

Please cite the Published Version

Lončarić, M, Stojanović, N, Rac-Justament, A, Coopmans, K, Majhen, D, Humphries, JD, Humphries, MJ and Ambriović-Ristov, A (2023) Talin2 and KANK2 functionally interact to regulate microtubule dynamics, paclitaxel sensitivity and cell migration in the MDA-MB-435S melanoma cell line. Cellular and Molecular Biology Letters: an international journal, 28 (1). 56 ISSN 1425-8153

DOI: <https://doi.org/10.1186/s11658-023-00473-6>

Publisher: BMC

Version: Published Version

Downloaded from: <https://e-space.mmu.ac.uk/632760/>

Usage rights:  [Creative Commons: Attribution 4.0](https://creativecommons.org/licenses/by/4.0/)

Additional Information: This is an open access article published in Cellular and Molecular Biology Letters, by BMC.

Enquiries:

If you have questions about this document, contact openresearch@mmu.ac.uk. Please include the URL of the record in e-space. If you believe that your, or a third party's rights have been compromised through this document please see our Take Down policy (available from <https://www.mmu.ac.uk/library/using-the-library/policies-and-guidelines>)

RESEARCH LETTER

Open Access



Talin2 and KANK2 functionally interact to regulate microtubule dynamics, paclitaxel sensitivity and cell migration in the MDA-MB-435S melanoma cell line

Marija Lončarić¹ , Nikolina Stojanović¹ , Anja Rac-Justament¹ , Kaatje Coopmans¹ , Dragomira Majhen¹ , Jonathan D. Humphries² , Martin J. Humphries³ and Andreja Ambriović-Ristov^{1*}

*Correspondence:
Andreja.Ambriovic.Ristov@irb.hr

¹ Laboratory for Cell Biology and Signalling, Division of Molecular Biology, Ruđer Bošković Institute, Zagreb, Croatia

² Department of Life Science, Manchester Metropolitan University, Manchester, United Kingdom

³ Wellcome Centre for Cell-Matrix Research, Faculty of Biology, Medicine and Health, University of Manchester, Manchester, United Kingdom

Abstract

Background: Focal adhesions (FAs) are integrin-containing, multi-protein structures that link intracellular actin to the extracellular matrix and trigger multiple signaling pathways that control cell proliferation, differentiation, survival and motility. Microtubules (MTs) are stabilized in the vicinity of FAs through interaction with the components of the cortical microtubule stabilizing complex (CMSC). KANK (KN motif and ankyrin repeat domains) family proteins within the CMSC, KANK1 or KANK2, bind talin within FAs and thus mediate actin-MT crosstalk. We previously identified in MDA-MB-435S cells, which preferentially use integrin $\alpha\text{V}\beta\text{5}$ for adhesion, KANK2 as a key molecule enabling the actin-MT crosstalk. KANK2 knockdown also resulted in increased sensitivity to MT poisons, paclitaxel (PTX) and vincristine and reduced migration. Here, we aimed to analyze whether KANK1 has a similar role and to distinguish which talin isoform binds KANK2.

Methods: The cell model consisted of human melanoma cell line MDA-MB-435S and stably transfected clone with decreased expression of integrin αV (3 αV). For transient knockdown of talin1, talin2, KANK1 or KANK2 we used gene-specific siRNAs transfection. Using previously standardized protocol we isolated integrin adhesion complexes. SDS-PAGE and Western blot was used for protein expression analysis. The immunofluorescence analysis and live cell imaging was done using confocal microscopy. Cell migration was analyzed with Transwell Cell Culture Inserts. Statistical analysis using GraphPad Software consisted of either one-way analysis of variance (ANOVA), unpaired Student's *t*-test or two-way ANOVA analysis.

Results: We show that KANK1 is not a part of the CMSC associated with integrin $\alpha\text{V}\beta\text{5}$ FAs and its knockdown did not affect the velocity of MT growth or cell sensitivity to PTX. The talin2 knockdown mimicked KANK2 knockdown i.e. led to the perturbation of actin-MT crosstalk, which is indicated by the increased velocity of MT growth and increased sensitivity to PTX and also reduced migration.

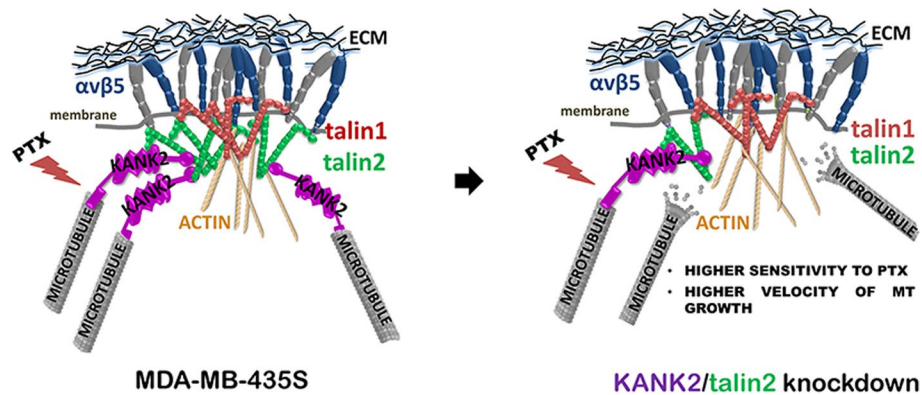
Conclusion: We conclude that KANK2 functionally interacts with talin2 and that the mechanism of increased sensitivity to PTX involves changes in microtubule dynamics.



These data elucidate a cell-type-specific role of talin2 and KANK2 isoforms and we propose that talin2 and KANK2 are therefore potential therapeutic targets for improved cancer therapy.

Keywords: Focal adhesion, Talin1, Talin2, KANK1, KANK2, Cortical microtubule stabilizing complex

Graphical Abstract



Background

Integrins are cell surface adhesion receptors composed of α and β subunits, forming 24 different $\alpha\beta$ heterodimers that connect cells either to other cells or to the extracellular matrix (ECM) [1, 2]. Binding of ECM ligands results in the formation of several functionally and structurally different classes of integrin adhesion complexes (IACs). These include nascent adhesions, focal adhesions (FAs) and fibrillar adhesions, which serve as a link to the actin cytoskeleton [3], hemidesmosomes, which link to intermediate filaments [4], and reticular adhesions (RAs), which are not associated to the cytoskeleton [5]. IACs activate a range of signaling pathways, which in turn control different physiological processes such as cell survival, differentiation, proliferation, migration and metabolism [2].

Focal adhesions are the most extensively studied IACs involved in the conversion of extracellular chemical and mechanical cues into biochemical signals [6–9]. Microtubules (MTs) regulate both FA assembly [10, 11] and turnover during cell migration [12, 13] by interacting with proximally-located cortical microtubule stabilizing complexes (CMSCs) [14] or plasma membrane-associated platforms (PMAPs) [15]. These complexes contain liprins α and β [16], ELKS (proteins rich in amino acids E, L, K and S) [17], kinesin family member KIF21A [18], plus-end tracking protein EB1, CLASPs (CLIP-associating proteins) and LL5 β [11, 16, 18, 19], as well as KN motif and ankyrin repeat domains (KANKs) family proteins [14, 20–22], all of which have been detected in biochemically-isolated IACs using mass spectrometry (MS)-based proteomics [23, 24]. KANKs, which bind directly to talins, are essential for connecting the CMSC to integrins [14, 20], but has not yet been established which isoforms contribute to the connection and whether their binding is cell type-specific.

Talins bind directly to the cytoplasmic domain of β integrin subunits via their FERM domain. The integrin-talin connection, supported further by binding of vinculin to talin, links integrins to actin [25]. Two isoforms, talin1 and talin2, are expressed by two separate genes but share a highly conserved structure composed of a head domain and 13 rod domains [26]. Knockout of talin1, but not talin2, is lethal in mice [27, 28]. KANKs bind to the R7 rod domain of talin, anchoring and stabilizing MTs in close proximity to FAs [14, 20]. The four KANK family isoforms (1–4) possess three distinct domains, a KANK N-terminal (KN) domain, a coiled-coil (CC) domain whose composition and number varies among the family members, and an ankyrin repeat domain (ANKRD), which consists of five ankyrin repeats [14, 20–22, 29]. The KN domain binds to talin, the CC domain to liprin- β 1 and ANKRD to KIF21A [18, 30, 31].

The composition of IACs of different cell lines (termed the adhesome) has been catalogued following biochemical isolation or in situ followed by proteomic analysis [32]. Isolated IACs have primarily been generated from cells seeded for short periods of time on fibronectin [33, 34], but also from cells in long-term culture where the cells secrete their own ECM, which is then used for binding of available integrin heterodimers. The latter method has shown that cells predominantly use a single integrin, whether it is organized as FAs and/or RAs [5, 24], or as hemidesmosomes [35, 36]. The advantage of using cell models in which IACs are already explored and are composed of preferentially one integrin heterodimer facilitates the research of the role of individual molecules in a specific integrin-forming IACs.

Our previous study showed that the melanoma cell line MDA-MB-435S preferentially uses integrin α V β 5 for adhesion to uncoated surfaces and identified key components of its adhesome [24]. Following integrin α V knockdown we observed increased sensitivity to MT poisons, paclitaxel (PTX) and vincristine, and decreased migration. α V β 5 was responsible for both effects [37]. By comparing the composition of IACs in MDA-MB-435S cells to cells in which integrin α V was stably knocked down, we identified the components of the integrin α V β 5 adhesome. Proteins decreased after α V knockdown included talin1, talin2 and all the components of CMSC, including KANK2, whose knockdown in MDA-MB-435S cells mimicked the effect of integrin α V knockdown and resulted in increased sensitivity to MT poisons and decreased migration. We concluded that KANK2 is a key molecule linking integrin α V β 5 IACs to MTs, enabling the actin-MT crosstalk important for both sensitivity to MT poisons and cell migration [24]. However, according to the (MS)-based proteomic analysis, these cells also express KANK1 at a low level. Furthermore, it was not possible to deduce which talin isoform plays a role in binding KANK1 or KANK2 proteins, and what is the exact mechanism of altered sensitivity to PTX.

The aim of the present work was to distinguish which talin isoform binds KANK2 and whether KANK1 has a similar role to KANK2. In addition, our aim was to further elucidate the mechanism of increased sensitivity to PTX upon integrin depletion. We show that KANK1 does not colocalize with α V β 5 FAs. Moreover, KANK1 is not altered after stable integrin α V knockdown and its knockdown does not change the sensitivity to PTX. We show that KANK2 and talin2 functionally interact thus enabling the actin-MT crosstalk important for both sensitivity to MT poisons and cell migration. We also show that the mechanism by which this occurs involves changes in microtubule dynamics.

Since talin2 or KANK2 knockdown mimicked the effect of integrin αV knockdown and resulted in increased sensitivity to PTX and reduced migration, talin2 and KANK2 are potential therapeutic targets for improved PTX therapy and reduced metastasis which might have a more uniform response than targeting integrins.

Materials and methods

Cell cultures and generation of stably transfected cells

The human melanoma cell line MDA-MB-435S (a spindle-shaped variant of the parental MDA-MB-435) was obtained from the American Type Culture Collection (ATCC). The isolation of the MDA-MB-435S-derived cell clone with decreased expression of integrin αV (3 αV clone) was described previously [24]. MDA-MB-435S and 3 αV cell populations with fluorescently labelled end-binding protein 3 (EB3) were generated by stable transfection of Dendra2-EB3-7 plasmid containing the photoconvertible fluorescent protein Dendra2 fused with EB3 (EB3-Dendra2) (Addgene plasmid #57715) using Lipofectamine 2000 (Thermo Fisher Scientific) and selection with 1.2 mg/mL geneticin (Sigma-Aldrich). Cells were grown in DMEM (Invitrogen) supplemented with 10% (v/v) FBS (Invitrogen) (DMEM-FBS) at 37 °C with 5% CO₂ (v/v) in a humidified atmosphere.

