

## Beta-catenin Mediates Soft Tissue Contracture in Clubfoot

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**Abstract** The contracted tissues from clubfeet resemble tissues from other fibroproliferative disorders such as palmar fibromatosis. Beta-catenin-mediated signaling is a crucial pathway controlling the fibroproliferative response in many fibroproliferative disorders. To determine if beta-catenin signaling plays a role in clubfoot, contracted and less contracted tissues from clubfeet were studied using Western analysis to determine the protein level of beta-catenin. Primary cell cultures were established from these tissues, and they were treated with either lithium to increase beta-catenin or Dickkopf-1 to inhibit beta-catenin. RNA was extracted from the cells and analyzed to determine how beta-catenin regulates expression of Type III collagen, an extracellular matrix protein upregulated in contracted clubfoot tissue. There was a more than twofold increase in beta-catenin protein in the contracted tissues. Treatment with either lithium or Dickkopf-1 showed Type III collagen RNA expression positively correlated with the protein level of beta-catenin. These data support the concept that beta-catenin-mediated signaling plays an

important role regulating contracture in clubfeet. Because pharmacologic agents are under development to block this signaling pathway, such drugs could be used in cases of severe stiffness to improve range of motion or to decrease the need for radical surgical approaches.

### Introduction

Clubfoot is one of the most common musculoskeletal anomalies in newborns [3]. Although a genetic cause is suggested, because it tends to run in families [27], data from early amniocentesis suggest oligohydramnios as a cause [35]. In either case, osseous deformities [39], muscle abnormalities [19, 22, 45], and arrested fetal development [18] are all hypothesized to play a role in its pathogenesis. Manipulation and casting has become the most frequently used treatment for clubfoot with generally good results reported [32, 34]. Long-term study of clubfoot shows feet that maintain a good range of motion and are not contracted have better function [10, 11, 15, 23].

Ultrastructural study of the contracted tissue in clubfeet shows cells from the medial side of clubfeet contain microfilaments, identical in nature to those described in palmar fibromatosis [36]. Cells from the contracted tissue express Type III collagen at high levels [18, 25]. Type III collagen is expressed in healing wounds, hypertrophic scar tissues, and in contracted tissues in conditions such as palmar fibromatosis [12, 33, 37]. Cells from palmar fibromatosis and the contracted tissues in clubfoot express a number of growth factors such as platelet-derived growth factor and transforming growth factor beta [25]. However, expression of these factors may not be the primary cause of the contracture, but instead may be secondary to activation of another signaling pathway.

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Work over the past decade demonstrates that beta-catenin-mediated signaling plays a central role regulating fibroblast activity [6, 7, 9, 21, 38]. This was initially suggested by the finding that mutations activating beta-catenin cause the fibroproliferative tumor aggressive fibromatosis [1, 42], but more recent data show hyperplastic wounds and palmar fibromatosis are also associated with activation of this pathway. Indeed, pharmacologic agents that activate beta-catenin signaling will cause hyperplastic wounds in mice [9].

Beta-catenin is a key member of the canonical Wnt signaling pathway. This pathway is activated by specific circulating Wnt ligands. In the absence of a Wnt ligand, beta-catenin is processed for degradation by phosphorylation by a multiprotein complex consisting of APC, GSK3-beta, and Axin. Phosphorylation of beta-catenin targets it for ubiquitin-mediated degradation at the proteasome. In the presence of Wnt signaling proteins, stabilized beta-catenin can translocate to the nucleus, bind to the members of the Tcf-Lef family of transcription factors to form a transcriptional activation complex, and induce expression of target genes [2, 10, 28, 31, 41] (Fig. 1). In the case of aggressive fibromatosis, specific mutations cause activation of beta-catenin independent of Wnt ligands [1, 42]. In the case of palmar fibromatosis, beta-catenin is activated in the absence of an identified mutation [30, 44].

An adjuvant treatment to reduce stiffness would result in a better outcome for patients with clubfeet. This might be useful in those rare feet that are resistant to manipulation and casting. Such adjuvant therapy may improve motion and function after treatment. Although agents to block beta-catenin signaling are not yet available, there are several

approaches to reduce its activity that are under investigation [14, 29]. As such, if beta-catenin is indeed found to play a major role in clubfoot, once such agents become available, they can be applied to this condition.

