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p63 expression in human tumors and normal tissues: a tissue microarray study on 10,200 tumors

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Abstract

Background: Tumor protein 63 (p63) is a transcription factor of the p53 gene family involved in differentiation of several tissues including squamous epithelium. p63 immunohistochemistry is broadly used for tumor classification but published data on its expression in cancer is conflicting.

Methods: To comprehensively catalogue p63 expression, tissue microarrays (TMAs) containing 12,620 tissue samples from 115 tumor entities and 76 normal tissue types were analyzed.

Results: p63 expression was seen in various normal tissues including squamous epithelium and urothelium. At least occasional weak p63 positivity could be detected in 61 (53%) of 115 different tumor types. The frequencies of p63 positivity was highest in squamous cell carcinomas irrespective of their origin (96–100%), thymic tumors (100%), urothelial carcinomas (81–100%), basal type tumors such as basal cell carcinomas (100%), and various salivary gland neoplasias (81–100%). As a rule, p63 was mostly expressed in cancers derived from p63 positive normal tissues and mostly not detectable in tumors derived from p63 negative cancers. However, exceptions from this rule occurred. A positive p63 immunostaining in cancers derived from p63 negative tissues was unrelated to aggressive phenotype in 422 pancreatic cancers, 160 endometrium cancers and 374 ovarian cancers and might be caused by aberrant squamous differentiation or represent stem cell properties. In 355 gastric cancers, aberrant p63 expression occurred in 4% and was linked to lymph node metastasis ($p = 0.0208$). Loss of p63 in urothelial carcinomas - derived from p63 positive urothelium - was significantly linked to advanced stage, high grade ($p < 0.0001$ each) and poor survival ($p < 0.0001$) and might reflect clinically relevant tumor dedifferentiation.

Conclusion: The high prevalence of p63 expression in specific tumor types makes p63 immunohistochemistry a suitable diagnostic tool. Loss of p63 expression might constitute a feature of aggressive cancers.

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Introduction

Tumor protein 63 (p63) is a transcription factor of the p53 gene family encoded by the TP63 gene located at chromosome 3q28. p63 regulates the activity of a multitude of genes involved in growth and development of the ectoderm and derived structures and tissues, such as basal layer keratins and cell cycle control genes [1]. Accordingly, p63 expression is found in basal cell layers of various organs, squamous epithelial cells of many organs and urothelium [1–4]. p63 (syn. TAp63) is closely related to p40 (syn. Δ Np63) as both proteins represent isoforms of the p63 gene with distinct molecular functions [3]. While “full length” p63 (TAp63) activates p53 target genes such as p21 or BAX [4], the shorter transcript p40 (Δ Np63) inhibits activation of p53 and “full length” p63 [4–6].

In diagnostic pathology, the consistently high expression of p63 in specific cell and tissue types is exploited for diagnostic purposes. For example, p63 immunohistochemistry (IHC) is commonly used to mark cell types with critical impact on cancer diagnosis such as basal cells in prostatic and breast glands. Together with other antibodies, p63 is also routinely used for tumor type determination, for example distinguishing squamous cell carcinoma from adenocarcinoma in lung biopsies, or urothelial carcinomas from renal cell carcinoma in tumors arising in the kidney as well as determining the tumor origin of metastases from unknown primary tumors. Since the first description of p63 antibodies, more than 2000 studies have evaluated p63 expression by IHC in various tumors leading to quite discrepant p63 positivity rates in a number of tumor entities [2, 7–77]. For example, the fraction of p63 positive cases ranged from 0 to 77% in small cell lung cancer [19, 78], from 50 to 100% in squamous cell lung cancer [22, 52], 0 to 84% in Merkel cell carcinoma [30, 56], 0 to 82% in papillary thyroid carcinoma [36, 74], 1.4 to 100% in colorectal adenocarcinoma [9, 45], 0 to 100% in urothelial carcinoma [41, 76], 0 to 100% in mucinous ovarian carcinoma [7, 79], and from 0 to 25% in endometrioid ovarian carcinoma [7, 79]. These conflicting data are likely to be caused by the use of different antibodies, immunostaining protocols, and criteria to determine p63 positivity in these studies.

To better understand the relative importance of p63 expression in different tumor types and normal tissues, a comprehensive study analyzing a large number of neoplastic and non-neoplastic tissues under highly standardized conditions is needed. We thus analyzed p63 expression in more than 12,000 tumor tissue samples from 115 different tumor types and subtypes as well as 76 non-neoplastic tissue types by IHC in a tissue microarray (TMA) format.

