

Research Paper



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The Roles of GSK-3β in Regulation of Retinoid Signaling and Sorafenib Treatment Response in Hepatocellular Carcinoma

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Abstract

Rationale: Glycogen synthase kinase- 3β (GSK- 3β) plays key roles in metabolism and many cellular processes. It was recently demonstrated that overexpression of GSK- 3β can confer tumor growth. However, the expression and function of GSK- 3β in hepatocellular carcinoma (HCC) remain largely unexplored. This study is aimed at investigating the role and therapeutic target value of GSK- 3β in HCC.

Methods: We firstly clarified the expression of GSK-3 β in human HCC samples. Given that deviated retinoid signalling is critical for HCC development, we studied whether GSK-3 β could be involved in the regulation. Since sorafenib is currently used to treat HCC, the involvement of GSK-3 β in sorafenib treatment response was determined. Co-immunoprecipitation, GST pull down, *in vitro* kinase assay, luciferase reporter and chromatin immunoprecipitation were used to explore the molecular mechanism. The biological readouts were examined with MTT, flow cytometry and animal experiments.

Results: We demonstrated that GSK-3 β is highly expressed in HCC and associated with shorter overall survival (OS). Overexpression of GSK-3 β confers HCC cell colony formation and xenograft tumor growth. Tumor-associated GSK-3 β is correlated with reduced expression of retinoic acid receptor- β (RAR β), which is caused by GSK-3 β -mediated phosphorylation and heterodimerization abrogation of retinoid X receptor (RXR α) with RAR α on RAR β promoter. Overexpression of functional GSK-3 β impairs retinoid response and represses sorafenib anti-HCC effect. Inactivation of GSK-3 β by tideglusib can potentiate 9-*cis*-RA enhancement of sorafenib sensitivity (tumor inhibition from 48.3% to 93.4%). Efficient induction of RAR β by tideglusib/9-*cis*-RA is required for enhanced therapeutic outcome of sorafenib, which effect is greatly inhibited by knocking down RAR β .

Conclusions: Our findings demonstrate that GSK-3 β is a disruptor of retinoid signalling and a new resistant factor of sorafenib in HCC. Targeting GSK-3 β may be a promising strategy for HCC treatment in clinic.

Key words: Retinoid Receptor; GSK-3β; Hepatocellular Carcinoma; Target Therapy

Introduction

Glycogen Synthase Kinase-3 (GSK-3) is a proline-directed serine/threonine kinase with two ubiquitously expressed homologous isoforms, GSK-3 α and GSK-3 β , in mammal cells [1, 2]. GSK-3 β

is constitutively active in resting cells essential for cellular metabolism and homeostasis in normal tissues. GSK- 3β is known to suppress tumor growth as it negatively regulate Wnt/ β -catenin, PI3K/AKT

and many other oncogenic pathways [3, 4]. Inversely, GSK-3 β can be phosphorylated and inactivated by AKT and sequestrated in multi-vesicular endosomes by Wnt signalling. However, many tumors highly express functional GSK-3β in irrespective of AKT and Wnt activation [5, 6]. Overexpression of GSK-3β can facilitate tumor growth and there are considerable evidence to show that GSK-3^β is rationally a therapeutic target [7, 8]. Down-regulation of GSK-3 β by siRNA and microRNA-129 or pharmacological inhibition of GSK-3β with lithium chloride, indirubins, 9-ING-41 and 9-ING-87 can inhibit tumor cell growth and metastasis [9-12].

Retinoids are very important for hepatic homeostasis, which effects are mediated by retinoic acid receptors (RAR α , RAR β and RAR γ) and retinoid X receptors (RXR α , RXR β and RXR γ) [13]. Deregulated metabolism of retinoids and altered expression of their receptors are implicated in HCC development and progression [14, 15]. GSK-3\beta can inhibit RARa-dependent differentiation of myeloid leukemia [16, 17]. Paradoxically, GSK-3β protects RXRα from calpain-mediated truncation in certain solid tumors [18]. Implication of GSK-3ß in retinoid signaling and HCC development need further explore. We characterized here RXRa as a direct substrate for GSK-3β. GSK-3β phosphorylates RXRa and impairs its activation of RAR^β promoter. Clinically, GSK-3\beta is overexpressed and associated with RAR^β reduction in a majority of HCC. RAR^β mediates retinoid action but is frequently silenced during carcinogenesis [14]. Thus, GSK-3β may confer HCC through interfering RARβ-mediated retinoid signalling. This prompted us to further determine whether targeting GSK- 3β /RAR β could be of therapeutic significance in HCC.

Sorafenib, a multi-kinase inhibitor, is currently used to treat HCC [19, 20]. However, its therapeutic resistance remains a significant problem in clinic [21, 22]. Interestingly, sorafenib can stimulate GSK-3 β activity *in vitro* and *in vivo*. We demonstrated that GSK-3 β regulation of RAR β is involved in sorafenib resistance in HCC.

Materials and Methods

Antibodies and Reagents

Antibodies include: anti-GSK-3β (D5C5Z, #12456), anti-phospho-GSK-3β (Ser9,5B3) (#9323) anti-Ki-67 (8D5, #9449) anti-Cleaved Caspase-3 (Asp175) (#9661) from Cell Signaling Technology (CST); anti-phospho-serine/threonine (ab15556) and anti-phospho-serine (ab9332) from Abcam; anti-Flag[®] M2 from Sigma-Aldrich; anti-RARB (A1603) from ABclonal; anti-PARP1 (66520-1-Ig) and anti-His 1231

(66005-1-Ig) from Proteintech; anti-rabbit and anti-mouse secondary antibodies conjugated to horseradish peroxidase, anti-rabbit secondary antibodies conjugated to Cy3 from Invitrogen; anti-RXRa (AN197, sc-774; D20, sc-553), anti-RARa (C20, sc-551), anti-GFP (B-2,sc-9996), anti-c-Myc (9E10, sc-40), anti-β-actin (H196, sc-7210) and anti-GAPDH (FL-335, sc-25778) from Santa Cruz Biotechnology. Chemicals and other regents are: lipofectamin 2000 from Invitrogen; enhanced chemiluminescence (ECL), protein A/G agarose from ThermoFisher Scientific; sorafenib, BIO, SB415286 and tideglusib from Selleck Chemicals; 3-(4,5-Dimethyl-2thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT), 4,6-Diamidino-2-phenylindole (DAPI), 9-cis-RA, isopropyl-1-thio-b-D-galactopyranoside (IPTG) and LiCl from Sigma-Aldrich; cocktail of proteinase phosphatase inhibitors and from Roche: Dual-Luciferase Assay System Kit from Promega.

