# Rac1b recruits Dishevelled and β-catenin to Wnt target gene promoters independent of Wnt3A stimulation

VAIJAYANTI V. PETHE $^{1,2\ast}, \ \text{GEORGE S. CHARAMES}^{1\cdot3\ast} \ \text{and} \ \text{BHARATI BAPAT}^{1\cdot3}$ 

<sup>1</sup>Samuel Lunenfeld Research Institute, Mount Sinai Hospital, Toronto, Ontario, M5T 3L9;

<sup>2</sup>Department of Pathology and Laboratory Medicine, Mount Sinai Hospital, Toronto, Ontario, M5G 1X5; <sup>3</sup>Department of Laboratory Medicine and Pathobiology, University of Toronto, Toronto, Ontario M5S 1A8, Canada

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Abstract. We previously reported a functional interaction between aberrant Wnt signaling and Rac1/Rac1b GTPases in tumorigenesis. In this study, we further investigated the mechanistic role of nuclear Rac1b. Using chromatin immunoprecipitation (ChIP) studies, we show that Rac1b resides at the promoters of Wnt target genes, c-Myc and Cyclin D1, in HCT116 cells with aberrant Wnt pathway. In HEK293T cells with intact Wnt signaling, Rac1b is tethered to these same gene promoters independent of Wnt3A stimulation and is further observed to recruit Dishevelled and  $\beta$ -catenin in the absence of Wnt3A stimulation. Our studies suggest a novel transcriptional co-activator role of Rac1b in  $\beta$ -catenin/TCF-mediated transcription.

# Introduction

Rac1 GTPase is one of the members of RhoGTPase proteins that are essential regulators of various cellular processes such as maintenance of actin cytoskeleton, mitogenesis, activation of kinase cascade and transcription (1-3). Aberrant Rac1 activation is associated with tumorigenesis (4) Rac1b GTPase, an alternatively spliced variant of Rac1 gene, is preferentially over-expressed in a subset of breast and colon tumors (4-6). The Rac1b protein contains an in-frame insertion of 19 amino acids between Rac1 residues 75 and 76 immediately preceding the Switch II region (7). As a result of this inclusion within a key functional region, several differences exist between Rac1 and Rac1b *In vitro* and *in vivo* studies have demonstrated that Rac1b possesses an impaired intrinsic GTPase activity, and consequently behaves as a constitutively active form of Rac1

*Correspondence to:* Dr Bharati Bapat, Samuel Lunenfeld Research Institute, Mount Sinai Hospital, Toronto, Ontario M5T 3L9, Canada E-mail: bapat@lunenfeld.ca

\*Contributed equally

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(5,7-9). Over-expression of Rac1b was recently attributed to the enhanced expression of exon 3b inclusion splice factor ASF/SF2 (10). Unlike Rac1 GTPase, Rac1b does not lead to lamellipodia formation, or the activation of PAK1, AKT1, or c-Jun-NH2kinase activities (11). However, Rac1b retains the ability to stimulate the NFkB pathway and was shown to induce transcriptional stimulation of a consensus NFkB promoter in a luciferase reporter construct (12,13). Rac1b is unable to bind to many regulators of Rho family GTPases, although it displays enhanced binding to SmgGDS, RACK1, and p120 catenin, proteins involved in cell-cell adhesion, motility, and transcriptional regulation (14). Rac1b signaling is essential for tumor cell viability and survival (11). Intriguingly, Raclb also cooperates functionally with B-Raf V600E mutations to sustain colorectal cell viability suggesting that the two proteins constitute an alternative survival pathway to oncogenic K-ras in colon tumors (15).