Materials and methods

Tissue microarrays (TMAs)

Our normal tissue TMA was composed of 8 samples from 8 different donors for each of 76 different normal tissue types (608 samples on one slide). The cancer TMAs contained a total of 12,620 primary tumors from 115 tumor types and subtypes. Detailed histopathological data on grade, pT and pN status were available from 1708 cancers (stomach, pancreas, ovarian, endometrium, urinary bladder). Clinical follow up data were only available from 254 patients who had undergone cystectomy for muscle invasive (pT \geq 2) urinary bladder cancer. In these patients the median follow-up time was 14 (range 1–77) months. Clinical follow up data from patients with carcinomas of the stomach, pancreas, ovarian, and endometrium are not available. The composition of both normal and cancer TMAs is described in detail in the results section. All samples come from the archives of the Institutes of Pathology of the University Hospital of Hamburg, Germany, Clinical Center Osnabrueck, Germany, and the Academic Hospital Fuerth, Germany. No other information beyond the histological tumor type was available for these samples. Tissues were fixed in 4% buffered formalin and then embedded in paraffin. For TMA manufacturing a tumor containing donor block (at least 70% tumor cells in a sufficiently large area) of each patient was selected. Per tumor block/patient one TMA tissue spot (diameter: 0.6 mm) was transferred to an empty recipient TMA block. The use of remnants of archived diagnostic tissues for manufacturing of TMAs and their analysis for research purposes as well as patient data analysis has been approved by local laws (HmbKHG, §12) and by the local ethics committee (Ethics commission Hamburg, WF-049/09). All work has been carried out in compliance with the Helsinki Declaration.

Immunohistochemistry

Freshly cut TMA sections were immunostained on one day and in one experiment. Slides were deparaffinized and exposed to heat-induced antigen retrieval for 5 min in an autoclave at 121 °C in pH 7.8 buffer. Primary antibody specific for p63 (clone 7B4, dilution 1:100) was applied at 37 °C for 60 min. Bound antibody was then visualized using the EnVision Kit (Agilent, CA, USA) according to the manufacturer's directions. For tumor tissues, the percentage of nuclear positive neoplastic cells was estimated, and the staining intensity was semiquantitatively recorded (0, 1+, 2+, 3+). For statistical analyses, the staining results were categorized into four groups. Tumors without any staining were considered negative. Tumors with 1+ staining intensity in \leq 70% of cells and 2+ intensity in \leq 30% of cells were considered weakly positive. Tumors with 1+ staining intensity in $>$ 70% of

cells, 2+ intensity in 31–70%, or 3+ intensity in $\leq 30\%$ were considered moderately positive. Tumors with 2+ intensity in $> 70\%$ or 3+ intensity in $> 30\%$ of cells were considered strongly positive.

Statistics

Statistical calculations were performed with JMP 14 software (SAS Institute Inc., NC, USA). Contingency tables and the χ^2 -test were performed to search for associations between p63 and tumor phenotype. Survival curves were calculated according to Kaplan-Meier. The Log-Rank test was applied to detect significant differences between groups. A significant difference was assumed for a p -value of ≤ 0.05 .

Results

Technical issues

A total of 10,200 (80.8%) of 12,620 tumor samples were interpretable in our TMA analysis. Non-interpretable samples either lacked unequivocal tumor cells or were lost from the TMA during the technical procedures. A statistical bias, which could potentially result from exclusion of non-interpretable samples, is highly unlikely in our study as non-interpretable samples were evenly distributed across all pathological diagnoses (Supplementary Table 1). On our normal tissue TMA, a sufficient number of samples was always interpretable per tissue type to determine the normal tissue p63 expression.

