



A Survey of Telomerase Activity in Human Cancer

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Research on the association of the ribonucleoprotein enzyme, telomerase, with human cancer has expanded rapidly in recent years. Essentially all major types of cancer have been screened and the presence of telomerase activity has been detected in the vast majority of cases. In this article we provide a summary, in table form, of the current data. © 1997 Elsevier Science Ltd. All rights reserved.

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INTRODUCTION

OVER ESSENTIALLY the last 2 years, screening of most types of human cancers has established a very strong association between the presence of telomerase activity and malignancy, making the enzyme one of the most common tumour markers ([1–3] and the articles in this Special Issue by N.W. Kim, pages 781–786; K.-F. Norrback and G. Roos, pages 774–780). Telomerase is a specialised reverse transcriptase that synthesises telomeric DNA and thus contributes to the maintenance of functional telomeres (see the article in this Special Issue by G.B. Morin, pages 750–760). In humans, telomerase activity in association with stable telomeres appears to be restricted to the germ-line and possibly to cycling stem cells in self-renewing somatic tissues (see articles by S.E. Holt *et al.*, pages 761–766; K.-F. Norrback and G. Roos, pages 774–780). More differentiated but proliferation competent somatic cells do not express detectable telomerase activity and indeed undergo progressive telomere shortening with cell division. In most cancers and in stem cells of renewal tissues, telomerase activity levels generally correlate with the proliferation state of the cells. The presence of the enzyme is almost always required for unlimited proliferation (immortality) whereas its absence may dictate a finite lifespan (senescence). Studies with experimentally transformed and immortalised human cells support this hypothesis (see article by T.M. Bryan and R.R. Reddel, pages 767–773).

Consequently, a requirement for telomerase in tumorigenesis is predicated on the assumption that tumour growth depends on, among other factors, the acquisition of the immortal phenotype by the malignant cells. This article summarises the current literature on the presence of telomerase in human tumours. Although the fast pace of

research in this field may render the information rapidly outdated, we hope that the present survey of over 3500 tumour and control samples will aid basic researchers and clinical oncologists in evaluating the role of telomerase in cancer, and in designing future studies that will eventually determine the usefulness of the enzyme as a diagnostic or prognostic marker or as a target for cancer therapy.

We have restricted the data in the present article to papers that have been published or are in press up to December 1996 and have excluded abstracts and presentation at meetings. Cancers have been combined into tables according to tissue of origin or body site and, for most cancer types, results from several studies have been pooled. In the case of giant cell tumours of bone and of chordoma, the current availability of only single studies of very small size did not warrant presentation in a table format. Results in those studies indicated that telomerase activity was present in 5/5 giant cell tumours of bone [4] and 1/2 chordomas [5].

With very few exceptions, tumour samples were assayed by the PCR-based TRAP protocol ([6], also see article by N.W. Kim, pages 781–786). Since this assay detects telomerase activity in as few as 1–10 positive cells and 0.01% positive cells in a mixed population, the chances that tumour samples containing admixtures of normal and malignant cells would score as negative is greatly reduced. For this reason, negative samples from the few studies that used the less sensitive assay without PCR amplification of telomerase products, have been disregarded. As mentioned above, most normal somatic tissues lack detectable telomerase activity, making discrimination between positive and negative samples an easy task. In the case of cancers arising from normal tissues that express the enzyme, however, we considered positive only tumour samples that had telomerase levels significantly higher than the matched control tissue.

Samples for which relative levels of activity were not given have not been included. One caveat that pertains to tumour samples with very weak activity is the possible contamination with haematopoietic stem cells and inflammatory cells (e.g. activated lymphocytes), since these cells express low levels of the enzyme that may increase upon proliferation (see articles by S.E. Holt *et al.*, pages 761–766 and by K.-F. Norrback and G. Roos, pages 774–780). This latter fact also renders the

issue of positive and negative haematological malignancies, such as leukaemias, the most difficult to resolve, since matched control samples are not available and basal levels of activity vary among individuals in an age-dependent manner.

The following ten tables summarise the current data on telomerase activity in various types of human cancer. The reader is advised to refer to the original articles for details and discussion.

Table 1. *Head/neck and lung tissue*

Pathology	No. positive/ No. tested	% positive	Ref.
Normal oral mucosa (or rinses)	1/39	(3)	[7,8]
Oral rinses from HNSCC*	14/44	(32)	[7]
Premalignant lesions	25/46	(54)	[7–9]
Head and neck squamous cell carcinoma	112/130	(86)	[3,7–10]
Non-small cell lung carcinoma	98/125	(78)	[11]
Small cell lung carcinoma	15/15	(100)	[3,11]
Adjacent tissue	3/68	(4)	[11]

*Head and neck squamous cell carcinoma.

Table 2. *Gastrointestinal tract and pancreas*

Pathology	No. positive/ No. tested	% positive	Ref.
Gastric metaplasia ^a /adenoma	4/15	(27)	[12]
Gastric carcinoma	72/85	(85)	[12,13]
Adjacent tissue	4/86	(5)	[12,13]
Colorectal adenoma*	20/44	(45)	[3,12,14,15]
Colorectal carcinoma†	123/138	(89)	[3,12,14–16]
Adjacent and normal tissues ^a	58/231	(25)	[3,12,14,17]
Benign pancreatic lesion‡	0/11	(0)	[18]
Pancreatic carcinoma	41/43	(95)	[18]
Adjacent tissue	5/36	(14)	[18]
Pancreatic brushings			
Benign	0/4	(0)	[18]
Carcinoma	8/8	(100)	

^aVery weak activity relative to tumour samples, comparable to activity level in normal mucosa. Includes adenomatous polyps from familial adenomatous polyposis (FAP) patients, tubular adenomas, sporadic polyps with and without carcinoma.

