

Up-Regulation of PTHrP and Bcl-2 Expression Characterizes the Progression of Osteochondroma towards Peripheral Chondrosarcoma and Is a Late Event in Central Chondrosarcoma

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SUMMARY: Chondrosarcomas are malignant cartilage-forming tumors arising centrally in bone (central chondrosarcoma) or within the cartilaginous cap of osteochondroma (peripheral chondrosarcoma). For hereditary multiple osteochondromas, two responsible genes, *EXT1* and *EXT2*, have been cloned. Their recently elucidated role in heparan sulfate biosynthesis and Hedgehog diffusion leads to the hypothesis that *EXT* inactivation affects fibroblast growth factor (FGF) and Indian Hedgehog (Ihh)/parathyroid hormone-related peptide (PTHrP) signaling, two important pathways in chondrocyte proliferation and differentiation. The expression of PTHrP, PTHrP-receptor, Bcl-2, FGF2, FGFR1, FGFR3, and p21 is investigated by immunohistochemistry in osteochondromas ($n = 24$) and peripheral ($n = 29$) and central ($n = 20$) chondrosarcomas. Ihh/PTHrP and FGF signaling molecules are mostly absent in osteochondromas. Although no somatic *EXT* mutations were found in sporadic osteochondromas, the putative *EXT* downstream targets are affected similarly in sporadic and hereditary tumors. In chondrosarcomas, re-expression of FGF2, FGFR1, PTHrP, Bcl-2, and p21 is found. Expression levels increase with increasing histological grade. Up-regulation of PTHrP and Bcl-2 characterizes malignant transformation of osteochondroma because PTHrP and Bcl-2 expression is significantly higher in borderline and grade I peripheral chondrosarcomas compared with osteochondromas. In contrast, up-regulation of PTHrP and Bcl-2 seems to be a late event in central cartilaginous tumorigenesis because expression is mainly restricted to high-grade central tumors. (*Lab Invest* 2000, 80:1925–1933).

Chondrosarcoma of bone is a malignant tumor characterized by the formation of cartilage and occurring principally in adults in the third to sixth decade. It is the second most frequent primary malignant bone tumor after osteosarcoma. The majority of chondrosarcomas (75%) are located within the medullary cavity (central chondrosarcoma), whereas 15% develop within the cartilage cap of a pre-existing osteochondroma (peripheral chondrosarcoma) (Mulder et al, 1993; Springfield et al, 1996). Although there are no apparent cytonuclear differences between central and peripheral chondrosarcomas, we previously demonstrated different genetic mechanisms to be operative in the origins of central and peripheral chondrosarcomas (Bovée et al, 1999a). Both subtypes are histologically classified into three grades, correlating with prognosis (Evans et al, 1977). Recurrent chondrosarcomas may exhibit a higher grade of malignancy than the original neoplasm (Bjornsson et al, 1998; Evans et al, 1977), suggesting that tumors may progress in grade.

Malignant transformation of solitary osteochondroma is low (< 1%), whereas it is estimated to occur in 1% to 5% of cases of hereditary multiple exostoses (HME), a familial skeletal disorder with an autosomal dominant mode of inheritance (Schmale et al, 1994; Wicklund et al, 1995). HME is genetically heterogeneous, and at present, two responsible genes, *EXT1* and *EXT2*, located respectively at 8q24 and 11p11-p12, have been isolated (Ahn et al, 1995; Stickens et al, 1996; Wuyts et al, 1996). We have previously demonstrated *EXT1* to be operative as a tumor-suppressor gene during hereditary osteochondroma development according to Knudson's two-hit model (Bovée et al, 1999b). Surprisingly, no somatic *EXT1* cDNA alterations have been found in sporadic tumors (Bovée et al, 1999b).

The human *EXT1* and *EXT2* genes both encode endoplasmic reticulum-localized type II transmembrane glycoproteins that possess or are tightly associated with glycosyltransferase activities involved in heparan sulfate polymerization (Lind et al, 1998; McCormick et al, 1998; Simmons et al, 1999). It was recently suggested that EXTL2, a related protein, is the key enzyme for the chain initiation of heparan sulfate (Kitagawa et al, 1999), after which an *EXT1*/*EXT2* complex performs chain polymerization (Kitagawa et al, 1999; McCormick et al, 1999, 2000).

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Heparan sulfate proteoglycans (HSPG) are large macromolecules composed of heparan sulfate glycosaminoglycan chains linked to a protein core, which participate in cell signaling pathways (McCormick et al, 1999). For instance, HSPGs are required for high-affinity interactions of fibroblast growth factor (FGF) with its receptor (FGFR) (Erlebacher et al, 1995; Goldfarb, 1996). Furthermore, an *EXT1* homolog (*tout-velu*, *Ttv*) in *drosophila*, which is also involved in heparan sulfate biosynthesis, is required for diffusion of an important segment-polarity protein called Hedgehog (Hh), a homolog of mammalian Indian Hedgehog (Ihh) (Bellaïche et al, 1998; The et al, 1999; Toyoda et al, 2000).