p63 in normal tissues

A strong nuclear p63 immunostaining was seen in squamous epithelium irrespective of its origin, peripheric germinative cells of sebaceous glands, urothelium, thymic epithelial cells, myoepithelial cells in breast, parotid, submandibular and sublingual glands, basal cells in prostate, seminal vesicle, and respiratory epithelium, cytotrophoblast of first trimester and mature placenta. The staining intensity was slightly decreasing from basal cells to the surface cell layer in squamous epithelia and in urothelium. A mild staining was seen in some lymphocytes and in high endothelial venules in lymph nodes. Representative images are given in Fig. 1. p63 staining was absent in aorta/intima, aorta/media, heart (left ventricle), striated muscle, skeletal muscle, skeletal muscle/tongue, uterus/myometrium, muscular wall of the GI-tract, renal pelvis and bladder, corpus spongiosum of the penis, ovarian stroma, fat, red and white pulp of the spleen, antrum and corpus of the stomach, mucosa and lamina propria of the stomach duodenum, ileum, appendix, colon descendens, rectum, and gallbladder, epithelium of the gallbladder, liver, pancreas, bone marrow, Brunner gland of the duodenum, cortex and medulla of the kidney, epididymis, testis, glands of the bronchus, endocervix, proliferative and secretory endometrium, mucosa of the fallopian tube, corpus luteum and follicular cyst of the ovary, adrenal gland, parathyroid, thyroid gland, stratum molecular and neuronorum of the cerebellum, grey and white

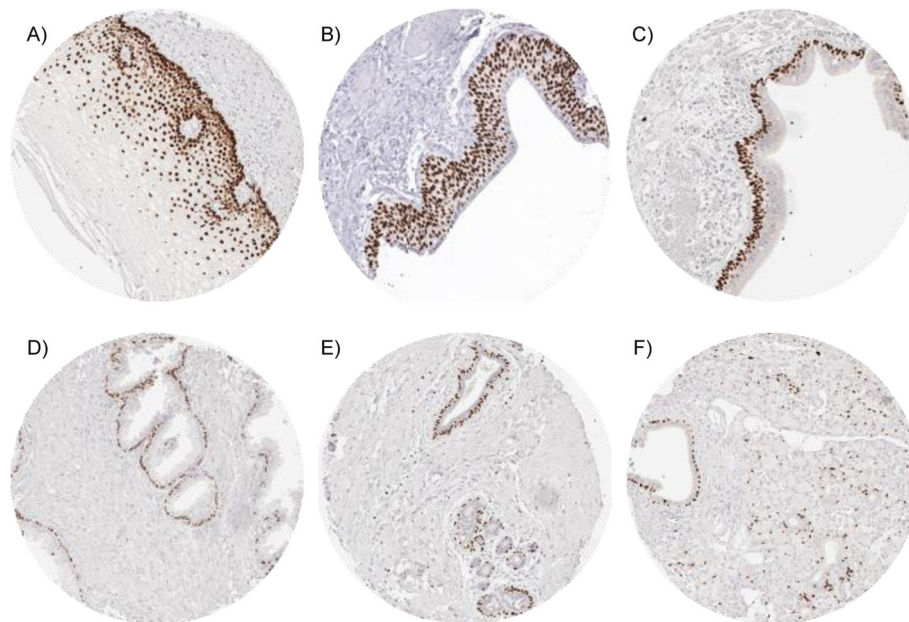


Fig. 1 p63 expression in normal tissues. Strong p63 immunostaining is seen in squamous epithelium of the esophagus (a) urothelium of the urinary bladder (b), basal cells of respiratory epithelium (c) and of the prostate (d) and myoepithelial cells of the breast (e) and of the salivary glands (f)

cerebrum, and posterior and anterior lobe of the pituitary gland.

p63 in neoplastic tissues

Nuclear immunostaining was observed in 1940 (19.0%) of 10,200 interpretable tumors with 13.4% showing strong, 3.0% moderate and 2.6% weak staining intensity. At least an occasional weak p63 positivity could be detected in 61 of 115 (53.4%) different tumor types and tumor subtypes and 37 (32.2%) tumor types and tumor subtypes had at least one tumor exhibiting strong positivity. Representative images of p63 positive tumors are shown in Fig. 2. The highest frequencies of p63 positivity were seen in squamous cell carcinomas irrespective of their origin, thymic tumors, urothelial cancers and basal type tumors such as basal cell carcinomas and various salivary gland neoplasia. A detailed description of the immunostaining results is given in Table 1 and Fig. 3.

p63 expression, tumor phenotype and prognosis

p63 immunostaining was not associated with parameters of disease aggressiveness in 422 pancreatic carcinomas, 160 endometrium cancers, and 374 ovarian cancers (Table 2). In a cohort of 355 gastric carcinomas, p63 positivity was seen in 4% and was linked to nodal metastasis ($p = 0.0208$; Table 2). In urinary bladder cancer, reduced or absent p63 immunostaining was

related to advanced stage and high-grade categories ($p < 0.0001$, Table 2) and reduced survival ($p < 0.0001$; Fig. 4).

