



Small molecules inhibiting the nuclear localization of YAP/TAZ for chemotherapeutics and chemosensitizers against breast cancers



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ABSTRACT

YAP and TAZ oncoproteins confer malignancy and drug resistance to various cancer types. We screened for small molecules that inhibit the nuclear localization of YAP/TAZ. Dasatinib, statins and pazopanib inhibited the nuclear localization and target gene expression of YAP and TAZ. All three drugs induced phosphorylation of YAP and TAZ, and pazopanib induced proteasomal degradation of YAP/TAZ. The sensitivities to these drugs are correlated with dependence on YAP/TAZ in breast cancer cell lines. Combinations of these compounds with each other or with other anti-cancer drugs efficiently reduced cell proliferation of YAP/TAZ-dependent breast cancer cells. These results suggest that these drugs can be therapeutics and chemosensitizers for YAP/TAZ-dependent breast cancers.

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1. Introduction

YAP and TAZ are transcriptional co-activators involved in tissue growth and stem cell maintenance in normal tissue through binding to TEADs. They are regulated through phosphorylation by the Hippo pathway, leading to inhibition of the nuclear translocation and the proteasomal degradation [1]. The Hippo pathway is activated by cellular density [2–4], soluble factors via GPCR [5,6], and actin cytoskeleton organization [7,8]. YAP and TAZ activations are also implicated in the tumorigenesis and malignancy of various cancers including breast [9], colon [10], lung [11], liver [12], and mesothelioma [13]. TAZ is associated with the maintenance of breast cancer stem cells and drug resistance [14]. Therefore, YAP and TAZ play causative roles in carcinogenesis and cancer progression, and inactivation of YAP and TAZ by small molecules is a promising strategy for therapeutics of various cancers with their activation [15].

Abbreviations: YAP, yes-associated protein; TAZ, transcriptional coactivator with PDZ-binding motif; TEAD, TEA domain family member; GPCR, G-protein coupled receptor; FDA, food and drug administration; NF2, neurofibromin 2; HMG-CoA, hydroxymethylglutaryl-coenzyme A; CTGF, connective tissue growth factor

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In drug discovery, one of the successful strategies is the exploitation of established drugs that have already been approved for treatment of other cancers or non-cancerous diseases (i.e., drug repositioning, drug repurposing, or indication switch). The major advantage of this approach is that the pharmacokinetic, pharmacodynamics and toxicity profiles of these drugs are well known, making their rapid shifts to clinical trials possible [16,17].

In this study, we screened for small molecules which inactivate YAP and TAZ from drugs with known targets for the drug repositioning against breast cancer. We found that dasatinib, statins, and pazopanib inhibited their nuclear localization and TEAD-dependent transcription, and induced YAP/TAZ phosphorylation. Pazopanib induced proteasomal degradation of YAP and TAZ. Furthermore, we explored the possibility of chemotherapy with them for breast cancer, and found that the sensitivities to these compounds are correlated with the dependence on YAP and TAZ in these cell lines. Combination of the YAP/TAZ inhibitors or of those with anti-cancer agents efficiently suppressed the breast cancer cell growth. Our findings thus opened the window for the application of dasatinib, statins, and pazopanib, clinically used drugs, for breast cancers with activation of YAP and TAZ.

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2. Materials and methods

2.1. Cell culture and treatments

MDA-MB-231, MDA-MB-453, HBC-4, HBC-5, MCF-7, BSY-1, ZR-75-1, and SKBR-3 breast cancer cell lines were maintained in RPMI-1640 containing 10% FBS and penicillin/streptomycin. HEK293 was maintained in Dulbecco's modified Eagle medium containing 10% FBS and penicillin/streptomycin. Dasatinib was purchased from JS Research Chemicals Trading Co. Fluvastatin, doxorubicin, and paclitaxel were purchased from Wako Pure Chemicals. Geranylgeranyl diphosphate (GGPP), farnesyl diphosphate (FPP), GGTI-286, and FTI-276 were purchased from Sigma. MG-132 was purchased from Merck Millipore, and pazopanib from ChemieTek.

2.2. Antibodies

For immunofluorescence, 1/200 rabbit anti-YAP antibody (H-125, Santa Cruz), 1/400 rabbit anti-YAP/TAZ antibody (D24E4, Cell Signaling), 1/200 phalloidin-AlexaFluor 594 (Life Technologies), and 1/1000 anti-rabbit IgG-AlexaFluor 488 conjugate (Life Technologies) were used. For immunoblot, 1/3000 rabbit anti-YAP antibody (H-125, Santa Cruz), 1/3000 rabbit anti-TAZ antibody (#2149, Cell Signaling), 1/3000 rabbit anti-YAP/TAZ antibody (D24E4, Cell Signaling), 1/10,000 mouse anti-GAPDH antibody (6C5, Millipore), 1/5000 anti-mouse IgG-HRP (GE Healthcare), and 1/5000 anti-rabbit IgG-HRP (GE Healthcare), were used. Antibodies for Western blot were diluted in Can Get Signal reagents (Toyobo). Western blot using standard SDS-PAGE gel or gels containing Phos-tag-acrylamide (SuperSep Phos-tag, Wako) was performed as previously described [18].