The orderly proliferation and maturation of chondrocytes in longitudinal columns within the normal growth plate (Fig. 1A) is regulated by a delicate paracrine feedback loop involving both Ihh and FGF signaling (Fig. 1, B and C) (van der Eerden et al, 2000). Chondrocytes in transition from the proliferative to the hypertrophic zone start to secrete Ihh, which binds to its receptor, Patched (Ptc). This results in increased secretion of parathyroid hormone-related peptide (PTHrP) via an as yet incompletely resolved mechanism. Binding of PTHrP to its receptor at the late-proliferating chondrocytes (Erlebacher et al, 1995) inhibits further differentiation by a direct or indirect up-regulation of the anti-apoptotic protein Bcl-2. This results in fewer Ihh-producing cells, which closes the feedback loop (Amling et al, 1997). Thus, PTHrP regulates the pace of chondrocyte differentiation by delaying the progression of chondrocytes towards the hypertro-

phic zone, allowing longitudinal bone growth (van der Eerden et al, 2000).

Ihh signaling is repressed when the FGF-receptor 3 (FGFR3), which is expressed at the proliferative zone of the growth plate, is activated by a presently unknown ligand (Erlebacher et al, 1995; Goldfarb, 1996; Naski et al, 1998). Furthermore, FGFR3 inhibits chondrocyte proliferation via induction of expression of p21^{WAF1/CIP1}, which is an inhibitor of the cell cycle (Sahni et al, 1999). In contrast to FGFR3, FGF2 (basic FGF) is the most potent mitogen for chondrocytes in vitro and stimulates extracellular matrix synthesis via its receptor (eg, FGF-receptor 1) (Iwamoto et al, 1991; Kato and Iwamoto 1990).

The stratified zones of the growth plate are less well regulated but can sometimes still be recognized in osteochondroma. The recent elucidation of the function of the *EXT* genes led us to the hypothesis that inactivation of both copies of *EXT(1)* in normal cells of the growth plate (Bové et al, 1999b) will result in altered heparan sulfate expression at the cell surface of chondrocytes (McCormick et al, 1998), which will in turn affect FGF signaling by preventing high-affinity binding of FGF to its receptor (Erlebacher et al, 1995; Goldfarb, 1996). Furthermore, because the *EXT1* homolog in *drosophila* is required for diffusion of Hedgehog (Bellaïche et al, 1998; The et al, 1999; Toyoda et al, 2000), *EXT1* inactivation in humans may also affect Ihh/PTHrP signaling by preventing the diffusion of Ihh. We have therefore investigated the protein expression of these putative downstream effectors of EXT by immunohistochemistry in a well-characterized group of osteochondromas and chondrosarcomas.

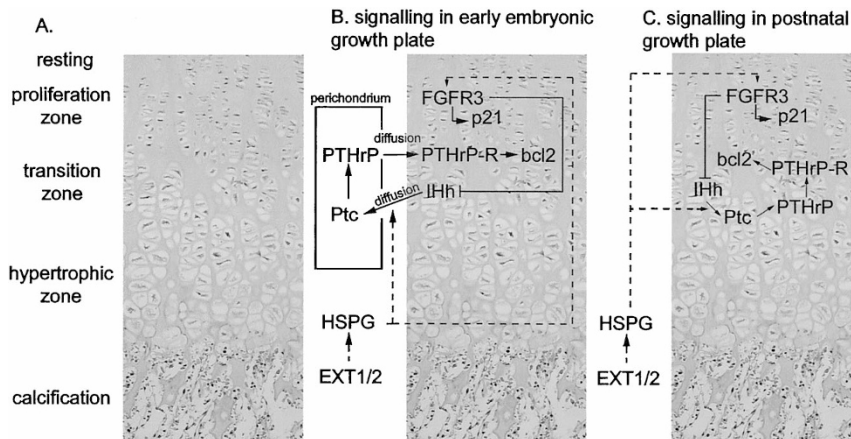


Figure 1.