All the experiments in our study were done with cells that were seeded on uncoated surfaces. The reason for this is that we published previously that MDA-MB-435S cells preferentially use integrin $\alpha V\beta 5$ for adhesion and that, in such conditions these cells become more sensitive to paclitaxel upon integrin αV or KANK2 knockdown [24].

Transient siRNA transfection

For transient siRNA transfection experiments, cells (4×10^5) were seeded in 3.5 cm Petri dishes and transfected 24 h later, using Lipofectamine RNAiMax (13778150, Thermo Fisher Scientific), with 25 nM of control (Silencer™ Select Negative Control No. 1 siRNA, Ambion) or gene-specific siRNA for KANK1 (target sequence: CAGAGAAGG ACATGCGGAT), KANK2 (target sequence: ATGTCAACGTGCAAGATGA), Talin1 (target sequence: TGAATGTCCTGTCAACTGCTG) or Talin2 (target sequence: TTT CGTTTTTCATCTACTCCTT) [14], all purchased from Sigma. Successful knockdown was validated by SDS-PAGE and western blotting (WB) using specific antibodies and matched labelled secondary antibodies. The primary and secondary antibodies are listed in Additional file 9: Table S1.

Isolation of IACs

Integrin adhesion complexes were isolated from cells cultivated for 48 h as previously described [24, 38]. Briefly, cells were washed with DMEM-HEPES and incubated with Wang and Richard's reagent (DTBP, 6 mM, Thermo Fisher Scientific) for 10 min. DTBP was quenched with 0.03 M Tris-HCl (pH 8) and cells were lysed using modified RIPA buffer (50 mM Tris-HCl, pH 7.6; 150 mM NaCl; 5 mM disodium EDTA, pH 8; 1% (w/v) Triton X-100, 0.5% (w/v) SDS, 1% (w/v) sodium deoxycholate). Cell bodies were removed by high-pressure washing with tap water for 10 s and remaining adhesion complexes were collected by scraping into adhesion recovery solution (125 mM Tris-HCl,

pH 6.8; 1% (w/v) SDS; 150 mM dithiothreitol). Samples containing isolated IACs were acetone-precipitated and further processed for WB analysis [33].

SDS-PAGE and western blotting

Total cell lysates were obtained from 3.5 cm Petri dishes in 200 μ L RIPA buffer supplemented with protease inhibitor cocktail (ThermoFisher). Samples for SDS-PAGE were collected by scraping. Samples containing an equal amount of protein were mixed in 6 \times Laemmli loading buffer (375 mM Tris-HCl (pH 6.8), 30% (w/v) glycerol, 12% (w/v) SDS, 0.02% (w/v) bromophenol blue, 12% (v/v) 2-mercaptoethanol) to reach a final 1 \times concentration, sonicated and heated for 5 min at 96 °C. Isolated IACs were prepared for SDS-PAGE by solubilization in 2 \times Laemmli loading buffer and heating for 20 min at 70 °C while shaking (1000 rpm). All samples were loaded onto pre-casted gradient gel (4–15% Mini-PROTEAN TGX) (Bio-Rad), separated by SDS-PAGE and semi-dry transferred to nitrocellulose (Bio-Rad). The membrane was blocked in 5% (w/v) non-fat dry milk or 5% (w/v) bovine serum albumin (BSA, Carl Roth), and incubated overnight with the appropriate antibodies, followed by incubation with horseradish peroxidase coupled secondary antibody. The primary and secondary antibodies are listed in Additional file 9: Table S1. Detection was performed using chemiluminescence (PerkinElmer) and documented with Uvitec Alliance Q9 mini (BioSPX b.v.). Blots were quantified using ImageJ.

Cell survival analysis

An MTT (3-((4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) (Millipore) assay was used to determine sensitivity of cells to PTX (Sigma-Aldrich). Briefly, 24 h after seeding in 96-well tissue culture plates ($1\text{--}1.2 \times 10^4$ cells/well) cells were treated with different concentrations of PTX. Seventy-two hours later, the absorbance of MTT-formazan product dissolved in dimethyl sulfoxide was measured with a microplate reader (Awareness Technology, Inc.) at 600 nm. Measured absorbance was proportional to the number of viable cells.

Confocal microscopy and live cell imaging

For immunofluorescence analysis (IF), cells were plated on coverslips in a 24-well plate (3.5×10^4 cells/well). After 48 h, cells were fixed using ice-cold methanol for 10 min or 2% (v/v) paraformaldehyde for 12 min followed by permeabilization with 0.1% (v/v) Triton X-100 for 2 min, and stained with protein specific primary antibodies for 1 h, followed by incubation with conjugated secondary antibodies for 1 h. The primary and secondary antibodies are listed in Additional file 9: Table S1. Actin stress fibers were stained with rhodamine phalloidin (Cell Signaling Technology). Cells were mounted with DAPI Fluoromount-G (SouthernBiotech) and fluorescence and respective IRM images were acquired using an inverted confocal microscope (Leica TCS SP8 X, Leica Microsystems) with the HC PL APOCS2 63 \times /1.40 oil-immersion objective, zoom set at 2 \times . Images were analyzed using LAS X software 3.1.1 (Leica Microsystems) and ImageJ. For actin stress fibers quantification, only those fibers that ended with FAs, marked by integrin β 5 staining, were identified as stress fibers and

quantified. For microtubule quantification, only microtubules within 5 μm from the cell edge (identified through IRM images) were quantified. All images were taken with the focus adjusted to the adhesion sites of cells at the upper surface of glass coverslip.

Time-lapse live cell microscopy of MDA-MB-435S-EB3 and 3 α V-EB3 cells was performed 48 h upon seeding in a 4-chamber 35 mm Cellvis glass bottom dish (4.5×10^4 cells/chamber). In experiments with transient knockdown or PTX treatment, cells were imaged 48 h after transfection or 24 h after treatment with PTX. Images were taken every 13 s for five min per cell using HC PL APO CS2 63 \times /1.40 oil-immersion objective on the Leica TCS SP8 X microscope (Leica Microsystems), 3.5 \times zoom. Obtained movies were analyzed with ImageJ software (Manual tracking plugin).

Cell migration

Serum-starved (24 h) cells (8×10^4) were placed in Transwell Cell Culture Inserts (pore size, 8 μm) (Corning) in DMEM containing 0.1% (w/v) BSA and left to migrate toward 10% (v/v) FBS in DMEM as a chemoattractant. After 22 h, cells that remained on the upper side of the inserts were removed with cotton-tipped swabs. Inserts were fixed in 4% paraformaldehyde for 15 min followed by staining with 1% (w/v) crystal violet in PBS for 90 min. Cells on the underside of the inserts were photographed using Olympus BX51TF microscope (five images/sample). The number of cells was determined using ImageJ software (Multi-point tool).

Cell proliferation assay

To determine cell proliferation upon knockdown of each talin and KANK, Click-iT assay was used according to the manufacturer's instructions (Thermo Fisher Scientific, United States). Cells were seeded in a 6-well plate (3.5×10^5 cells/well) and upon 24 h transfected with either control siRNA or specific siRNA. After 24 h, cells were trypsinized and seeded back in the same well to mimic steps during MTT assay. Next day, two hours before harvesting, modified thymidine analog EdU (5-ethynyl-2'-deoxyuridine, final concentration 10 μM) was added. Cells were collected, fixed with 4% (w/v) paraformaldehyde, permeabilized with saponin, stained with Alexa-Fluor 488 azide (in the presence of CuSO_4) and analyzed by flow cytometry. Flow cytometry was performed using BD FACSCalibur (BD Biosciences, United States). Data were analyzed using FCS Express (De Novo Software, United States). To determine the proliferation rate, the frequencies of the proliferative (EdU+) cells were compared.

Statistical analysis

GraphPad Prism version 9.0.0 (GraphPad Software) was used to analyze data. Data obtained from IF, WB, proliferation and migration assay were analyzed by one-way analysis of variance (ANOVA) with Dunnett's multiple comparison or by unpaired Student's *t*-test, and expressed as histograms depicting mean \pm standard deviation (SD); violin plots or scatter plots with marked median; ns denotes not significant; * denotes $p < 0.05$; ** denotes $p < 0.01$; *** denotes $p < 0.001$; **** denotes $p < 0.0001$. Data from time-lapse live cell microscopy were analyzed by ordinary one-way ANOVA with Šidák's multiple comparisons test, with a single pooled variance and shown as violin plots or scatter plots with marked median; ns denotes not significant; * denotes $p < 0.05$; ** denotes

$p < 0.01$; *** denotes $p < 0.001$; **** denotes $p < 0.0001$ or by unpaired Student's *t*-test, and expressed as mean \pm SD; ns denotes not significant; * denotes $p < 0.05$; ** denotes $p < 0.01$; *** denotes $p < 0.001$; **** denotes $p < 0.0001$. Data obtained from MTT experiments were analyzed by two-way ANOVA with Šidák's multiple comparisons test, with a single pooled variance and expressed as mean \pm SD; ns denotes not significant; * denotes $p < 0.05$; ** denotes $p < 0.01$; *** denotes $p < 0.001$; **** denotes $p < 0.0001$.

Results

Talin isoform specific regulation of integrin $\alpha V\beta 5$ and KANK localization

To determine the location of talin isoforms within integrin $\alpha V\beta 5$ FAs, we analyzed the distribution of talin1, talin2 and integrin $\beta 5$ using IF. Figure 1A shows that both talin1 and talin2 colocalize with integrin $\beta 5$, thus confirming MS data that both talins are constituents of integrin $\alpha V\beta 5$ IACs [24]. Although KANK1 and KANK2 are similar in structure [22], they displayed different subcellular localizations in MDA-MB-435S cells. KANK1 was primarily restricted to the cell edge, in regions that were rich in talin1- and talin2-positive FAs (Fig. 1B). Conversely, KANK2 localized to talin1 and talin2-positive FAs both at the cell edge and in the center of the cell (Fig. 1C).

We have previously shown that KANK2 is a key molecule linking integrin $\alpha V\beta 5$ IACs to MTs, enabling the actin-MT crosstalk required for both sensitivity to MT poisons and cell migration. MDA-MB-435S cells also express KANK1, but it was not clear from our MS data whether there was a difference in KANK1 expression after integrin αV knockdown [24]. Therefore, we analyzed KANK1 expression in 3 αV cells, obtained from MDA-MB-435S cells following stably decreased expression of integrin αV , by WB of total cell lysates, and found that its level was not significantly different from that in MDA-MB-435S cells (Fig. 2A, B; Additional file 1: Fig. S6). In addition, we performed knockdown of talin1 or talin2 and showed that neither KANK1 level (Fig. 2A,B; Additional file 1: Fig. S6) nor its localization changed (Additional file 1: Fig. S1A, B). Most significantly, knockdown of KANK1 did not alter the sensitivity of MDA-MB-435S cells to PTX (Additional file 1: Fig. S1C).

KANK2 expression was reduced in the 3 αV clone compared to MDA-MB-435S cells [24] (Fig. 2A, B; Additional file 1: Fig. S6). To identify the isoform dependence of talin binding to KANK2, we analyzed the expression of KANK2 in MDA-MB-435S cells transiently transfected with talin1- or talin2-specific siRNA, using WB. Knockdown of talin 1 decreased the level of talin2 and KANK2, while knockdown of talin2 did not change the level of KANK2 (Fig. 2A, B; Additional file 1: Fig. S6). Simultaneous knockdown of both talins resulted in a comparable level of KANK2 as observed upon talin1 knockdown (Fig. 2A, B; Additional file 1: Fig. S6).