HCC samples

HCC samples (tumors and para-tumor tissues) were collected from The 174th Hospital affiliated to Xiamen University. The tumors were histologically diagnosed as described [23, 24]. All the use of human samples and study protocols were approved by the Hospital Ethics Committee. All patients signed an informed consent form in prior to sample collection. The clinical data were provided in Table S1.

Cell culture and transfection

HepG2 (HB-8065) and HEK293T (CRL-11268) were purchased from ATCC, while SMMC-7721, Bel-7402, and QGY-7703 from Institute of Biochemistry and Cell Biology (SIBS, CAS). All cell lines were obtained between 2008 and 2013 and authenticated by the vendors. The newly received cells were expanded and aliquots of less than 10 passages were stored in liquid nitrogen. All cell lines were kept at low passage, returning to original frozen stocks every 6 months. During the course of this study, cells were thawed and passaged within 2 months in each experiment. QGY-7703 was cultured in RPMI-1640 medium, while other cell lines were grown in Dulbecco's Modified Eagle's Medium. The cultured cells were supplemented with 10% fetal bovine serum. Sub-confluent cells with exponential growth were used throughout the experiments. Transfections carried by were using out Lipofectamine 2000 according to manufacturer's instructions.

Generation of stable lines

 $GSK-3\beta$ stable lines were generated with retroviral vectors. Briefly, HEK293T cells were

transfected with PCDH-puro-GSK-3 β together with envelope plasmid VsVg (addgene, #8454) and packaging plasmid psPAX2 (addgene, #12260). Retroviral supernatant was harvested at 48 h after initial plasmid transfection and then infected various HCC cell lines. Stable cell pools were selected with 1 µg/ml puromycin (Amresco). The expression efficiency was determined by Western blotting and RT-PCR.

MTT assay and flow cytometry

Cell viability was performed with MTT method as described [15]. For apoptotic analysis, control and treated cells were harvested and washed with precooled PBS twice. The cells were then stained with Annexin V-FITC and propidium iodide (PI) at room temperature for 15 min in the dark. Apoptotic cells were quantitated with flow cytometer analysis (Thermo, Attune NxT).

siRNA synthesis

The siRNA sequences for RAR β (AUA UUC UUC UGA AUA CUU CUG), for GSK-3 β (GUA UUG CAG GAC AAG AGA UdTdT), and non-targeting sequence (UUC UCC GAA CGU GUC ACG UTT) were synthesized by Sangon (Shanghai).

$RXR\alpha$ mutant constructs and $\beta RARE$ reporter

pCMV-myc vector was used to clone a set of plasmids: N-terminal truncation of RXR α (Δ N20, Δ N60, Δ N80, Δ N100) and its point mutations (S49A, S66A, T53A, S78A, T82A, and T82D). Luciferase reporter plasmid of pGL6- β RARE was constructed with RAR β gene's retinoic acid response element sequence (5'-GGG TAG GGT TCA CCG AAA GTT CAC TCG-3').

Dual-luciferase reporter assays

Cells were co-transfected with pGL6- β RARE firefly luciferase reporter constructs, renilla luciferase expression vector (renilla), and HA-RAR α /myc-RXR α in the presence or absence of GSK-3 β . The cells were treated with 1 μ M 9-*cis*-RA combined with or without 5 μ M tideglusib for 20 h. Cell lysates were measured for luciferase activities. The fluorescence intensity was detected in Multiskan Spectrum (PerkinElmer, USA). The renilla luciferase activity was used to normalize for transfection efficiency.

Co-immunoprecipitation

HCC cells and tumor tissues were lysed and sonicated in 500 μ L lysis buffer containing 150 mM NaCl, 100 mM NaF, 50 mM Tris-HCl (pH 7.6), 0.5% NP-40 and 1 mM PMSF. The lysates were incubated with antibodies against endogenous or tags of ectopic proteins and purified with protein A/G beads. For

detection of GSK-3 β -associated RXR α phosphorylation, HCC tumor lysates were subjected to two rounds of immunoprecipitation (IP). The first round of IP (IP1) was performed with anti-RXR α (D20), which product was then subjected to secondary IP (IP2) with anti-GSK-3 β antibody. The lysates or IP2 samples were separated by 10% SDS-PAGE and blotted with anti-p-S/T antibody.

GST pull down

The constructs of pGEX-4T-1-GST-RXRa and pET-22b-His-GSK-3ß were transformed into E. coli. BL21 and E. coli. BL21 (DE3) respectively. Both cells were induced with 1 mM IPTG at 25°C for 16 h. Bacterial pellets were sonicated and centrifuged. The supernatants were mixed with Glutathione-Sepharose (Thermo Scientific) and Ni-NTA agarose (Qiagen), for purification of GST-RXRα and His-GSK-3 β , respectively. Immobilized GST-RXRa protein was incubated with 1 µg His-GSK-3β protein in 500 µl of 150 mM NaCl, 100 mM NaF, 50 mM Tris-HCl, pH 7.6, 0.5% NP-40 and 1 mM PMSF at 4 °C for 2 h. After thoroughly washing, bound proteins were eluted with elution buffer (20 mM Tris-HCl, pH 8.0; 10 mM GSH; 1 mM PMSF; 5 mM DTT) and processed for SDS-PAGE.