A, Light-micrograph of a normal postnatal growth plate that is characterized by the orderly proliferation and maturation of chondrocytes in longitudinal columns, forming stratified zones of resting, proliferative, prehypertrophic/maturing, and hypertrophic cartilage. Ossification begins with invasion of calcified hypertrophic cartilage by capillaries accompanied by apoptosis of terminal hypertrophic chondrocytes, resorption of cartilage matrix, and deposition of bone matrix by osteoblasts. B, Growth signaling in the early embryonic growth plate is depicted within the photograph. Chondrocytes in the transition zone secrete Indian hedgehog (Ihh), which has to diffuse to the lateral perichondrium where its receptor, Patched (Ptc), is expressed. Binding will induce up-regulation of parathyroid hormone-related peptide (PTHrP) at the apical perichondrium, which then diffuses to its receptor localized at late-proliferating chondrocytes. Via up-regulation of Bcl-2, further differentiation of late proliferating chondrocytes is inhibited, resulting in fewer Ihh-producing cells, which closes the feedback loop. Activation of fibroblast growth factor receptor-3 (FGFR3), expressed at the proliferative zone, by an as yet unknown ligand inhibits chondrocyte proliferation via up-regulation of p21^{WAF1/CIP1} and represses Ihh signaling. C, Growth signaling in the postnatal growth plate. The Ihh/PTHrP feedback loop is now confined to the growth plate. Ihh will bind Ptc in the hypertrophic zone, stimulating PTHrP expression. PTHrP then binds its receptor in the prehypertrophic zone, up-regulating Bcl-2. Defective or absent EXT proteins leading to altered or absent heparan sulfate proteoglycans (HSPG) expression at the cell surface may affect high-affinity binding at the FGFR3 receptor and may disturb Ihh diffusion towards the apical perichondrium (early embryonic growth plate) or to its neighboring cells (postnatal growth plate).

Results

Immunohistochemistry

Expression of FGF2, FGFR1, p21, PTHrP, and Bcl-2 was found in 25% to 83% chondrosarcomas. In contrast, only 0% to 41% of osteochondromas expressed these antigens (Table 1). Examples are shown in Figures 2 and 3. To the extent that the stratified zones that are found in the normal growth plate could still be recognized in positive osteochondroma cases, no spatial distribution of protein expression was found. The expression of the PTHrP-receptor and FGF-receptor 3 was almost equally distributed over osteochondromas and chondrosarcomas.

Some immunostained sections could not be evaluated because of the loss of attachment to the glass slides or the absence of staining of an internal positive control. In the latter situation, prolonged decalcification may have altered the configuration of the antigen, and these cases were therefore excluded from the analysis. This was mainly a problem in osteochondromas containing a lot of bony material requiring long decalcification. Results of all cases that could be evaluated are summarized in Table 1.

Statistical Analysis

Osteochondroma versus Peripheral Chondrosarcoma. The expression of FGF2, FGFR1, p21, PTHrP, and Bcl-2 was found in a significantly higher percentage of peripheral chondrosarcomas compared with osteochondromas (Table 1). A further comparison between the group of osteochondromas and a group of exclusively borderline and grade I peripheral chondrosarcomas revealed a significantly higher percentage expressing PTHrP ($p = 0.003$) and Bcl-2 ($p = 0.001$) (Fisher's Exact Test) in the latter group (borderline and grade I peripheral chondrosarcomas)(Fig. 3). PTHrP expression was found in 7 of 12 borderline and grade I chondrosarcomas, but in only 1 of 18 osteochondromas, resulting in a sensitivity of 58% and a specificity of 94%. Five of 9 borderline and grade I

chondrosarcomas expressed Bcl-2 compared with none of 19 osteochondromas, resulting in a sensitivity of 56% and a specificity of 100%.

Peripheral versus Central Chondrosarcoma. Expression of PTHrP and Bcl-2 was also significantly higher in peripheral compared with central chondrosarcomas ($p = 0.031$ and $p = 0.017$, respectively). Remarkably, in central chondrosarcomas, expression of PTHrP and Bcl-2 was mainly restricted to high-grade cases (Table 2, Fig. 3), with the exception that one borderline central chondrosarcoma expressed both proteins.

Sporadic versus Hereditary Cases. There were no differences in expression of any molecule tested between HME and non-HME peripheral tumors (Fisher's Exact test). Considering the level of expression, however, only Bcl-2 expression was stronger in non-HME cases compared with HME cases ($p = 0.031$, Chi-square test, linear by linear). For the level of expression of the other factors, no differences were found. Furthermore, within the group of osteochondromas, there were no differences in expression between patients before and after the age of 15 years or between male and female patients.

Correlation with Prognosis. The expression levels of FGF2 ($p = 0.039$), FGFR1 ($p = 0.000$), FGFR3 ($p = 0.043$), p21 ($p = 0.002$), PTHrP ($p = 0.039$), and Bcl-2 ($p = 0.011$) increased with increasing histological grade (Chi square test, linear by linear). Only the expression of p21 was correlated with a shorter disease-free survival ($p = 0.0198$, Log Rank Test), although the effect of this parameter on disease-free survival was not independent of histological grading ($p = 0.2443$, p21; $p = 0.0520$, histological grade, Cox regression analysis).