Next, the talin isoform dependency of MDA-MB-435S cell adhesion was analyzed by knockdown with control, talin1- or talin2-specific siRNA. Figure 2C shows that knockdown of talin1 (Fig. 2A, B; Additional file 1: Fig. S6), led to disruption of $\alpha V\beta 5$ FAs and a simultaneous reduction of talin2-positive structures (Fig. 2C, D). This conclusion was supported by a reduction in the cell spreading area (Additional file 1: Fig. S2), with only a few remaining $\alpha V\beta 5$ FAs containing talin1 (Fig. 2D first row), and shown by reduced expression of talin1 (Fig. 2A, B; Additional file 1: Fig. S6). It should be noted that after knockdown of talin1 there is some residual talin2 that colocalizes with integrin $\beta 5$, which

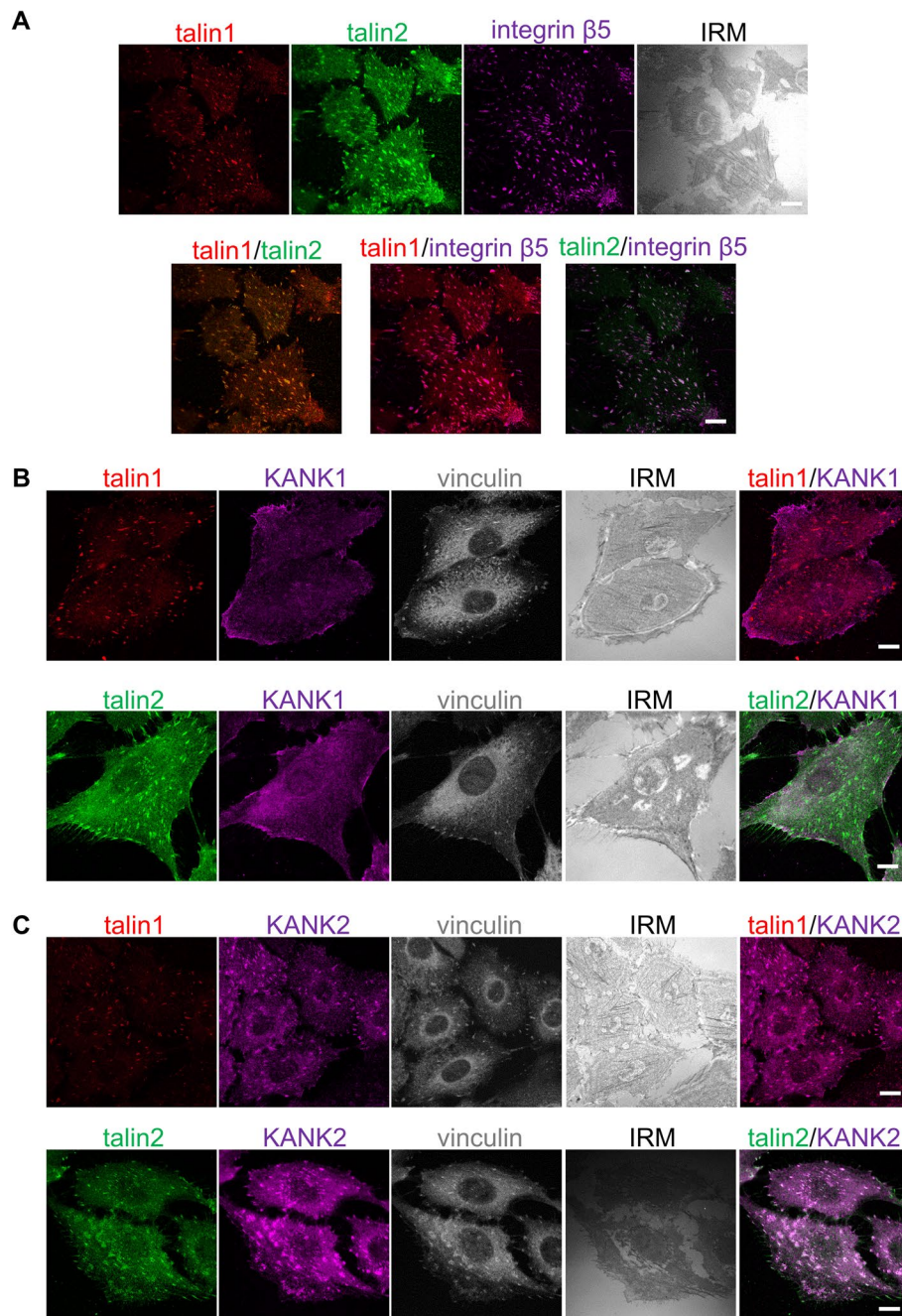


Fig. 1 KANK2, unlike KANK1, is localized within α V β 5 FAs. **A** Talin1 and talin2 are localized within integrin α V β 5 FAs. Forty-eight hours after seeding MDA-MB-435S cells were methanol fixed, and stained with anti-talin1 antibody followed by Alexa-Fluor IgG1 555-conjugated antibody (red), anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green) and anti-integrin β 5 antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) and interference reflection microscopy (IRM) images were taken. **B, C** KANK2, unlike KANK1, localizes in talin1 and talin2-positive FAs. Forty-eight hours after seeding MDA-MB-435S cells were methanol fixed and stained with anti-talin1 or anti-talin2 antibody followed by Alexa-Fluor 546-conjugated antibody (red) or Alexa-Fluor 488-conjugated antibody (green) and anti-KANK1 or anti-KANK2 antibody followed by Alexa-Fluor 555-conjugated antibody or Alexa-Fluor 488-conjugated antibody (shown in magenta). Vinculin was visualized using conjugated anti-vinculin Alexa Fluor 647 antibody (shown in grey) and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m

is very likely a part of RAs and whose presence in MDA-MB-435S cells was observed previously [5, 24]. Upon talin2 knockdown, colocalized integrin α V β 5 FAs and talin1 were still present (Fig. 2C) and cell spreading was not affected (Additional file 1: Fig. S2) despite the highly efficient knockdown of talin2 (Fig. 2A, B; Additional file 1: Fig. S6), demonstrated as disappearance of talin2-positive structures (Fig. 2C). This morphological analysis was supported by a quantitative assessment of the FA number, where talin1-, talin2- or β 5-positive structures decreased upon talin1 knockdown, but the number of talin1- and β 5-positive structures didn't change upon talin2 knockdown (Fig. 2D first row). These data, together with decreased FA size upon talin1 knockdown, measured by analyzing talin1-, talin2- or β 5-positive structures, and increased FA size upon talin2 knockdown, measured by analyzing talin1- and β 5-positive structures but not by analyzing talin2-positive structures (Fig. 2D second row), indicates that talins are not functionally redundant.

We next performed IF analysis of KANK2 following either talin1 or talin2 knockdown. Again, talin1 knockdown reduced cell spreading (Additional file 1: Fig. S2), with only a few remaining structures containing talin1, talin2 and KANK2 (Fig. 3A, B) consistent with their reduced level as measured by WB (Fig. 2A, B; Additional file 1: Fig. S6). Of note, we observed colocalization of talin2 and KANK2 remaining in structures lacking talin1 which is reminiscent of RAs [5, 24]. However, upon talin2 knockdown, there was no change in the expression (Fig. 2A, B; Additional file 1: Fig. S6) or the appearance of KANK2 which colocalizes with talin1 (Fig. 3A). A statistical analysis of the number of KANK2-positive structures per cell confirmed that only talin1 knockdown had a significant effect (Fig. 3B).

In conclusion, our data show that MDA-MB-435S cells in long term culture assemble FAs containing talin1, talin2 and KANK2 but not KANK1. In addition, these data indicate that talins are not functionally redundant. Namely, talin1 led to disruption of α V β 5 FAs and a simultaneous reduction of talin2- and KANK2 positive structures while reducing their total protein levels in the cell. In contrast, knockdown of talin2 did not affect either the protein levels or the localization of integrin α V β 5 FA, talin1, or KANK2.

(See figure on next page.)

Fig. 2 Talin1, unlike talin2, is necessary for formation of integrin α V β 5 FAs. **A** Talin1 knockdown, unlike talin2, reduces the level of KANK2 in whole cell lysates, while KANK1 level does not change upon either talin1 or talin2 knockdown. WB analysis of talin1, talin2, KANK1 or KANK2 in MDA-MB-435S cells transfected with either control, talin1, talin2 or a combination of talin1 and talin2-specific siRNAs and in the 3aV clone with decreased expression of integrin α V. Forty-eight hours after transfection total cell lysates were collected and WB analysis was performed. The results presented are representative of three independent experiments with similar results. **B** Quantification of data presented in (A). Histogram data are plotted as mean \pm SD ($n \geq 3$) relative to expression in MDA-MB-435S cells transfected with control siRNA that was set as 1 (indicated by a dotted line). Data were analyzed by one-way ANOVA with Dunnett's multiple comparison. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **C** Knockdown of talin1 leads to disruption of α V β 5 FAs. Forty-eight hours after transfection with either control siRNA, talin1 or talin2-specific siRNA, MDA-MB-435S cells were methanol fixed, and stained with anti-talin1 antibody followed by Alexa-Fluor IgG1 555-conjugated antibody (red), anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green) and anti- β 5 antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m. **D** Quantification of data presented in (C). Violin plots (number of structures/cell) and scatter plots with median marked in size (size of structures/cell) represents measurements of > 45 cells. Data were analyzed by one-way ANOVA with Dunnett's multiple comparison. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$