Aberrant activation of the Wnt signaling pathway is one of the major causes of colorectal cancer (CRC) (16,17). Wnt signaling regulates expression of a wide range of target genes that direct cell proliferation, migration, establishment of cell polarity, development, specification of cell fate and adult stem cell proliferation (18-22). In the canonical Wnt signaling pathway, the core protein,  $\beta$ -catenin, plays a dual role in the cell, by participating in the maintenance of cadherin-mediated cell-cell contacts, and as a transcriptional co-activator in the nucleus. The list of potential co-activators present at the promoters of Wnt target genes is increasing, however, their specific roles are not well understood. Dishevelled plays a key role as a molecular branch point between the canonical and non-canonical Wnt signaling pathways. Dishevelled also exists in the nucleus and its nuclear localization is required for Dishevelled function in Wnt/ $\beta$ -catenin signaling (23,24). Gan *et al* (25) have shown the involvement of Dishevelled (Dvl-3) in the transcriptional regulation of a Wnt target gene, c-Myc, in SW480 colon cancer cells.

Our group previously reported cross-talk between canonical Wnt signaling and Rac1/Rac1b GTPases (26,27). We demonstrated that over-expression of the splice variant Rac1b facilitates tumour progression by enhancing Dvl-3-mediated Wnt pathway signaling and induction of Wnt target genes (26). We observed that Rac1b downregulated endogenous E-cadherin expression and decreased cell-cell adhesion of HCT116 colorectal cancer

Abbreviations: Dvl-3, Dishevelled; TCF, T cell factor; ChIP chromatin immunoprecipitation

cells and this effect was further augmented by combined action of Raclb and Dvl-3. The classical mechanism of Raclb function occurs through activation at the plasma membrane; however, it is becoming increasingly apparent that the oncogenic properties of Raclb are also mediated by its entry into the nucleus. Mutation of the nuclear localization signal sequence in the C-terminal polybasic region (PBR) of Raclb, however, resulted in a complete loss of Raclb stimulatory effects on TCF-mediated Wnt target gene transcription and the suppressive effects seen on cell adhesion, indicating the importance of nuclear and membrane localization of Raclb. In this study, we investigated the mechanistic role of nuclear Raclb GTPase in Wnt signaling mediated tumorigenesis.

# Materials and methods

Cell culture. HCT116 parental cells (ATCC, Manassas, VA) were cultured in McCoy's 5A media supplemented with 10% fetal bovine serum (FBS). HCT116 cells with stable-inducible expression of Rac1b were sub-cultured and maintained in McCoy's 5A media supplemented with 10% FBS, 10  $\mu$ g/ml blasticidine and 150  $\mu$ g/ml zeocine. 293T cells (ATCC) were cultured in ATCC-formulated DMEM medium supplemented in 10% FBS. L (L cells transfected only with pGKneo) and Wnt3A expressing cells (L cells transfected with pGK Wnt3A), a kind gift of Dr Liliana Attissano (University of Toronto, ON, Canada), were maintained in DMEM supplemented with 10% FBS. Wnt3A expressing cells were additionally supplemented with 400  $\mu$ g/ml G418 (Invivogen, San Diego, CA).

Antibodies. Antibodies for co-immunoprecipitation and ChIP studies were as follows: Raclb (Millipore, Billerica, MA), Monoclonal Anti-Flag M2 (Sigma, St. Louis, MO),  $\beta$ -catenin (H-102), Dvl-3 (4D3) and TCF 4 (H125) were all from Santa Cruz Biotechnology, Santa Cruz, CA.

Antibodies for immunoblotting were as follows: Flag-tag (M2-anti-FLAG-HRP) from Sigma, Rac1b (Millipore, Billerica, MA),  $\beta$ -catenin (Transduction Laboratories, Franklin Lakes, NJ), topoisomerase II (Oncogene Research Products, Boston, MA), Paxillin (Transduction Laboratories, BD Biosciences, Franklin Lakes, NJ),  $\beta$ -actin (Sigma-Aldrich).

*Plasmids*. Mammalian Rac1b expression construct (Rac1bpcDNA3-Flag N) was a kind gift of Dr M.R. Ahmadian, Max-Planck Institute, Heidelberg, Germany. pCDN<sup>™</sup> 3.1 (Invitrogen, Carlsbad, CA) was used as empty vector control for transient transfections.