In vitro kinase assays

GFP-RXR α was expressed in and purified from HepG2 cells with immunoprecipitation (IP) using anti-GFP antibody. The cell lysates and IP products were incubated with bacterially purified His-GSK-3 β protein in a kinase reaction buffer (pH 7.5, 20 mM Tris-HCl, 10 mM MgCl₂ and 100 mM ATP) at 37°C for 45 min. The reactions were stopped by boiling the samples in loading buffer for 10 min and then separated with 10% SDS-PAGE. GSK-3 β -induced RXR α phosphorylation was detected by anti-phospho-ser/thr (p-S/T) antibody.

Chromatin immunoprecipitation assay

Cells were cross-linked with 0.75% formaldehyde in PBS for 10 min and sonicated in lysis buffer (50 mM HEPES-KOH, pH 8.0, 140 mM NaCl, and 1% TritonX-100). Immunoprecipitation of the chromatin was performed with anti-RXRa (D20) or normal rabbit IgG in 1× dilution buffer (1.0% Triton-X-100, 2 mM EDTA, 150 mM NaCl, and 20 mM Tris-HCl, pH 8.0). Immunocomplexes were purified with A/G agarose beads (Thermo Scientific) and incubated with RNase A (0.5 mg/ml) at 65°C overnight to remove RNA contamination and reverse formaldehyde-induced cross-linking. DNA fragments were purified with a DNA purification kit (Axygen Incorporation, China) and subjected to real-time PCR

analysis. The primers for PCR amplification were: 5'-GCT CTG TGA GAA TCC TGG GA-3' (Forward) and 5'-TGC CTC TGA ACA GCT CAC TT-3' (Reverse) located between -124~+37 bp on RAR β promoter.

RT-PCR

Total RNA was extracted by Trizol (Transgen Biotech). Complementary DNA was synthesized using FastQuant RT Kit (TianGen, Beijing, China). PCR experiments were performed with $2 \times \text{Hieff}^{\text{TM}}$ PCR Master Mix (Yeasen, Shanghai, China) following manufacturer's protocol. The primers for RAR β mRNA transcription were: 5'-TCT CAG TGC CAT CTG CTT AAT CTG-3' (Forward) and 5'-CCA GCA ATG GTT CTT GTA GCT TAT C-3' (Reverse); for GAPDH mRNA transcription: 5'-AGG TCG GAG TCA ACG GAT TT-3' (Forward) and 5'-TGA CGG TGC CAT GGA ATT TG-3' (Reverse).

Animal experiments

Male BALB/c nude mice were injected with HepG2/3 β cells (2×10⁶ cells) subcutaneously in the posterior flanks and treated with 10 mg/kg sorafenib, 2 mg/kg 9-cis-RA, and 5 mg/kg tideglusib every other day at Day 3 of post-implantation. After three weeks of treatment, the mice were sacrificed. The tumors and various organ tissues were collected for further analysis. Tumor volume was measured twice weekly with a caliper. Tumor samples were immunoblotted with antibodies against RAR β , GSK-36, p-GSK-36 and GAPDH. Paraffin sections were immunostained using antibodies against Ki-67 and cleaved caspase 3 with DAB Detection Kit (Polymer) (MXB biotechnologies, Fuzhou, China). The expression of p-GSK-3ß and RARß was determined with fluorescent immunostaining. The sections were co-stained with DAPI and detected by Laser Scanning Confocal Microscope (Zeiss). The study protocols were approved by the Institutional Animal Care and Use Committee of University of Xiamen University.

Statistical analysis

Data were represented as mean \pm standard deviation (SD) or median \pm SEM. The statistical significances of differences were determined using an analysis of variance or Student t test. A P value of <0.05 was considered as significant. All data were acquired in at least three independent experiments.

Results

GSK-3 β is overexpressed and associated with RAR β reduction in HCC

To clarify the role of GSK-3 β in HCC, we firstly

collected HCC samples (n=18) to examine GSK-3β expression. Our results showed that GSK-3ß was upregulated in 66.7% of tumors (≥1.5-fold increase) compared to adjacent liver tissues (Fig. 1A, 1B and Table S1). This was consistent with other reports [25, 26]. In most tumors, increased GSK-3ß expression remained significant active (with low Ser⁹ phosphorylation level) (Fig. 1A). We noted that high GSK-3^β was not correlated to downregulation of total and nuclear β -catenin (Fig. S1A and B), suggesting that the tumor-suppressing effect of GSK-3ß via Wnt/ β -catenin was lost in HCC. To study whether GSK-3β could confer HCC growth, we overexpressed or knocked down GSK-38 with various HCC cell lines. We showed that overexpression of GSK-3^β could strongly promote colony-forming capability of HCC cells, while siRNA-mediated downregulation of GSK-3^β resulted in reduced colony formation (Fig. S1C). The role of GSK-3^β in HCC was further strengthened in two HepG2/siβ clones (Fig. S1D). Importantly, GSK-3β-mediated tumor growth and proliferation was confirmed in *in vivo* experiment (Fig. S1E-G).

Interestingly, tumor-associated GSK-3β was inversely correlated with RARβ expression (Fig. 1A and C). To further evaluate the possible clinical relevance of GSK-3β/RARβ, we analyzed GEPIA (Gene Expression Profiling Interactive Analysis) database (http://gepia.cancer-pku.cn/). Agreement with our results, GSK-3β was higher and RARβ lower in HCC than adjacent liver tissue (Fig. 1D), which phenomenon was also observed in many other (Supplementary malignant tumors Fig. S2). Kalpen-Meier survival plot showed that the patients with high GSK-3 β had a shorter overall survival (OS) than those with low GSK-3 β (Fig. 1E). Together, our results suggest that GSK-36 play a role in RAR6 regulation and HCC development.