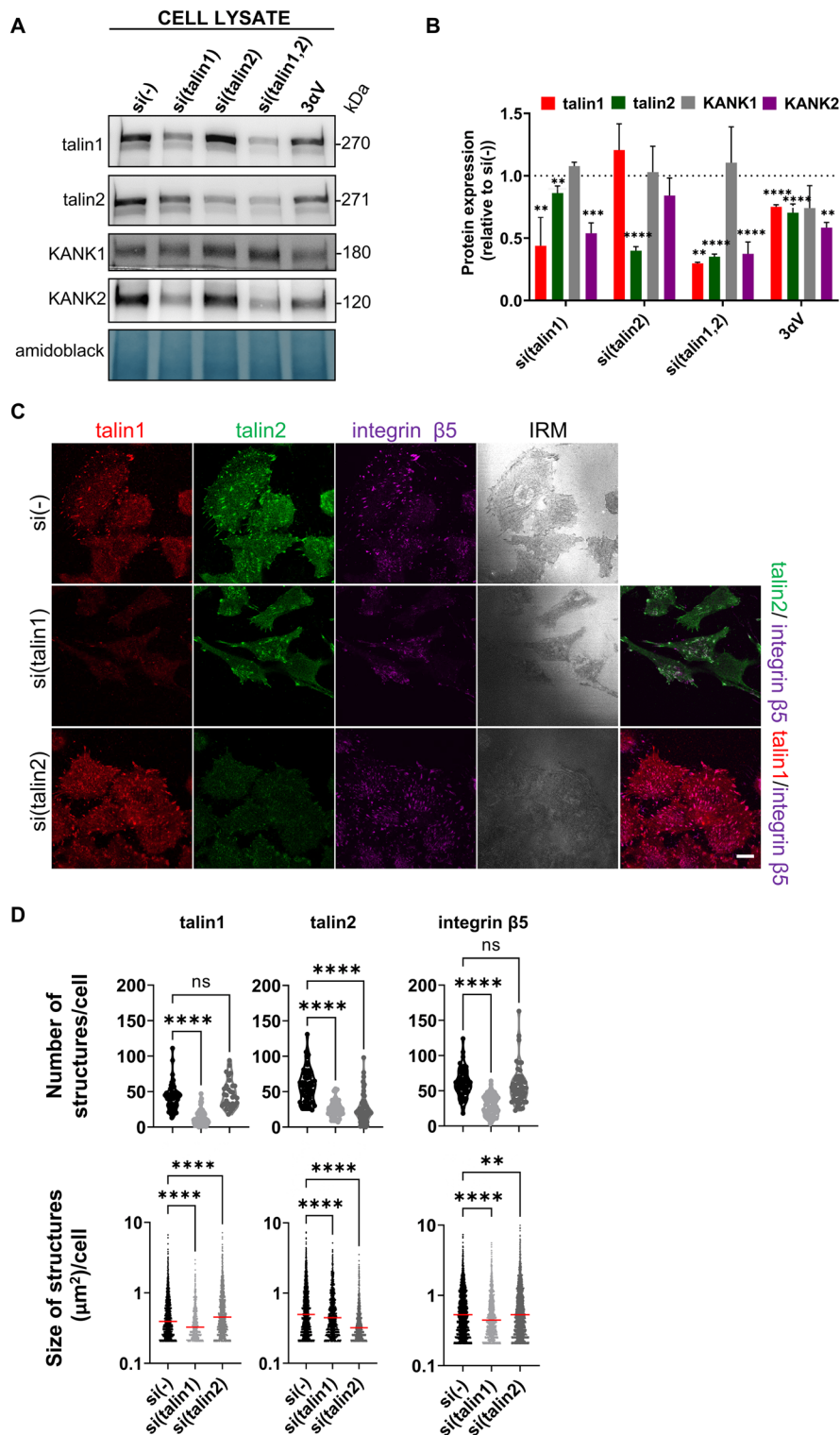


Fig. 2 (See legend on previous page.)

Talin isoform specific regulation of integrin $\alpha\beta 5$ and KANK in isolated IACs

Talins are located in FAs, while KANK2 bridges FAs to the CMSCs [14, 18, 20]. In our experimental system, the expression of CMSC components, including KANK2, depends on

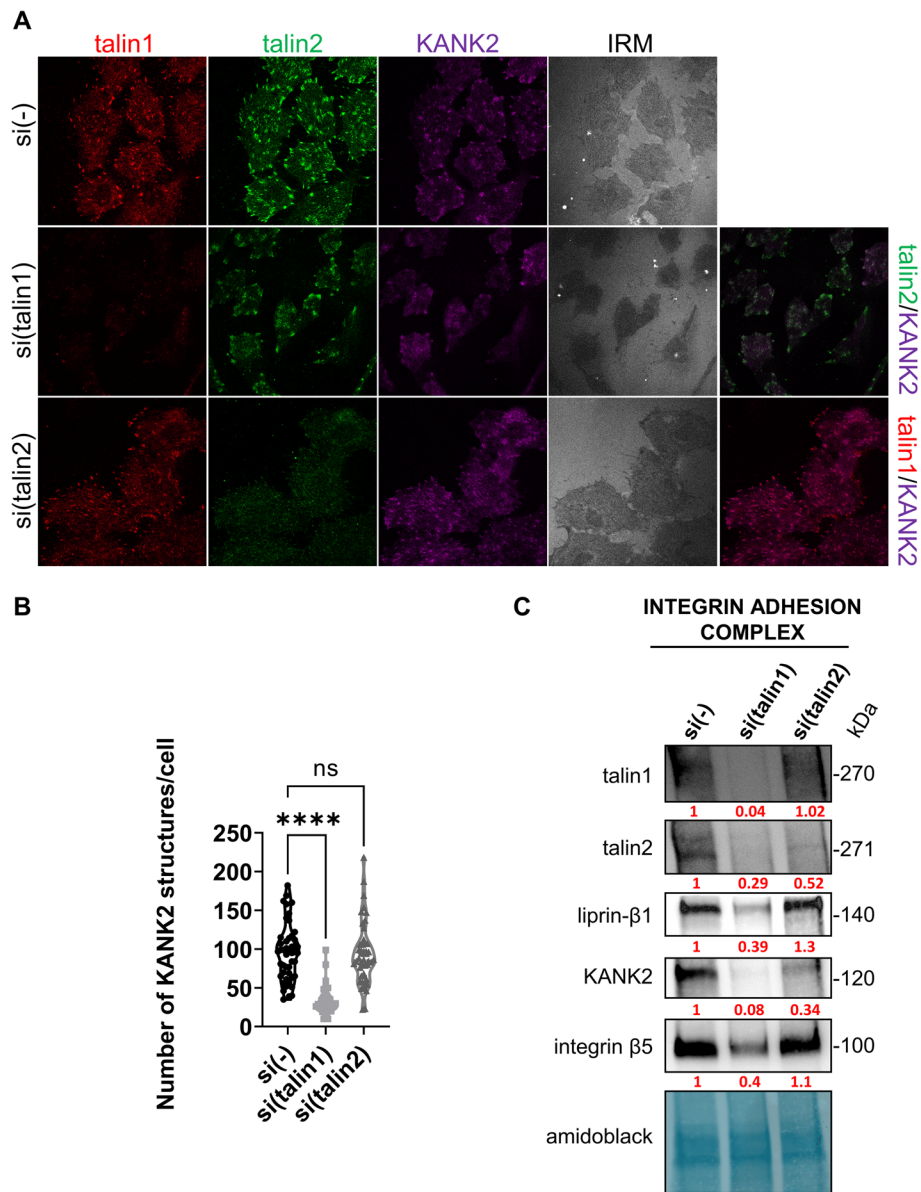


Fig. 3 Talin2 functionally interacts with KANK2. **A** Knockdown of talin1, unlike talin2, leads to changes in KANK2 localization. Forty-eight hours after transfection with either control siRNA, talin1 or talin2-specific siRNA, MDA-MB-435S cells were methanol fixed and stained with anti-talin1 antibody followed by Alexa-Fluor IgG1 555-conjugated antibody (red), anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green), anti-KANK2 antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μm. **B** Quantification of data presented in (A). Violin plot represents measurements of > 30 cells. Data were analyzed by one-way ANOVA with Dunnett’s multiple comparison. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **C** WB analysis of talin1, talin2, liprin-β1, KANK2 or integrin β5 in IACs isolated from MDA-MB-435S cells transfected with either control siRNA, talin1 or talin2-specific siRNA. Forty-eight hours after transfection, IACs were isolated and WB analysis was performed. The results presented are representative of two independent experiments yielding similar results. Numbers below WB scan represent expression of target proteins after talin1 or talin2 knockdown as relative to expression of target proteins in cell transfected with control siRNA set as 1

α V β 5 FAs, which is evident from MS analysis of isolated IACs upon knockdown of integrin α V [24], but also from WB and IF analysis upon talin1 knockdown, which leads to the reduction of KANK2 (Figs. 2A,B, 3A, B; Additional file 1: Fig. S6). Since talin2 knockdown results in an augmentation in FA size (Fig. 2D, second row), we hypothesized that KANK2 might bind talin2 and therefore, after talin2 knockdown, the connection between FAs and CMSC might be partially disrupted. Our attempts to demonstrate a direct link between talin1 or talin2 and KANK2 by coimmunoprecipitation failed. However, it was previously demonstrated, using in vitro fluorescence polarization assay, that both talins (1 and 2) can bind both KANKs (1 and 2) through interaction of KN domain with talin R7 domain [14]. Therefore, we sought to demonstrate a functional interaction between talins and KANK2 within α V β 5 FAs in MDA-MB-435 s cells. To test this, we performed knockdown of either talin1 or talin2, crosslinked and isolated IACs and analyzed the levels of talin1, talin2, integrin β 5, KANK2 and one of the main components of CMSC, liprin- β 1. As expected, knockdown of talin1 reduced the levels of integrin β 5, talin2, KANK2 and liprin- β 1 while knockdown of talin2 did not affect levels of β 5 or talin1 from FAs (Fig. 3C, Additional file 1: Fig. S7), which is in line with our previous data (Fig. 2D). However, despite the fact that upon talin2 knockdown, KANK2 did not change its localization (Fig. 3A) or expression level in total cell lysate (Fig. 2A ,B; Additional file 1: Fig. S6), talin2 knockdown reduced the level of crosslinked KANK2 in samples of isolated IACs (Fig. 3C, Additional file 1: Fig. S7), suggesting that talin2 and KANK2 might be interaction partners. In other words, talin2 knockdown disrupted the link between FA and CMSC. The level of liprin- β 1 did not change upon talin2 knockdown (Fig. 3C, Additional file 1: Fig. S7) indicating that the formation of CMSCs does not depend on talin2. This conclusion is further supported by monitoring the localization of liprin- β 1, upon talin2 or KANK2 knockdown. Knockdown of talin2 did not change the localization of liprin- β 1 (Additional file 1: Fig. S3A). However, knockdown of KANK2 reduced the amount of liprin- β 1 (Additional file 1: Fig. S3B). Therefore, we conclude that the binding of KANK2 to the CMSC is essential for its stabilization and specific localization near FAs.

Talin isoform specific regulation of the actin and MT cytoskeleton

MDA-MB-435S cells contain a high level of stress fibers [37]. KANK2 knockdown in MDA-MB-435S cells does not alter cell size or stress fiber level, but alters MT appearance [24]. We therefore analyzed changes in the actin and MT cytoskeleton upon talin1 or talin2 knockdown using IF. Knockdown of talin1 led to reduced cell spreading area (Additional file 1: Fig. S2) and a significant loss of stress fibers (Fig. 4A). Knockdown of talin2 did not alter cell area (Additional file 1: Fig. S2), but increased the level of stress fibers (Fig. 4A) confirmed by the quantification of confocal images (Fig. 4B). We also observed a complete breakdown of the MT network upon talin1 knockdown (Fig. 4A) which is in line with the disruption of integrin α V β 5 FAs (Fig. 2C). Upon talin2 knockdown, the MT network was still present, but was denser than the control (Fig. 4A) and resembled the MT network upon KANK2 knockdown [24]. We also performed quantitative MT image analysis within 5 μ m from the cell edge and showed a slightly decreased MT content (Fig. 4C). These data provide further evidence that talin2 and KANK2 are potential interaction partners.