2.3. Screening of the inhibitors inhibiting nuclear localization of YAP

MDA-MB-231 cells (10,000–15,000 cells) were inoculated in a nuclear imaging plate (Corning) and 24 h later, 10 μ M chemicals in a SCADS inhibitor kit (provided by the Screening Committee of Anticancer Drugs, Japan) was added and incubated for 6 h. Cells were fixed with 4% paraformaldehyde and immunostained with anti-YAP antibody as described below.

2.4. Immunofluorescence and imaging

Paraformaldehyde-fixed cells were permeabilized with 0.3% TritonX-100 in PBS and blocked with 3% FBS in PBST for 30 min. They were then incubated with anti-YAP antibody at 4 °C overnight and washed with PBS three times. Cells were incubated with anti-rabbit IgG-Alexa Fluor 488 for 1 h at room temperature, and washed with PBS three times. For confocal microscopy, cells were mounted in Prolong Gold reagent containing 10 μ g/ml Hoechst 33342 (Life Technologies). When appropriate, cells were stained with phalloidin-Alexa Fluor 594 (Life Technologies) prior to mounting. Images were obtained with an FV1000-D confocal microscope equipped with a 40 \times objective lens using FV10-ASW software (Olympus). For screening of small molecules, PBS containing 10 μ g/ml Hoechst 33342 was added and images were obtained by IN Cell Analyzer 2000 (GE Healthcare) using a 40 \times objective lens.

2.5. Reporter assay

8xGTIIIC-luciferase was obtained from Addgene (Addgene #34615). MDA-MB-231 cells were transfected with 1.4 μ g of 8xGTIIIC-luciferase plasmid and 50 ng of pRL-CMV (Promega) by using lipofectamine 2000 (Life Technologies) or Viafect

transfection reagent (Promega) as previously described [7]. After 24 h, cells were treated with chemicals for 18 h. Luciferase activities were measured using the Dual-Luciferase Reporter Assay System (Promega) according to the manufacturer's instructions.

2.6. Quantitative real time PCR

Total RNA was prepared with ISOGEN (Nippon Gene) and cDNA was synthesized with ReverTra Ace qPCR RT master mix with gDNA remover (TOYOBO). Real time PCR was performed with QuantiTect SYBR Green PCR Master Mix (Qiagen) using the Eco Real Time PCR system (Illumina). The sequences of PCR primers for CTGF and GAPDH were as follows; CTGF-F: AGGAGTGGGTGTGTGACGA, CTGF-R: CCAGGCAGTTGGCTCTAATC, GAPDH-F: AGCCACATCGCTC-AGACAC, GAPDH-R: GCCCAATACGACCAAATCC.

2.7. RNAi

siRNA transfection was performed using Lipofectamine RNAiMAX (Life Technologies) according to the manufacturer's instructions. The siRNA sequence is listed as followed; siYAP: GACAUCUUCUGGUCAGAGAUU, siTAZ: ACGUUGACUUAGG-AACUUUUU [14].

2.8. MTT assay

MTT assay was performed as previously described [18]. 3000–10,000 cells suspended in RPMI-1640 containing 1% FBS were seeded on 96 well plates. Fifteen μ l of medium containing drugs was added, and cells were incubated for 4 days.

2.9. Colony formation assay

MDA-MB-231 or MCF-7 cells (1000–2000 cells per well) were seeded on 24 well plates and treated with inhibitors for 10 days. Cells were fixed with 4% formaldehyde and stained with 0.5% crystal violet.

3. Results

3.1. Dasatinib, fluvastatin, and pazopanib inhibit the function of YAP/TAZ transcriptional co-activator

To identify drugs targeting YAP and TAZ, we performed image-based screening for small molecules which inhibit their nuclear localization using MDA-MB-231 breast cancer cell line. MDA-MB-231 harbors homozygous mutation in NF2, which positively regulates the Hippo pathway; therefore, YAP and TAZ are constitutively activated in this cell line [19]. We screened about 400 chemicals with known targets from SCADS inhibitor kit (provided by the Screening Committee of Anticancer Drugs, Japan) which consists of classical anti-cancer agents, kinase inhibitors, metabolic pathway inhibitors, and signaling pathway inhibitors, including FDA-approved drugs [20]. MDA-MB-231 cells were treated with chemicals and nuclear localization of YAP was evaluated by immunofluorescence. Chemicals which induced nuclear exclusion of YAP were treated as positive. We found that thiazovivin, dasatinib, lovastatin, cucurbitacin I, and pazopanib inhibited the nuclear localization of YAP (Fig. 1A). Among them, dasatinib, statins, and pazopanib are approved as clinically used drugs, and we therefore analyzed them further. We found that dasatinib, fluvastatin, and pazopanib inhibited nuclear localization of YAP and TAZ in the nanomolar to micromolar range, although the effect of pazopanib on the nuclear localization of YAP/TAZ was relatively weak compared to the other two drugs (Fig. 1B). They also