Correlation between the Expression of the Different Proteins. Considering the delicate paracrine growth signaling in which the investigated molecules are normally involved (Fig. 1), a correlation between the level of expression of the separate molecules investigated was tested using the Chi square test, linear by linear. Indeed, for most proteins the level of expression strongly correlated with the level of expression of

Table 1. Results of Immunohistochemistry

Antigen	Osteochondroma		Osteochondroma vs peripheral CS p value ^b	Peripheral CS		Peripheral vs central CS p value ^b	Central CS	
	pos ^a	%		pos	%		pos	%
FGF-2	2/17	12	p = 0.044	12/27	44	p = 0.767	9/18	50%
FGFR-1	9/22	41	p = 0.017	21/27	78	p = 0.721	15/18	83%
FGFR-3	4/21	19	p = 0.517	8/28	29	p = 0.737	4/19	21%
p21	5/20	25	p = 0.009	17/26	65	p = 0.539	10/19	53%
PTHrP	1/18	6	p = 0.000	23/29	79	p = 0.031	9/19	47%
PTHrP-R	6/19	32	p = 0.543	11/26	42	p = 0.540	6/20	30%
Bcl-2	0/19	0	p = 0.000	15/24	63	p = 0.017	5/20	25%

FGF-2, fibroblast growth factor 2; FGFR-1, fibroblast growth factor receptor 1; PTHrP, parathyroid hormone-related peptide; PTHrP-R, PTHrP receptor.

Results of the immunohistochemical analysis are shown for osteochondromas, peripheral chondrosarcomas (CS), and central chondrosarcomas.

^a pos: the number of positive tumors/total number of tumors that could be evaluated.

^b Results within the group of osteochondromas are compared to results within the group of peripheral chondrosarcomas for each antigen tested using the Fisher's Exact Test. Similarly, results within the group of peripheral chondrosarcomas are compared to the results within the group of central chondrosarcomas. Significant p values are shown in bold. For PTHrP and Bcl-2, differing in expression between central and peripheral chondrosarcomas, results per tumor grade are shown in Table 2.

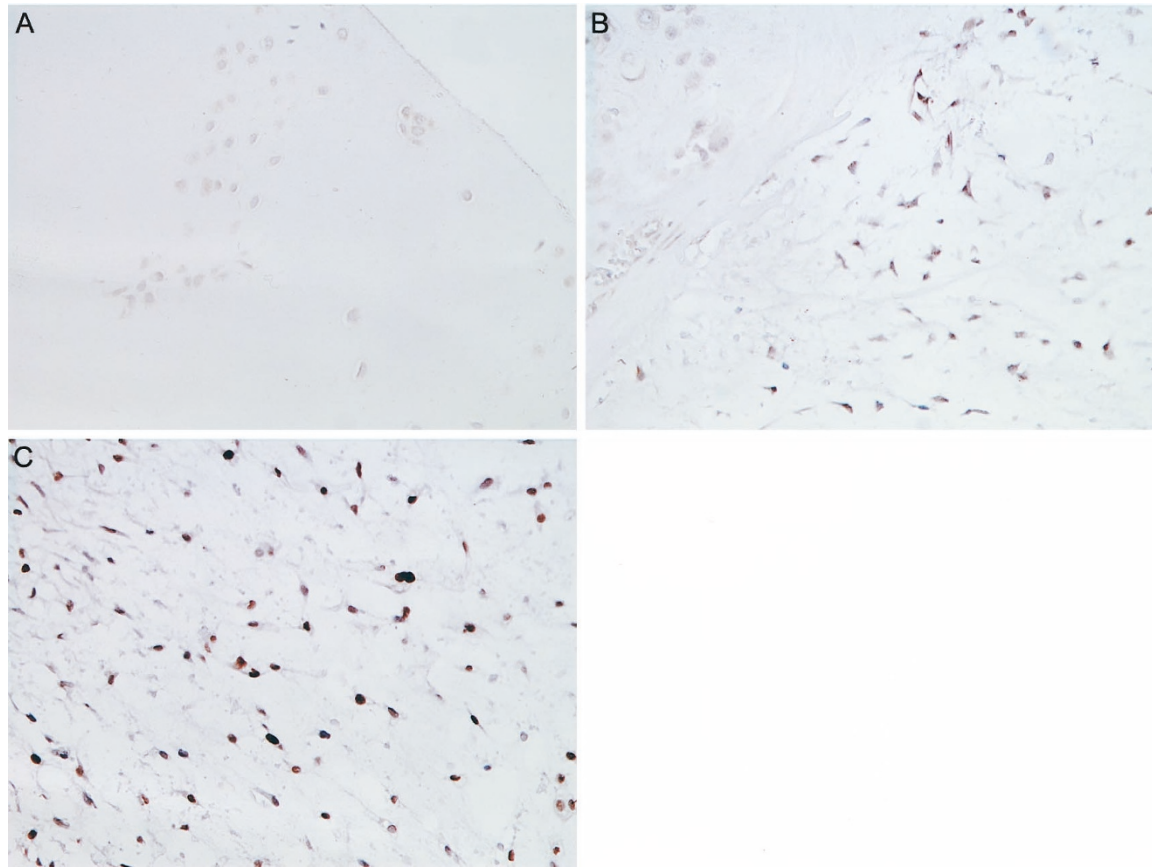


Figure 2.