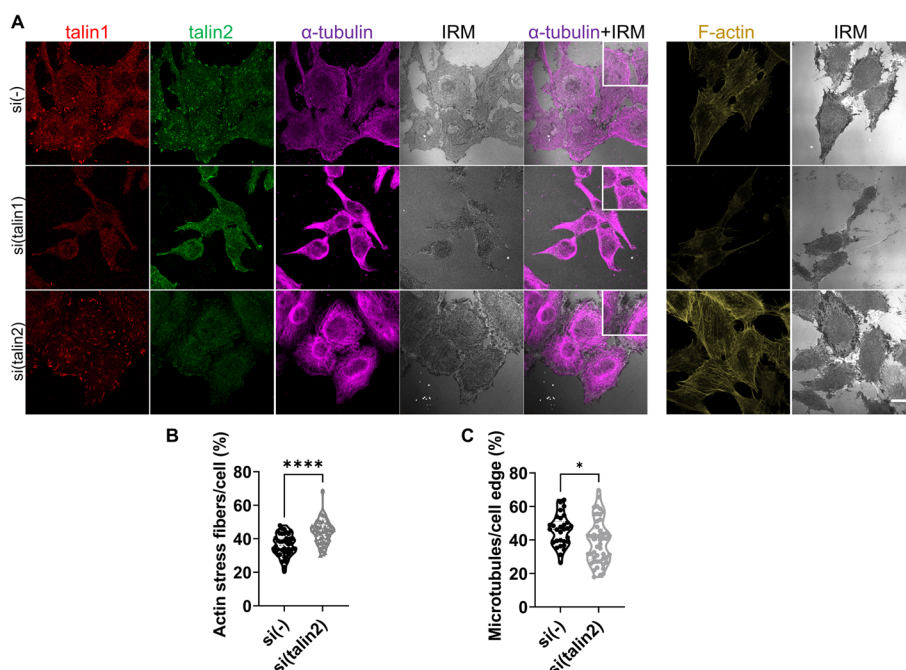


Fig. 4 Talin1 or talin2 knockdown affects actin and MT cytoskeleton. **A** Visualization of α -tubulin and F-actin upon knockdown of talin1 or talin2. Forty-eight hours after transfection with either control, talin1 or talin2-specific siRNA, MDA-MB-435S cells were fixed with methanol (for α -tubulin visualization) or PFA followed by permeabilization with Triton X-100 (for F-actin visualization), and stained with anti-talin1 antibody followed by Alexa-Fluor IgG1 555-conjugated antibody (red), anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green), anti- α -tubulin antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) and IRM images were taken. For F-actin visualization cells were incubated with Alexa-Fluor 488 conjugated phalloidin (shown in gold). Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m. **B, C** Quantification of data presented in (A). Violin plots represents measurements of >30 cells. Data were analyzed by unpaired Student's *t*-test. ns, not significant; **P* < 0.05; ***P* < 0.01; ****P* < 0.001; *****P* < 0.0001

Breaking the link between CMSCs and FAs leads to increased velocity of microtubule growth

As the results above suggest that KANK2 might be an interaction partner of talin2, thus linking the integrin α V β 5 FAs to the CMSCs, we hypothesized that the mechanism of PTX sensitization might be changes in MT dynamics. In order to test this hypothesis, we measured the velocity of MT growth upon knockdown of talins and KANKs. To visualize changes in the velocity of MT growth, we isolated a population of MDA-MB-435S cells and 3 α V cell clone transfected with fluorescent EB3 (435S-EB3 and 3 α V-EB3). Since fluorescent EB3 (EB3-Dendra2) binds to the growing tip of MTs, growing MTs could be followed during time-lapse live cell microscopy [39]. The 435S-EB3 and 3 α V-EB3 cell populations retained their initial characteristics after isolation (Additional file 1: Fig. S4). The 3 α V-EB3 cells expressed lower amounts of talin1, integrin β 5 and KANK2 compared to 435S-EB3 (Additional file 1: Fig. S4A, B) and were more sensitive to PTX compared to 435S-EB3 cells (Additional file 1: Fig. S4C).

Using live cell imaging of fluorescent EB3, we showed that the velocity of MT plus end growth in 3 α V-EB3 was approximately 1.6 times faster than in 435S-EB3 cells (Fig. 5A; Additional file 8: Fig. S8; Additional file 2: Movie S1, Additional file 3: Movie S2). Knockdown of talin1, talin2, KANK2 or integrin β 5 had the same effect, while knockdown

of KANK1 had no effect (Fig. 5B; Additional file 8: Fig. S8; Additional file 2: Movie S1; Additional file 3: Movie S2; Additional file 4: Movie S3; Additional file 5: Movie S4; Additional file 6: Movie S5; Additional file 7: Movie S6). We conclude that knockdown of integrin α V, integrin β 5, talin1, talin2 or KANK2 interferes with either integrin α V β 5 FA formation (integrin α V, integrin β 5 or talin1) or breaks the link between the MT cytoskeleton and integrin α V β 5 FAs (talin2 or KANK2). These results therefore further support the conclusion that talin2-KANK2 interaction is functional in connecting FAs and CMSCs and that its loss leads to changes in MT dynamics.

The increased sensitivity to paclitaxel correlates with increased velocity of microtubule growth

PTX is among the most widely used chemotherapeutic drugs which binds to and stabilizes MTs. PTX causes concentration-dependent inhibition of MT dynamics and perturbation of mitosis [40]. Considering that 435S-EB3 and 3 α V-EB3 cells show different sensitivity to PTX and different velocities of MT growth, we exposed both cells to an equitoxic or equimolar dose of PTX and measured the velocity of MT growth using time-lapse imaging. A decrease in the velocity of MT growth was observed in both 435S-EB3 and 3 α V-EB3 cells with both concentrations of PTX with respect to the nontreated control (Fig. 5C). However, in 3 α V-EB3 cells, which demonstrated a 1.6-fold

(See figure on next page.)

Fig. 5 Disruption of crosstalk between CMSCs and FAs leads to increased velocity of microtubule growth and increased sensitivity to PTX. **A** Knockdown of α V integrin subunit increases velocity of microtubule growth. Quantification of time-lapse live cell microscopy data of 435S-EB3 cells and 3 α V-EB3 cell clone with decreased expression of integrin α V. Violin plot represents measurements of > 450 analyzed microtubules, ($n = 3$) relative to velocity of MT growth in MDA-MB-435S cells that was set as 1. Data were analyzed by unpaired Student's *t*-test. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. Still images of Additional file 2 : Movie S1 and Additional file 3: S2 represent tracking of one microtubule tip in 104 s through five frames. **B** 435S-EB3 cells transfected with either talin1, talin2, KANK2 or β 5-specific siRNA, but not KANK1 siRNA, showed a significant increase in velocity of MT growth compared to cells transfected with control siRNA. Quantification of time-lapse live cell microscopy data. Violin plot represents measurement of > 250 analyzed microtubules, ($n \geq 2$) relative to velocity of MT growth in MDA-MB-435S cells transfected with control siRNA that was set as 1. Data were analyzed by one-way ANOVA with Šidák's multiple comparisons test. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **C** PTX treatment decreases MT growth velocity much stronger in 3 α V-EB3 cells compared to 435S-EB3 cells. Quantification of time-lapse live cell microscopy data of 435S-EB3 cells and 3 α V-EB3 clone upon treatment with equitoxic and equimolar concentrations of PTX. Scatter plot with median marked in red represents measurements of > 200 analyzed microtubules, ($n = 3$) relative to velocity of MT growth in MDA-MB-435S cells that was set as 1. Data were analyzed by one-way ANOVA with Šidák's multiple comparisons test. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **D** Cell proliferation decreases upon knockdown of talin1, but not talin2, KANK1 or KANK2. Cell proliferation was measured using ClickIT Edu assay upon transfection with either control, talin1, talin2, combination of talin1 and talin2, KANK1 or KANK2-specific siRNA. Histogram represents measurements of > 30 cells, plotted as mean \pm SD ($n = 2$). Data were analyzed by one-way ANOVA with Dunnett's multiple comparison. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **E** Talin2 knockdown increases sensitivity to PTX in MDA-MB-435S cells. Twenty-four hours upon transfection, cells were seeded in 96-well plates and 24 h later treated with different concentrations of PTX. Cytotoxicity was measured by MTT assay. Data were analyzed by two-way analysis of variance (ANOVA) with Šidák's multiple comparisons test, with a single pooled variance. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. ($n = 3$). **F** Talin2 knockdown decreases migration in MDA-MB-435S cells. Serum starved (24 h) cells, transfected previously with either control or talin2-specific siRNA were seeded in Transwell cell culture inserts and left to migrate for 22 h toward serum. Cells on the underside of the inserts were stained with crystal violet, photographed, and counted. Scale bar = 100 μ m. **F** Histogram data represents averages of five microscope fields of three independently performed experiments, plotted as mean \pm SD. Data were analyzed by unpaired Student's *t*-test. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$

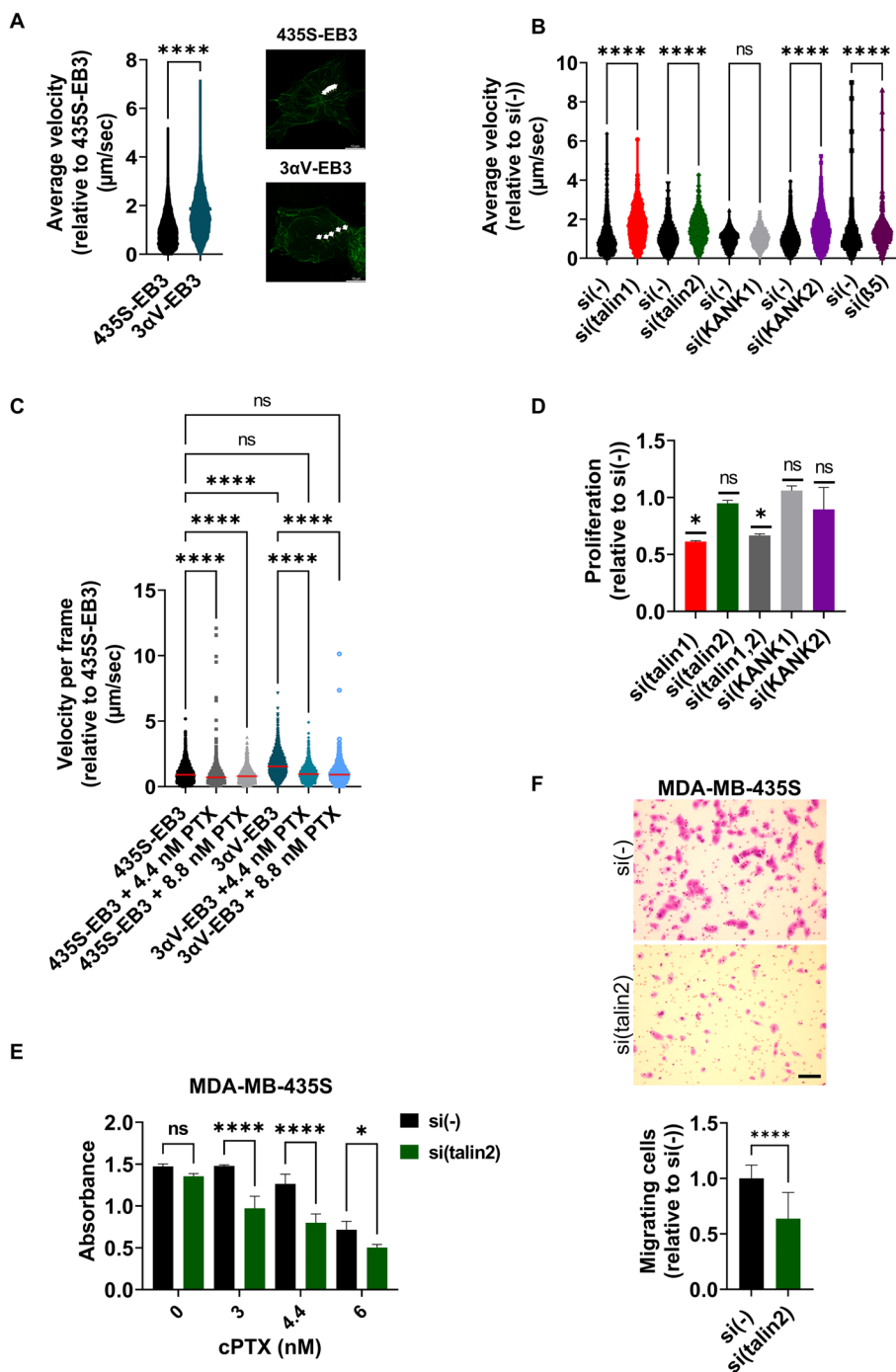


Fig. 5 (See legend on previous page.)

increased velocity compared to 435S-EB3 cells (Fig. 5A), the reduction was much more pronounced (Fig. 5C).