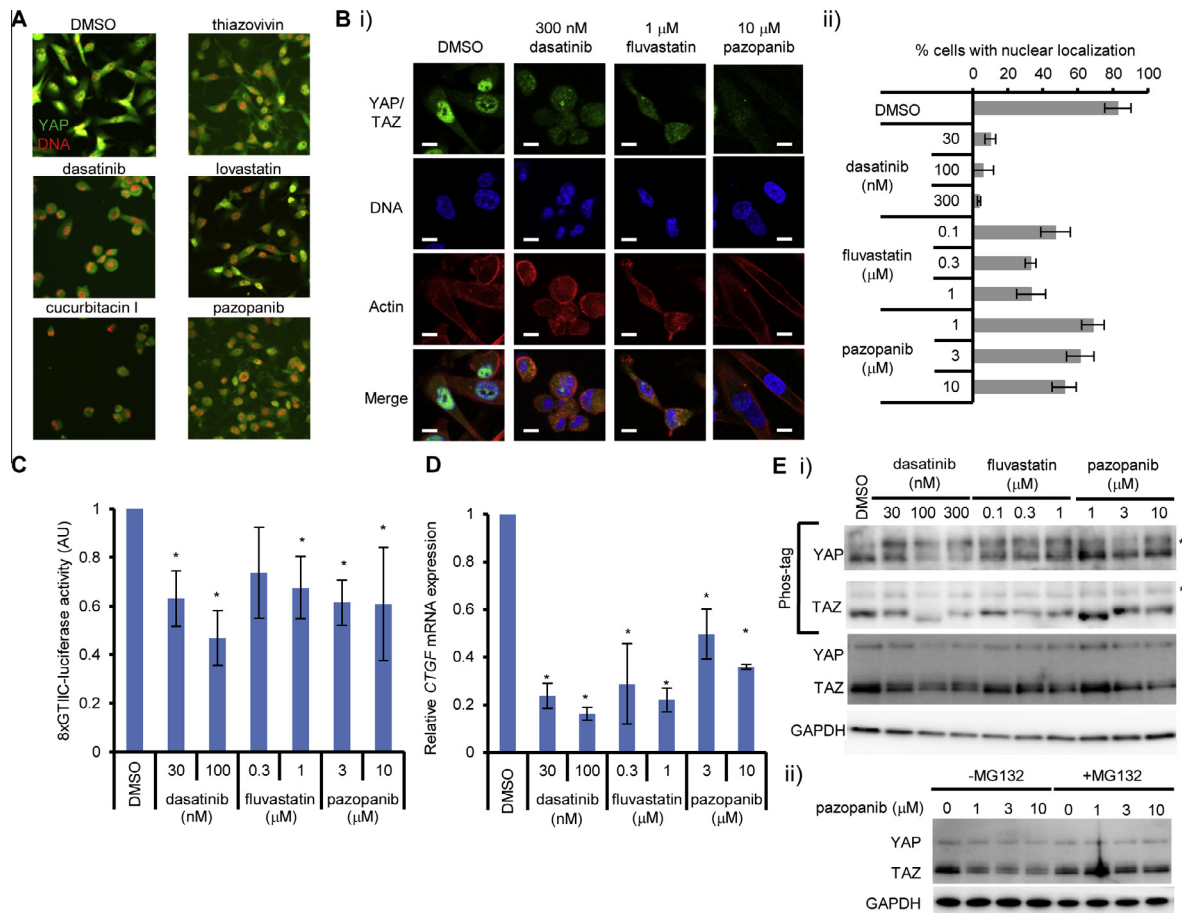


Fig. 1. Dasatinib, fluvastatin, and pazopanib inhibit the function of YAP/TAZ transcriptional co-activator. (A) Identification of small molecule agents which inhibit the nuclear localization of YAP. MDA-MB-231 cells were treated with a chemical library for 8 h. YAP was immunostained. Red: DNA, green: YAP. Each agent was used at 10 μM. (B) Dasatinib, fluvastatin, and pazopanib inhibited the nuclear localization of YAP and TAZ. MDA-MB-231 cells were treated with inhibitors for 8 h and YAP and TAZ were visualized by immunofluorescence. (i) Representative images of immunofluorescence. Bar represents 10 μm. (ii) Cells with nuclear localization of YAP and TAZ were counted. At least 75 cells were counted for each sample. Data represents mean and standard deviation from three independent experiments. (C) Inhibition of TEAD-dependent promoter activity by dasatinib, fluvastatin, and pazopanib. MDA-MB-231 cells were transfected with 8xGT11C-luciferase TEAD reporter plasmid with pRL-CMV control plasmid, and treated with inhibitors overnight. Relative luciferase activity (8xGT11C-luciferase/*Renilla* luciferase) was measured. Data represents mean and standard deviation from at least three independent experiments. * $P < 0.02$. (D) Inhibition of endogenous target genes of YAP and TAZ by dasatinib, fluvastatin, and pazopanib. MDA-MB-231 cells were treated with inhibitor for 18 h and total RNA was prepared. *CTGF* mRNA was quantified by quantitative RT-PCR. Data represents mean and standard deviation from at least three independent experiments. * $P < 0.02$. (E) Dasatinib, fluvastatin, and pazopanib activate the Hippo pathway. (i) Induction of phosphorylation of YAP and TAZ by dasatinib, fluvastatin and pazopanib. MDA-MB-231 cells were treated with inhibitors for 8 h. Whole cell extract was analyzed by SDS-polyacrylamide gel containing Phos-tag acrylamide. Asterisks show phosphorylated YAP and TAZ. Data is representative of at least three independent experiments. (ii) Degradation of YAP and TAZ by pazopanib is proteasome-dependent. MDA-MB-231 cells were treated with pazopanib for 8 h in the presence or absence of 10 μM MG132. Whole cell extract was analyzed by Western blot. Data is representative of at least three independent experiments.