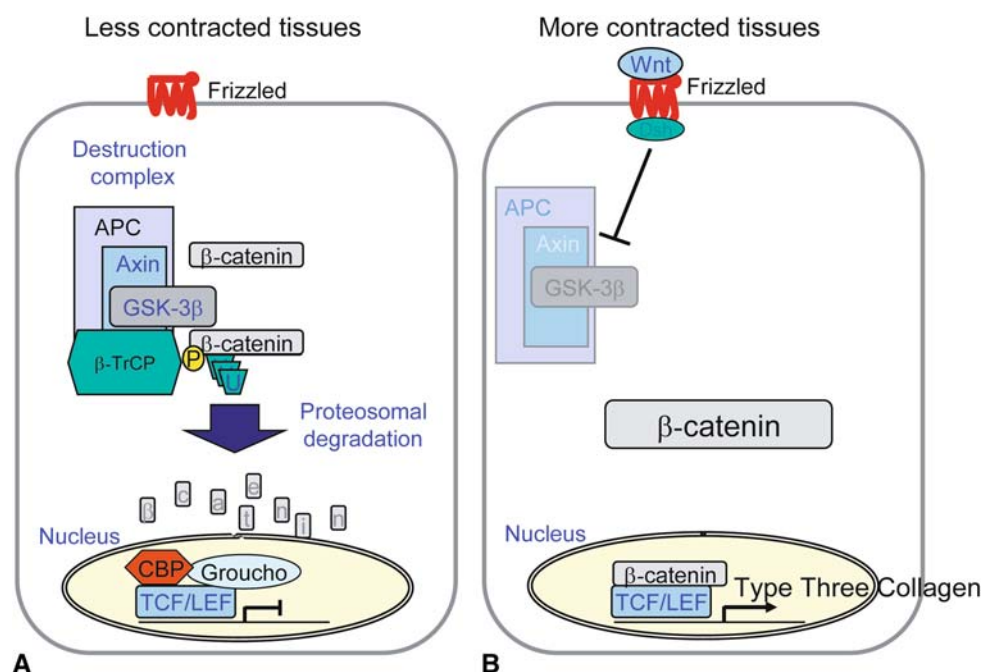
We therefore asked whether beta-catenin signaling is activated in the contracted tissues in clubfeet and if this signaling pathway regulates the expression of genes important in contracture formation.

## Materials and Methods

We studied tissues from patients undergoing surgery for clubfeet. These tissues were the same ones that were used in a previous investigation of growth factors in patients with clubfeet [25]. Contracted tissues were obtained from the medial aspect of the talonavicular joint, and less contracted tissues were obtained from the plantar surface of the calcaneocuboid joint. Sufficient materials from 10 feet were available for study. We obtained institutional approval for the study, and all patients completed informed consent to participate.

We processed tissue as soon as possible after surgical excision as previously reported [6, 8, 24, 25]. A portion of each sample was cryopreserved in liquid nitrogen vapor, and this was used for protein analysis. We used another portion of the tissue, from the more contracted side of the clubfoot, to establish primary cell cultures. Cultures were established by mechanical and enzymatic dissociation (trypsin and collagenase) in culture medium (Dulbecco's modified Eagle's medium) with 10% fetal calf serum.

**Fig. 1A–B**  $\beta$ -catenin regulation and function in less contracted and more contracted clubfoot tissues. **(A)** When Wnt signaling is quiescent, in less contracted tissues, a multiprotein complex phosphorylates amino terminal serine and threonine residues, resulting in  $\beta$ -catenin degradation by a ubiquitin-mediated pathway. **(B)** Wnt activation, as is the case in contracted clubfoot tissues, inhibits the ubiquitin-mediated degradation of  $\beta$ -catenin, elevating  $\beta$ -catenin protein level. Stabilized  $\beta$ -catenin translocates to the nucleus and binds members of the tcf-lef family, resulting in transcriptional activation. One of the targets in clubfoot is Type III collagen.



We extracted protein from samples by homogenization in lysis buffer (1% sodium dodecyl sulfate [SDS], 10 mM Tris-HCl, pH 7.4) followed by centrifugation at 12,000 g for 5 minutes. Equal amounts of total protein were electrophoresed on an SDS polyacrylamide gel, transferred to a polyvinylidene difluoride membrane, and stained to verify equal amounts of transferred protein from each sample. Western blot was performed using an antibody to activated beta-catenin (R & D Systems, Minneapolis, MN), Phospho, and total GSK-3-beta (Cell Signaling Technology, Danvers, MA). Hybridization was carried out overnight at 4°C and detected using an antimouse IgG-horseradish peroxidase secondary antibody and chemiluminescence. The membranes were stripped and reprobed with an antibody to GAPDH (R & D Systems) as an additional protein loading control.