Since we have previously shown that KANK2 knockdown sensitizes cells to PTX [24], and have demonstrated here a functional link between talin2 and KANK2 and their effect on MT dynamics, we tested the effects of talin2 knockdown on sensitivity of MDA-MB-435S cells to PTX by measurement of cell survival following the treatment with PTX. It

has to be emphasized that neither talin2 nor KANK2 knockdown altered cell proliferation, thus indicating that the observed sensitization is the result of altered MT dynamics. Cell proliferation was affected only by talin1 knockdown (Fig. 5D), consistent with the fact that it leads to the destruction of FAs. Talin2 knockdown increased sensitivity to PTX compared to cells transfected with control siRNA (Fig. 5E) and reduced cell migration (Fig. 5F). Since cell migration is a process driven by the cytoskeleton [41] it was not surprising that talin2 knockdown also reduced MDA-MB-435S cell migration as was observed for KANK2 knockdown [24].

In conclusion, the data presented here support a functional interaction between KANK2 and talin2 that maintains the link between FAs and CMSCs, affecting actin-MT crosstalk and regulating sensitivity to MT poison PTX and cell migration.

Discussion

MTs can affect FA dynamics and functions through trafficking and signaling functions, but it is also clear that there is a physical and functional link between MTs and FAs. The importance of this actin-MT crosstalk has been recognized in several physiological processes, survival and migration being some of them [41, 42]. In this paper, we have shown that the loss of interaction between $\alpha V\beta 5$ FAs and MTs leads to increased sensitivity to PTX and that the mechanism by which this occurs is a change in the velocity of MT growth. Our results identify, in melanoma cell line MDA-MB-435S, talin2 from integrin $\alpha V\beta 5$ FAs and KANK2 from CMSCs, as functional partners regulating MT dynamics and sensitivity to PTX. Specifically, stable integrin αV knockdown in 3 αV -EB3 cells, which reduces talin1, talin2 and KANK2 levels, increased the velocity of MT growth and sensitivity to PTX as compared to 435S-EB3 cells. Importantly, upon PTX treatment, a reduction of the velocity of MT growth is much more pronounced in 3 αV -EB3 than in 435S-EB3 cells thus correlating to the sensitivity to PTX. In MDA-MB-435S cells, knockdown of talin2 or KANK2 mimics the effect of integrin αV knockdown, i.e. increased velocity of MT growth and increased sensitivity to PTX, thus providing evidence that KANK2 functionally interacts with talin2.

Because knockdown of talin1 disrupts and effectively removes all links between the proteins within and associated to $\alpha V\beta 5$ FAs, while knockdown of talin2 does not affect the formation or number of $\alpha V\beta 5$ FA, and because, in our hands, co-immunoprecipitation experiments between talins and KANK2 failed, we couldn't conclude which talin physically binds KANK2. However, we show that talin2 and KANK2 functionally interact by analyzing isolated IACs after talin2 knockdown in which the level of cross-linked KANK2 is significantly reduced. It has been shown *in vitro* that talin1 and talin2 can bind KANK2 through interaction with talin R7 domain [14]. However, to our knowledge, direct link between endogenous talins and KANK2 was not shown using co-immunoprecipitation. The localization of KANK1 [14] or KANK2 [24] within CMSC has been demonstrated but the cause of these differences is not known. The functional interaction of talin2 and KANK2 is very likely cell type specific affected by unknown factor/s that should be investigated in the future.

The knockdown of integrin αV [37], KANK2 [24] and talin2 (this work) not only increased sensitivity to PTX but also decreased cell migration. Similarly, in HeLa cells, the loss of KANK1 reduced cell motility [14] while in cultured mouse podocytes

depletion of both, KANK1 or KANK2, resulted in reduced migration [43]. Conversely, in mouse fibroblasts KANK2 does not localize within CMSC but to the lateral border of FAs (FA belt) where it activates talin, reduces force transduction across integrins and induces central adhesion formation. This adhesion sliding correlates with reduced cell migration speed i.e. KANK2 depleted mouse fibroblast cells showed higher migration velocities [20]. Therefore, KANK proteins play a dual role in the cell: they can play a role in MT targeting [14] or in force transmission, but these KANK2 complexes function separately [20]. We can assume that KANK localization and interaction partners potentially determine the way it will affect cell migration.

We showed in our MDA-MB-435S cells that KANK1 is not a part of the CMSC associated with integrin $\alpha V\beta 5$ FAs because KANK1 level does not change after knockdown of either talin1, talin2 or integrin αV , nor does KANK1 knockdown affect cell sensitivity to PTX or change the velocity of MT growth. KANK1 function and potential interaction partners should be further investigated.

Currently little is known about the isoform dependence of talin-KANK binding. In HeLa cells KANK1 binds to talin1 and controls cortical organization of CMSC components [14] while in the fibrosarcoma cell line HT1080 KANK1 or KANK2 knockdown mimicked the effects of nocodazole-induced MT depolymerization i.e. uncoupling MTs from adhesion complexes [44]. KANK2 in mouse fibroblasts is present in mature integrin $\alpha 5\beta 1$ FAs where it regulates force transmission in a CMSC-independent manner [20].

Talin1 and talin2 mRNAs display different tissue distributions and a reciprocal relationship between their expression [45]. Even though the isoforms share 76% protein sequence their role in cells is not fully redundant [46–48] and the mechanism of interplay of the isoform expression is still not known [26]. Our results support cell-specific and $\alpha V\beta 5$ integrin-specific ratio of talin1 and talin2 in FAs. We have previously shown that integrin $\alpha V\beta 5$ FAs, used preferentially for adhesion in MDA-MB-435S cells, contain both talins [24]. Interestingly, using the same cell line, Qi and colleagues [46] showed that talin1 is required for small FA formation, whereas talin2 is responsible for larger FA assembly. However, they plated cells on fibronectin and therefore $\alpha V\beta 5$ may not have been the primary receptor used by cells. In several cell types talin1 was concentrated in peripheral FAs while talin2 was observed in both FAs and fibrillar adhesions [49]. In our case both talin isoforms, 1 and 2, were present in $\alpha V\beta 5$ FAs. We showed that talin1 has a role in formation of $\alpha V\beta 5$ FAs since knockdown of talin1, unlike talin2, led to the disruption of $\alpha V\beta 5$ FAs. It would follow that this disruption would result in a decrease of other proteins within the FAs. Since we have previously shown that talin2 and KANK2 are also a part of the integrin $\alpha V\beta 5$ FAs [24], it was to be expected that knockdown of talin1 would lead to a decrease in the level of talin2 and KANK2. Besides, that means that talin2 alone is not sufficient for successful integrin $\alpha V\beta 5$ FA formation, which is consistent with the fact that knockout of talin1, but not talin2, is lethal in mice [27, 28]. This conclusion is supported by the fact that in our MDA-MB-435S cells the proliferation was strongly affected by talin1 knockdown (down to 60%) but not upon talin2 knockdown (down to 95%). It should be noted that talin1 cannot replace the role of talin2 in maintaining actin-MT crosstalk, because after talin2 knockdown in IAC isolates KANK2 levels are reduced and the velocity of MT growth is increased, thus

indicating uncoupling of MTs from FAs. Together, these results confirm that talins are not functionally redundant.

In our MDA-MB-435S cells the size of talin1 and $\beta 5$ -positive structures upon talin2 knockdown increased, without affecting the number of $\alpha V\beta 5$ FAs, indicating the change in FA dynamics. The knockdown of the other functional partner involved in actin MT-crosstalk, KANK2, also slightly increased talin2-positive structures (Additional file 1: Fig. S5) thus supporting a talin2-KANK2 functional linkage. In addition, we observed an increased amount of stress fibers upon talin2 knockdown which is not the consequence of talin1 replacing talin2, since WB analysis of talin1 in either total cell lysates or IACs showed that its level did not change upon talin2 knockdown. Conversely, KANK2 knockdown did not alter the amount of stress fibers [24]. The absence of changes in amount of stress fibers upon KANK2 knockdown still does not exclude functional involvement of KANK2 in actin cytoskeleton dynamics. Indeed, it has been shown in the human fibrosarcoma cell line HT1080 that uncoupling of FAs and CMSCs, by depleting KANKs or disrupting MT stability, leads to dispersion of CMSCs and augmentation of FAs. The release of the RhoA GEF ARHGEF2/GEF-H1 from MTs resulted in increased RhoA-mediated actomyosin contractility, reinforcement of the stress fiber-associated FAs and decreased FA turnover [44].

KANK proteins are generally localized to the periphery of FAs [14, 20] and complex stability measurements combined with cell biological experiments suggested that shear-force stretching promotes KANK1 localization to the periphery of FAs [21]. Since talin2 knockdown did not affect the formation and number of $\alpha V\beta 5$ FAs we assume that in our cell model FAs are formed preferentially through involvement of talin1 while talin2 is recruited to the outer rim of FAs later, enabling the link to KANK2 protein from CMSC and establishing actin-MT crosstalk. The talins are mechanotransducers due to their ability to undergo force dependent structural rearrangements [26]. It has been shown in vitro that cyclin-dependent kinase-1 (CDK1) regulates IACs through binding within the R8 domain of talin1 and phosphorylation at the end of R7 rod domain leading to significant reduction in binding of KANK to R7 [50]. We hypothesize that in our cell model, talin1, shown to be engaged in IAC formation, is likely phosphorylated by CDK1 and therefore has a low KANK binding affinity. Instead, talin2, which we have shown is not essential for FA formation, is the one sterically accessible and maintains an affinity to KANK2.

We have shown that talin2 knockdown, which disconnects FAs from CMSCs, does not affect either KANK2 or liprin- $\beta 1$ localization. Nevertheless, upon KANK2 knockdown, the liprin- $\beta 1$ signal was dispersed and almost lost. Indeed, the clustering of CMSC components depends on interactions of its components, primarily KANK and liprins $\alpha 1$ and $\beta 1$ and possibly other interactions [15, 18, 19, 51]. Our results are in line with results obtained by Bouchet and colleagues [14] in HeLa cell model in which the localization of CMSC clusters depends on the KANK1-liprin- $\beta 1$ connection.