*Preparation of L/Wnt3A conditioned media*. For the preparation of L and Wnt3A conditioned media, cells were split from the original flask and seeded at 1:10 dilution using DMEM (with 10% FBS) in 15-cm culture dishes. The first batch of the media was harvested after 4 days. For the second batch, we replenished the media and incubated the cells for additional 3 days. Media for both batches were centrifuged at 1000 g for 10 min and saved at 4°C.

Generation of stable inducible cells. Rac1b-pcDNA3-Flag N was used as a template for generation of inducible plasmid

construct. The Raclb ORF including the N-terminal-Flag tag was amplified by polymerase chain reaction (PCR) with oligonucleotide primers to introduce 5'-*Kpn*I and 3'-*Eco*RI restriction sites, respectively. The following forward and reverse primers were used: forward: 5'-CTTAAGCTTGGTACCTAGC CACCATGGACTACAAGGACGACGACGATGACAAG-3' and reverse primer sequence used was 5'-AGGAAGAGAAAAT GCCTGCCTGCTGTTGTAAGAATTCTGCAGATAT CCCAGC-3'. The resulting PCR amplicon was digested with *Kpn*I and *Eco*RI, purified, and was subcloned in to *KpnI-Eco*RI digested pCDNA 4/TO/myc-His (frame B) parent vector (Invitrogen). The expression of Raclb in the newly generated construct was detected and confirmed after induction with doxycyclin at different time-points using Flag-HRP antibody.

T-REx system, a tetracycline regulated mammalian expression system (Invitrogen) was used for generation of a stable inducible cell line according to the manufacturer's instructions. The HCT116/TR6 blasticidine resistant cell line stably expressing the repressor was made as described previously (28). HCT116/TR cells were then transfected with 6  $\mu$ g of Fsp I-linearized Rac1b inducible construct in 6-cm dishes using Lipofectamine 2000 transfection agent (Invitrogen). Forty-eight hours after transfection, the cells were re-seeded into 15-cm plates at 1:10 dilution in medium containing 5  $\mu$ g/ml of blasticidine and 100  $\mu$ g/ml of zeocine. To screen resistant foci, cells were treated with either vehicle (water) or doxycyclin (0.01  $\mu$ g/ml) and protein lysates were harvested after 24 h and analyzed by Western blotting. Positive clones were maintained in blasticidine and zeocine.

Cell fractionation, co-immunoprecipitation and Western blotting. Whole cell lysates were harvested in RIPA buffer and co-immunoprecipitation analysis of HCT116 uninduced and induced whole cell lysates was performed as described (27). Whole cell lysates (500  $\mu$ g) were subjected to immunoprecipitation using 3  $\mu$ g antibody. Immunocomplexes were resolved on 10% SDS-PAGE gel, and analyzed by Western blotting using indicated antibodies. Cytoplasmic and nuclear extracts were prepared according to the manufacturer's instructions using NE-PER<sup>®</sup> Kit (Pierce, Rockford, IL).

Chromatin immunoprecipitation (ChIP) assay. ChIP assays were performed using the ChIP-IT<sup>™</sup> Express Enzymatic kit (Active Motif, Carlsbad, CA, USA) according to the manufacturer's instructions. Briefly, ~1x10<sup>7</sup> HCT116 cells in 15-cm dishes were fixed for 10 min with 1% formaldehyde and then chromatin was prepared using ChIP-IT Express Enzymatic kit protocol. Chromatin was sheared with the Enzymatic Shearing Cocktail for 10 min at 37°C to generate optimally sheared chromatin fragments in the range of 200-1500 bp. Sheared and homogenized chromatin was immunoprecipitated using  $3 \mu g$ of indicated antibodies. Chromatin samples were then eluted, subjected to cross-link reversal, treated with proteinase K. Recovered DNA was analyzed by PCR using primer pairs that flank Tcf-Binding-Elements (TBEs) in C-Myc or Cyclin D1 promoters. The sequences of the primers are as described previously (28). The negative control primers flank a region of genomic DNA of GAPDH gene that is devoid of TBEs. The GAPDH primers were: forward: 5'-ATGGTTGCCACTGG GGATCT-3' and reverse: 5'-TGCCAAAGCCTAGGGGA