Discussion

The results of our study on 12,620 tissues show that p63 expression is largely limited to few normal tissues including squamous epithelium, urothelium, thymic epithelial cells and basal/myoepithelial cells of various epithelial organs. The fact that p63 expression in these cells is usually strong but completely undetectable in other tissues fits well with the known role of p63 in driving embryonal cellular evolution towards specific cell types. The S-shaped curve displaying the ranking order of p63 positive tumors demonstrates, that frequent and intensive p63 immunostaining is predominantly seen in these few cancers that appear to be derived from p63 positive normal cell types. The most commonly positive cancers include squamous cell carcinomas irrespective of their origin, thymic tumors, urothelial carcinomas and basal type tumors such as basal cell carcinomas and various salivary gland neoplasias.

However, in the majority of p63 positive tumor types, p63 expression was not seen in all cases. Absence of detectable p63 immunostaining in a tumor derived from a p63 expressing normal tissues may in some instances reflect inefficient immunostaining in

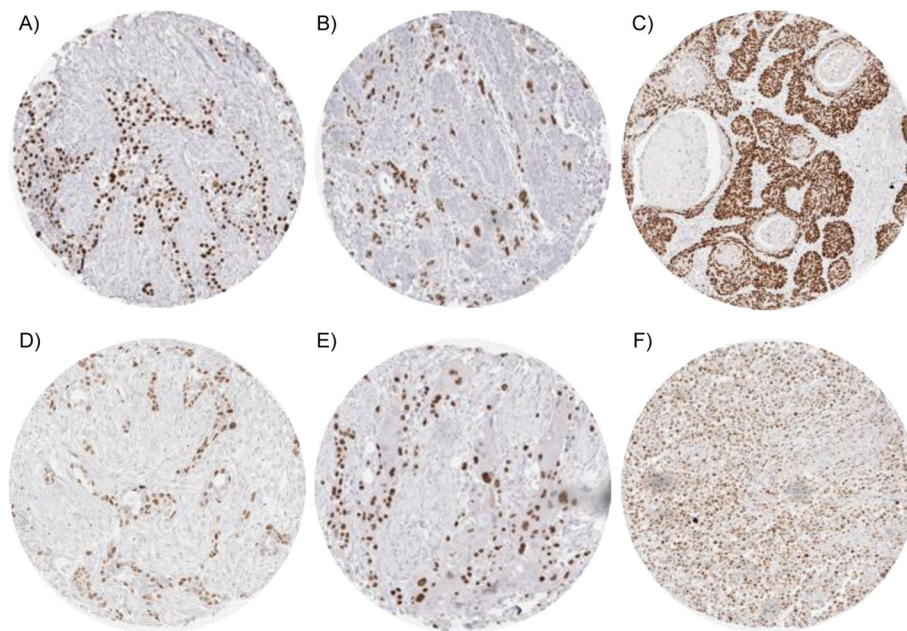


Fig. 2 p63 expression in cancerous tissues. Strong p63 immunostaining is seen in invasive urothelial carcinoma (a) gastric cancer, diffuse type (b), anal squamous cell carcinoma (c), gastric cancer, intestinal type (d), and adenocarcinoma of the pancreas with focal squamous differentiation (e). Moderate intensity p63 staining is seen in a diffuse large B-cell lymphoma (f)