Identification of RXR α as GSK-3 β substrate

RARβ expression was subjected to transcriptional regulation by RXRa [27]. We thus investigated whether GSK-3 β could directly interact with and regulate RXRa function. Indeed, our results showed that GSK-3β and RXRa were coimmunoprecipitated mutually in HCC tissues (Fig. 2A) and HepG2 hepatoma cells (Fig. 2B). When co-transfection, GSK-3 β could interact with RXRa in HEK293T cells (Fig. 2C). Interestingly, the interaction between GSK-3β and RXRα could be abrogated when GSK-3^β was inactivated by LiCl (Fig. 2C). Further, cell-free experiments showed that GST-RXRa could pull down His-GSK-3β demonstrating their direct association (Fig. 2D).

We could consistently detect an upper-shifted band of RXR α in GSK-3 β dose-dependent manner in various HCC cell lines and HEK293T cells (Fig. 3A and Fig. S3A), which could be specifically erased by calf intestinal alkaline phosphatase (CIP) (Fig. 3B) and inhibited by GSK-3 β inhibitors, LiCl, SB415286 and BIO (Fig. 3C and D), indicating that GSK-3 β could induce RXR α phosphorylation. Enhanced RXR α phosphorylation was seen in GSK-3 β /S9A (mutation with enhanced enzyme activity), while no RXRa phosphorylation event in GSK-3 β /K85R (a kinase-dead mutant) (Fig. 3E). Using anti-phosphor-Ser/Thr (p-S/T) antibody, we confirmed that GSK-3 β could induce RXRa phosphorylation in HepG2 cells (Fig. 3F) and HCC tumor tissues (Fig. 3G). *In vitro* kinase assays showed that His-GSK-3 β could effectively phosphorylate GFP-RXRa protein in the presence of 100 μ M ATP (Fig. 3H). Taken together, we



Figure 1. GSK-3 β is overexpressed and associated with reduced RAR β expression in HCC. (A) The surgical samples were collected from 18 patients with HCC. GSK-3 β , p-GSK-3 β /Ser⁹ and RAR β expression were detected by western blotting in tumor (T) and adjacent liver (L) tissues. GAPDH was used as loading control. (B) The protein levels were calculated and normalized to GAPDH based on grey values using Quantity One software (Bio-Rad). The median levels of GSK-3 β and RAR β were compared between tumor and nontumor tissues. *p* values are shown. (C) The correlation of GSK-3 β with RAR β was determined by Pearson Correlation Analysis. (D) The clinical relevance of GSK-3 β and RAR β in HCC were further analysed with GEPIA by using TCGA-LIHC database (http://gepia.cancer-pku.cn/). The median levels of GSK-3 β and RAR β were compared between tumors (n=369) and liver tissues (n=160); **p*<0.05 and ***p*<0.01 (Tumor vs Liver). (E) Overall survival (OS) rate analysis. The cut-off level of GSK-3 β Ratio) =1.6, p(HR)=0.0089.

demonstrated that RXR α is a direct substrate of GSK-3 β .

We thus proceeded to map GSK-3 β -mediated phosphorylation site on RXR α . Deletion mutation analysis showed that GSK-3 β could phosphorylate RXR α / Δ N20, Δ N40 and Δ N60, which effect was impaired in Δ N80 and lost in Δ N100, indicating that the putative phosphorylation site is located between 60~100 aa (Fig. 4A). To identify the phosphorylation

site, we introduced Ala point mutation into these putative sites. Our results showed that RXRα phosphorylation by GSK-3β was kept at S49A and S66A, but abolished in S78A and S78A-containing mutations (Fig. 4B and C), thus identifying that Ser⁷⁸ is the site for phosphorylation by GSK-3β. Since GSK-3β recognizes sequence motif in the context of S/T-X-X-X-S/T, Thr⁸² was expected as a priming phosphorylation site. We thus also introduced Asp





Figure 2. GSK-3 β interacts with RXR α . (A)(B) The interaction between GSK-3 β and RXR α was determined via co-IP. The tumor tissues (A) and HepG2 cells (B) were lysed and incubated with anti-RXR α (D20 or \triangle N197) or anti-GSK-3 β antibodies or nonspecific IgG. The IP samples was immunoblotted by anti-RXR α and anti-GSK-3 β antibodies. (C) HEK-293T cells were transfected with myc-RXR α in the presence or absence of GFP-GSK-3 β . The cells were treated with vehicle or 3 mM LiCl for 3 h. The cell lysates were incubated with anti-myc tag or with nonspecific IgG. For A, B and C, the inputs were loaded with 5% of each whole-cell lysate. (D) GST-pull down. GST and GST-tagged RXR α were purified with glutathione agarose beads and incubated with His-GSK-3 β . The membranes were detected with anti-His antibody. GST and GST-RXR α expressions were indicated in coomassie blue staining.

mutation into Ser⁷⁸ (S78D) and Thr⁸² (T82D) to mimic their phosphorylation. As a result, GSK-3 β -mediated RXR α phosphorylation was abolished in S78A and T82A, but retained in S78D and T82D (Fig. 4D). The upper-shifted band seen in S78D was due to its acidic carboxylic group-contained aspartic acid, which was unrelated to the activity of GSK-3 β [28]. Thus, Thr⁸² phosphorylation primes RXR α for subsequent phosphorylation of Ser⁷⁸ by GSK-3 β . This phosphorylation event was confirmed with anti-p-S/T antibody (Fig. 4E and data not shown).