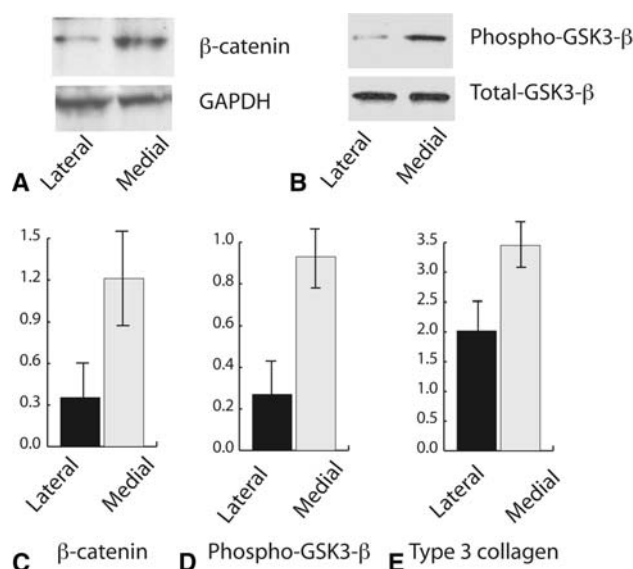
To determine the role of beta-catenin in these cells, they were treated with lithium, which activates beta-catenin-mediated signaling [4, 9], or with Dickkopf-1 (DKK-1), which inhibits canonical Wnt signaling [4, 5, 20, 43]. We performed in vitro experiments with either 50 mM LiCl (Sigma, St Louis, MO) or an equal concentration of NaCl as a control to activate signaling. Cells were treated with 200 multiplicity of infection Ad DKK1-HA in serum-free media using identical conditions as previously reported [4, 5, 20, 43]. As a control, we used the same adenovirus but expressing an empty vector. After 1.5 hours, cells were returned to regular 10% fetal bovine serum media and left to grow overnight. RNA and protein were extracted and used for expression studies and Western analysis to determine that beta-catenin levels did indeed change. In additional cell cultures, we added bromodeoxyuridine (BrDU) to the cells overnight and the relative proportion of cells incorporating BrDU was measured using a commercially available colorimetric assay (Roche, Indianapolis, IN).

Previous work suggests Type III collagen is upregulated in contracted tissues in clubfeet and as such is a marker of tissue contracture [25, 36]. We used real-time polymerase chain reaction to determine differences in Type III collagen expression between cultures, using GAPDH as a house-keeping control, using previously reported primers and conditions [16].

Means, standard deviations, and 95% confidence intervals were calculated for the data. We used two-way Student's t-test to compare comparisons between data sets (Microsoft Excel; Microsoft, Redmond, WA).

## Results

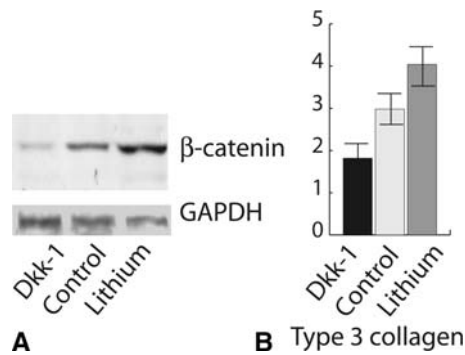
We found Type III collagen, a gene expressed at high levels in contracted tissues, was expressed at a higher ( $p = 0.006$ ) level in the more contracted (medial) tissues



