

# Acromegaly: Re-thinking the cancer risk

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**Abstract** Acromegaly is characterized by sustained elevation of circulating growth hormone (GH) and insulin-like growth factor I (IGF-I), and is clearly associated with increased morbidity and overall mortality mainly due to cardiovascular, metabolic, and respiratory diseases. Although cancer-related mortality varies widely amongst retrospective studies, it appears to be consistently elevated mainly in patients with uncontrolled disease. We review individual tumor types including neoplasms of the colon, breast, prostate, and thyroid where *in vitro*, animal studies, and studies in non-acromegalic cancer patients have established a role for the GH/IGF-I axis in tumor progression and possibly initiation. We highlight deficiencies in data in acromegalic patients where the evidence is less convincing. Instead, we explore the hypothesis that acromegaly, independent of hormone secretion, is a disease that heralds genetic and/or epigenetic alterations predisposing to cancer risk elsewhere.

**Keywords** Acromegaly · Growth hormone/insulin-like growth factor-I axis · Cancer risk · Genetic susceptibility

## 1 Cancer mortality in acromegalic patients

Retrospective studies focusing on mortality in acromegaly have shown that 9–50% of deaths in acromegalic patients are attributable to cancer related complications (average 24% in 5,024 patients with acromegaly) (Table 1). While earlier studies by Wright et al. [1], Alexander et al. [2], Nabarro [3], Bengtsson et al. [4], and Etxabe et al. [5] demonstrated a 2–3 fold increased overall mortality for acromegalic patients, cancer related mortality rates were either not increased [6] or increased only in females (2–3 fold, [1, 3, 4] or in males (5–7 fold, [2, 5]). If the cancer entities were documented, they covered a wide array of different malignancies but cancers of the breast, colon, prostate, and hematological system were the most frequent types. Most of the subsequent studies focusing on mortality analyzed overall rates in comparison to sex- and age-matched population based on historical controls. It is important to note that more recent studies examined mortality rates in relationship to post-treatment growth hormone (GH) and /or insulin-like growth factor I (IGF-I) levels. Such studies were able to demonstrate that mortality rates were comparable to population-based controls if post-treatment GH levels (either nadir or random) were below threshold cut-offs suggestive of “cure” or “in-remission” and were elevated in patients with persistent disease [7–14]. At least two studies also found a correlation between post treatment IGF-I levels and mortality rates [9, 13] while two others [12, 14] did not detect an effect of IGF-I levels on outcome. The authors of these studies suggest that increased mortality risk in acromegalic patients is restricted mainly to those with elevated post-treatment GH levels. It is of interest to note that the largest cohort of patients studied by Orme et al. [10] as well as those studied by Holdaway et al. [13] demonstrated increased mortality rates for all

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**Table 1** Retrospective studies focusing on mortality in acromegalic patients

Author	Number of patients/deceased patients/cancer related deaths	Prevalence of cancer related deaths in %	SMR	Conclusion for overall mortality and cancer related mortality	Types of cancer (Number of patients)	Analysis of mortality rates related to GH/IGF-I levels or persistent disease
Wright et al. 1970 [1]	194/55/10	18	Overall: SMR 1.93. Cancer related: SMR 1.39 (F: 2.05)	Overall: increased. Cancer related: increased in females	Colon (2), lung (3), stomach (2), thyroid (2), fibrosarcoma (1)	NA
Alexander et al. 1980 [2]	164/54/7	16	Overall: SMR 3.97. Cancer related: SMR 1.94 (M: 4.62)	Overall: increased. Cancer related: increased in males	NA	NA
Nabarro 1987 [3]	256/47/11	23	Overall: SMR 1.26 (F: 1.68). Cancer related: SMR 0.96	Overall: increased in females. Cancer related: not increased	NA for mortality	NA
Bengtsson et al. 1988 [4]	166/62/15	24	Overall: SMR not listed. Cancer related: SMR 2.68 (F: 3.3)	Overall: increased. Cancer related: increased in females	Breast (3), lung (2), kidney (1), stomach (1), prostate (1), leukemia (3), other (4)	NA
Eixabe et al. 1993 [5]	74/10/5	50	Overall: M: SMR 3.2. Cancer related: M: SMR 7.1	Overall: increased in males. Cancer related: increased in males	Lung (2), pancreas (1), prostate (1), osteosarcoma (1)	Mortality increased irrespective of cured vs. persistent disease
Bates et al. 1993 [7]	79/28/11	39	Overall: SMR 2.63	Overall: increased	NA	Mortality not increased in patients in remission
Rajasooriya et al. 1994 [8]	151/32/3	9	Overall: SMR not listed. Cancer related: SMR 1	Overall: increased. Cancer related: not increased	Breast (1), colon (1), chondrosarcoma (1), melanoma (1), mesothelioma (1)	Increased mortality with higher post treatment GH levels

Swearingen et al. 1998 [9]	149/12/4	33	Overall: SMR 1.16	Overall: not increased. Cancer related: NA	Leukemia (1), prostate (1), astrocytoma (1), other (1)	Mortality increased in persistent disease; mortality not increased in patients in remission
Orme et al. 1998 [10]	1,362/366/83	23	Overall: SMR 1.60. Cancer Related: SMR 1.16	Overall: increased. Cancer related: not increased	Breast (13), lung (13), colon (13), rectum (3), other (42)	Increased overall and cancer related mortality with higher post treatment GH
Drange et al. 2000 [189]	176/9/2	22	Overall: SMR 0.85	Overall: not increased. Cancer related: NA	NA	NA
Beaugreard et al. 2003 [11]	103/18/9	50%	Overall: remission SMR 0.88. Persistent disease SMR 4.82	Overall: increased. Cancer related: NA	Lung (5), ovary (1), leukemia (1), other (2)	Mortality increased in persistent disease; mortality not increased in patients in remission
Ayuk et al. 2004 [12]	419/95/21	22	Overall: SMR 1.26. Cancer related: SMR 0.91	Overall: increased. Cancer related: not increased	NA	Increased mortality with higher post treatment GH but not IGF-I levels
Holdaway et al. 2004 [13]	208/72/17	24	Overall: SMR 1.87. Cancer related: SMR 0.91	Overall: increased. Cancer related: not increased	Breast (2), colon (8), stomach (1), pancreas (1), lymphoma (1), other (4)	Increased mortality with higher post treatment GH and IGF-I levels
Mestron et al. 2004 [190]	1,219/56/9	16	NA	NA	NA for mortality	NA
Kauppinen-Mäkelin et al. 2005 [14]	334/56/12	21	Overall: SMR 1.16	Overall: not increased. Cancer related: NA	NA	Increased mortality with higher post treatment GH but not IGF-I levels
Total	5,054/972/219	23	Overall: SMR 2.05. Cancer related: 1.37			

