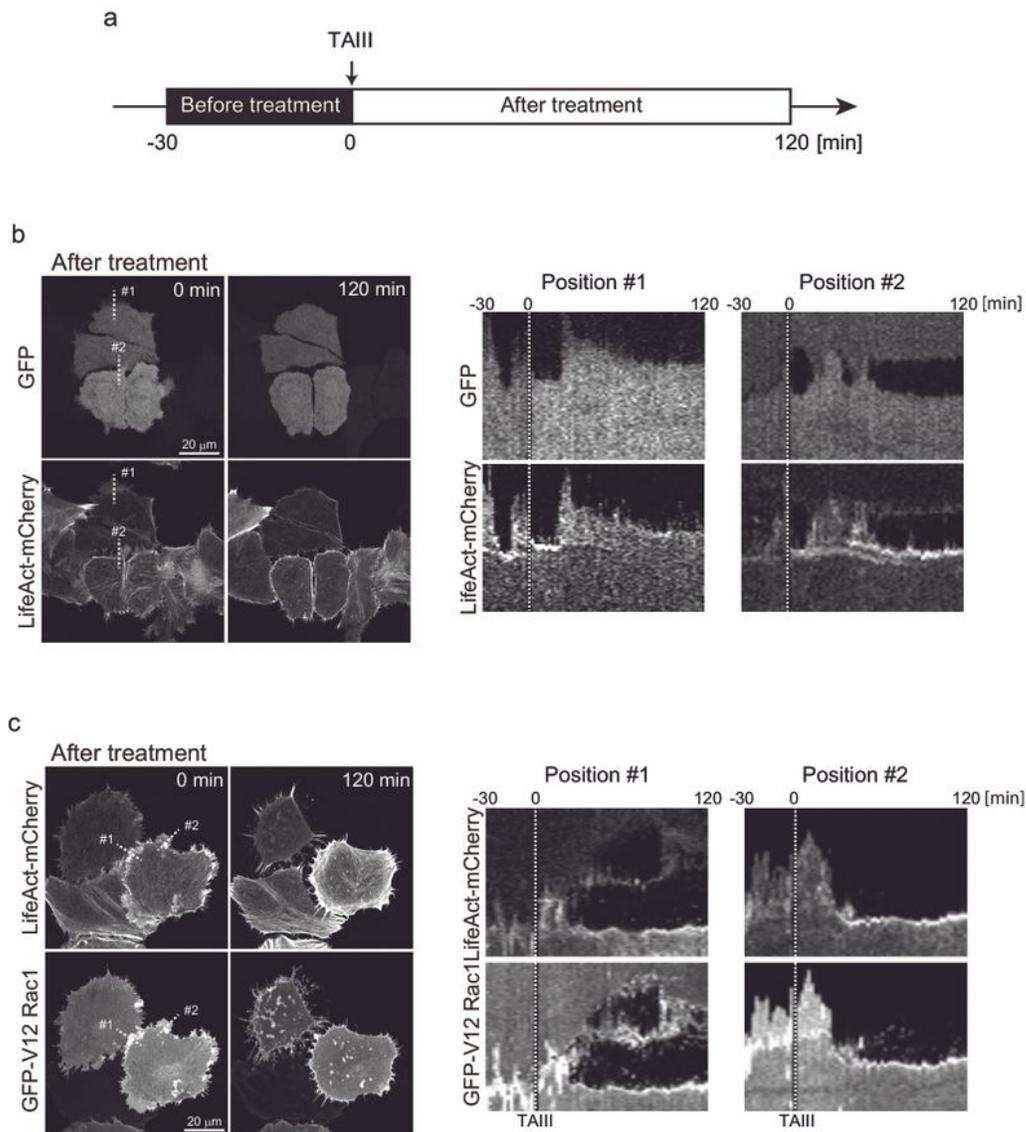


Figure 4 TAlII interfered with membrane dynamics regulated by Rac.