Figure 4. Identification of RXR α phosphorylation sites by GSK-3 β . (A) HEK293T cells were transfected with Flag-GSK-3 β in the presence of myc-RXR α or its various N-terminal truncated forms (Δ N20, Δ N40, Δ N60, Δ N80 and Δ N100). The cells were lysed and blotted with anti-Flag, anti-myc and anti- β -actin antibodies. (B) HEK-293T cells were transfected with Flag-GSK-3 β in the presence of RXR α or its point mutations including S49A, S66A, S78A, and T82A (alone or in combination). The blots were detected with anti-myc, anti-Flag and anti- β -actin antibodies. (C) HepG2 cells were transfected Flag-GSK-3 β combined with myc-RXR α or myc-RXR α /S78A. The cells were treated with or without 20 mM LiCl for 3 h. RXR α expression was detected with anti-myc tag or D20 or Δ N197 antibodies. GSK-3 β and its Ser⁹ phosphorylation were examined with anti-Flag and anti-Ser⁹ antibodies. GAPDH expression was detected with anti-myc tag or D20 or Δ N197 antibodies. GAPDH expression was used as loading control. (D) HepG2 cells were transfected with excert or Flag-GSK-3 β in combination with myc-RXR α or its mutants (S78A, S78D, T82A, T82D) for 24 h. The blots were determined with anti-myc, anti-Flag, and anti-GAPDH antibodies. (E) HepG2 cells were transfected with anti-myc, anti-Flag and anti-GAPDH antibodies. (E) HepG2 cells were transfected with anti-myc, anti-Flag and anti-GAPDH antibodies. (E) HepG2 cells were transfected with anti-myc, anti-Flag, and anti-GAPDH antibodies. (E) HepG2 cells were transfected with anti-myc, anti-Flag, and anti-GAPDH antibodies. (E) HepG2 cells were transfected with anti-myc, anti-Flag and anti-SA plus myc-RXR α or the determined with anti-myc, anti-Flag and anti-GAPDH antibodies. (E) HepG2 cells were transfected with flag-GSK-3 β plus myc-RXR α or myc-RXR α phosphorylation w

GSK-3 β inhibits RXR α transcriptional activity and impairs retinoid signaling

A luciferase reporter containing β RARE sequence was constructed. This reporter was 9-*cis*-RA inducible in the presence of RXR α , which could be inhibited by overexpression of GSK-3 β (Fig. 5A). When Ser⁷⁸ was mutated into Ala⁷⁸ (RXR α /S78A), GSK-3 β -mediated β RARE inhibition could be reactivated by 9-*cis*-RA (Fig. 5A). 9-*cis*-RA could also restore β RARE activity *via* GSK-3 β inhibition by tideglusib (Fig. 5B). Such phenomenon was reproducible in SMMC7721 and Bel-7402 (Fig. S3B and S3C).

We then investigated whether GSK-3 β could modulate the heterodimeric capacity of RXR α . As expected, 9-*cis*-RA could induce RXR α interaction with RAR α . When overexpression of GSK-3 β , 9-*cis*-RA-induced RXR α :RAR α complex was dismantled (Fig. 5C and D). Conversely, the complex could be reassembled when GSK-3 β was knocked down by specific siRNA (Fig. 5D) or inactivated by tideglusib (Fig. 5E, Fig. S3D and S3E).

RXR α could also form dimer with itself. We thus also study the effect of GSK-3 β on 9-cis-RA-induced TREpal luciferase reporter activity, which expression was driven by RXR α :RXR α . We showed that overexpression of GSK-3 β could strongly inhibit the formation of RXR α homodimer, which could be rescued when GSK- β was inactivated by tideglusib (Fig. S3F and S3G). Thus, our results demonstrated that GSK-3 β could impair the dimeric capacity and transcriptional activity of RXR α .

RAR β and p21, two direct target genes of RXR α [29, 30], could be induced by 9-*cis*-RA. Such induction was inhibited when overexpression of GSK-3 β (Fig. 5F and Fig. S4A). GSK-3 β -mediated silence of RAR β and p21 could be relieved when GSK-3 β was inactivated by tideglusib or LiCl (Fig. 5G and Fig. S4C). In contrast, 9-*cis*-RA and LiCl alone only played minor role in modulating the expression of p27, Cyclin D1 and Cyclin B1, all of which do not contain RXR α binding sites on their promoters (Fig. S4C). Interestingly, combined treatment of 9-*cis*-RA and LiCl could synergistically induce expression of these genes (Fig. S4C).

RXRα could bind to RARβ and p21 promoters in the presence of 9-*cis*-RA, while overexpression of GSK-3 β inhibited this activity (Fig. 5H and Fig. S4B). 9-*cis*-RA could reactivate RARβ promoter silenced by GSK-3β when combining with tideglusib (Fig. 5H). Consequently, RARβ expression was induced by co-treatment of 9-*cis*-RA and tideglusib (Fig. 5I). Biologically, overexpression of GSK-3 β rendered HCC cells resistant to 9-*cis*-RA (Fig. 5J). When GSK-3 β was inactivated by tideglusib, the sensitivity of 9-*cis*-RA was re-established in GSK-3 β stably transfected cells. Combination of 9-*cis*-RA and tideglusib effectively induced RAR β expression, which was required for their synergy to inhibit HCC cell growth. Knocking down RAR β largely blocked the synergistic anticancer activity of 9-*cis*-RA/tideglusib (Fig. 5J).

GSK-3β contributes to sorafenib resistance

Sorafenib was shown to activate GSK-3 β [31–33]. We then decided to study the role of GSK-3 β in the action of sorafenib. Although sorafenib could not affect GSK-3ß expression, it extensively increased GSK-3β activity (reduced Ser⁹ phosphorylation) in both HepG2 (Fig. 6A and B) and HepG2/3 β (Fig. 6D and E). Interestingly, 9-cis-RA could reverse sorafenib-induced GSK-3ß activation (increased Ser9 phosphorylation) in cells with low endogenous GSK-3β (HepG2) (Fig. 6A, 6B and Fig. S5A), but failed to do so in GSK-3 β overexpressed cells (HepG2/3 β) (Fig. 6D and E). Consistently, 9-cis-RA could not affect phosphorylation RXRα when GSK-3β was overexpressed (Fig. S5B). Sorafenib-induced PARP cleavage was enhanced when RAR β was induced by 9-cis-RA (Fig. 6A and C).