SMR = standardized mortality ratio, F = female, M = male, NA = not analyzed

cancers only in acromegalic patients with higher post-treatment GH levels while cancer-related mortality for all acromegalic patients regardless of treatment outcome was not increased compared to population controls. Orme found a 1.8 fold increased mortality risk of all cancers, a 4.6 fold increased mortality risk of colon cancer and a 2.9 fold increased mortality risk of breast cancer if random serum GH levels were higher than 10 ng/ml. On the other hand there was no significant relationship between duration of acromegaly and cancer related mortality in these patients.

It should also be noted that many authors have argued that most deaths amongst acromegalic patients are attributable to cardiovascular disease (60%) followed by respiratory causes (25%). Correspondingly, analyses of mortality determinants in patients with persistently elevated GH levels show that hypertension and heart disease, and not cancer, are the strongest predictors of mortality [15, 16].

Cancer related mortality data, on the other hand, does not reflect cancer incidence or cancer morbidity. This along with the potential bias of cause of death reporting from autopsy examinations, we will argue that studies focusing on mortality are not likely to be helpful in answering the question of cancer risk.

## 2 Overall cancer incidence in acromegalic patients

Unfortunately, studies focusing on cancer incidence or cancer prevalence in acromegaly do not provide conclusive evidence of increased cancer risk (Table 2). Most studies report a moderately (1.5 to 4-fold) increased relative risk for acromegalic patients to develop tumors, mainly of the colon/rectum, breast, prostate, thyroid, and hematological system [3–24] while other studies do not confirm increased tumor prevalence or incidence [6, 10].

The study by Mustacchi and Shimkin, for example, analyzing 223 patients treated for acromegaly from 1937 to 1955 [6] did not find an increased cancer incidence compared to population tumor incidence rates (13 vs. 10 malignancies, respectively). In 1982, Klein et al. [17] studied 44 patients and reported 11 cases of cancer 4 of which involved the colon. Although the observed rates were not compared to expected rates, the control group was comprised of 29 patients with chromophobic pituitary adenomas. Subsequently, Nabarro [3] reported in his personal series an increased cancer prevalence in women, mainly due to breast cancer (11 observed vs. 2.6 expected) while men showed no difference in cancer incidence rates. Brunner et al. [18] were able to demonstrate an increased risk only for colon cancer in 52 acromegalic patients compared to population tumor incidence rates (standardized incidence ratio, SIR, of 4.7 for colon cancer). Barzilay et al. [19], Cheung & Boyages [21], Popvich et al. [22], and

Higuchi et al. [23] described an increased cancer incidence in their smaller series of acromegalic patients (with patient numbers varying between 44 and 220) compared to population tumor incidence rates (SIR 2.45, SIR 2.5, SIR 3.39, SIR3.53, respectively) but the statistical power of such studies has been questioned [10].

The studies by Ron et al. [20], Orme et al. [10], and Baris et al. [24], analyzed large retrospective epidemiological cohort data. Ron et al. reviewed hospital records of 1,041 acromegalic male patients admitted to the United States Veteran Affairs (VA) hospitals and after excluding 87 patients with a diagnosis of cancer prior to the diagnosis of acromegaly documented 116 cases of cancers (including 13 colon cancers) and an increased tumor incidence compared to tumor incidence rates from 3.7 million first admissions to VA hospitals (SIR 1.6, SIR for colon cancer 3.08). This study was criticized for the inclusion of 27 cases of pituitary tumors as cancer manifestations; also, the prevalence of acromegaly greatly out-numbered that in the general population. Instead of using population tumor incidence rates to calculate the expected numbers of tumors the authors utilized non-acromegalic hospitalized patients as controls.

The important study by Orme et al. analyzed a large cohort of 1,239 acromegalic patients, yielding 16,778 person-years at risk. After excluding 16 patients with the diagnosis of cancer before the diagnosis of acromegaly 79 cancers were observed in the patient group while 104 were expected from calculations utilizing data from the national cancer register for England and Wales (SIR 0.76). Nevertheless, sub-analysis of the incidence of different tumor types revealed a non-significant increased incidence of colon cancers in acromegalic patients (SIR 1.68).

Baris et al., on the other hand, found an increased risk for all cancers in 1,634 patients from Sweden and Denmark compared to population tumor incidence rates (SIR 1.5). Risks were increased for all gastroenteropancreatic (GEP) cancers (as well as colon and rectum separately), kidney cancers, thyroid, and breast in females while prostate cancer was not significantly increased [24].

Epidemiologic analyses of cancer incidence/prevalence and cancer mortality in patients with acromegaly is hampered by several factors. Acromegaly is a rarely diagnosed disease with a reported prevalence of 40 to 90 cases per million population, which makes it difficult for a single center to gather sufficient numbers of patients for statistical analysis [25]; this may lead to biased reporting by specialized centers limiting generalization to the larger community. Secondly, as the main causes of death are cardiovascular and respiratory events, patients with uncontrolled acromegaly might succumb before developing recognizable cancer. Indeed, nearly 50% of patients in studies from the 1940s–1970s died before the age of 60 years. In this regard it remains to be shown conclusively

if current standards will change cancer incidence in the acromegalic population.