Figure 1. Raclb is expressed 7 h post-induction of HCT116 cells with 0.01  $\mu$ g/ml of doxycyclin. HCT116 cells with stable inducible expression of Raclb were plated on 6-cm dishes. Next day, the cells were induced with doxycyclin and incubated for the time-points as indicated. Whole cell lysates were harvested in RIPA buffer. Raclb expression was detected with Flag antibody. Equal protein loading was confirmed with  $\beta$ -actin expression.

AGA-3'. The PCR products were run on 2.5% agarose gel and visualized with ethidium bromide.

# Results

Subcellular distribution of Rac1b, Dvl-3 and  $\beta$ -catenin in HCT116 and HEK293T cells. HCT116 cells harbour consti-

tutively active Wnt signaling due to β-catenin mutation and do not express endogenous Raclb. To address the nuclear role of Rac1b in these cells, we first generated an HCT116 colorectal cancer cell line with stable and doxycyclin-inducible expression of flag-tagged Rac1b and verified its expression by Western blotting. We determined conditions for optimal expression of Rac1b by performing time point and doxycyclin induction dose curves. We observed robust expression of Raclb as early as 7 h post-induction with doxycyclin and at a concentration as low as 0.01  $\mu$ g/ml with no observable expression in cells treated with a drug vehicle control (Fig. 1). To assess the subcellular distribution of Rac1b in stable inducible HCT116 colon cancer cells, we isolated cytoplasmic and nuclear extracts from cells with and without Rac1b induction. The purity of the nuclear and cytoplasmic fractions was verified by probing the Western blots with antibodies specific for nuclear and cytoplasmic marker proteins, topoisomerase II and paxillin, respectively. Robust Rac1b expression was found along with endogenous Dvl-3 and  $\beta$ -catenin in the cytoplasm as well as the nucleus (Fig. 2A). We observed that Rac1b does not affect endogenous Dvl-3 and β-catenin protein profiles since Raclb induction did not alter the expression or cytonuclear distribution of these proteins.

To investigate whether activation of the Wnt signaling could alter the subcellular expression and/or distribution of Rac1b in a cell system with intact canonical Wnt pathway, we treated human embryonic kidney 293T (HEK293T) cells with Wnt3A conditioned medium (WCM) in order to stimulate the canonical Wnt signaling pathway. Control L-cell conditioned medium (LCM) is devoid of secreted Wnt3A protein and was used as a negative control. We first confirmed the ability of Wnt3A conditioned medium to stimulate the canonical Wnt pathway in HEK293T cells by performing the classical  $\beta$ -catenin stability assay as described previously (28). HEK293T cells do not express endogenous Rac1b (Fig. 2B). Following



Figure 2. Raclb co-localizes with endogenous  $\beta$ -catenin and Dvl-3 in cytoplasm and the nucleus and does not modulate their expression. (A) HCT116 cells with stable-inducible expression of Raclb were treated with (I) and without (U) 0.01  $\mu$ g/ml doxycyclin for 15 and 24 h. Uninduced and induced cells were then separated into cytoplasmic (CE) and nuclear (NE) fractions and subjected to Western blotting. Blots were assessed for Raclb, endogenous  $\beta$ -catenin and Dvl-3 expression. Blots were stripped and re-probed with topoisomerase II and Paxillin antibodies to assess the purity of nuclear and cytoplasmic fractions respectively. (B) HEK293 T cells were plated and transfected with empty vector (pCDN 3.1) and Raclb expressing plasmid constructs (Raclb-pcDNA3-Flag N). Next day the medium was removed and cells were treated with L-CM (LCM) and Wnt3A-CM (WCM) for 24 h. Cytoplasmic (C) and nuclear (N) fractions were analyzed by Western blotting using Dvl-3,  $\beta$ -catenin and antibody specific to Raclb. The blots were also probed with topoisomerase II and Paxillin to check the purity of nuclear and cytoplasmic fractions, respectively.