inhibited the YAP/TAZ-TEAD-dependent reporter activity (Fig. 1C). We found that they also reduced the transcripts of *CTGF*, whose transcription is dependent on YAP/TAZ (Fig. 1D). Dasatinib and fluvastatin are known to change actin dynamics, and this leads to phosphorylation of YAP/TAZ [19]. Dasatinib inhibits SRC and affects YAP nuclear localization [21]. Statins inhibit HMG-CoA reductase and lead to impaired geranylgeranylation of RHOA, resulting in the inactivation of YAP/TAZ as reported recently [22,23]. We reproduced the geranylgeranyl-dependent inactivation of YAP/TAZ by fluvastatin, as the simultaneous addition of geranylgeranyl diphosphate (GGPP), but not farnesyl diphosphate (FPP), with fluvastatin canceled the effect of fluvastatin (Supplementary Fig. S1). We also found that the direct inhibition of protein geranylgeranylation recapitulated phosphorylation of YAP (Supplementary Fig. S1). These results confirmed the RHOA-dependent inactivation of YAP/TAZ by statins [22,23].

Pazopanib inhibits VEGFR and PDGFR signaling. VEGF signaling induces activation of RHOA in cervical cancer cells as well as vascular endothelial cells [24,25]. Inhibition of VEGF signaling is known to impair the activation of SRC and FAK, leading to impaired activation

of RHOA. PDGF signaling also induced activation of RHOA [26]. MDA-MB-231 is known to express VEGFR2, a VEGFR family member [27] and PDGFR [28]. We hypothesized that the phosphorylation of YAP/TAZ accounted for the inhibition of YAP/TAZ nuclear localization by pazopanib. Therefore, we examined whether pazopanib induced the phosphorylation of YAP and TAZ, and found the increase in the ratio of phosphorylated YAP and TAZ by this drug by pazopanib similar to that by dasatinib and fluvastatin (Fig. 1E i), suggesting that pazopanib also induces YAP/TAZ phosphorylation. We further noted that treatment of cells with pazopanib reduced the total amount of YAP/TAZ in the cell. This reduction was canceled by a proteasome inhibitor, MG-132, (Fig. 1E ii) suggesting that pazopanib facilitates degradation of YAP/TAZ by the ubiquitin-proteasome system as previously described [29].

3.2. YAP/TAZ-dependent breast cancer cell lines are sensitive to dasatinib, fluvastatin, and pazopanib

YAP and TAZ activation is frequently observed in various cancers and is associated with resistance to anti-cancer agents,

invasiveness, stemness, and poor prognosis [9,14,30–32]. We hypothesized that dasatinib, statins, and pazopanib are effective for YAP/TAZ-dependent breast cancer cells and therefore addressed whether they suppressed the breast cancer cell growth. The response to these drugs was different among cell lines, and we found that MDA-MB-231 was most sensitive to these inhibitors. On the other hand, some cell lines (MCF-7, for example), and HEK293 (human embryonic kidney cell), were resistant to them (Fig. 2A). Examining the level of YAP/TAZ-dependent transcription in these cell lines, we found that MDA-MB-231 showed the highest YAP/TAZ-dependent transcriptional activity among eight breast cancer cell lines (Fig. 2B), which is consistent with the *NF2* mutation in this cell line [19]. MDA-MB-231 expressed higher YAP and TAZ, and phosphorylation of YAP and TAZ was lower than that in other cell lines (Fig. 2C). We hypothesized that the activation and dependence of YAP and TAZ is one determinant of the effectiveness of these YAP/TAZ inhibitors and examined the colony formation assay using cells treated with YAP and TAZ siRNA. Colony formation was dramatically reduced when both YAP and TAZ were depleted in MDA-MB-231 (Fig. 2D). Therefore, cell growth of MDA-MB-231 is YAP/TAZ-dependent. On the other hand, some cell lines which are resistant to the agents, including MCF-7, are also resistant to silencing of YAP and TAZ by siRNA (Fig. 2D), suggesting

MCF-7 is a YAP/TAZ-independent cell line. In summary, we concluded that there was a correlation between the dependence of YAP/TAZ in breast cancers and the sensitivities to dasatinib, fluvastatin, and pazopanib.