a. Schematic experimental procedure for time-lapse imaging analysis in HeLa cells treated with TAlII. b. HeLa cells transfected with GFP and LifeAct-mCherry expression plasmids. At 16 h post-transfection, the cells were subjected to time-lapse imaging analysis in the presence of TAlII treatment (10 μ M). Left: representative images of transfected cells after TAlII treatment (0 and 120 min); scale bar: 20 μ m. Right: kymograph analyses at dotted lines in the left panel. c. HeLa cells transfected with GFP-tagged constitutively active Rac1 (Val12) and LifeAct-mCherry expression plasmids were subjected to time-lapse imaging analysis as in (b). Left: representative images of transfected cells after TAlII treatment (0 and 120 min); scale bar: 20 μ m. Right: kymograph analyses at dotted lines in the left panel.

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Figure 4

TAlII interferes with membrane dynamics regulated by Rac. a. Schematic experimental procedure for time-lapse imaging analysis in HeLa cells treated with TAlII. b. HeLa cells transfected with GFP and LifeAct-

mCherry expression plasmids. At 16 h post-transfection, the cells were subjected to time-lapse imaging analysis in the presence of TAlII treatment (10 μ M). Left: representative images of transfected cells after TAlII treatment (0 and 120 min); scale bar: 20 μ m. Right: kymograph analyses at dotted lines in the left panel. c. HeLa cells transfected with GFP-tagged constitutively active Rac1 (Val12) and LifeAct-mCherry expression plasmids were subjected to time-lapse imaging analysis as in (b). Left: representative images of transfected cells after TAlII treatment (0 and 120 min); scale bar: 20 μ m. Right: kymograph analyses at dotted lines in the left panel.