Another important consideration is that acromegaly study patients were included in cancer incidence estimates only if they presented with an initial diagnosis of acromegaly. Patients presenting with a malignancy as the primary diagnosis have been traditionally excluded (see Table 2). In the case of the Orme et al. study 16 patients with cancer and subsequent diagnosis of acromegaly were excluded accounting for 17% of all cancer patients. Ron et al. excluded 87 patients with an initial diagnosis of a malignancy before diagnosis of acromegaly (49% of all patients with cancer). Although this exclusion is justified to evaluate the effects of increased GH and/or IGF-I levels on cancer development it may obscure the alternate hypothesis; namely that acromegaly and cancer share a common unifying non-endocrine etiology.

We conclude that the data generated from retrospective studies suggest an increased overall tumor incidence/prevalence in patients with acromegaly but additional prospective studies are required to resolve the significance of these observations.

## 2.1 Colorectal cancer

Colorectal tumors constitute the most frequent malignancy associated with acromegaly. The summary risk ratio for all colorectal cancer estimated from the three largest retrospective population-based studies by Ron et al. [20], Orme et al. [10], and Baris et al. [24] is 2.04 (95% CI: 1.32, 3.14) suggesting a modestly increased risk [26]. Prospective studies utilizing colonoscopy or flexible sigmoidoscopy with barium examination have been performed to evaluate the risk of colon polyps and colorectal cancer and are listed in Table 3. Most of these studies included only a small number of patients and the controls vary between in-house non-acromegalic patients, e.g. subjects with irritable bowel syndrome, and models generated from screening colonoscopy and/or autopsy studies. Three of the four larger studies including more than 100 acromegalic patients each demonstrate an increased risk of colon adenomas with an adenoma prevalence of 22 to 26% [27–29] while two of them show an increased risk for colon carcinoma with a prevalence of 4 to 5% [28, 29]. The study by Renehan et al. on the contrary, did not confirm either the increased prevalence of adenomas or carcinomas in acromegalic patients if compared to either literature data of 2,559 bowel examinations in autopsy studies or 810 bowel examinations performed in screening colonoscopy studies [30]. Overall, the data presented in colonoscopy studies are problematic mainly because of inadequate controls (e.g. autopsy studies tend to present a higher frequency of adenomas in the asymptomatic general population due to “optimal” exami-

nation conditions; predominance of males in some colonoscopy screening studies also leads to a higher incidence of adenomas/carcinomas in the general population; age and environmental influences are important factors which have not been taken into account when comparing prevalence in acromegalic vs. controls). Therefore, the data suggesting an increased prevalence/incidence of colon adenomas in acromegalic patients have to be interpreted with caution. The available population-based data are more consistent. Renehan et al. [26] estimated in their meta-analysis that acromegaly is associated with a moderate (2-fold) increase in colorectal cancer risk. This estimate is increasingly accepted by other investigators and consensus working groups [15, 26, 31–36].

There is furthermore evidence that colorectal neoplasia in acromegaly has different characteristics compared with that in the general population. Adenomatous lesions are more likely to be right sided (35 to 68% of the lesions) [27, 30, 34]; they tend to be larger and more dysplastic [28, 34, 35]; the adenomas are often multiple [27]. When resected there is a 25 to 41% chance of recurrence within 3 years and this seems to increase with higher GH/IGF-I levels [29, 36].

In patients with acromegaly the length of the colon and sigma is generally greater than in controls, and epithelial cells of sigmoid crypts present a slightly enhanced pattern of proliferation which is positively correlated to circulating IGF-I levels and an decreased apoptotic activity [30, 37, 38]. It has been speculated that GH/IGF-I excess in acromegaly increases epithelial cell proliferation and decreases apoptosis leading to greater susceptibility to accumulation of genetic defects resulting in colon cancer [16, 39]. Abnormalities in lymphocyte subset patterns in the colon mucosa of acromegalic patients and increased serum and intra-colonic unconjugated deoxycolic acids may represent an alternate pathophysiological mechanism of colonic neoplasia. The latter has been associated with the development of colon cancer in the general population and may also contribute to a tumor favorable environment in acromegalic patients [35, 40, 41].

### 2.1.1 The role of GH/IGF-I in tumor initiation and progression

Circumstantial evidence supporting a role of GH/IGF-I in the development of adenomas and cancers of the colon arises from cell and animal models and data in non-acromegalic patients with colon cancer.

The actions of GH are mediated by the GH receptor, which is ubiquitously expressed in colonic epithelium [42]. GH is mitogenic and anti-apoptotic in many tissues [43, 44], but there is relatively little data on its action in colonic tissue. Instead, IGF-I-based studies have demonstrated that

**Table 2** Retrospective studies focusing on cancer prevalence/incidence in acromegalic patients

Author	Number of patients/cancers	Cancer prevalence/incidence	SIR	Conclusion for cancer prevalence/incidence	Types of cancer (number of patients)	Patient inclusion criteria
Mustacchi & Shimkin 1957 [6]	223/13	6	SIR 1.33. F: SIR 1.34. M: SIR 1.31	Not increased <sup>a</sup>	Breast (2), lung (3), FGT (2), leukemia (1) skin (3), osteosarcoma (1), angiosarcoma (1)	All acromegalic patients included (regardless if cancer was diagnosed prior to acromegaly)
Klein et al. 1982 [17]	44/11	25	Tumor prevalence in controls 10%	Increased <sup>b</sup>	Colon (4), breast (2), kidney (1), thyroid (1), parathyroid (1), FGT (2), thymoma (1)	All acromegalic patients included
Nabarro 1987 [3]	256/26	10	F: SIR 1.87 (SIR for breast cancer 4.23). M: SIR 1	Increased in females <sup>a</sup>	Breast (11), lung (3), GEP (2), prostate (1), thyroid (1), ovary (2), kidney (1), hematological (2)	All acromegalic patients included
Ritchie 1990 [124]	125/15	12	NA	Suggestive of increase <sup>a</sup>	Breast (2), lung (3), GEP (7), kidney (1), bladder (2)	All acromegalic patients included
Brunner et al. 1990 [18]	52/7	14	SIR for colon cancer 4.7	Increased for colon cancer <sup>a</sup>	Breast (1), lymphoma (1), FGT (2), colon (2), adrenal (1)	Only patients with primary diagnosis of acromegaly
Barzilay et al. 1991 [19]	87/7	8	SIR 2.45	Increased <sup>a</sup>	Breast (1), sarcoma(1), thyroid (2), ovary (1), colon (1), leukemia (1)	Only patients with primary diagnosis of acromegaly
Ron et al. 1991 [20]	1,041/89	9	SIR 1.23. SIR for colon 3.08	Increased for gastrointestinal tumors including colon <sup>b</sup>	Lung (22), colon (13), other GEP (14), skin (3), thyroid (1), male genital and urinary (15), brain (5), hematological (7), other (9)	Only patients with primary diagnosis of acromegaly (87 patients excluded with primary diagnosis of cancer)