Integrin expression is frequently disturbed in tumor cells and therefore integrins represent attractive therapeutic targets for arresting tumor growth, reducing resistance to chemo- or radiotherapy, or attenuating invasion and metastasis [52]. We have previously identified, in melanoma MDA-MB-435S cells, integrin αV as a potential therapeutic target whose knockdown increased sensitivity to PTX and reduced metastasis [24, 37]. However, targeted integrin therapy has proven to be ineffective in the clinic [52, 53]. We

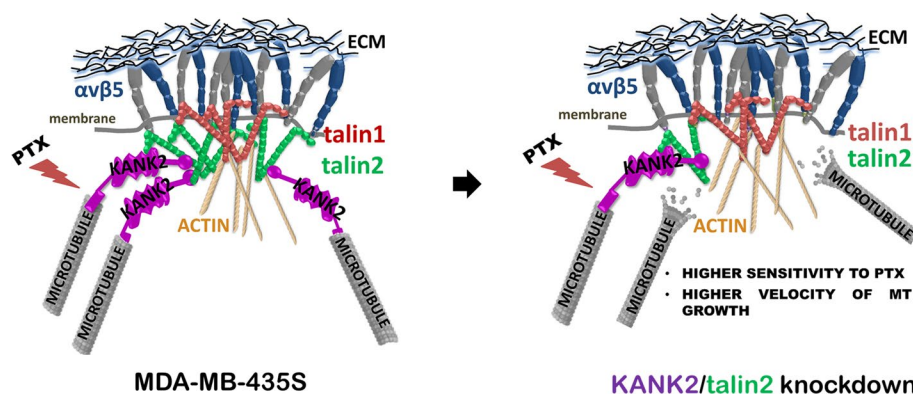


Fig. 6 Schematic representation of the functional link between talin2 from $\alpha V\beta 5$ FAs and KANK2 from CMSC. Disruption of the link, either by KANK2 or talin2 knockdown, results in increased sensitivity to PTX and decreased migration of MDA-MB-435S cells

therefore suggest here, that besides KANK2, which we already previously identified as a protein linked to integrin $\alpha V\beta 5$ -mediated sensitivity to PTX [24], talin2 might also be a potential therapeutic target because its knockdown improves PTX therapy and reduces metastasis, and may have a more uniform response than targeting integrins. Disruption of a functional link between talin2 and KANK2 isoforms, either by KANK2 or talin2 knockdown, alters MT dynamics, i.e., increases the velocity of MT growth, thereby increasing sensitivity to PTX and decreasing migration. Indeed, few data from the literature confirm that talin2 is a potential target for tumor therapy because of its role in cell migration, invasion and metastasis. It has previously been shown that talin2 binds much more strongly to the $\beta 1$ -integrin tail than talin1 and that interaction of talin2 with integrins is required to generate traction force, which in turn drives invadopodium-mediated matrix degradation [46, 54]. In the breast cancer cell line MDA-MB-231, the loss-of-function of talin1 has been shown to increase chemosensitivity to docetaxel [55] while talin2 knockdown inhibited growth, migratory capacity and invasiveness and promoted apoptosis [54, 56]. Both talin1 and talin2 correlate with the malignancy potential of the human hepatocellular carcinoma MHCC-97 L cells [57]. Conversely, talin2 is downregulated in clear cell renal cell carcinoma (ccRCC) tissues and cells, and its overexpression in ccRCC cell lines inhibits cell growth and metastasis [58]. It remains to be discovered which factors influence the role of talin2 in tumors.

In conclusion, we show in melanoma cell line MDA-MB-435S the different subcellular localizations of KANK1 and KANK2, different roles of talin1 and talin2, and functional link between talin2 and KANK2 affecting MT dynamics, sensitivity to PTX and cell migration (Fig. 6). The different subcellular localizations of KANK isoforms 1 and 2 likely are caused by yet unknown interaction partners which should be further investigated.

Abbreviations

ECM	Extracellular matrix
IAC	Integrin adhesion complex
FA	Focal adhesion
RA	Reticular adhesion
MT	Microtubule
CMSC	Cortical microtubule stabilizing complex

PMAP	Plasma membrane-associated platform
ELKS	Proteins rich in amino acids E, L, K and S
KIF21A	Kinesin family member KIF21A
CLASP	CLIP-associating protein
KANK	KN motif and ankyrin repeat domain
MS	Mass spectrometry
KN	KANK N-terminal
CC	Coiled-coil
ANKRD	Ankyrin repeat domain
PTX	Paclitaxel
EB3	End-binding protein 3
SDS-PAGE	Sodium dodecyl sulphate–polyacrylamide–gel electrophoresis
WB	Western blotting
HEPES	Hydroxyethylpiperazine ethanesulfonic acid
DTBP	Dimethyl 3,3'-dithiobispropionimidate, Wang and Richard's reagent
MTT	3-((4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide
IRM	Interference reflection microscopy
IF	Immunofluorescence analysis
CDK1	Cyclin-dependent kinase-1
ANOVA	One-way analysis of variance
ccRCC	Clear cell renal cell carcinoma

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s11658-023-00473-6>.

Additional file 1: Fig. S1. KANK1 localization is not affected by talin1 or talin2 knockdown. (A, B) Talin2, but not talin1, knockdown slightly affects KANK1 appearance. Forty-eight hours after transfection with either control, talin1 or talin2-specific siRNA, MDA-MB-435S cells were methanol fixed and stained with anti-talin1 or anti-talin2 antibody followed by Alexa-Fluor 546-conjugated antibody (red) or Alexa-Fluor 488-conjugated antibody (green), respectively. KANK1 was further visualized by anti-KANK1 antibody followed by Alexa-Fluor 555-conjugated antibody or Alexa-Fluor 488-conjugated antibody (shown in magenta). Finally, vinculin was visualized using conjugated anti-vinculin Alexa Fluor 647 antibody (shown in grey) and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m. (C) KANK1 knockdown does not affect sensitivity to PTX. Sensitivity of cells transfected with either control or KANK1-specific siRNA to PTX was measured by MTT assay. Twenty-four hours upon transfection, cells were seeded in 96-well plates and 24 h later treated with different concentrations of PTX. Data were analyzed by two-way analysis of variance (ANOVA) with Šidák's multiple comparisons test, with a single pooled variance; ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **Fig. S2.** Cell area decreases upon knockdown of talin1, but not talin2. Quantification of data presented in Fig. 2. Violin plot represents measurements of > 30 cells ($n = 2$). Data were analyzed by one-way ANOVA with Dunnett's multiple comparison. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **Fig. S3.** The CMSC protein liprin- β 1 loses its organization upon KANK2, but not upon talin2 knockdown. (A, B) Forty-eight hours after transfection with talin2 or KANK2-specific siRNA, MDA-MB-435S cells were fixed with methanol and stained with anti-liprin- β 1 antibody followed by Alexa-Fluor 546-conjugated antibody, anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green) and anti-KANK2 antibody followed by Alexa-Fluor 647-conjugated antibody and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m. **Fig. S4.** Verification of the MDA-MB-435S and 3aV cell model expressing fluorescently labelled EB3. (A, B) Clone 3aV-EB3 shows decreased expression of talin1, integrin β 5 and KANK2 as compared to MDA-MB-435S-EB3 cells. Forty-eight hours after seeding cells were methanol fixed and stained with anti-talin1 followed by Alexa-Fluor 546-conjugated antibody (red), anti-KANK2 antibody or anti- β 5 antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) and IRM images were taken. Analysis was performed using TCS SP8 Leica. Scale bar = 10 μ m. (C) Clone 3aV-EB3 demonstrates increased sensitivity to PTX as compared to parental MDA-MB-435S-EB3 cells. Twenty-four hours upon seeding in 96-well plates cells were treated with different concentrations of PTX. Cytotoxicity was measured by MTT assay. Data were analyzed by two-way analysis of variance (ANOVA) with Šidák's multiple comparisons test, with a single pooled variance; ns denotes not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$ ($n = 3$). **Fig. S5.** Knockdown of KANK2 increases talin2-positive FAs size. (A) Forty-eight hours after transfection with either control or KANK2-specific siRNA, MDA-MB-435S cells were methanol fixed and stained with anti-talin2 antibody followed by Alexa-Fluor IgG2b 488-conjugated antibody (green) and anti-KANK2 antibody followed by Alexa-Fluor 647-conjugated antibody (magenta) IRM images were taken. (B) Quantification of data presented in (a). Scatter plot with median marked in red represents measurements of ≥ 30 cells, ($n = 2$). Data were analyzed by unpaired Student's t -test. ns, not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; **** $P < 0.0001$. **Fig. S6.** Full images of the blots in Fig. 2A. Images were obtained using Uvitec Alliance Q9 mini, which directly scanned membranes developed with ECL reagents. **Fig. S7.** Full images of the blots in Fig. 3C. Images were obtained using Uvitec Alliance Q9 mini, which directly scanned membranes developed with ECL reagents.

Additional file 2: Movie S1. Time-lapse live cell microscopy of MDA-MB-435S cells with fluorescent EB3 (435S-EB3 cells) used for measurement of velocity of MT growth.

Additional file 3: Movie S2. Time-lapse live cell microscopy of clone 3aV cells with fluorescent EB3 (3aV-EB3 cells) used for measurement of velocity of MT growth.

Additional file 4: Movie S3. Time-lapse live cell microscopy of MDA-MB-435S cells with fluorescent EB3 (435S-EB3 cells) transfected with control siRNA used for measurement of velocity of MT growth.

Additional file 5: Movie S4. Time-lapse live cell microscopy of MDA-MB-435S cells with fluorescent EB3 (435S-EB3 cells) transfected with talin1-specific siRNA used for measurement of velocity of MT growth.

Additional file 6: Movie S5. Time-lapse live cell microscopy of MDA-MB-435S cells with fluorescent EB3 (435S-EB3 cells) transfected with talin2-specific siRNA used for measurement of velocity of MT growth.

Additional file 7: Movie S6. Time-lapse live cell microscopy of MDA-MB-435S cells with fluorescent EB3 (435S-EB3 cells) transfected with KANK2-specific siRNA used for measurement of velocity of MT growth.

Additional file 8: Fig. S8. Still images of Additional file 2–Additional file 8: Movie S1–S6. Images were obtained using Image J manual tracking tool. Each arrow represent position of one microtubule tip through 104 s. Images were captured every 26 s.

Additional file 9: Table S1. List of used antibodies and dyes.

Acknowledgements

We thank Marina Sutalo of the Laboratory for Cell Biology and Signalling for technical assistance.

Author contributions

Material preparation, data collection and analysis were performed by [ML], [NS], [ARJ], [KK], [DM], and [AA]. The first draft of the manuscript was written by [AA] and [ML], corrections were made by [NS], [DM], [JDH] and [MJH]. All authors read and approved the final manuscript. [AA], [ML] and [NS] contributed to the study conception and design.

Funding

This work was supported by the Croatian Science Foundation Project (Grant No IP-2019-04-1577 to AA-R) and Cancer Research United Kingdom (Grant C13329/A21671 to MH). ML received research fellowship from British Scholarship Trust (UK Charity 1031428).

Availability of data and materials

All relevant data could be obtained from the corresponding author.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors have no relevant financial or non-financial interests to disclose.

Received: 31 March 2023 Accepted: 27 June 2023

Published online: 17 July 2023

References

1. Hynes RO. Integrins: bidirectional, allosteric signaling machines. *Cell*. 2002;110(6):673–87.
2. Green HJ, Brown NH. Integrin intracellular machinery in action. *Exp Cell Res*. 2019;378(2):226–31.
3. Burridge K, Guilluy C. Focal adhesions, stress fibers and mechanical tension. *Exp Cell Res*. 2016;343(1):14–20.
4. Walko G, Castanon MJ, Wiche G. Molecular architecture and function of the hemidesmosome. *Cell Tissue Res*. 2015;360(3):529–44.
5. Lock JG, Jones MC, Askari JA, Gong XW, Oddone A, Olofsson H, et al. Reticular adhesions are a distinct class of cell-matrix adhesions that mediate attachment during mitosis. *Nat Cell Biol*. 2018;20(11):1290–302.
6. Geiger B, Yamada KM. Molecular architecture and function of matrix adhesions. *Cold Spring Harb Perspect Biol*. 2011;3(5):a005033.
7. Winograd-Katz SE, Fassler R, Geiger B, Legate KR. The integrin adhesome: from genes and proteins to human disease. *Nat Rev Mol Cell Biol*. 2014;15(4):273–88.
8. Wolfenson H, Lavelin I, Geiger B. Dynamic regulation of the structure and functions of integrin adhesions. *Dev Cell*. 2013;24(5):447–58.
9. Zaidel-Bar R, Itzkovitz S, Ma'ayan A, Iyengar R, Geiger B. Functional atlas of the integrin adhesome. *Nat Cell Biol*. 2007;9(8):858–67.
10. Zhou X, Li J, Kucik DF. The microtubule cytoskeleton participates in control of beta2 integrin avidity. *J Biol Chem*. 2001;276(48):44762–9.
11. Ng DH, Humphries JD, Byron A, Millon-Fremillon A, Humphries MJ. Microtubule-dependent modulation of adhesion complex composition. *PLoS ONE*. 2014;9(12):e115213.
12. Ezratty EJ, Partridge MA, Gundersen GG. Microtubule-induced focal adhesion disassembly is mediated by dynamin and focal adhesion kinase. *Nat Cell Biol*. 2005;7(6):581–90.