Biologically, sorafenib resistance was observed in various GSK-3 β stable cell lines compared to their vector-transfected counterparts (Fig. 6F and Fig. S6A). Sorafenib response was reestablished in HepG2/3β when GSK-3ß activity was inhibited by tideglusib. The enhancement of sorafenib response regarding its anti-proliferation activity (Fig. 6F) and anti-colony formation (Fig. S6B) were achieved by co-treatment of 9-cis-RA and tideglusib, which combination could effectively induce RAR^β expression and PARP cleavage (Fig. 6G). When RARB was silenced by specific siRNA, the dose-dependent effect of sorafenib on inducing PARP cleavage was inhibited even in the presence of 9-cis-RA/tideglusib (Fig. 6G). We then used flow cytometry to evaluate the synergy of 9-cis-RA/tideglusib on enhancing the apoptotic effect of sorafenib. Sorafenib could alone induce 22.1% apoptotic cell death, which effect was promoted to 50.8% by combining with 9-cis-RA/tideglusib. The enhancement of 9-cis-RA/tideglusib on sorafenib apoptotic response was impaired when silencing RAR_β (Fig. 6H, 6I and Fig. S6C). Thus, our results demonstrated that GSK-3\beta-mediated RAR^β inhibition was responsible for sorafenib resistance in HCC cells.



Figure 5. GSK-36 impairs retinoid signaling. (A) Reporter assays. HepG2 and HepG2/36 cells were transfected with myc-RXRa or myc-RXRa/S78A. The cells were co-transfected with pGL6-BRARE, Renilla, and HA-RAR of or 24 h and then treated vehicle or 1 µM 9-cis-RA for 20 h. The fluorescence intensities were determined with Multiskan Spectrum (PerkinElmer). Renilla luciferase activity was used to normalize for transfection efficiency. ** p<0.01 (vs respective control). (B) HepG2 cells were transfected with Flag-GSK-3β or Flag vector, in combination with HA-RARa, myc-RXRa, pGL6-βRARE and Renilla for 24 h. The cells were treated with vehicle or 1µM 9-cis-RA for 20 h. HepG2/3β stable cells were transfected with pGL6-βRARE, Renilla, HA-RARα and myc-RXRα. After 24 h transfection, the cells were pretreated with vehicle or with different concentrations of tideglusib (2 µM, 5 µM) for 1 h followed by 1µM 9-cis-RA for 20 h. Luciferase activities were similarly detected. **p<0.01 (vs respective control); ##p<0.01 (GSK-3β vs mock transfection). (C) HepG2 cells were transfected with vector or Flag-GSK-3β in combination with myc-RXRa and HA-RARa for 36 h. The cells were treated with or without 1 µM 9-cis-RA for 6 h. The lysates were immunoprecipitated with anti-myc tag and blotted with anti-HA and anti-myc antibodies. (D) HepG2/3β cells were transfected with siGSK-3ß or scramble (500 pmol in 10 cm dish) for 48 h. The cells were then treated with or without 1 µM 9-cis-RA for 6 h. Co-IP was performed with anti-RAR α and blotted with anti-RAR α and anti-RXR α antibodies. (E) HepG2/3 β cells were pretreated with 5 μ M tideglusib for 1 h and then treated with vehicle or 1 μ M 9-cis-RA for 6 h. The cell lysates were immunoprecipitated with anti-RARa and blotted with anti-RARa and anti-RXRa (D20) antibodies. For (C), (D), and (E), the inputs were detected with 5% of whole cell lysates. (F)(G) RARB mRNA expression. HepG2 cells were transfected with vector or Flag-GSK-3B for 24 h (F). HepG2/3B cells were pretreated with 5 µM tideglusib for 1 h (G). Both HepG2 and HepG2/3β cells were treated with 1 µM 9-cis-RA or vehicle for 24 h. RARβ and GAPDH transcripts were detected with RT-PCR. (H) CHIP assays. HepG2 cells were transiently transfected with Flag-GSK-3β, while HepG2/3β stable cells were pretreated with 5 µM tideglusib for 1 h. Both cell lines were treated with vehicle or 1 µM 9-cis-RA for 20 h. The Chromatin DNA was purified and immunoprecipitated with anti-RXR (D20) antibody or nonspecific IgG. The IPs were subjected to RT-PCR analysis by using specific RARB promoter primers as indicated in Materials and Methods. (I) Different GSK-3B stable cell lines were pretreated with 5 µM tideglusib and then treated with vehicle or with increasing concentrations of 9-cis-RA for 20 h. The lysates were blotted with anti-RARB, anti-Flag, anti-Ser⁹ GSK-3B and anti-GAPDH antibodies. (J) HepG2 and HepG2/3β cells were pretreated with 5 µM tideglusib and then exposed to vehicle or increasing concentrations of 9-cis-RA for 48 h. HepG2/3β cells were also transfected with RARβ siRNA or scramble siRNA and then subjected to similar treatments. The cell proliferation was detected with MTT. *p<0.01 (vs respective control).



Annexin V

Figure 6. GSK-3β induces sorafenib resistance. (A) HepG2 cells were treated with vehicle or 2 μ M 9-*cis*-RA in the presence or absence of 5 μ M sorafenib for different time intervals. The lysates were blotted with antibodies against PARP, GSK-3β, p-GSK-3β, RARβ and GAPDH. (B) GSK-3β activities (GSK-3β/p-GSK-3β) were analysed by normalizing to GAPDH based on band grey values. (C) RARβ expression was eemiquantified based on band grey intensities. (D) HepG2/3β cells were treated alone or in combination with 2 μ M 9-*cis*-RA, $\beta\mu$ M tideglusib and 5 μ M sorafenib for different time intervals. Expression levels of GSK-3β, p-GSK-3β, RARβ and GAPDH were determined with respective anti-bodies. (E) The role of 9-*cis*-RA on sorafenib-induced GSK-3β activities was analysed based on IB band grey intensities derived in (D). For (B), (C), and (E), *p<0.05, *p<0.01 (vs respective controls). (F) HepG2/3β cells were treated alone or in combination with 2 μ M 9-*cis*-RA, $\beta\mu$ M tideglusib and 5 μ M sorafenib for 48 h. The cell growth was examined with MTT assays. **p<0.01 (vs control in HepG2/3β), ##p<0.01 (HepG2/3β vs HepG2), \$p<0.01 (vs control in HepG2/3β). (G)(H)(I) HepG2/3β cells were transfected with RARβ siRNA or scramble siRNA. The cells were then treated alone or combined with 2 μ M 9-*cis*-RA and 5 μ M tideglusib in the presence of absence of sorafenib (with increasing concentrations) for 48 h. The lysates were documented using immunoblot with anti-PARP, anti-RARβ and anti-GAPDH antibodies (G) Apoptotic induction was analysed with Flow Cytometry through Annexin V-FITC/PI staining (H). Representative results were provided (I) *p<0.05 and *p<0.01 (vs respective control); ##p<0.01 (RARβ siRNA vs scramble). C, control; T, tideglusib; S, corafenib; S9c, co-treatment with sorafenib and 9-*cis*-RA; S9cT, combination of sorafenib, 9-*cis*-RA and tideglusib; 9cRA, 9-*cis*-RA.