A, FGFR1 expression within osteochondroma. The remaining cartilaginous cap (not shown) has low cellularity without FGFR1 expression, whereas this part of the cartilaginous cap, with increased cellularity, demonstrates FGFR1 expression, suggesting local progression. B, Grade III peripheral chondrosarcoma demonstrating cytoplasmic FGFR3 expression with moderate intensity, found in 50% to 75% of the tumor cells. C, Grade III peripheral chondrosarcoma demonstrating strong nuclear p21 expression, found in 50% to 75% of the tumor cells.

other proteins. For instance, strong Bcl-2 expression was correlated with a strong expression of all other molecules (p value range: 0.000–0.007, Chi-square test, linear by linear). A correlation was lacking only between FGF2 and p21 ($p = 0.051$), FGFR3 and p21 ($p = 0.394$), PTHrP and PTHrP-R ($p = 0.249$), and FGFR3 and PTHrP-R ($p = 0.069$).

Discussion

We studied the expression of seven proteins that are known to be involved in normal chondrocyte growth and differentiation (Fig. 1) and that are predicted to be affected by EXT inactivation in cartilaginous tumors. We previously demonstrated inactivation of both copies of *EXT1* in hereditary osteochondromas (Bovée et al, 1999b). The involvement of EXT1 and EXT2 proteins in heparan sulfate biosynthesis (McCormick et al, 1998) and Hedgehog diffusion (Bellaïche et al, 1998) was shown recently. This leads to the hypothesis that EXT inactivation will affect both Ihh/PTHrP signaling by preventing the diffusion of Ihh (Bellaïche et al, 1998; The et al, 1999; Toyoda et al, 2000) and FGF signaling by preventing HSPG-mediated high-affinity binding of FGF to its receptor (Erlebacher et al, 1995; Goldfarb, 1996) (Fig. 1).

Table 2. Results of Bcl-2 and PTHrP Immunohistochemistry in Chondrosarcomas, According to Histological Grade

	Peripheral CS		Central CS	
	pos ^a	%	pos ^a	%
PTHrP				
Borderline	0/2	0	1/3	33
Grade I	7/10	70	4/9	44
Grade II	9/10	90	3/5	60
Grade III	7/7	100	1/1	100
Metastasis	—	—	0/1	0
Bcl-2				
Borderline	0/1	0	1/3	33
Grade I	5/8	63	0/9	0
Grade II	5/8	63	2/6	33
Grade III	5/7	71	1/1	100
Metastasis	—	—	1/1	100

^a pos: the number of positive tumors/total number of tumors that could be evaluated.

We demonstrate that the growth regulators involved in both Ihh/PTHrP and FGF signaling are absent, or only lowly expressed in a minority of cells, in osteo-

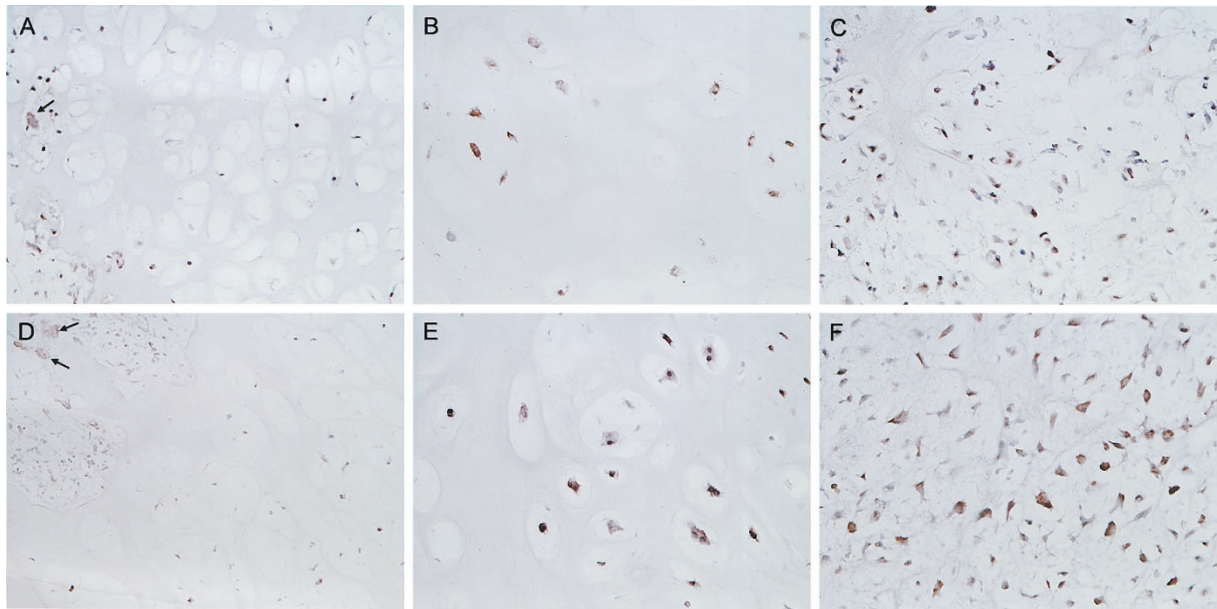


Figure 3.