Figure 3. Raclb interacts with endogenous  $\beta$ -catenin, Dvl-3 and TCF-4. HCT116 cells with stable inducible expression of Flag-tagged Raclb were treated with (I) and without (U) 0.01 µg/ml Doxycyclin for 24 h. Whole cell lysates were prepared in RIPA buffer and quantitated for total protein. Whole cell lysates (~500 µg total protein) from both uninduced and induced cells were immunoprecipitated (IP) with Raclb, Dvl-3,  $\beta$ -catenin and TCF-4 antibodies and the immunocomplexes were analyzed by immunoblotting (IB) with Flag antibody. IgG antibody was used as negative control.

ectopic expression of Rac1b in HEK293T cells treated with LCM- and WCM, Rac1b was found in both cytoplasmic and nuclear compartments. We observed that activation of Wnt signaling following WCM treatment of HEK293T cells did not modulate the expression levels of Rac1b (as detected by Rac1b-specific antibody) nor did it affect the cytonuclear distribution of Rac1b, indicating that Wnt signaling activation does not regulate Rac1b in these cells.

Rac1b interacts with endogenous  $\beta$ -catenin, Dvl-3 and TCF-4. Our observation of co-localization of Rac1b,  $\beta$ -catenin, and Dishevelled (Dvl3) in HCT116 stable inducible cells prompted us to investigate if these proteins also interact with each other. Whole cell lysates from HCT116 cells with and without Rac1b induction were immunoprecipitated with antibodies to Dvl-3,  $\beta$ -catenin and TCF-4. As shown in Fig. 3, Rac1b was found to co-precipitate with Dvl-3,  $\beta$ -catenin and TCF-4 suggesting that Rac1b is involved in complex formation with these proteins. These observations are similar to our previous findings where the active form of Rac1 GTPase, V12Rac1, was shown to interact with  $\beta$ -catenin in the HCT116 cells. Raclb binds to the Wnt target gene promoters in HCT116 cells and recruits Dishevelled (Dvl3) and  $\beta$ -catenin to these promoters independent of Wnt3A stimulation in HEK 293 T *cells*. We previously showed that Rac1b expression contributes to elevated transcription of a subset of Wnt target genes in colorectal cells (26). Using TopFlash reporter assays and Cyclin D1 promoter-reporter assays, we established that Rac1b participates in further activation of Wnt signaling in cells with constitutively active Wnt pathway. In addition, with specific knockdown of Raclb using siRNA in HT29 colon cancerderived cells, we observed a decrease in the expression of Cyclin D1 transcript. Given its presence in the nucleus and its role as a co-activator in TopFlash reporter assays, we sought to determine if Rac1b itself is present at the  $\beta$ -catenin/TCF-4 promoter complex in a subset of Wnt target genes in HCT116 cells (Fig. 4A). To address this, we conducted chromatin immunoprecipitation (ChIP) assays to examine the in vivo association of Rac1b with TCF-binding elements (TBEs) located within the promoters of two well-known canonical Wnt transcriptional targets, c-Myc and Cyclin D1. We observed that Raclb is indeed at the promoters of Wnt target genes in colon cancer cells with constitutively active Wnt signaling. We and others have previously shown Dvl-3, \beta-catenin and TCF-4 to bind *c*-Myc and Cyclin D1 promoters in vivo in colon cancer cells (25,28). However, to our knowledge, this is the first report demonstrating the presence of Raclb at the promoter of endogenous Wnt targets in HCT116 cells.