Table 1 p63 immunostaining in human cancers

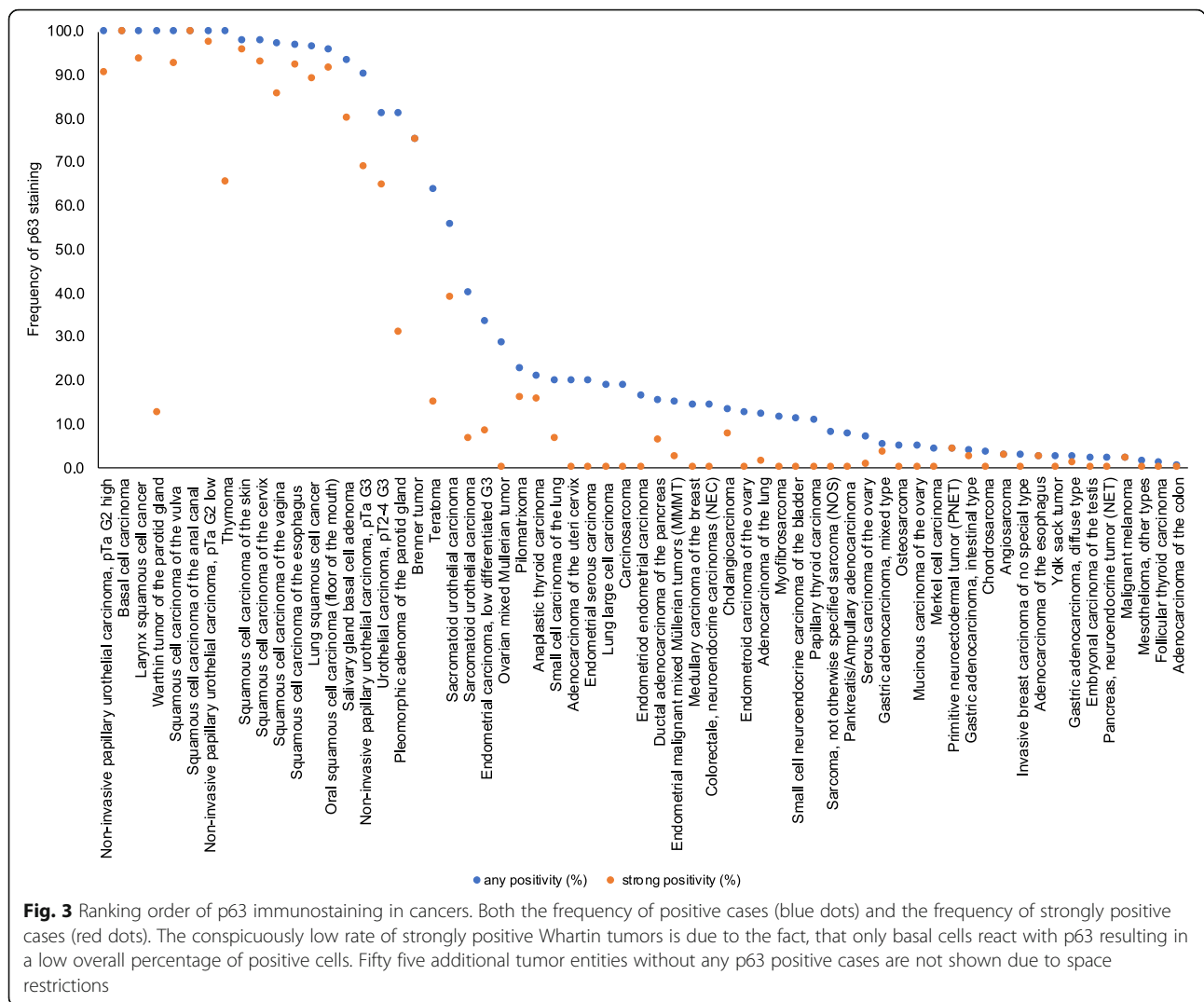
	Tumor type	TMA (n)	p63 immunohistochemistry					
			analyzable (n)	negative (%)	weak (%)	moderate (%)	strong (%)	positive (%)
Tumors of the skin	Pilomatrixoma	35	31	77.4	0.0	6.5	16.1	22.6
	Basal cell carcinoma	48	45	0.0	0.0	0.0	100.0	100.0
	Benign nevus	29	22	100.0	0.0	0.0	0.0	0.0
	Squamous cell carcinoma of the skin	50	47	2.1	2.1	0.0	95.7	97.9
	Malignant melanoma	48	43	97.7	0.0	0.0	2.3	2.3
	Merkel cell carcinoma	46	45	95.6	0.0	4.4	0.0	4.4
Tumors of the head and neck	Squamous cell carcinoma of the larynx	100	88	9.1	11.4	15.9	63.6	90.09
	Oral squamous cell cancer	50	47	4.3	0.0	4.3	91.5	95.7
	Oral squamous cell carcinoma (floor of the mouth)	49	48	0.0	6.3	81.3	12.5	100.0
	Pleomorphic adenoma of the parotid gland	15	15	6.7	6.7	6.7	80.0	93.3
	Warthin tumor of the parotid gland	127	83	3.6	4.8	2.4	89.2	96.4
	Basal cell adenoma of the salivary gland	31	16	81.3	18.8	0.0	0.0	18.8
Tumors of the lung, pleura and thymus	Adenocarcinoma of the lung	250	189	87.8	10.6	0.0	1.6	12.2
	Squamous cell carcinoma of the lung	6	5	100.0	0.0	0.0	0.0	0.0
	Small cell carcinoma of the lung	20	15	80.0	6.7	6.7	6.7	20.0
	Malignant mesothelioma	39	39	100.0	0.0	0.0	0.0	0.0
	Mesothelioma, other types	76	67	98.5	0.0	1.5	0.0	1.5
	Thymoma	29	29	0.0	13.8	20.7	65.5	100.0
Tumors of the female genital tract	Squamous cell carcinoma of the vagina	48	35	2.9	2.9	8.6	85.7	97.1
	Squamous cell carcinoma of the vulva	50	41	0.0	0.0	7.3	92.7	100.0
	Squamous cell carcinoma of the cervix	50	43	2.3	2.3	2.3	93.0	97.7
	Adenocarcinoma of the cervix uteri	50	45	80.0	15.6	4.4	0.0	20.0
	Endometrioid endometrial carcinoma	236	206	83.5	6.8	9.7	0.0	16.5
	Endometrial serous carcinoma	82	60	80.0	11.7	8.3	0.0	20.0
	Endometrial malignant mixed Müllerian tumors (MMMT)	48	40	85.0	7.5	5.0	2.5	15.0
	Endometrial carcinoma, high grade G3	13	12	66.7	0.0	25.0	8.3	33.3
	Endometrial clear cell carcinoma	8	4	100.0	0.0	0.0	0.0	0.0
	Endometrial stromal sarcoma	12	12	100.0	0.0	0.0	0.0	0.0
	Endometrioid carcinoma of the ovary	115	102	87.3	3.9	8.8	0.0	12.7
	Serous carcinoma of the ovary	567	496	92.9	4.2	2.0	0.8	7.1
	Mucinous carcinoma of the ovary	97	81	95.1	2.5	2.5	0.0	4.9
	Clear cell carcinoma of the ovary	54	48	100.0	0.0	0.0	0.0	0.0
	Ovarian malignant mixed Müllerian tumors (MMMT)	47	42	71.4	9.5	19.0	0.0	28.6
	Brenner tumor	9	8	25.0	0.0	0.0	75.0	75.0
Tumors of the breast	Invasive breast carcinoma of no special type	126	67	97.0	1.5	1.5	0.0	3.0
	Lobular carcinoma of the breast	123	97	100.0	0.0	0.0	0.0	0.0
	Medullary carcinoma of the breast	15	14	85.7	7.1	7.1	0.0	14.3
	Tubular carcinoma of the breast	18	15	100.0	0.0	0.0	0.0	0.0
	Mucinous carcinoma of the breast	22	16	100.0	0.0	0.0	0.0	0.0
	Phyllodes tumor of the breast	50	36	100.0	0.0	0.0	0.0	0.0