A, Low-power view of a sporadic osteochondroma demonstrating weak PTHrP expression found in only 1% to 25% of the tumor cells, which is therefore scored as negative. Note the strong staining of the osteoclast, used as an internal positive control. B, Grade I peripheral chondrosarcoma secondary to a sporadic osteochondroma demonstrating PTHrP expression with moderate intensity, found in 25% to 50% of the tumor cells. C, In central chondrosarcoma, PTHrP expression was mainly restricted to high-grade tumors. A grade III central chondrosarcoma is shown with staining of moderate intensity, found in 50% to 75% of the tumor cells. D, Low-power view of an osteochondroma from a patient demonstrating hereditary multiple exostoses syndrome with weak Bcl-2 expression found in only 1% to 25% of the tumor cells, which is therefore scored as negative. Note strong positive staining of the osteoclast, used as internal positive control. E, Same grade I peripheral chondrosarcoma as shown in B, clearly demonstrating moderate Bcl-2 expression, found in 50% to 75% of the tumor cells. F, Same grade III central chondrosarcoma as shown in C, clearly demonstrating strong Bcl-2 positive staining in 75% to 100% of the tumor cells.

chondromas. Although both the PTHrP receptor and the FGF receptor 1 are detected in 32% and 41% of osteochondromas, respectively, their ligands PTHrP and FGF2 are detected in only 6% and 12% of osteochondromas, respectively. Others demonstrate the absence of FGF-2 expression in enchondromas, whereas all chondrosarcomas studied are positive with varying intensity (Uria et al, 1998). Investigations in *drosophila* demonstrate that defects in the *EXT1* homolog *tout velu* specifically affect Hedgehog diffusion, whereas other HSPG-dependent pathways, such as FGF or Wingless signaling, are unaffected (The et al, 1999). Our data however suggest that in human osteochondromas, Ihh/PTHrP and FGF signaling are affected equally, and often simultaneously. Alternatively, because osteochondromas cease growing when the growth plates close, it may be that with puberty most of these growth-signaling molecules are down-regulated in osteochondroma in a physiological manner under hormonal control. However, we did not find differences in (the level of) expression of all factors in osteochondromas before and after the age of 15.

In a previous study, we failed to detect any somatic mutations in *EXT1* or *EXT2* in sporadic osteochondromas and chondrosarcomas, despite loss of heterozygosity (LOH) at the *EXT1* locus at 8q24 (Bovée et al, 1999b). So far, only one somatic *EXT1* mutation in a sporadic chondrosarcoma has been described in the literature (Hecht et al, 1997). These data may imply that another mechanism is operative in sporadic compared with hereditary osteochondroma development. In the present study, we show, however, that the

putative downstream effectors of EXT are affected similarly in sporadic and hereditary osteochondromas and peripheral chondrosarcomas, because we did not find any differences in expression. This suggests that, in sporadic osteochondroma development, other genetic aberrations affecting the same EXT pathway are operative.

Interestingly, re-expression of most signaling molecules in chondrosarcomas is demonstrated, because the expression of FGF2, FGFR1, p21, PTHrP, and Bcl-2 is significantly higher in peripheral chondrosarcomas compared with osteochondromas. Furthermore, the levels of expression of all factors, apart from the PTHrP receptor, increase with increasing histological grade. This re-expression of growth factors normally involved in cartilage growth and differentiation is probably responsible for the proliferation and cell survival in chondrosarcomas. Whether this re-expression after malignant transformation is EXT-dependent or not remains to be elucidated. However, because of the fact that in hereditary osteochondromas EXT is inactivated irreversibly (Bovée et al, 1999b), the re-expression is most probably induced by other signals. Perhaps, growth stimulation becomes autocrine instead of paracrine, avoiding extracellular HSPG-dependent signaling.

The distinction between benign tumors and those of low-grade malignancy is considered difficult at both the clinico-radiological (Geirnaerd et al, 1997) and histological levels (Mirra et al, 1985), and so far the diagnosis is usually based on a combination of clinical, radiological, and histological findings. We demon-

strate expression of PTHrP and Bcl-2 in a significantly higher percentage of borderline/grade I peripheral chondrosarcomas compared with osteochondromas. This suggests that up-regulation of the expression of these two proteins characterizes malignant transformation of osteochondroma. Although the sensitivity is relatively low (58% and 56%), a specificity of 94% (PTHrP) and 100% (Bcl-2) is found because expression, especially of Bcl-2, is mostly absent in osteochondromas. Positive Bcl-2 immunostaining seems, therefore, to be indicative of malignancy. Additional studies, including double stainings and other methodologies like mRNA in situ hybridization, should be performed on a larger series of osteochondromas and low-grade chondrosarcomas to determine whether the detection of Bcl-2 and/or PTHrP can be used in the differential diagnosis between benign and low-grade malignant peripheral chondrosarcomas in the future.