Next, we examined if the binding of Rac1b to these promoters is dependent on Wnt stimulation of HEK293T cells by performing chromatin immunoprecipitation (ChIP) studies. We had already established that WCM treatment of HEK293T cells did not modulate the expression levels of ectopic Rac1b (as detected by Rac1b-specific antibody) nor did it have any effect on cyto-nuclear distribution of Rac1b (Fig. 2B). As expected, Wnt3A stimulation of HEK293T cells leads to an increase in the amount of both Dvl-3 and  $\beta$ -catenin proteins



Figure 4. (A) Rac1b binds to the promoter of Wnt target genes in HCT116: HCT116 cells with stable inducible expression of Flag-tagged Rac1b were treated with (I) and without (U) 0.01  $\mu$ /ml Doxycyclin for 24 h. Chromatin was isolated as described in Materials and methods and subjected to immunoprecipitation with Flag, Dvl-3,  $\beta$ -catenin and TCF-4 antibodies. IgG was used as a negative control. Enriched chromatin was PCR amplified using the primers that flank TCF binding sites (TBEs) in C-*Myc* or *Cyclin D1* promoters. GAPDH primers as described in Materials and methods were used as negative controls. Input represents 10% of chromatin used for immunoprecipitation. (B) Rac1b binds to the promoter of Wnt target genes in Wnt-independent manner and recruits  $\beta$ -catenin and Dvl-3 to the *c*-*Myc* and *Cylin D1* promoters in the absence of Wnt3A stimulation of HEK293T cells: HEK293 T cells were plated and transfected with empty vector (pCDN 3.1) and Rac1b expressing plasmid constructs (Rac1b-pcDNA3-Flag N). Next day the medium was removed and cells were treated with L-CM (LCM) and Wnt3A-CM (WCM) for 24 h. For ChIP assay, chromatin was subjected to immunoprecipitation using Dvl-3,  $\beta$ -catenin and Flag antibodies. Enriched chromatin was analyzed by PCR using primers that flank the TBEs in Wnt target genes, *c*-*Myc* and *Cyclin D1*. GAPDH primers as described in Materials and methods were used as negative controls. Input represents 10% of chromatin used for immunoprecipitation.



Figure 5. Proposed model for the nuclear role of Rac1b in HCT116/HEK293T cells. Upon Wnt3A stimulation in a normal cell (left panel),  $\beta$ -catenin and Dvl-3 proteins are recruited to the promoter of canonical Wnt target genes. In cancer, nuclear expression of Rac1b (center panel), leads to the recruitment of  $\beta$ -catenin and Dishevelled to these promoters, independent of Wnt3A stimulation. Rac1b expression in the presence of Wnt3A stimulation (right panel) results in a synergistic increase in transcription of canonical Wnt target genes.

present at the *c-myc* and *Cyclin D1* promoters (Fig. 4B) in the absence of Raclb. Intriguingly, we observed that when Raclb is present, it binds to the promoters of endogenous Wnt target genes, *c-myc* and *Cyclin D1*, independent of Wnt3A stimulation. Furthermore, Raclb recruits Dvl-3 and  $\beta$ -catenin to these same promoters independent of Wnt3A stimulation. Taken together, these observations lend further support for a novel role of Raclb in the nucleus.

### Discussion

In the present study, we explored the nuclear role of Rac1b in a colon cancer cell line with aberrant Wnt signaling due to constitutive  $\beta$ -catenin expression (HCT116), as well as in human embryonic kidney cells (HEK293T) with intact Wnt signaling as a model base system. While it is well known that the classical mechanism of Rac1b function occurs through activation in the cytoplasm and/or at the plasma membrane, emerging evidence indicates that the oncogenic properties of Raclb are also mediated by its entry into the nucleus. After establishing cross talk between RaclGTPase and wnt signaling, we reported for the first time that ectopically expressed constitutively active mutant of Rac1, V12 Rac1GTPase is associated with a subset of Wnt-responsive gene promoters in vivo. In addition, endogenous Tiam 1, a Rac1-specific activator, associated with nuclear TCF-binding sites to form a complex of Rac-Tiam-\beta-catenin-TCF to enhance Wnt target gene transcription. In extension of work previously published by our group and after establishing the nuclear localization of Raclb, we were interested in exploring if Raclb is recruited to these promoters. We show that Raclb indeed associates with the  $\beta$ -catenin-TCF complex *in vivo* in colon cells with inherently dysregulated Wnt signaling. This makes us believe that, aside from the nuclear  $\beta$ -catenin-TCF complex, which is a prerequisite for transcription, a myriad of proteins may act as transcriptional co-activators consequently leading to enhanced dysregulated expression of endogenous oncogenic targets. Besides its over-expression in a subset of colon and breast tumors, our work demonstrates an alternate mechanistic role of nuclear Raclb that may further enhance tumor progression.