3.3. Combination of dasatinib, fluvastatin and pazopanib efficiently reduced the viability of MDA-MB-231

Although dasatinib, fluvastatin, and pazopanib can inactivate YAP and TAZ in breast cancers, their doses were relatively high compared to clinically relevant ones. Higher blood concentration of these drugs can be a risk factor for adverse effects. Reduced concentrations of them to cancer cells are relevant in terms of drug safety. Their use in combination is a feasible approach for this. We first combined the dasatinib, fluvastatin, and pazopanib for MDA-MB-231 cell line, and found that their combinations efficiently reduced viability of the cells (Fig. 3A). Although emergence of the colonies was not largely affected, the colony sizes were efficiently reduced by the combination of these inhibitors (Fig. 3B). Then again, single agents or their combinations did not reduce colony formation of MCF-7, YAP/TAZ-independent cell line (Fig. 3B). These results suggested that combinations of these agents are one of the efficient strategies against YAP/TAZ-dependent cell lines.

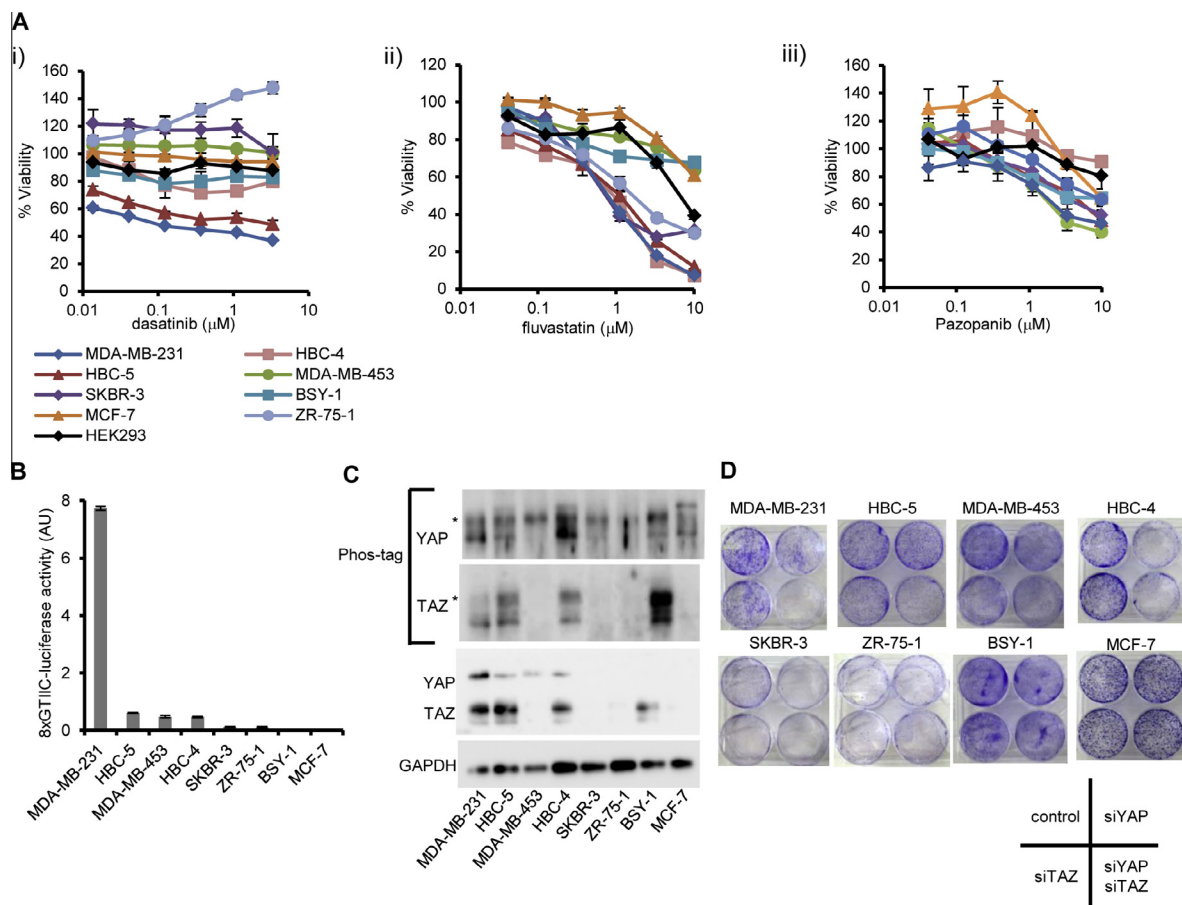


Fig. 2. YAP/TAZ-dependent breast cancer cell lines are sensitive to dasatinib, fluvastatin, and pazopanib. (A) Sensitivity of breast cancer cell lines to (i) dasatinib, (ii) fluvastatin, and (iii) pazopanib. Cells were treated with drugs for 4 days. Viability was measured by MTT assay. Data represents mean and standard deviation from triplicate. Data is representative of at least three independent experiments. HEK293 (black line) was used as normal cell control. (B) Activation states of YAP/TAZ-dependent transcription in breast cancer cell lines. Cells were transfected with 8xGT1C-luciferase and pRL-CMV vector. Promoter activity was measured 2 days after transfection. Data represents mean and standard deviation from triplicate. Data is representative of at least two independent experiments. (C) Expression of YAP and TAZ in breast cancer cell lines. Asterisks show phosphorylated YAP and TAZ. (D) Dependence of cell growth on YAP and TAZ. Cells were transfected with siRNA for YAP and TAZ and were diluted in 6 well plates. Colony formation was measured by crystal violet. Data is representative of at least two independent experiments.