Cheung & Boyages 1997 [21]	50/7	14	SIR 2.5. F: SIR 4.17. M: SIR 1.25	Increased <sup>a</sup>	Breast (2), kidney (2), colon (1), parotid (1), prostate (1)	Only patients with primary diagnosis of acromegaly (1 patient excluded with cancer)
Orme et al. 1998 [10]	1,239/79	6	SIR 0.76	Not increased <sup>a</sup>	Breast (14), lung (6), colon (12), rectum (4), thyroid (1), others (42)	Only patients with primary diagnosis of acromegaly (16 patients excluded with cancer)
Popovich et al. 1998 [22]	220/23	9	SIR with in-house control 3.21. SIR with population control 3.39	Increased <sup>a,b</sup>	Breast (4), ovary (1), cervix (4), kidney (2), colon (2), pancreas (2), skin (2), thyroid (3), hematological (3)	Only patients with primary diagnosis of acromegaly
Higuchi et al. 2000 [23]	44/5	11	M: SIR 3.53	Increased in males <sup>a</sup>	Colon (1), stomach (1), bladder (1), thyroid (2)	Only patients with primary diagnosis of acromegaly
Baris et al. 2002 [24]	1,634/177	11	SIR 1.52. SIR colon 2.6, rectum 2.5, thyroid 3.7, kidney 3.2, bone 13.8, brain 2.7	Increased <sup>a</sup>	Colon (23), other GEP (36), hematological (9), lung (14), breast (20), FGT (9), skin (13), prostate (13), kidney (12), other (28)	Only patients with primary diagnosis of acromegaly
Mestron et al. 2004 [190]	1,219/90	7	NA	NA	Breast (23), colon (15), lung (5), gastric (4), other (43)	Only patients with primary diagnosis of acromegaly
Total	6,234/549	9	SIR 1.78			If all patients with primary cancer included: total number of patients with cancer 653 (11%)

SIR = standardized incidence ratio, F = female, M = male, NA = not analyzed, FGT = female genital tract, GEP = gastroenteropancreatic

<sup>a</sup>Compared to population tumor incidence rates

<sup>b</sup>Compared to in-house controls

**Table 3** Retrospective and prospective studies screening for colorectal neoplasia in acromegalic patients

Author	Number of patients/ control participants	Adenoma prevalence		Cancer prevalence		Comments on controls	Conclusion for prevalence rate adenoma/carcinoma
		Acromegaly	Controls	Acromegaly	Controls		
Klein et al. 1982 <sup>b</sup> [17]	17/-	5 (29)	NA	2 (12)		NA specifically for colon cancer	Suggestive of increase for adenoma/carcinoma
Iuarte et al. 1984 <sup>a,b</sup> [191]	12/-	2 (17)	-	3 (25)	-	Population cancer incidence rates	Increased/increased
Pines et al. 1985 <sup>a</sup> [192]	48/-	2 (4)	-	3 (6)	-	Population cancer incidence rates	Increased/increased
Brunner et al. 1990 <sup>a</sup> [18]	29/-	3 (10)	-	2 (7)	-	Population cancer incidence rates	Increased/increased
Ezzat et al. 1991 <sup>b</sup> [193]	23/-	8 (35)	-	0 (0)	-	Compared with literature	Increased/not increased
Ladas et al. 1994 <sup>b</sup> [194]	54/-	5 (9)	-	0 (0)	-	Compared with literature	Not increased/not increased
Vasen et al. 1994 <sup>b</sup> [34]	49/57	11 (22)	5 (9)	0 (0)	0 (0)	In-house non-acromegalic patients	Increased/not increased
Delhougne et al. 1995 <sup>b</sup> [27]	103/138	23 (22)	11 (8)	0 (0)	0 (0)	In-house non-acromegalic patients	Increased/not increased
Colao et al. 1997 <sup>b</sup> [35]	50/318	11 (22)	18 (6)	1 (2)	4 (1)	In-house non-acromegalic patients	Increased/not increased
Jenkins et al. 1997 <sup>b</sup> [28]	129/-	34 (26)	-	6 (5)	-	Models from screening colonoscopy/population	Increased/increased
Renehan et al. 2000 <sup>b</sup> [30]	115/-	14 (12)	-	3 (3)	-	Models from autopsy/ screening colonoscopy	Not increased/not increased
Terzolo et al. 2005 <sup>b</sup> [29]	235/233	55 (23)	34 (15)	10 (4)	2 (1)	In-house non-acromegalic patients	Increased/increased
Matano et al. 2005 <sup>b</sup> [195]	19/76	8 (42)	13 (17)	3 (16)	2 (3)	In-house non-acromegalic patients	Increased/increased
Total (prospective studies)	806/822	176 (22)	81 (10)	28 (4)	8 (1)		

Values in parentheses are percentages

NA = not analyzed

<sup>a</sup> Retrospective study

<sup>b</sup> Prospective study

both normal and tumoral colorectal cancer cell lines express the IGF1-R, and that IGF-I induces proliferation and anti-apoptotic activity in colorectal cancer cell lines [45–47].