13. Ezratty EJ, Bertaux C, Marcantonio EE, Gundersen GG. Clathrin mediates integrin endocytosis for focal adhesion disassembly in migrating cells. *J Cell Biol.* 2009;187(5):733–47.
14. Bouchet BP, Gough RE, Ammon YC, van de Willige D, Post H, Jacquemet G, et al. Talin-KANK1 interaction controls the recruitment of cortical microtubule stabilizing complexes to focal adhesions. *Elife.* 2016;5: e18124.
15. Astro V, Chiaretti S, Magistrati E, Fivaz M, de Curtis I. Liprin-alpha1, ERC1 and LL5 define polarized and dynamic structures that are implicated in cell migration. *J Cell Sci.* 2014;127(17):3862–76.
16. Astro V, Tonoli D, Chiaretti S, Badanai S, Sala K, Zerial M, et al. Liprin-alpha1 and ERC1 control cell edge dynamics by promoting focal adhesion turnover. *Sci Rep.* 2016;6:33653.
17. Hida Y, Ohtsuka T. CAST and ELKS proteins: structural and functional determinants of the presynaptic active zone. *J Biochem.* 2010;148(2):131–7.
18. van der Vaart B, van Riel WE, Doodhi H, Kevenaar JT, Katrukha EA, Gumy L, et al. CFEOM1-associated kinesin KIF21A is a cortical microtubule growth inhibitor. *Dev Cell.* 2013;27(2):145–60.
19. Lansbergen G, Grigoriev I, Mimori-Kiyosue Y, Ohtsuka T, Higa S, Kitajima I, et al. CLASPs attach microtubule plus ends to the cell cortex through a complex with LL5beta. *Dev Cell.* 2006;11(1):21–32.
20. Sun Z, Tseng HY, Tan S, Senger F, Kurzawa L, Dedden D, et al. Kank2 activates talin, reduces force transduction across integrins and induces central adhesion formation. *Nat Cell Biol.* 2016;18(9):941–53.
21. Yu M, Le SM, Ammon YC, Goult BT, Akhmanova A, Yan J. Force-dependent regulation of talin-KANK1 complex at focal adhesions. *Nano Lett.* 2019;19(9):5982–90.
22. Tadjan A, Samarzija I, Humphries JD, Humphries MJ, Ambriovic-Ristov A. KANK family proteins in cancer. *Int J Biochem Cell Biol.* 2021;131: 105903.
23. Byron A, Askari JA, Humphries JD, Jacquemet G, Koper EJ, Warwood S, et al. A proteomic approach reveals integrin activation state-dependent control of microtubule cortical targeting. *Nat Commun.* 2015;6:6135.
24. Paradzik M, Humphries JD, Stojanovic N, Nestic D, Majhen D, Dekanic A, et al. KANK2 links alphaVbeta5 focal adhesions to microtubules and regulates sensitivity to microtubule poisons and cell migration. *Front Cell Dev Biol.* 2020;8:125.
25. Klapholz B, Brown NH. Talin—the master of integrin adhesions. *J Cell Sci.* 2017;130(15):2435–46.
26. Gough RE, Goult BT. The tale of two talins—two isoforms to fine-tune integrin signalling. *FEBS Lett.* 2018;592(12):2108–25.
27. Monkley SJ, Zhou XH, Kinston SJ, Giblett SM, Hemmings L, Priddle H, et al. Disruption of the talin gene arrests mouse development at the gastrulation stage. *Dev Dyn.* 2000;219(4):560–74.
28. Debrand E, Conti FJ, Bate N, Spence L, Mazzeo D, Pritchard CA, et al. Mice carrying a complete deletion of the talin2 coding sequence are viable and fertile. *Biochem Biophys Res Commun.* 2012;426(2):190–5.
29. Chen NP, Sun Z, Fassler R. The Kank family proteins in adhesion dynamics. *Curr Opin Cell Biol.* 2018;54:130–6.
30. Pan W, Sun K, Tang K, Xiao Q, Ma C, Yu C, et al. Structural insights into ankyrin repeat-mediated recognition of the kinesin motor protein KIF21A by KANK1, a scaffold protein in focal adhesion. *J Biol Chem.* 2018;293(6):1944–56.
31. Weng Z, Shang Y, Yao D, Zhu J, Zhang R. Structural analyses of key features in the KANK1.KIF21A complex yield mechanistic insights into the cross-talk between microtubules and the cell cortex. *J Biol Chem.* 2018;293(1):215–25.
32. Chastney MR, Lawless C, Humphries JD, Warwood S, Jones MC, Knight D, et al. Topological features of integrin adhesion complexes revealed by multiplexed proximity biotinylation. *J Cell Biol.* 2020;219(8).
33. Robertson J, Jacquemet G, Byron A, Jones MC, Warwood S, Selley JN, et al. Defining the phospho-adesome through the phosphoproteomic analysis of integrin signalling. *Nat Commun.* 2015;6:6265.
34. Horton ER, Byron A, Askari JA, Ng DHJ, Millon-Fremillon A, Robertson J, et al. Definition of a consensus integrin adhesion and its dynamics during adhesion complex assembly and disassembly. *Nat Cell Biol.* 2015;17(12):1577–87.
35. Tadjan A, Humphries JD, Samarzija I, Stojanovic N, Zha J, Culjak K, et al. The tongue squamous carcinoma cell line Cal27 primarily employs integrin alpha6beta4-containing type II hemidesmosomes for adhesion which contribute to anticancer drug sensitivity. *Front Cell Dev Biol.* 2021;9: 786758.
36. Humphries JD, Zha J, Burns J, Askari JA, Below CR, Chastney MR, et al. Pancreatic ductal adenocarcinoma cells employ integrin alpha6beta4 to form hemidesmosomes and regulate cell proliferation. *Matrix Biol.* 2022;110:16–39.
37. Stojanovic N, Dekanic A, Paradzik M, Majhen D, Ferencak K, Ruscic J, et al. Differential effects of integrin alphaV knockdown and cilengitide on sensitization of triple-negative breast cancer and melanoma cells to microtubule poisons. *Mol Pharmacol.* 2018;94(6):1334–51.
38. Jones MC, Humphries JD, Byron A, Millon-Fremillon A, Robertson J, Paul NR, et al. Isolation of integrin-based adhesion complexes. *Curr Protoc Cell Biol.* 2015;66:9.8.1–9.8.15.
39. Stepanova T, Slemmer J, Hoogenraad CC, Lansbergen G, Dortland B, De Zeeuw CI, et al. Visualization of microtubule growth in cultured neurons via the use of EB3-GFP (end-binding protein 3-green fluorescent protein). *J Neurosci.* 2003;23(7):2655–64.
40. Jordan MA, Wilson L. Microtubules as a target for anticancer drugs. *Nat Rev Cancer.* 2004;4(4):253–65.
41. Seetharaman S, Etienne-Manneville S. Cytoskeletal crosstalk in cell migration. *Trends Cell Biol.* 2020;30(9):720–35.
42. Seetharaman S, Etienne-Manneville S. Microtubules at focal adhesions—a double-edged sword. *J Cell Sci.* 2019;132(19): jcs232843.
43. Gee HY, Zhang FJ, Ashraf S, Kohl S, Sadowski CE, Vega-Warner V, et al. KANK deficiency leads to podocyte dysfunction and nephrotic syndrome. *J Clin Invest.* 2015;125(6):2375–84.
44. Rafiq NBM, Nishimura Y, Plotnikov SV, Thiagarajan V, Zhang Z, Shi S, et al. Publisher Correction: a mechano-signalling network linking microtubules, myosin IIA filaments and integrin-based adhesions. *Nat Mater.* 2019;18(6):638–49.
45. Senetar MA, McCann RO. Gene duplication and functional divergence during evolution of the cytoskeletal linker protein talin. *Gene.* 2005;362:141–52.
46. Qi L, Jafari N, Li X, Chen Z, Li L, Hytonen VP, et al. Talin2-mediated traction force drives matrix degradation and cell invasion. *J Cell Sci.* 2016;129(19):3661–74.
47. Zhang X, Jiang G, Cai Y, Monkley SJ, Critchley DR, Sheetz MP. Talin depletion reveals independence of initial cell spreading from integrin activation and traction. *Nat Cell Biol.* 2008;10(9):1062–8.

48. Manso AM, Okada H, Sakamoto FM, Moreno E, Monkley SJ, Li R, et al. Loss of mouse cardiomyocyte talin-1 and talin-2 leads to beta-1 integrin reduction, costameric instability, and dilated cardiomyopathy. *Proc Natl Acad Sci U S A*. 2017;114(30):E6250–9.
49. Praekelt U, Kopp PM, Rehm K, Linder S, Bate N, Patel B, et al. New isoform-specific monoclonal antibodies reveal different sub-cellular localisations for talin1 and talin2. *Eur J Cell Biol*. 2012;91(3):180–91.
50. Gough RE, Jones MC, Zacharchenko T, Le S, Yu M, Jacquemet G, et al. Talin mechanosensitivity is modulated by a direct interaction with cyclin-dependent kinase-1. *J Biol Chem*. 2021;297(1): 100837.
51. Astro V, de Curtis I. Plasma membrane-associated platforms: dynamic scaffolds that organize membrane-associated events. *Sci Signal*. 2015;8(367):re1.
52. Bergonzini C, Kroese K, Zweemer AJM, Danen EHJ. Targeting integrins for cancer therapy—disappointments and opportunities. *Front Cell Dev Biol*. 2022;10: 863850.
53. Samarzija I, Dekanic A, Humphries JD, Paradzik M, Stojanovic N, Humphries MJ, et al. Integrin crosstalk contributes to the complexity of signalling and unpredictable cancer cell fates. *Cancers (Basel)*. 2020;12(7):1910.
54. Li L, Li X, Qi L, Rychahou P, Jafari N, Huang C. The role of talin2 in breast cancer tumorigenesis and metastasis. *Oncotarget*. 2017;8(63):106876–87.
55. Singel SM, Cornelius C, Batten K, Fasciani G, Wright WE, Lum L, Shay JW. A targeted RNAi screen of the breast cancer genome identifies KIF14 and TLN1 as genes that modulate docetaxel chemosensitivity in triple-negative breast cancer. *Clin Cancer Res*. 2013;19(8):2061–70.
56. Liang Y, Chen H, Ji L, Du J, Xie X, Li X, Lou Y. Talin2 regulates breast cancer cell migration and invasion by apoptosis. *Oncol Lett*. 2018;16(1):285–93.
57. Fang KP, Dai W, Ren YH, Xu YC, Zhang SM, Qian YB. Both Talin-1 and Talin-2 correlate with malignancy potential of the human hepatocellular carcinoma MHCC-97 L cell. *BMC Cancer*. 2016;16:45.
58. Cai J, Huang Z, Zhou J, Wu W, Ye Y. TLN2 functions as a tumor suppressor in clear cell renal cell carcinoma via inactivation of the Wnt/ β -catenin signaling pathway. *Transl Androl Urol*. 2022;11(1):39–52.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