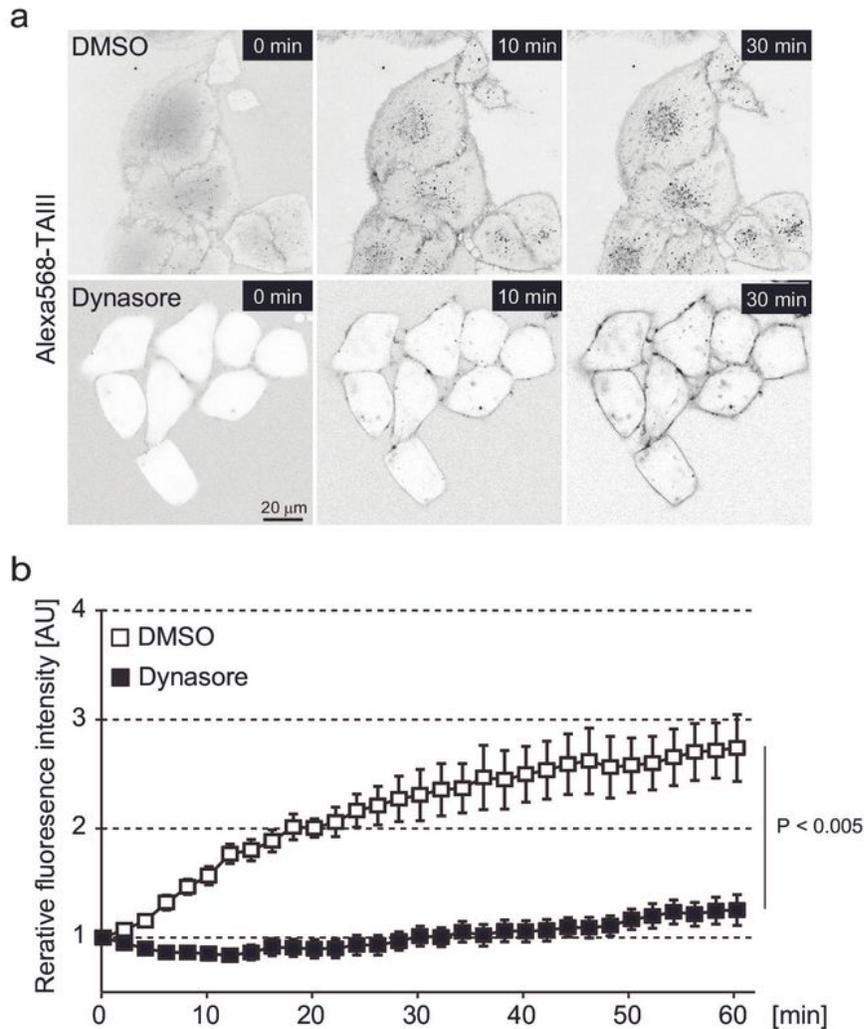


Figure 5 TAlII was internalised dependent on dynamin.

a. HeLa cells were treated with DMSO or dynasore (10 μ M) for 6 h and further treated with Alexa Fluor 568 labelled-TAlII. Then, time-lapse imaging analysis was performed for 60 min. Representative images (0, 10 and 30 min after TAlII treatment) are presented. Scale bar: 20 μ m. b. Quantification of signals intensity of internalised labelled-TAlII. Intensity of Alexa Fluor 568 normalized against levels at onset of 0 min. Data are presented as the mean \pm s.e.m. ($n > 6$). $p < 0.005$ at 60 min.

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Figure 5

TAlII is internalised dependent on dynamin. a. HeLa cells were treated with DMSO or dynasore (100 μ M) for 6 h and further treated with Alexa Fluor 568 labelled-TAlII. Then, time-lapse imaging analysis was performed for 60 min. Representative images (0, 10 and 30 min after TAlII treatment) are presented. Scale bar: 20 μ m. b. Quantification of signals intensity of internalised labelled-TAlII. Intensity of Alexa Fluor 568 normalized against levels at onset of 0 min. Data are presented as the mean \pm s.e.m. ($n > 6$). $p < 0.005$ at 60 min.