Animal studies involving the over expression of GH or IGF-I distinguish two different murine phenotypes of hypersomatotropism [48, 49]. The selectively over expressed GH transgene is associated with both high GH and IGF-I levels while the IGF-I transgenic mouse has high IGF-I but low GH levels [48, 49]. Although both models exhibit biochemical and phenotypical features of acromegaly, the IGF-I transgenic mouse exhibits an increased bowel length with highly proliferative colonic crypt cells and decreased apoptosis, whereas no proliferative or apoptotic changes were present in the crypt cells of the GH transgenic model [48–51]. Intriguingly evidence to date suggests that IGF-I transgenic mice do not show an increased incidence of bowel cancer while GH transgenics do so [52]. A liver-

specific gene-deletion of IGF-I results in reduced circulating IGF-I levels and a reduction in cancer growth and metastases in three cancer models, one for colon cancer and two for breast cancer [53]. Serum IGF-I concentrations vary between individuals across the general population, and it has been postulated that this may impact upon cancer risk [54]. A recent meta-analysis of a number of prospective studies on colon cancer risk has pooled the data and estimated a summary odds ratio of 1.58 (CI:1.11–2.27) for the association of the uppermost categories of serum IGF-I concentration compared to the lowermost categories [55].

In summary, there appears to be an increased prevalence of colorectal neoplasms in acromegaly where regular screening and polypectomy is advisable. Total colonoscopy should be performed as acromegalic patients more frequently develop cancers in the ascending and transverse colon. Predisposing features include multiple skin tags, a



positive family history, increasing age, and persistently elevated GH and IGF-I levels [56].

Remaining questions include the age and frequency at which colonoscopy should be performed. As suggested by different authors we agree that colonoscopic screening is recommended to all newly diagnosed patients and that thereafter the American Cancer Society guidelines for patients with increased risk are applicable (colonoscopy every 5 years if no cancer or polyps are detected, with more frequent follow-up if any lesions are detected, depending on the number, size, and histologic features of the lesions) [33, 57–59]. While some have recommended even more intensive screening and follow-up programs [59], others have argued that acromegalic patients should not receive special screening or follow-up treatment as the modest increase in relative risk for colon cancer does not warrant special screening or surveillance measures. In fact guidelines for familial colorectal cancer screening suggest that colonoscopic surveillance is only warranted for a lifetime risk of 1 in 10 or greater [60, 61].

## 2.2 Breast cancer

Determining the incidence of breast cancer in women with acromegaly is even more difficult to ascertain than of other tumors as the restriction to the female sex nearly halves the number of patients at any single centre. Only one study by Nabarro [3] has reported a 4-fold higher risk for breast cancer, while subsequent studies failed to confirm a significantly increased incidence in patients with acromegaly [10, 24].

In contrast to that of colon cancer, *in vitro* data, animal studies, and epidemiologic studies in non-acromegalic patients have implicated the GH/IGF-I axis in the development of breast cancer. The GH-R is expressed in human mammary tumors (predominantly in the epithelial cell but also to some extent in the stromal cells) and human breast cancer cell lines [62]. The additional finding of GH protein expression in mammary tumors as well as normal tissue appears to suggest a possible autocrine/paracrine role for GH on tumor development and growth [63]. In line with this hypothesis, studies in MCF-7 mammary cancer cells have shown that blockade of GH-R by a specific GH antagonist abrogates cell proliferation [64].

The IGF-IR is also expressed on primary cultures of malignant breast epithelial cells as well as in several estrogen-dependent and, at lower levels, in estrogen-independent cell lines and IGF-I causes marked proliferation of breast cancer cells which can be inhibited by anti-IGF-I-R antibody [65–67]. A paracrine as well as an autocrine IGF-I/IGF-IR loop has been demonstrated in stromal cell—breast cancer cell interactions and in breast cancer cell lines [68–70]. Early studies involving chronic administration of

GH in female rats demonstrated an increased development of breast tumors over the long term [71]. In addition, transgenic mice over-expressing human GH [72] or IGF-I [73] have displayed an increased incidence of mammary tumors. Conversely, *lit/lit* mice with a defective GH-releasing hormone receptor and thus decreased GH/IGF-I levels show a significant growth inhibition of transplanted human breast cancer cells [74].

Furthermore, epidemiologic studies in non-acromegalic patients with breast carcinoma have suggested that these patients have significantly higher serum GH and IGF-I levels compared to individuals without carcinoma [75, 76] and that high IGF-I tertiles alone predict a 4.6 fold higher risk, and in combination with a low IGFBP3, a 7.3 fold higher risk for premenopausal but not postmenopausal breast cancer [77]. The meta-analysis by Renehan et al. [55] estimated a summary odds ratio of 1.93 (CI:1.38–2.69) for the association of the uppermost categories of serum IGF-I concentration compared to the lowermost categories in premenopausal breast cancer, but did not find an association between IGF-I levels and postmenopausal breast cancer. Unexpectedly, the authors found a positive association between IGFBP-3 and premenopausal breast cancer.

We conclude that until further studies are available to evaluate the true incidence, it is advisable to offer periodic mammography to women over the age of 50, as in the general population.

## 2.3 Prostate cancer

Even though acromegaly is associated with benign prostate hyperplasia as well as micro- and macro-calcifications regardless of age or gonadal status [78–79], the incidence of prostate cancer in patients with acromegaly is not significantly elevated and may even be reduced [10, 20–24]. It has been argued that this reduction may be seen because the disease is limited to elderly men and only recently men with acromegaly have survived long enough to participate in epidemiologic studies [80]. Again, *in vitro*, animal and non-acromegalic prostate cancer patient studies have implicated the involvement of the GH/IGF-I axis in prostate cancer.

Primary cultures of tumor prostate epithelial cells, as well as androgen sensitive and androgen insensitive prostate tumor cell lines all express significant levels of IGF-IR [81, 82]; increased expression of the IGF-IR is associated with progression of androgen-sensitive prostate cancer cell lines to androgen independence [83, 84]; treatment with IGF-I causes cell proliferation which is independent of the presence of androgens in the androgen-insensitive but dependent on androgens in the androgen-sensitive cell lines and which is inhibited by IGF-IR antisense oligonucleotide transfection [82, 85]. In concor-

dance with these observations are the findings that injection of PA-III prostate adenocarcinoma cells transfected with an IGF-IR anti-sense construct into rats resulted in a 90% decrease in tumor growth compared to control transfected cells [86]. Not only prostate epithelial cells but also stromal cells derived from benign prostatic hyperplasia express IGF-IR mRNA and IGF-I stimulation increases stromal cell density by 80%, an effect abrogated by co-incubation with IGF-IR neutralizing antibodies providing a possible explanation for the above mentioned association of acromegaly with benign prostate hyperplasia [87].