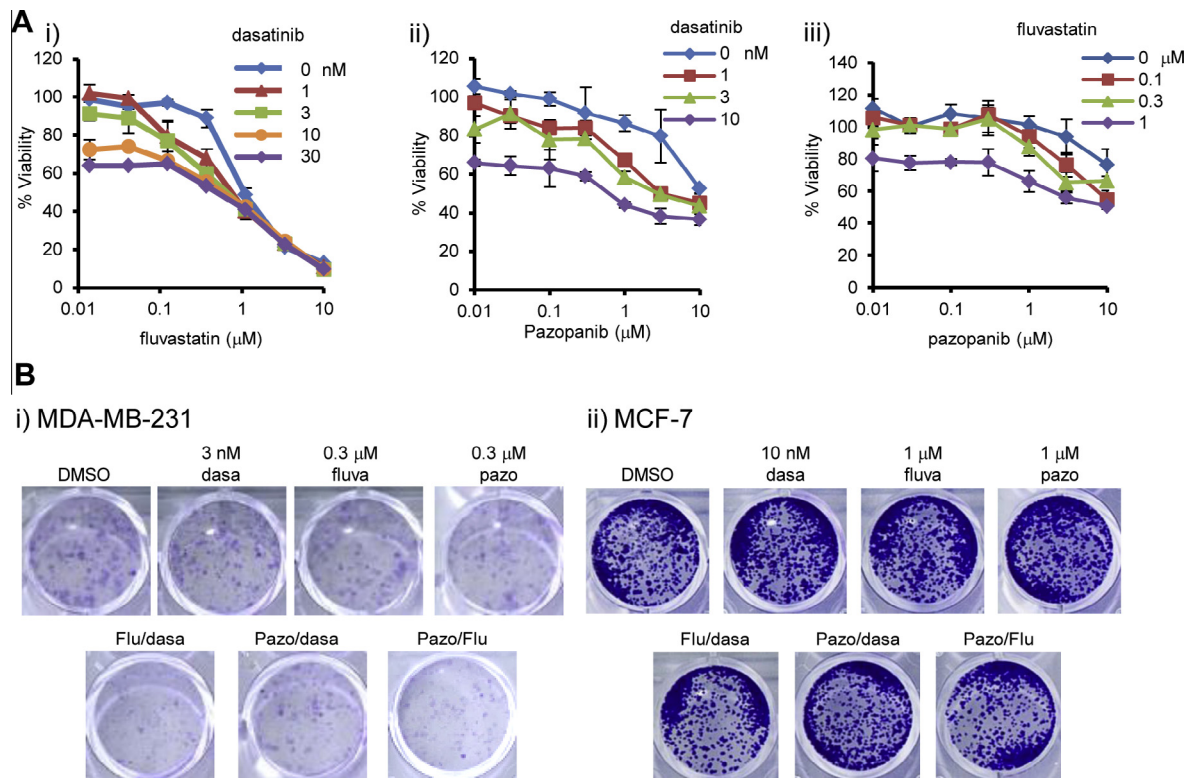


Fig. 3. Combination of dasatinib, fluvastatin and pazopanib efficiently reduced the viability of MDA-MB-231. (A) Combination therapy using dasatinib, fluvastatin, and pazopanib for MDA-MB-231. (i) Fluvastatin in combination with dasatinib. (ii) Pazopanib with dasatinib. (iii) Pazopanib with fluvastatin. Cells were treated with inhibitors for 4 days and viability was measured by MTT assay. Data represents mean and standard deviation of triplicate. Data is a representative of at least three independent experiments. (B) Colony formation reduced by the combination of dasatinib, fluvastatin, and pazopanib in MDA-MB-231. (i) MDA-MB-231 and (ii) MCF-7 cells were treated with the combination of inhibitors (dasa: dasatinib, fluva: fluvastatin, pazo: pazopanib) for 7 or 10 days. Colony formation was determined by staining with crystal violet. Data is representative of three independent experiments.

3.4. Dasatinib, fluvastatin, and pazopanib sensitize YAP/TAZ-dependent cancer cells to doxorubicin and paclitaxel

TAZ confers the resistance to paclitaxel and doxorubicin, and its depletion leads to sensitization of cells to these anti-cancer drugs in breast cancer cells [9,32]. Therefore, we hypothesized that dasatinib, fluvastatin, and pazopanib can act as chemosensitizers for YAP/TAZ-dependent breast cancer cells. MDA-MB-231 was then treated with paclitaxel or doxorubicin together with dasatinib, fluvastatin, or pazopanib. We found that some combination of YAP/TAZ inhibitors with these anti-cancer agents synergistically reduced cell viability (Fig. 4A). Colony formation was also significantly reduced by the combination therapies (Fig. 4B).