We previously demonstrated a diverging genetic mechanism to be operative in the origin of central and peripheral chondrosarcoma (Bové et al, 1999a). Peripheral chondrosarcomas are characterized by genetic instability as demonstrated by a broad range in DNA ploidy and a high percentage of loss of heterozygosity involving many different chromosomes. In contrast, central chondrosarcomas are mostly peridiploid and have limited LOH (Bové et al, 1999a). In the present study, expression of PTHrP and Bcl-2 was found in a significantly higher percentage of peripheral chondrosarcomas compared with central chondrosarcomas, suggesting that up-regulation of these two proteins is highly specific for malignant transformation during peripheral cartilaginous tumorigenesis. In contrast, in central chondrosarcoma, up-regulation of PTHrP and Bcl-2 seems to be a late event, because expression is mainly found in high-grade tumors. Amling et al (1998) reported a high co-expression of PTHrP and Bcl-2 only in grade II and grade III chondrosarcomas without subdivision in central and peripheral chondrosarcoma. These data suggest that, in

central cartilaginous tumorigenesis, up-regulation of PTHrP and Bcl-2 is a late event characterizing the progression from low-grade towards high-grade chondrosarcoma. Thus, these data substantiate that different genetic pathways are involved in the genesis of central and peripheral chondrosarcoma.

In conclusion, we investigated the protein expression of molecules that are normally involved in chondrocyte proliferation and differentiation and that are predicted to be affected by EXT inactivation. Our data suggest that indeed both IHH/PTHrP and FGF signaling are affected during osteochondroma development. Although we previously failed to detect any somatic *EXT* mutations in sporadic osteochondroma, we here demonstrate that the putative *EXT* downstream targets are affected similarly in sporadic and hereditary peripheral tumors. In chondrosarcomas, re-expression of most signaling molecules is found, and although the up-regulation of PTHrP and Bcl-2 seems to be a late event in central chondrosarcoma, malignant transformation of osteochondroma towards low-grade peripheral chondrosarcoma is characterized by the up-regulation of these two molecules.

Materials and Methods

Patient Data

Formalin-fixed, paraffin-embedded material was available for 73 tumors (osteochondroma, $n = 24$; peripheral chondrosarcoma, $n = 29$; and central chondrosarcoma, $n = 20$) from 64 patients. Six hereditary osteochondroma cases were retrieved from other laboratories using PALGA (Dutch National Information System for Pathology, Utrecht, The Netherlands), whereas all other cases originated from the files of the Leiden University Medical Center (Leiden, The Netherlands), a tertiary bone tumor referral center. Dedifferentiated, mesenchymal, juxtacortical, and clear-cell chondrosarcomas were excluded because of their distinctly different clinico-pathological features. Pa-

Table 3. Clinical and Tumor-Related Data

	Osteochondroma 20 patients (24 tumors)	Peripheral CS 26 patients (29 tumors)	Central CS 18 patients (20 tumors)
Male/female	15/5	14/12	6/12
Median age at diagnosis	20.7 yr (range 4–49)	39.8 yr (range 16–68)	37.6 yr (range 14–63)
Histological grade			
Borderline		2	3
Grade I		10	9
Grade II		10	6
Grade III		7	1
Metastasis			1
Hereditary multiple exostoses	11/18 ^a	7/20 ^a	—
Median follow-up	—	46 mo (range 13–128)	52 mo (range 16–192)

CS, chondrosarcoma.

^a Patients with inconclusive data are omitted.

tient data, shown in Table 3, were obtained by review of pathology specimens and reports, clinical charts, and radiographs. Central and peripheral chondrosarcoma were separated based upon accepted clinicopathological and radiological criteria (Huvos, 1991). For peripheral chondrosarcoma, a pre-existing osteochondroma or an identifiable stalk was documented either radiographically or by gross pathology. Multiplicity of osteochondromas, which was considered the criterion for HME because an unequivocal family history was not always available, could be assessed for 38 patients. Histological grading was performed according to the method of Evans et al (1977). Five tumors were diagnosed as borderline chondrosarcoma because they showed histological features that were not sufficient to diagnose grade I chondrosarcoma, although their X-rays showed features of an aggressive neoplasm (Unni, 1996).