Transgenic mice expressing human IGF-I in basal epithelial cells of prostate demonstrate a stepwise development of prostatic adenocarcinomas [88].

Finally, in retrospective population studies, prostate cancer is associated with IGF-I levels in the higher normal range [89, 90]. Matzoros et al. estimated an odds ratio of 1.9 for a serum IGF-I increment of 60 ng/ml and also demonstrated that in the presence of higher serum testosterone levels the relative risk for prostate cancer increased even further in regard to serum IGF-I increments [89]. A prospective study demonstrated that serum IGF-I levels in the upper quintile of the normal range were associated with a significantly increased 4-fold risk of developing prostate cancer [91] and also with a 5-fold increased risk of advanced stage prostate cancer [92]. The meta-analysis of Renehan et al. estimated a summary odds ratio of 1.83 (CI:1.03–3.26) for the association of the uppermost categories of serum IGF-I compared to the lowermost categories [55].

It is of interest to point out, that acromegaly is also associated with increased liver synthesis of IGFBP-3, which has IGF-independent (as well as IGF-dependent) pro-apoptotic and anti-tumoral effects at the local level. In non-acromegalic patients with prostate cancer, levels of IGFBP-3 are often found to be decreased in the serum and in prostatic tissue [93, 94]. Because IGFBP-3 is a substrate for PSA, a member of the kallikrein family of serine proteases, it has been postulated that rising levels of PSA during prostate cancer evolution facilitates disease progression by proteolytically cleaving IGFBP-3, thereby increasing bioavailable IGF-I at the cellular level [95]. As IGFBP-3 proteolysis is not elevated in patients with acromegaly it has been speculated that this might present a protective mechanism in these patients against the development of prostate cancer [16].

In conclusion, we suggest that men with acromegaly should be screened for prostate cancer as in the general population using serum PSA measurement, rectal examination and/or prostatic ultrasound annually. Patients with high IGF-I receiving replacement testosterone should be monitored with special attention even though the data, thus far, does not confirm an increased risk in acromegalic men.

## 2.4 Thyroid cancer

Acromegaly is clearly associated with diffuse and nodular goiter, the prevalence of which is dependent on the diagnostic methods ranging from 78–92% using ultrasound [96–102] and 11–87% using palpation [19, 99, 103]. Most studies show to some degree a positive relationship between GH/IGF-I levels and thyroid volume [96–98, 101]. The majority of patients have normal thyroid function; hyperthyroidism ranges between 4 and 14% [100–103].

Retrospective studies focusing on overall cancer incidence thyroid cancer constitutes 3.1% of malignancies in acromegaly. Larger studies examining the relative risk for cancer subtypes demonstrate a 4.3 increased [20], 2.5 increased [10], and 3.7 increased [24] relative risk for thyroid cancer, but it has to be taken into account that the patient numbers with thyroid cancer were very low (1 to 3 patients). Retrospective studies focusing on thyroid disease in acromegaly demonstrated either no thyroid cancer in 37 to 80 acromegalic patients [97, 98, 103], or an increased thyroid cancer prevalence compared to the general population [100, 102, 104]. The study by Tita et al. [102] reported a 60-fold increased prevalence of thyroid carcinoma in patients with acromegaly (5.6% of their 125 acromegalic patients had thyroid cancer compared to 0.1% of a general control population in iodine deficient area). Reported types of thyroid cancer include papillary, follicular-variant papillary and follicular carcinomas.

*In vitro* studies have demonstrated not only that IGF-I increases proliferation of porcine thyroid cells and rat FRTL-5 thyroid cells but also potentiates TSH-mediated thyroid cell proliferation [105]. Furthermore, the existence of a local autocrine IGF-I loop is suggested by the increased presence of IGF-IR in follicular and papillary cancer cells and tumor specimens [106, 107] and by the synthesis of IGF-I in follicular thyroid cells [108], papillary cell lines [109], and surgical thyroid specimens [110].

In view of the high prevalence of nodular goiter in acromegaly and the potential roles of IGF-I and IGFBP3 in malignancy we suggest that thyroid nodules in acromegalic patients should be monitored in accordance with the American Thyroid Association guidelines and that special emphasis should be placed on fine-needle aspiration of all non-functioning nodules 1 cm or greater in size [111].

## 2.5 Other tumors

As shown in Table 2 other tumor entities have been found in acromegalic patients. Similarly scattered case reports exist describing various tumors including brain tumors [112–114], osseous tumors [115], skin epidermoid tumors and melanomas [116], adrenal tumors [117–120], and

different lymphohematopoietic neoplasms (lymphoma, multiple myeloma, chronic myeloid or lymphocytic leukemia) [121–123].

In the early smaller studies it was also suggested that lung tumors occurred more frequently in patients with acromegaly [3, 6, 124], but the larger studies by Ron et al. [20], Orme et al. [10], and Baris et al. [24] did not confirm an increased prevalence. On the other hand, experimental studies have shown that normal and tumoral lung tissue, as well as small cell lung cancer (SCLC) and non-SCLC express the IGF-IR and produce IGF-I in an autocrine fashion [125–129]. Stimulation with IGF-I or over expression of the IGF-IR induces cell proliferation and increases the metastatic activity of lung cancer cells [130]. Furthermore, a large-scale genome-wide association study in UK Caucasians undertaken to identify susceptibility alleles for lung cancer identified variants in the GH/IGF-I pathway associated with the inherited predisposition to this cancer [131]. Additional evidence is provided by a case-control-study in non-acromegalic lung cancer patients, which demonstrated a positive association between IGF-I levels and lung cancer risk and an inverse relationship for IGFBP3 [132]. Conversely, in the meta-analysis by Renehan et al. IGF-I was not associated with an increased risk for lung cancer [55].