Chemosensitizing effects were not consistently observed when MCF-7, whose cell growth is YAP/TAZ-independent, was treated with these combinations (Fig. 4C). Colony formation of this cell line was also not affected by the combination of agents (Fig. 4D). These results suggested that dasatinib, fluvastatin, and pazopanib can be chemosensitizers for YAP/TAZ-dependent breast cancer cells.

4. Discussion

YAP and TAZ are emerging targets for breast cancer treatment. In this study we identified the clinically used drugs, dasatinib, statins, and pazopanib as small molecule agents which inhibit the nuclear localization of YAP/TAZ transcriptional co-activators. These agents are effective for YAP/TAZ-dependent breast cancer;

their combination increased the efficacy. Furthermore, they can also act as chemosensitizers of doxorubicin and paclitaxel. We proposed them as candidate chemotherapeutic agents for YAP/TAZ-dependent breast cancers.

Thiazovivin, cucurbitacin I, dasatinib, fluvastatin, and pazopanib inhibit nuclear localization of YAP/TAZ, and activate the Hippo pathway. Thiazovivin is a RHO-kinase inhibitor and modulates actin cytoskeletal dynamics. Inhibition of RHO-kinase is known to inhibit nuclear localization of YAP/TAZ [19], indicating the validity of our screening. Cucurbitacin I is a plant poison and indirectly affects actin dynamics [33]. Changes in the cytoskeletal dynamics can activate YAP/TAZ as previously reported [7]. Among them, we focused on three FDA-approved drugs. Dasatinib, a SRC inhibitor, changes actin dynamics, leading to decrease in nuclear localization of YAP and TAZ [21]. Recent papers have shown that statins inhibit nuclear localization of YAP/TAZ via reduction of geranylgeranylation of RHO family GTPases, followed by phosphorylation [22,23]. In addition to them, we found that pazopanib, a multikinase inhibitor, inhibited the nuclear localization of YAP/TAZ, leading to reduced transcription. Pazopanib also induced phosphorylation of YAP, similar to that by dasatinib or fluvastatin (Fig. 1E). Pazopanib is known as a multikinase inhibitor, and mainly targets VEGFR and PDGFR. Previous studies revealed that VEGFR2 and PDGFR are expressed in MDA-MB-231 [27,34–36], and that their signalings are involved in migration and cell growth [34,37]. In vascular endothelial cells, VEGFR signaling affects actin filament dynamics through the RHO/ROCK pathway in vascular endothelial cells [38]. Thus, inhibition of VEGFR2 in MDA-MB-231 may influence cytoskeletal rearrangements,