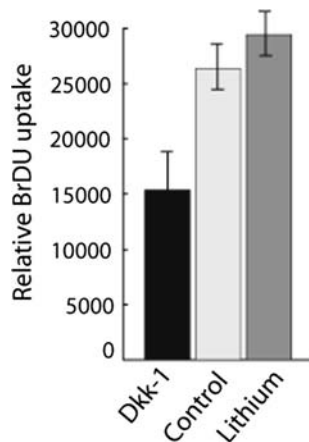
**Fig. 2A–E** Contracted clubfoot tissues have higher levels of beta-catenin protein. **(A)** Western analysis from a protein extract from a representative clubfoot sample shows a higher level of protein expression in the medial, more contracted tissues compared with lateral, less contracted tissues compared with GAPDH as a loading control. **(B)** Western analysis from a representative clubfoot sample shows a higher level of GSK-3-beta phosphorylation in the medial contracted tissues compared with the lateral, less contracted tissues compared with total GSK-3-beta. **(C)** Densitometry results from Western analysis for beta-catenin level from all 10 clubfeet are shown. Band density is given relative to GAPDH as a loading control. Results are shown as means and 95% confidence intervals. **(D)** Densitometry results from Western analysis for phosphorylated GSK-3-beta level from all 10 clubfeet are shown. Band density is given relative to total GSK-3-beta as a loading control. Results are shown as means and 95% confidence intervals. **(E)** Results of real-time polymerase chain reaction for RNA expression of Type III collagen compared with GAPDH showing a higher level of expression in the medial, more contracted tissues. Results are shown as means and 95% confidence intervals.

compared with the less contracted lateral tissues (Fig. 2). This confirms the medial tissues examined are indeed representative of contracted clubfoot tissues. Using Western analysis, we found a more than twofold increase ( $p = 0.003$ ) in activated beta-catenin in the medial (contracted) tissues compared with the lateral (less contracted) tissues. This increase was found in tissues from all 10 feet we studied (Fig. 2). We then examined GSK-3-beta, a protein that is phosphorylated with canonical Wnt signaling activation, and found a higher ( $p = 0.01$ ) level of phosphorylation in the medial capsular tissues (Fig. 2).

Using Western analysis, we found DKK-1 decreased ( $p = 0.002$ ) and lithium increased ( $p = 0.005$ ) beta-catenin in the cell cultures derived from the contracted clubfeet compared with controls (Fig. 3). Quantitative real-time polymerase chain reaction showed an increase ( $p = 0.02$ ) in expression of Type III collagen with lithium treatment and a decrease ( $p = 0.01$ ) in the expression level of Type III collagen with DKK-1 treatment (Fig. 3).



**Fig. 3A–B** Beta-catenin regulates expression of Type III collagen in the contracted tissues from clubfeet. (A) Western analysis shows treatment with DKK-1 decreases and treatment with lithium increases beta-catenin protein level compared with GAPDH as a loading control. (B) The level of RNA expression for Type III collagen as detected using real-time polymerase chain reaction shows a positive relationship between beta-catenin levels and Type III collagen expression. Results are shown as means and 95% confidence intervals.



**Fig. 4** Beta-catenin regulates proliferation in the contracted tissues from clubfeet. Proliferation is measured using BrdU incorporation in cell cultures from the contracted tissues from clubfeet. Results are shown as means and 95% confidence intervals for relative BrdU incorporation with a higher level indicating a higher proliferation rate. There is a positive relationship between beta-catenin level and proliferation rate.

BrdU incorporation, as a measure of proliferation, was assessed in the cell cultures subjected to the same treatments. There was an increase ( $p = 0.01$ ) in the relative uptake of BrdU with lithium treatment and a decrease ( $p = 0.03$ ) in uptake with DKK-1 treatment compared with controls (Fig. 4).

## Discussion

The contracted tissues from clubfeet share histologic and ultrastructural similarities with contracted tissues from

fibroproliferative conditions such as palmar fibromatosis [14, 29]. Beta-catenin-mediated signaling is a cell regulatory pathway that is activated in fibroproliferative conditions causing cell proliferation and the deposition of matrix components. Pharmacologic agents to modulate beta-catenin signaling activation are in development. As such, it is possible beta-catenin levels are elevated in the contracted tissues from clubfeet and therapeutically targeting this signaling pathway could be used to decrease contracture and improve outcome for children being treated for clubfeet. In this study, we tested the hypothesis that beta-catenin protein level was elevated in contracted tissues from clubfeet and that modulation of beta-catenin could decrease cell proliferation and the expression of extracellular matrix components in a way that might decrease the severity of the contracture.