### 3 Pathogenesis of tumor development in acromegaly

#### 3.1 The GH/IGF-I axis (endocrine/autocrine theory)

Endocrine GH secretion occurs in a pulsatile manner from the anterior pituitary gland to mediate somatic growth through membrane bound growth hormone receptor (GHR), a member of the cytokine receptor superfamily [133] (for review see [134]). Upon binding of GH receptor dimers undergo a conformational change [135] bringing two Janus kinases 2 (JAK2) which are associated with the intracellular domain of the receptor into close association that then autophosphorylate [136]. Subsequently, these kinases phosphorylate the GHR and activate the signal transducers and activators of transcription (STAT) proteins, in particular STAT5, which translocates to the nucleus and transactivates specific genes such as c-fos, insulin, and acid-labile subunit [137]. STAT 5b binding sites have also been demonstrated in the IGF-I promoter and shown to regulate IGF-I transcription in a GH-dependent manner [138, 139]. GH receptor activation is also mediated via the phosphatidylinositol 3'-kinase (PI3K) and Ras/MAPK pathways (for review see [140, 141]).

Elevated circulating GH levels, independent of IGF-I, have mitogenic and anti-apoptotic effects in many tissues through transcriptional regulation of different signaling

pathways (reviewed in [142]) including AKT [143], inhibition of the pro-apoptotic caspase-3 protein [144], and activation of nuclear factor (NF)- $\kappa$ B, PI3K [145] and p70S6K, a kinase involved in the control of cell proliferation and differentiation [146]. Negative regulators of GH signaling include suppressor of cytokine signaling (SOCS) -1, -2, -3 and CIS (cytokine inducible SH2-containing protein) [147]. Haplo-insufficiency of SOCS2 enhances the trophic actions of GH in the small intestine and promotes formation of colon polyps, thus suggesting that small variations in SOCS2 expression levels may significantly influence the impact of elevated GH in the intestine [148].

In addition to the classical endocrine actions of the GH/IGF-I axis, important paracrine and autocrine effects of locally produced GH and/or IGF-I are increasingly recognized. Autocrine GH increases MCF-7 cell proliferation, protects from apoptosis, promotes epithelial to mesenchymal transition and confers an invasive cell type [149, 150]. Forced expression of GH in MCF-10A cells, an immortalized but otherwise normal human mammary epithelial cell line, is sufficient for oncogenic transformation and tumor formation *in vivo* [151]. Importantly, GH administration was not transforming in this system highlighting the role of autocrine GH in this process.

The IGF system is comprised of three ligands (IGF-I, IGF-II, and insulin), multiple receptors (IGF-IR, IR, hybrids of IGF-IR and IR, and IGF-IIR), six IGF binding proteins (IGFBP), three families of IGFBP proteases, and nine growth-mediating factors (reviewed in [152]). IGF-I circulates in the serum bound to IGFBP3 and ALS as a complex, which acts both as a reservoir and regulator of local IGF-I distribution. IGF-I is released from IGFBP3 by proteolysis. The IGF-I cellular effects are achieved through the interaction of IGF-I with the IGF-IR, a transmembrane receptor tyrosine kinase. Ligand-binding to the IGF-IR results in a conformational change leading to transphosphorylation of one receptor subunit by the other. This is followed by recruitment of specific docking intermediates, including insulin-receptor substrate-1 (IRS-1), Shc and 14-3-3 proteins [153, 154]. These molecules link the IGF-IR to diverse signaling pathways, allowing induction of growth, transformation, differentiation and protection against apoptosis. IGF-IR activation or over expression is associated with an increased propensity for invasion and metastasis. This effect is mediated by multiple signaling intermediates including of IRS-1, which can disrupt E-cadherin from the actin cytoskeleton and promote  $\beta$ -catenin-mediated transcriptional activity [155]. Similarly, tumor cell motility and invasive potential are influenced by crosstalk between the IGF axis and integrins [156], and by IGF induced secretion of matrix metalloproteinases [157]. The finding that IGF-IR over expression confers an

invasive, metastatic phenotype in a murine model of pancreatic cancer highlights the importance of these functions [158]. Furthermore, IGF-IR inhibition is capable of inhibiting metastasis *in vivo* [159, 160].

We conclude that experimental and epidemiological studies provide robust evidence that elevated GH and/or IGF-I levels can contribute to tumor progression and possibly tumor initiation. However, elevated circulating levels of GH and IGF-I in acromegalic subjects may not necessarily lead to increased tumor development due to differences in the endocrine and autocrine/paracrine GH/IGF-I axis. Furthermore, elevated IGFBP3 which has proapoptotic and anti-proliferative effects, may alter tissue sensitivity to GH/IGF-I signaling [161]. On the other hand, a biological mechanism distinct from this frequently cited enhancement of the GH/IGF-I pathway may be operational in determining the risk of cancer in acromegaly.

### 3.2 The genetic/epigenetic theory

It is conceivable that factors that predispose to a GH-secreting pituitary tumor may also predispose to the development of different cancers. In this regard, a recent study utilizing the nationwide Swedish Family-Cancer Database to analyze familial risks for pituitary adenomas and associated tumors through parental and sibling probands is of interest [162]. While familial pituitary tumor syndromes were seldom (only three offspring-parent pairs showed concordant pituitary tumors), the results suggested an association between pituitary adenomas in the offspring and parental skin cancer (SIR 1.60; CI:1.13–2.21), leukemia (SIR 1.90; CI:1.31–2.66), and nervous system hemangiopericytomas (SIR 182, CI not listed). Among siblings there was a significant association between pituitary tumors and breast cancer (SIR 1.46; CI:1.02–2.01), while the risk of pituitary adenoma was marginally increased in individuals whose siblings were diagnosed with colorectal cancer (SIR 1.53; CI:0.69–2.92). The same authors reported earlier that pituitary tumors (SIR 2.2; CI:1.5–3.1) as well as other endocrine tumors (SIR 3.2; CI:1.82–4.96) were increased in patients first diagnosed with colorectal cancer [163, 164]. Similarly, the authors described an increased rate of second primary endocrine tumors but also non-endocrine neoplasia (including breast, renal, liver, nervous system cancers) after first endocrine tumors, even though the authors acknowledge, that some of these tumors occur in the setting of familial tumor syndromes, such as multiple endocrine neoplasia (MEN) 1/2, Neurofibromatosis, von-Hippel-Lindau disease [165].