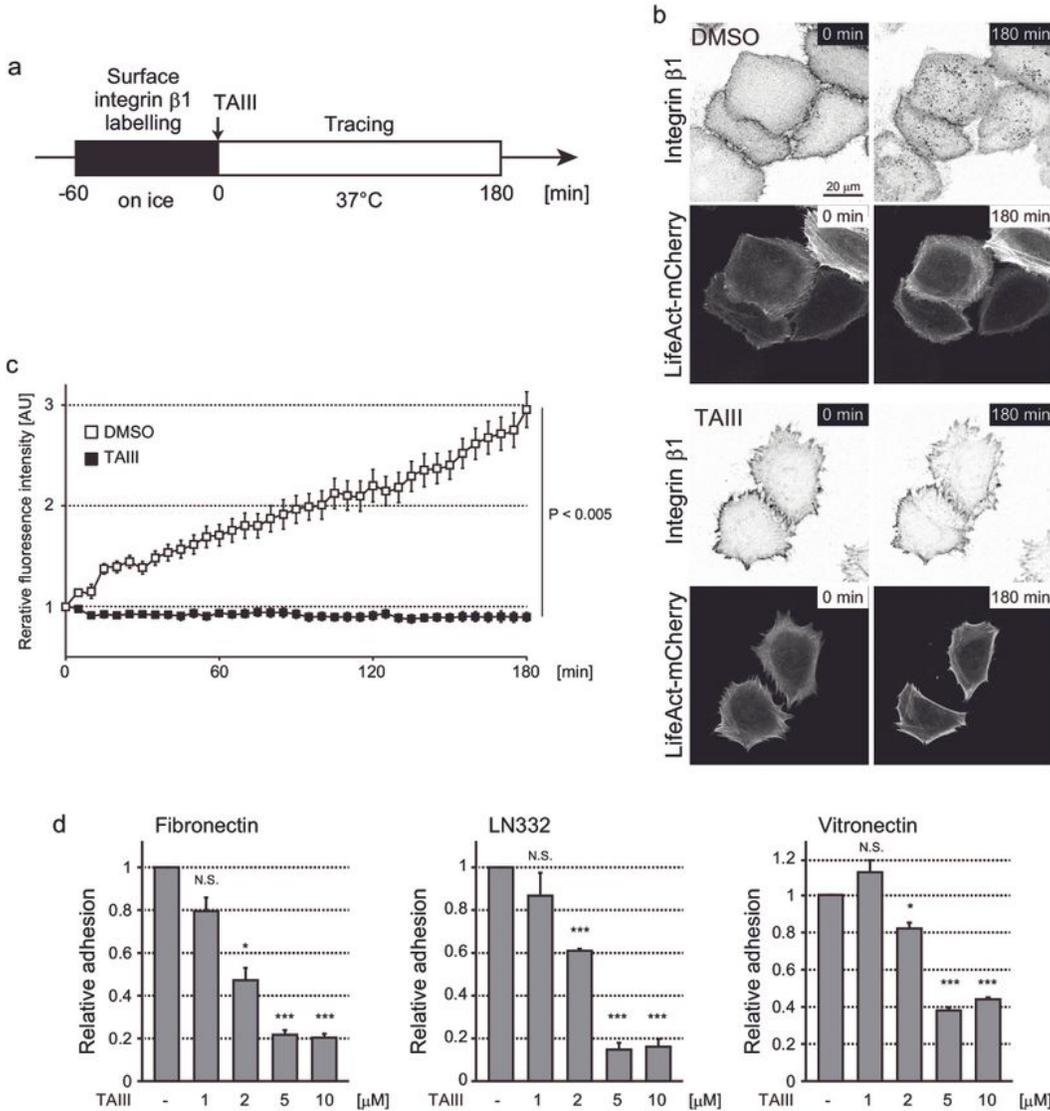


Figure 6 TAlII blocked integrin internalisation.

a. Schematic experimental procedure for the integrin internalisation assay. b. HeLa cells expressing LifeAct-mCherry plated onto fibronectin-coated dish were subjected to the integrin internalisation assay. Cells were analysed by time-lapse imaging analysis until 180 min after incubation at 37°C (tracing). Representative images of transfected cells after DMSO and TAlII treatment are shown. Scale bar: 20 μ m. c. Quantifications of signal intensity of internalised integrin $\beta 1$ after tracing. Intensity of Alexa Fluor 488 was normalized against levels at onset of tracing (0 min). Data are presented as the mean \pm s.e.m. ($n=20$). $p < 0.005$ at 60 min. d. HeLa cells were trypsinised and treated with TAlII for 30 min at the indicated concentrations, and then subjected to adhesion assay. Each column shows the mean \pm s.e.m. of three independent experiments. * $p < 0.05$, ** $p < 0.001$ and *** $p < 0.005$ compared with cells treated with DMSO.

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Figure 6

TAllI blocks integrin internalisation. a. Schematic experimental procedure for the integrin internalisation assay. b. HeLa cells expressing LifeAct-mCherry plated onto fibronectin-coated dish were subjected to the integrin internalisation assay. Cells were analysed by time-lapse imaging analysis until 180 min after incubation at 37°C (tracing). Representative images of transfected cells after DMSO and TAllI treatment are shown. Scale bar: 20 µm. c. Quantifications of signal intensity of internalised integrin β1 after tracing. Intensity of Alexa Fluor 488 was normalized against levels at onset of tracing (0 min). Data are presented as the mean ± s.e.m. (n=20). $p < 0.005$ at 60 min. d. HeLa cells were trypsinised and treated with TAllI for 30 min at the indicated concentrations, and then subjected to adhesion assay. Each column shows the mean ± s.e.m. of three independent experiments. * $p < 0.05$, ** $p < 0.001$ and *** $p < 0.005$ compared with cells treated with DMSO.

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