†Hereditary polyposis and non-polyposis cases, sporadic cases with and without mutator phenotype.

‡Cystadenoma, adenoma, cystic tumour, hyperplasia.

Table 3. *Hepatic tissue*

Pathology	No. positive/ No. tested	% positive	Ref.
Normal tissue	0/15	(0)	[19,20]
Non-malignant liver disease*	43/148	(29)	[19–22]
Hepatocellular carcinoma	149/173	(86)	[3,19–22]
Adjacent tissue	1/50	(2)	[19–22]

*Chronic/active hepatitis, cirrhosis; activity is very weak compared to carcinoma.

Table 4. Breast

Pathology	No. positive No. tested	% positive	Ref.
Normal tissue	0/15	(0)	[3,23,24]
Breast fibrocystic disease	0/34	(0)	[24,25]
Breast fibroadenoma	11/40	(28)	[24-26]
Breast carcinoma <i>in situ</i>	9/12	(75)	[26,27]
Breast carcinoma (ductal and lobular)	300/339	(88)	[3,23-25]
Adjacent tissue	4/85	(5)	[24,25]
Breast FNA* benign	4*/47	(9)	[24,25]
Breast FNA malignant	26/32	(81)	[24,25]

*FNA, fine needle aspirates. *Positive samples from patients with fibroadenoma with Ki-67 positive ductal epithelia.

Table 5. Female reproductive tract

Pathology	No. positive No. tested	% positive	Ref.
Ovary (fetal)	2/2	(100)	[3]
Ovary (adult)	1/3	(33)	[3,28]
Unfertilised egg	0/3	(0)	[29]
Normal myometrium/endometrium	0/18	(0)	[3,30]
Leiomyoma	0/14	(0)	[3,30]
Leiomyosarcoma	5/5	(100)	[3,30]
Endometrial adenocarcinoma	4/4	(100)	[30]
Cervical carcinoma	16/16	(100)	[30]
Vaginal carcinoma	3/3	(100)	[29]
Ovarian carcinoma	21/23	(91)	[28,30]

Table 6. Male reproductive tract

Pathology	No. positive No. tested	% positive	Ref.
Testis (fetal/adult)	3/3	(100)	[3,29]
Mature spermatozoa	0/3	(0)	[29]
Normal prostate	0/43	(0)	[3,31,32]
BPH* without carcinoma	1/20	(5)	[3,32]
BPH with carcinoma	4/35	(11)	[31,32]
High grade PIN†	3/5	(60)	[3]
Prostate carcinoma	52/58	(90)	[3,31,32]

*Benign prostatic hyperplasia. † Prostatic intra-epithelial neoplasia.

Table 7. Kidney/urinary tract

Pathology	No. positive No. tested	% positive	Ref.
Normal urothelium	0/45	(0)	[33-36]
Normal voided urine	3*/83	(4)	[35]
Dysplastic urothelium	3/7	(43)	[33-35]
Bladder carcinoma (all stages)	172/185	(92)	[33-36]
Bladder carcinoma (washings)	29/40	(73)	[34]
Bladder carcinoma (voided urine)	16/56	(29)	[34,35]
Renal carcinoma	95/115	(83)	[37,38]
Adjacent tissue	0/115	(0)	[37,38]
Wilms' tumour	6/6	(100)	[3]
Adjacent tissue	2/6	(33)	[3]

*Positive samples were from patients with benign prostatic hyperplasia.

Table 8. Neural tissue

Pathology	No. positive/ No. tested	% positive	Ref.
Normal retina/brain	0/9	(0)	[29,39,40]
Retinoblastoma*	17/34	(50)	[39]
Glioblastoma multiforme	45/60	(75)	[3,40,41]
Oligodendroglioma	19/19	(100)	[41]
Anaplastic astrocytoma	2/20	(10)	[41]
Meningioma			
Ordinary	5/30	(17)	
Atypical	12/13	(92)	[42]
Malignant	9/9	(100)	
Normal adrenal (newborn)	0/3	(0)	[43]
Ganglioneuroma	0/4	(0)	[43]
Neuroblastoma	99/105	(94)	[43]
Adjacent tissue	0/13	(0)	[43]

*Primary, unilateral or bilateral.

Table 9. Skin

Pathology	No. positive/ No. tested	% positive	Ref.
Normal epidermis	4/9*	(44)	[44]
Squamous cell carcinoma	15/18	(83)	[44]
Basal cell carcinoma	73/77	(95)	[44]
Melanoma	6/7	(86)	[44]

*Activity very weak compared to carcinoma.

Table 10. Haematological tissues*

Pathology	No. positive/ No. tested	% positive	Ref.
Myeloma	1/1	(100)	[45]
Lymphoma			
Low grade	12/14	(86)	[45]
High grade	16/16	(100)	[45,46]
Lymph nodes, benign	5/15	(33)	[45]
Tonsils, benign	23/23	(100)	[45]
MDS	4/6	(67)	[47]
CML			
Chronic	30/42	(71)	[48-50]
Early accelerated	1/3	(33)	[48]
Blast	21/21	(100)	[49,50]
CLL			
Early	2/14	(14)	[47]
Late	4/7	(57)	[47]
APL	1/1	(100)	[48]
ALL	4/5	(80)	[48,49]
AML	47/64	(73)	[47,48,51]

*Data are for samples with activity above basal levels of normal peripheral blood or bone marrow (see text for discussion of telomerase activity in haematopoietic cells). MDS, myelodysplastic syndrome; CML, chronic myeloid leukaemia; CLL, chronic lymphoid leukaemia; APL, acute promyelocytic leukaemia; ALL, acute lymphoblastic leukaemia; AML, acute myeloid leukaemia.

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