The larger studies analyzing cancer incidence in acromegalic patients listed in Table 2 typically excluded patients in their analysis with a first primary non-pituitary tumor followed by a second primary somatotroph adenoma.

Ron et al. [20] excluded 87 patients in whom a non-pituitary tumor preceded the diagnosis of acromegaly accounting for nearly 50% of patients with acromegaly and tumors. Similarly, Orme et al. [10] excluded 16 patients with primary non-pituitary tumors while many of the other studies did not list the number of patients excluded. If a common genetic-epigenetic susceptibility is significant in the development of malignant tumors as well as pituitary somatotroph adenomas these previously excluded patients should be included in the analysis. It is tempting to speculate that if all patients were to be included into the statistical analysis the risk of malignant tumors in patients with acromegaly (occurring before or after the diagnosis of acromegaly) is remarkably higher than the estimated SIRs listed in Table 2.

The data supporting second primary tumors after first primary pituitary adenomas is of a much smaller size compared to the data on first and second non-endocrine cancers (see overview [166]). Indeed the latter category predicts that heritable factors and environmental influences may be responsible for the increased occurrence of second primary tumors following primary pituitary adenomas. Thus, we will next discuss potential common genetic events that might predispose to the development of a pituitary somatotroph adenoma as well as other neoplasms.

The first potential candidate gene is the recently described aryl hydrocarbon receptor interacting protein (*AIP*) gene. *AIP* interacts with the aryl hydrocarbon receptor and heat shock protein 90 dimer where it is involved in mediating cellular responses to environmental toxins, such as dioxin [167]. Vierimaa et al. [168] identified germline mutations in this gene in 16% of all patients diagnosed with pituitary adenomas in a population-based series from Northern Finland. Many patients lacked a strong family history of pituitary adenomas, thus suggesting that *AIP* is a low penetrance tumor susceptibility gene. The prevalence of *AIP* germ-line mutations also varies in different clinical settings. In unselected sporadic pituitary tumors, the overall prevalence seems to be low as shown by studies in patients from the US [169], from Japan [170], and in a patient cohort from France, Belgium, and Italy [171]. In familial isolated pituitary adenomas *AIP* mutations occur more frequently (up to 15%) [172, 173]. While the numbers of patients with *AIP* mutation are too small to evaluate the risk of developing other tumors, the incidence of somatic *AIP* mutations was analyzed in a Finnish series of 373 colorectal cancers, 82 breast cancers, and 44 prostate cancers [174]. Only two missense mutations were identified in colorectal cancer but not in healthy controls and the authors suggested that somatic *AIP* mutations are not common in these tumor entities. To prove this hypothesis, germline *AIP* mutations will have to be examined in the

context of additional common somatic changes in the pituitary and other affected organs.

Finally, in common with multiple other tumors, epigenetic modification is a feature of pituitary tumorigenesis and methylation-associated silencing of cell cycle regulators have been reported by different groups (reviewed in [175]). For example, the p16 member of the INK4 family blocks cell cycle progression in the G1 phase by specific binding to cyclin-dependant kinase 4 and 6 (CDK 4/6). If p16 is inactivated CDK4/6 binds to cyclin D and phosphorylates Rb, which stimulates entry into the S phase and activates cell proliferation [176, 177]. Sporadic pituitary adenomas frequently show loss of p16 due to promoter hypermethylation [178–180]. Loss of p16 by aberrant promoter methylation is also a frequent event in many other tumors, including breast, prostate, colon, liver, and lung [181–184]. Epigenetic events targeting p16 promoter in acromegaly and a wide range of tumors raise the interesting possibility that common environmental factors, such as folate deficiency, may contribute to acromegaly and other tumors. Folate deficiency appears to increase the risk of developing several malignancies including colorectal cancer [185], and changes in folate metabolism have also been implicated in pituitary adenoma formation [186]. Folate deficiency has been associated with overall DNA hypomethylation but with promoter DNA hypermethylation, especially of the p16 gene promoter thus potentially leading to decreased p16 levels and enhanced cell proliferation [187, 118].

Although the epigenetic mechanisms discussed here are speculative it presents an alternative integrated approach of how environmental factors may impose on genetic events thus endangering one individual to multiple neoplastic diseases including pituitary adenomas.

#### 4 Conclusions

While the majority of epidemiologic studies in acromegalic patients indicate a modest increase of cancer risk, other studies clearly do not support this notion. The true incidence and prevalence of neoplasia in patients with acromegaly is, therefore, difficult to determine on the basis of currently published studies. Deficiencies in these studies include patient numbers, inclusion criteria, selected control populations, and statistical methodologies.

Increasing evidence from *in vitro* data, animal studies and studies in non-acromegalic patients continue to strengthen the role of the GH/IGF-I axis in tumor development. Such evidence should provide the impetus for more data gathering from large prospective and even retrospective studies relying on tumor registries. Given the increasingly recognized risk of secondary tumors in patients

with primary neoplasia, the search for common early genetic and/or epigenetic changes in acromegaly will prove critical in examining cancer risk. Until firmer evidence proves otherwise, acromegalic patients should receive tumor-screening programs for patients at increased risk of cancer. Maximal and sustained GH and IGF-I normalization should be sought to prevent and reduce co-morbidities and normalize life expectancy.

#### 5 Key unanswered questions

The main question if acromegaly is associated with an increased cancer risk still remains controversial. Therefore, it is also currently disputable, if and how patients with acromegaly should be screened for secondary neoplasia. Large scale epidemiologic studies are still required to answer this question. Prospective evaluation of patients with pre-existing conditions should be considered.

The endocrine as well as non-endocrine mechanisms responsible for increased cancer risk deserve a more balanced evaluation. In particular, further studies examining possible epigenetic and genetic variations including multiple tumors from the same patients are required to support the contribution of non-endocrine mechanisms in cancer risk.

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