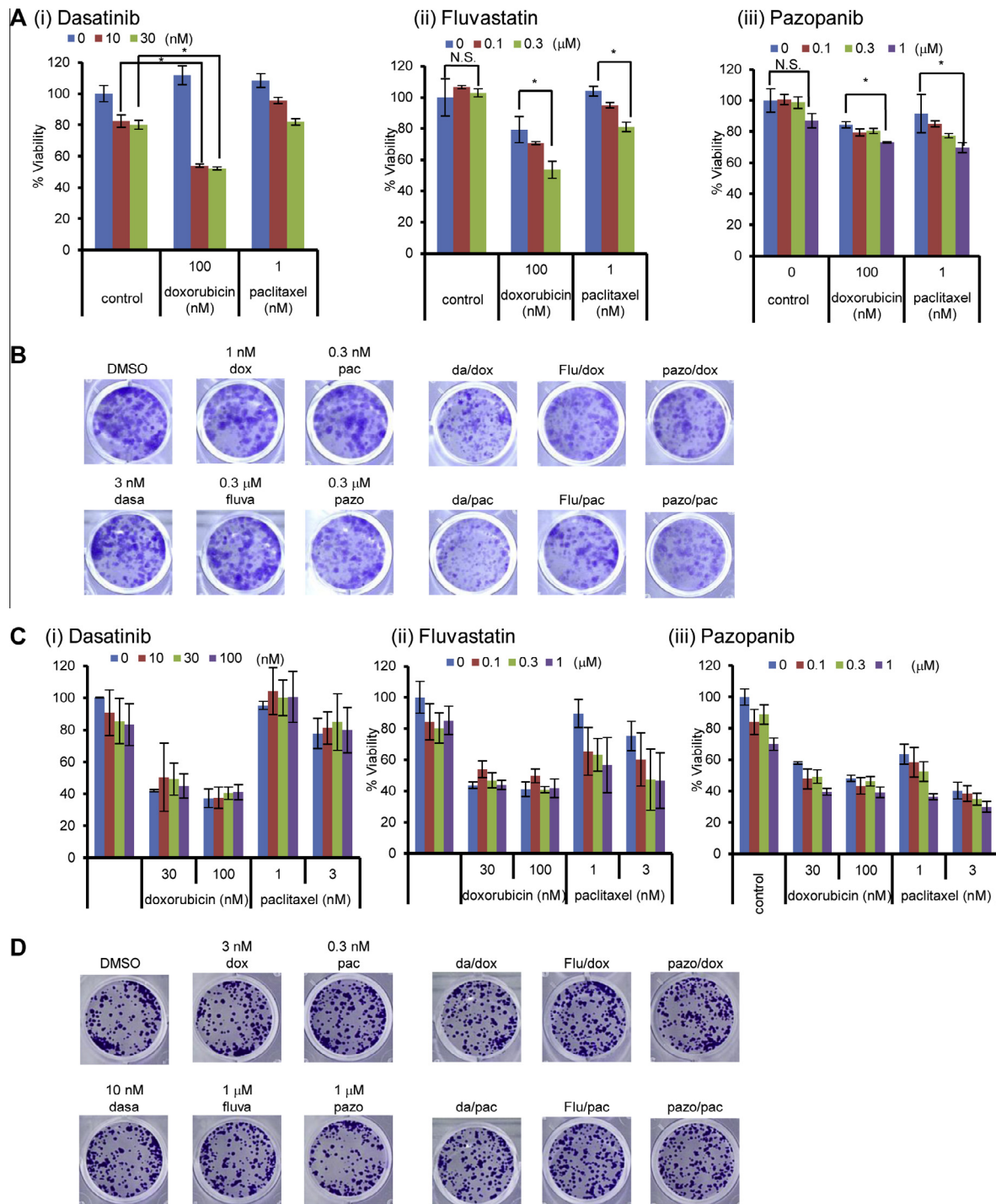


Fig. 4. Dasatinib, fluvastatin, and pazopanib sensitize YAP/TAZ-dependent cancer cells to doxorubicin and paclitaxel. (A) Combination therapy with doxorubicin and paclitaxel with (i) dasatinib, (ii) fluvastatin, or (iii) pazopanib. Cells were treated with dasatinib, fluvastatin, or pazopanib in combination with doxorubicin or paclitaxel for 4 days. Viability was measured by MTT assay. Data represents mean and standard deviation from triplicate, and data is representative of at least three independent experiments. $P < 0.02$. N.S.: not significant. (B) Colony formation of MDA-MB-231 cells treated with dasatinib, fluvastatin, and pazopanib in combination with doxorubicin or paclitaxel. MDA-MB-231 cells were treated with the combination of inhibitors (dasa: dasatinib, fluva: fluvastatin, pazo: pazopanib, dox: doxorubicin, pac: paclitaxel) for 10 days. Colony formation was determined by staining with crystal violet. Data is representative of two independent experiments. (C) Effect of (i) dasatinib, (ii) fluvastatin, and (iii) pazopanib as chemosensitizers in MCF-7, YAP/TAZ-independent cell line. Viability was measured by MTT assay. Data represents mean and standard deviation from triplicate, and data is representative of at least two independent experiments. (D) Colony formation of MCF-7 cells treated with dasatinib, fluvastatin, and pazopanib in combination with doxorubicin or paclitaxel. MCF-7 cells were treated with the combination of inhibitors (dasa: dasatinib, fluva: fluvastatin, pazo: pazopanib, dox: doxorubicin, pac: paclitaxel) for 7 days. Colony formation was determined by staining with crystal violet. Data is representative of three independent experiments.

resulting in Hippo pathway activation. Furthermore, pazopanib induced the proteasome-dependent degradation of YAP and TAZ, which was not prominent in the cells treated with dasatinib and

fluvastatin (Fig. 1E). Pazopanib more strongly induced proteasomal degradation than dasatinib or fluvastatin. Although further study is required to know the precise mechanisms of proteasomal

degradation of YAP/TAZ by pazopanib, it might exemplify a unique class of Hippo pathway activators.

We found that the efficacy of dasatinib, fluvastatin, and pazopanib is correlated with the dependency on YAP/TAZ in breast cancer cell lines. MDA-MB-231 exhibited higher YAP/TAZ-dependent transcriptional activity and its cell growth was YAP/TAZ-dependent (Fig. 2). This cell line was sensitive to these agents, whereas the cell line with low dependence on YAP/TAZ, MCF-7, for example, was resistant to them (Fig. 2E). We further showed that the combination of the identified compounds was effective for MDA-MB-231 (Fig. 3). YAP/TAZ expression is associated with poor prognosis in many cancers, including breast cancer [9,11,39]. YAP/TAZ is also associated with cancer stem cell traits in breast cancer cells [14]; therefore, these agents might be effective in YAP/TAZ-dependent cancers. We also found that dasatinib, fluvastatin, and pazopanib sensitize MDA-MB-231 to doxorubicin and paclitaxel. YAP/TAZ is associated with drug resistance to classical chemotherapeutics [9,32,40]. Combination therapies efficiently reduced the viability of MDA-MB-231 (Fig. 4). Drug resistance is a bottleneck for anti-cancer therapy, and reversing drug resistance to susceptibility is an attractive strategy. The agents identified in this study are effective against YAP/TAZ-dependent cell line (Fig. 2). In recent molecular targeted approaches in cancer therapeutics, the stratification of patients with biomarkers is important. Expression or activation status of YAP/TAZ might be a predictor for these drugs, and such stratification might be important for anti-YAP/TAZ therapy. Our findings can offer an alternative regimen for the treatment of breast cancer using clinically approved drugs.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.fob.2015.06.007>.

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