Antibodies and Controls

Details of the available antibodies and controls used are described in Table 4. The commercially available antibodies against IHH and its receptor, Ptc (Santa Cruz Biotechnology, Santa Cruz, California), were shown not to work reliably in our hands, either on fresh-frozen or formalin-fixed, paraffin-embedded material, and were therefore not used in this study. The FGFR-1 antibody was kindly provided by Dr. J. Walters (Oxford Brookes University, Oxford, United Kingdom), and the characteristics of this antibody have been described (Bovée et al, 1998; De-Boer et al, 1996). To avoid false-negative results, all antibodies used were tested for their susceptibility to formalin fixation. Immunohistochemistry was performed on appropriate tissues fixed for 1, 3, 7, and 40 days, respectively. None of the antibodies revealed diminished staining reactivity or staining sensitivity. Moreover, for most of the antibodies used, internal positive controls were present in the histological slides (Hughes, 1997; Hughes and Hall, 1993; Qu et al, 1995; Reed et al, 1995) (Table 4) to determine whether negative tumor cells are truly negative or whether prolonged decalcification might have altered the configuration of the antigen, resulting in a negative internal control. As negative controls, slides were incubated with mouse or rabbit immunoglobulin G (IgG) of a corresponding (iso-)type and concentration instead of with primary specific antibodies.

Immunohistochemistry

Immunohistochemical reactions were performed according to standard laboratory methods as described previously (Bovée et al, 1998). After deparaffinizing, rehydration, and blocking of endogenous peroxidase, antigen retrieval was performed as detailed in Table 4, followed by an overnight incubation with the primary antibodies. Biotin-labeled rabbit antimouse or swine antirabbit immunoglobulins and a biotinylated HRP-Streptavidin complex (DAKO, Glostrup, Denmark) were applied. Visualization was carried out in a diami-

Table 4. Details of the Purchase of Antibodies and Protocols Used

Antigen	Manufacturer	Mono/poly-clonal	Staining	Positive control	Internal positive control in bone tissue	Dilution	Antigen retrieval
FGF-2 (Clone 6)	Transduction Laboratories	Monoclonal	Nuclear	Normal tonsil and skin	Osteoblasts, blood vessel walls (Qu et al., 1995), mast cells (Reed et al., 1995)	1:125	Citrate
FGFR-1	Kindly provided	Monoclonal	Cytoplasmic	Normal skin	Osteoblasts, blood vessel walls (Hughes and Hall 1993)	1:2000	None
FGFR-3	Sigma	Polyclonal	Cytoplasmic	Normal skin	Striated muscle, vessel walls (Hughes 1997), connective tissue osteoblasts	1:2000	Trypsin
p21 ^{WAF1/CYP1} (Clone AB-1)	Calbiochem	Monoclonal	Nuclear	Normal colon	None (osteoblasts and vessels occasionally positive)	1:200	Citrate
PTHrP	Oncogene	Polyclonal	Cytoplasmic	Normal skin	None (osteoclasts occasionally positive)	1:25	Trypsin
PTH/PTHrP-Receptor Bcl-2 (Clone 124)	Babco Boehringer	Polyclonal Monoclonal	Cytoplasmic/nuclear Cytoplasmic	Normal skin Normal tonsil	Vessel walls, osteoblasts Osteoclasts, lymphocytes	1:500 1:100	Citrate Citrate

nobenzidine solution (Sigma, St. Louis, Missouri) with the addition of 0.01M Imidazole for FGFR-1. Hematoxylin was used for counterstaining the slides.

Evaluation and Scoring

Staining was evaluated by two observers independently. Scoring was performed as described (Bové et al, 1999a, 1999c; Detre et al, 1995; Elkhuizen et al, 1999) with some minor modifications. In brief, both staining intensity (0, no staining; 1, weak; 2, moderate; 3, strong intensity, as related to a positive internal control) and the percentage of positive tumor cells (0, 0%; 1, 1% to 24%; 2, 25% to 49%; 3, 50% to 74%; 4, 75% to 100% positive tumor cells) were evaluated, and scores of intensity and percentage positivity were added up. In general, a score greater than 3 was considered positive (Elkhuizen et al, 1999), with the exception of p21 and FGFR1, for which a score greater than 0 was considered positive. For instance, for PTHrP and FGF2, almost all tumors demonstrated variable expression leading to a cut-off at a score greater than 3, whereas expression of p21 and FGFR1 was completely absent in about half of the tumors, allowing a cut-off at a score greater than 0. The observers were blinded towards all clinicopathological data.

Statistical Analysis

Immunohistochemical results of peripheral chondrosarcomas were compared to osteochondromas as well as to central chondrosarcomas using Fisher's Exact test. In the same way, a comparison was made between the group of osteochondromas and a group of only borderline and grade I peripheral chondrosarcomas. Sensitivity and specificity were calculated for PTHrP and Bcl-2 to determine whether their expression could differentiate between osteochondroma and low-grade peripheral chondrosarcoma. Also, HME cases were compared to non-HME cases using Fisher's Exact Test. To test the prognostic value of the investigated molecules, a correlation between histological grade and the level of positivity was investigated using the Chi-square test, linear by linear. To test a correlation with disease-free survival, the Log Rank Test was performed for all variables separately. The Cox regression analysis was used to test whether certain parameters have an independent effect on disease-free survival. Correlation of the levels of expression of molecules with each other was tested using the Chi-square test, linear by linear. *p* values equal to or less than 0.05 were considered to be significant.

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