One limitation of this study is clubfoot samples were obtained from feet that underwent operative treatment almost always after a period of corrective casting. It is therefore possible the change in beta-catenin protein level is the result of the period of immobilization rather than the foot deformity itself. Although this possibility cannot be completely excluded, it is not likely to be the case, because all of the feet had at least a 6-week period without immobilization before operative treatment, the entire foot was immobilized in the cast, and the changes resulting from immobilization would likely be present throughout the foot. Furthermore, the medial tissues from fetal clubfeet have the same cytologic features characteristic of contracture and express matrix components associated with contracture at a similar level as medial tissues from surgically treated clubfeet [14, 29] suggesting changes associated with tissue contracture are not related to post-natal therapy. Another potential limitation is that using the current treatment approach, most clubfeet are treated with manipulation and casting; one could argue these are abnormally severe feet that were available for study. However, these samples were obtained several years ago, at a time when more than half of all clubfeet that presented to our institution were treated with open circumferential releases. In an ideal situation, fetal tissues would be used for such a study. Because of difficulties in obtaining appropriately preserved tissue from prenatal clubfeet for these analyses, it is impractical to test for differences in beta-catenin using fetal clubfoot tissues. The control tissues used in this study were from the less contracted tissues from clubfeet rather than from normal feet or other capsular tissues that were not related to a pathologic process. Obtaining such normal tissues is rather difficult and, as such, they are not included in this study. However, the level of beta-catenin protein is approximately the same as that found in normal control tissues from other sites in the body as previously reported in a number of studies [1, 6,

31, 38] suggesting the level of expression of the less contracted tissues is at approximately a baseline level for normal connective tissues.

We found the contracted capsular tissues in clubfoot were associated with increased protein levels of beta-catenin. Furthermore, regulating beta-catenin in cell cultures from the contracted tissues shows a positive correlation with expression of Type III collagen and proliferation rate with the protein level of beta-catenin. Because Type III collagen is an important extracellular component of the contracted clubfoot tissues, this suggests the beta-catenin signaling pathway plays an important role regulating contracture in clubfoot. Beta-catenin is also elevated in contracted tissues in a variety of other soft tissue fibroproliferative processes (eg, palmar fibromatosis) [2, 31]. Many of these processes are associated with myofibroblast cells and soft tissue contracture. The level of beta-catenin protein we found in the contracted clubfoot tissues is approximately the same as that found in palmar fibromatosis [38]. As such, it seems the regulation of canonical Wnt signaling through beta-catenin is a common factor in fibrosis. Although growth factors and extracellular matrix components also play a role in these processes, and because beta-catenin directly regulates transcriptional control of many of the genes important in fibrosis [9], it is likely a central regulator of this process. Because treatment with DKK-1 inhibited beta-catenin, it is likely that in clubfoot, a Wnt-ligand-dependent process activates beta-catenin.

Intriguingly, a genetic linkage study searching for the gene(s) predisposing to idiopathic clubfoot found a relationship between Wnt 7A and clubfoot [13]. Because Wnt ligands activate beta-catenin, it is possible this could play a role as a causative factor in clubfeet. However, a second study of genetic linkage showed Wnt 7A is not likely to be a causative gene for clubfoot [26]. Taken together, this suggests the beta-catenin-mediated fibrosis is more likely an end result rather than a causative factor in clubfoot. Despite this, it is possible genetic changes that alter the level of expression or activity of a Wnt ligand will alter the level of beta-catenin and thus the severity of clubfoot. As such, genes in the beta-catenin signaling pathway might play important roles regulating severity of contracture and response to treatment.

The data from this study raise several intriguing possibilities for the management of clubfoot, which could be explored as part of future work. For instance, levels of beta-catenin could be a prognostic factor, and lower levels may be related to less contracted tissues. Beta-catenin levels also may differ between idiopathic and syndromic or atypical clubfeet. They might also predict recurrence after successful manipulative treatment, the response to manipulative treatment, or a better response to anterior tibialis

transfers for recurrence. Data supporting these possibilities could be obtained from samples from patients undergoing surgery for relapsed clubfeet.

Although drugs that regulate beta-catenin are not yet available for clinical use, there is a great deal of research into developing such agents [14, 17, 40, 46]. In addition, screening of known compounds may identify drugs that modulate beta-catenin activity [40]. Such agents may be used as an adjuvant in the treatment of stiff or syndromic clubfeet or for feet in which there is a recurrence requiring additional surgeries. They may decrease contracture and increase the ease of manipulative correction and increase the posttreatment range of motion. As such agents become available, they would be good candidates for clinical trials in children with clubfeet.

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