Targeting GSK-3 β enhances the anticancer effect of sorafenib

The significance of GSK-3 β /RAR β involved in regulation of sorafenib treatment response was finally determined in vivo. Targeting GSK-3β could significantly inhibit tumor growth of subcutaneous or orthotopical xenografts (Fig. 7A and Fig. S6D). We found that inactivation of GSK-38 by tideglusib could significantly shrink tumor (22.3% inhibition), inhibit Ki-67 expression (14.2%) and induce caspase 3 activation (9.1%), further supporting that overexpression of GSK-3β might be a tumor promoter in HCC (Fig. 7A and B). Sorafenib treatment resulted in tumor inhibition by 48.3%, which effect could be largely enhanced to 93.4% by combining 9-cis-RA/ tideglusib (Fig. 7A). Consistently, sorafenib-induced Ki-67 inhibition and caspase 3 activation were greatly promoted by 9-cis-RA/tideglusib from 28.4% to 50.2% and from 12.4% to 30.3% respectively (Fig. 7B). 9-cis-RA was alone inefficient to improve the anti-tumor effect of sorafenib when GSK-3ß remained active (Fig. 7A, B and D). Combination of 9-cis-RA/tideglusib/sorafenib did not affect the mouse weight and change the normal histological characteristics of various tissues including liver, lung, kidney, heart, and spleen, demonstrating that this strategy has less toxic side effect (Fig. S7A and B). Mechanistically, tideglusib and 9-cis-RA/tideglusib could efficiently inactivate GSK-3β, but only combination could strongly induce RAR^β expression (Fig. 7C-E). Interestingly, RAR β expression by 9-cis-RA/tideglusib was extensively translocated into cytoplasm (Fig. 7C), implying that the nuclear export of RAR β is responsible for apoptotic induction and tumor growth inhibition. The involvement of GSK-3 β /RAR β in sorafenib action was summarized in Fig. 7F.

Discussion

Functional GSK-3 β is recently demonstrated to confer tumor development and poor prognosis in a wide range of solid tumors [34, 35]. It is thus widely attempted to design GSK-3 β inhibitor for cancer treatment [36-38]. However, the concern is its another important function in suppressing tumor growth [2]. Inhibition of active GSK-3 β , whether beneficially or detrimentally, is highly dependent on contextual environment and clinical settings [35, 39]. Disclosing the role and mechanism of GSK-3 β in tumor will help develop new therapeutic strategy. HCC is the fourth most common tumor worldwide but with very limited treatment options [24, 40]. The expression and therapeutic significance of GSK-3 β in HCC remain largely unexplored. Although the samples we examined are small, we could consistently demonstrate that GSK-3 β is increased in almost every tumor and upregulation of ≥1.5-fold is seen in 66.7% HCC (Fig. 1A, B and E). Increased GSK-3β is closely associated with shorter overall survival (OS) (Fig. 1E). Consistently, it was recently demonstrated that GSK-3 β is overexpressed in HCC and targeting GSK-3β can induce degradation of c-FLIPL, a master anti-apoptotic regulator [25]. Our study further showed that overexpression of GSK-3ß conferred HCC cell proliferation, colony formation and tumor development, while targeting GSK-3^β by tideglusib can significantly induce about 22.3% of growth inhibition in HepG2/3 β xenografts (Fig. 7A). Thus, we demonstrated that overexpression of functional GSK-3β supports HCC growth. Since overexpression of GSK-3β renders HCC resistant to certain chemotherapies like retinoid and sorafenib, the therapeutic significance of targeting GSK-3β may lie on its combination with other anticancer drugs.

Hepatocarcinogenesis is closely linked to impaired retinoid metabolism and altered retinoid receptors [15, 41, 42]. GSK-3 β is recently suggested to be a modulator of retinoid signaling as it strongly inhibits RARα-dependent myeloid leukemia differentiation in response to all-trans retinoic acid treatment [16, 17]. However, the roles of retinoid receptors in leukemia and solid tumors can be quite different. The therapeutic effects of retinoids are usually less efficacy in solid tumors than leukemia. The mechanism and implication of GSK-3β-mediated impairment of retinoid signaling in solid tumors have not been reported. We demonstrated here that overexpression of GSK-3\beta can inhibit RAR\beta expression and impair retinoid signaling in HCC (Fig. 1 and 5). RAR β expression is required for mediating retinoid action [30], but this protein is frequently down-regulated in HCC with poorly understood mechanism [43]. It was demonstrated that chromatin hypermethylation can impact negatively on RARB expression [44]. Interestingly, GSK-3 β was shown to play a fundamental role in maintaining DNA methylation [45]. There are currently no reports on GSK-3β regulation of RARβ. We demonstrated here that GSK-3 β -mediated RAR β inhibition is attributed to its direct inactivation of RXRa (Fig. 5D), suggesting that a functional RXR α is required for RAR β induction. We identified RXR α as a new substrate for GSK-3β. GSK-3β can directly interact with and phosphorylate RXRa at Ser78 within its N-terminal proline-directed context of S/T-X-X-S/T (Fig. 4). Such modification renders RXRa incapable of heterodimerizing with RAR α to activate retinoic acid response element on RAR β promoter (Fig. 5). Targeting GSK-3 β can recover the function of RXR α

(Fig. 5D) and promote retinoid-induced RAR β expression *in vitro* (Fig. 5G) and *in vivo* (Fig. 7D). Our

results thus disclosed a novel mechanism by which GSK-3 β regulates RAR β expression in HCC.