Wnt proteins constitute a family of secreted signaling molecules that regulate pathways essential for development, and its aberrant activation drives oncogenesis in a multitude of human cancers. The total expression levels of Dvl-3 and  $\beta$ -catenin are inherently elevated in HCT116 cells due to a  $\beta$ -catenin mutation that results in its stabilization, and consequently, the constitutive activation of the canonical Wnt pathway. It is well known that  $\beta$ -catenin is an essential nuclear effector of canonical Wnt signaling. Extracellular Wnt ligands activate Dishevelled, which is believed to function as a scaffold protein bridging the receptors and downstream components of What signaling pathway leading to activation of  $\beta$ -catenin target gene. Nuclear localization of Dishevelled is required for its function in the Wnt/β-catenin pathway. A subsequent study further demonstrated that, in addition to its cytoplasmic role in regulating the stability of  $\beta$ -catenin, Dishevelled could independently interact with c-Jun and  $\beta$ -catenin in the nucleus to mediate the formation of a transcription complex comprising

Dishevelled, c-Jun, β-catenin, and TCF. Recent studies have shown a novel role of Dishevelled in the canonical Wnt signaling pathway by mediating the transcriptional activity of the  $\beta$ -catenin/TCF complex. Gan *et al* (25) demonstrated that Dvl-3 is recruited to the promoter of Wnt target gene, *c-myc*, in a Wnt3A-dependent manner, whereas binding of TCF-4 to the promoter was independent of Wnt3A stimulation. Similarly, our previous work showed that Rac1 and the Rac1-specific activator Tiam1 are also components of transcriptionally active β-catenin/TCF-4 complexes at *c-Myc* and *Cyclin D1* promoters; and that the presence of Rac1 and Tiam1 within these complexes serves to enhance target gene transcription (28). We found that while Raclb is also present at the gene target promoters in vivo, this binding is not dependent on Wnt signaling. Further, Rac1b recruits Dvl-3 and β-catenin to these promoters independent of Wnt3A stimulation.

Based on our current data as well as previously published studies, we propose a model as illustrated in Fig. 5 to describe the novel role of Rac1b in the nucleus. Despite functional differences between Rac1 and Rac1b, both proteins are found to be present in complexes with TCF/ $\beta$ -catenin in colon cancer cells, and also bind the promoters of Wnt target genes. Dvl-3 and β-catenin are known to bind these Wnt target promoters only upon Wnt3A stimulation of HEK293T cells. We now demonstrate, for the first time, that Rac1b is associated with the promoters of Wnt target genes, c-Myc and Cyclin D1, independent of Wnt activation. Further, similar to Wnt3A stimulation, Raclb also leads to the recruitment of Dvl-3 and  $\beta$ -catenin to these promoters, independent of Wnt activation. Using a promoter reporter assay, we have previously shown that, in HEK293T cells, Rac1b augments Dvl-3-mediated activation of TCF/LEF-dependent transcription. Co-expression of Rac1b and Dv1-3 in HEK293T cells caused a striking transcriptional activation of TOPFlash and Cyclin D1-luc constructs compared to expression of individual proteins. Taken together, we speculate that it is this novel nuclear role for Raclb in co-operation with Wnt components that leads to augmented transcription of Wnt target genes, and subsequently tumorigenesis. The data presented here suggests the existence of an additional alternative mechanism of Rac1b mediated activation of Wnt signaling, and underscores the molecular complexity associated in the signaling networks leading to cancer. We believe that this new mechanism further extends our understanding of functional role of nuclear Rac1b in cell signaling.

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