Figure 7. Animal Experiments. (A) HepG2/3 β cells (2 × 10⁶ cells) were transplanted subcutaneously into mice's posterior flanks. 3 days later, the mice were intraperitoneally injected with 10 mg/kg sorafenib, 2 mg/kg 9-cis-RA, and 5 mg/kg tideglusib, alone or combination every other day. Tumors were collected at Day 21. The tumor volume were evaluated every 3 days after treatment. The tumor weight were measured. (B) Immunostaining of Ki-67 expression and caspase 3 cleavage were detected with respective antibodies (left panel), which expression were also quantified (right panel). *p<0.05 and **p<0.01 (vs respective control). (C) Tumor paraffin sections were immunostained with anti-p-GSK-3 β and anti-RAR β antibodies and detected with anti-rabbit cy3 conjugated secondary antibodies. DAPI counterstaining was provided. (D) Tumors were lysed for IB analysis for detection of RAR β , GSK-3 β , p-GSK-3 β . GAPDH expression was served as control. (E) Quantification of sorafenib-induced GSK-3 β activities was analysed as described in Figure 6B. (F) Mechanistic scheme. The synergy of 9-cis-RA/tideglusib on enhancing the anti-HCC action of sorafenib was summarized. C, control; T, tideglusib; S, sorafenib; 9cRA, 9-cis-RA; S9c, treatment with sorafenib and 9-cis-RA; S9CT, combination of sorafenib, 9-cis-RA and tideglusib.

Deregulation of RAR β -mediated retinoid signaling by GSK-3 β may at least partially explain why clinical trials of some classical retinoids like β -retinoic acid have no proven benefit in HCC [46]. Interestingly, clinical trial of acyclic retinoid, a synthetic analog of retinoids that target at phosphorylated RXR α , revealed a promising effect in reducing the incidence rate of secondary HCC by about 20% [47, 48].

Sorafenib, a multi-kinase targeted anti-cancer drug, is being widely used to treat HCC [19, 20] but with significant treatment resistance. We thus asked if GSK-3β-mediated RARβ inhibition could impact on sorafenib treatment response. We found that sorafenib can extensively activate GSK-3β both in vitro (Fig. 6A-D) [32] and in tumor microenvironment (Fig. 7E). Since GSK-3 β is highly expressed in HCC, sorafenib treatment will generate abundantly hyperactive GSK-3β. Overexpression of functional GSK-3β strongly inhibits sorafenib action as indicated in various GSK-3ß stable liver cell lines vs their vector-transfected counterparts (Fig. 6F and Fig. S6A). 9-cis-RA cannot alone induce RARB expression in GSK-3 β stable cell lines, in which RAR β is silenced by GSK-3 β . Targeting GSK-3 β by tideglusib can greatly potentiate 9-cis-RA activation of RARβ-dependent Importantly, reactivation of RARβsignaling. signaling that is inhibited dependent bv overexpression of GSK-3β returns profoundly unexpected sorafenib treatment outcome (tumor inhibition raised sharply from 48.3% to 93.4%) (Fig. 7A). In this study, we only used low dose of tideglusib in animal experiment by considering that GSK-3 β is normally a critical regulator of cell metabolism and homeostasis. In addition, tideglusib has been demonstrated to have fewer side effects under phase II trial in Alzheimer's disease treatment [49, 50]. On the other hand, normal tissues are resistant to 9-cis-RA-induced cytoxicity [51]. Combination of tideglusib and 9-cis-RA do not exacerbate deleterious effect of sorafenib in liver and other normal tissues (Fig. S7B).

In summary, our findings suggest that HCC may take advantage of GSK-3 β overexpression to support its growth possibly through interfering RAR β -mediated retinoid signalling. The discovery of GSK-3 β /RAR β in sorafenib treatment response may help design improved strategy to overcome the significant treatment resistant problem of sorafenib in clinic.

Abbreviations

HCC: hepatocellular carcinoma; GSK-3: Glycogen Synthase Kinase-3; RAR: retinoic acid receptor; RXR: retinoid X receptor; RARE: retinoic acid response element; TREpal: thyroid hormone response element; 9cRA: 9-cis-RA; CIP: calf intestinal alkaline phosphatase; RT-PCR: reverse transcriptionpolymerase chain reaction; siRNA: small interfering 3-(4,5-Dimethyl-2-thiazolyl)-2,5-RNA; MTT: diphenyl-2H-tetrazolium bromide; DAPI: 4.6-Diamidino-2-phenylindole; IPTG: isopropyl-1-thio-b-D-galactopyranoside; Co-IP: co-immunoprecipitation; CHIP: chromatin immunoprecipitation assay; GEPIA: gene expression profiling interactive analysis; OS: overall survival.

Supplementary Material

Supplementary figures and table. http://www.thno.org/v10p1230s1.pdf

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Author Contributions

Conception and design: JZ, HZ, SZ. Acquisition, analysis and interpretation of data: SZ, HZ, JT, HZ, JL, FL, KC, WL, ST, WG and YZ. Primary samples collection and purification and validation: WL, SZ. Animal model analysis and data collection: SZ, JT. Writing, and revision of the manuscript: JZ, SZ, HZ. Critical discussion and edition of the manuscript: JZ, AW, XZ. All authors have read and approved the manuscript.

Competing Interests

The authors have declared that no competing interest exists.

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