Table 1 p63 immunostaining in human cancers (Continued)

	Tumor type	TMA (n)	p63 immunohistochemistry					strong (%)	positive (%)
			analyzable (n)	negative (%)	weak (%)	moderate (%)			
Tumors of the digestive system	Adenomatous polyp, low-grade dysplasia	50	41	100.0	0.0	0.0	0.0	0.0	
	Adenomatous polyp, high-grade dysplasia	50	43	100.0	0.0	0.0	0.0	0.0	
	Adenocarcinoma of the colon	1882	1743	99.6	0.3	0.1	0.0	0.4	
	Adenocarcinoma of the small intestine	10	5	100.0	0.0	0.0	0.0	0.0	
	Gastric adenocarcinoma, diffuse type	176	153	97.4	1.3	0.0	1.3	2.6	
	Gastric adenocarcinoma, intestinal type	174	151	96.0	0.7	0.7	2.6	4.0	
	Gastric adenocarcinoma, mixed type	62	56	94.6	1.8	0.0	3.6	5.4	
	Adenocarcinoma of the esophagus	83	75	97.3	0.0	0.0	2.7	2.7	
	Squamous cell carcinoma of the esophagus	75	63	3.2	1.6	3.2	92.1	96.8	
	Squamous cell carcinoma of the anal canal	50	35	0.0	0.0	0.0	100.0	100.0	
	Cholangiocarcinoma	50	38	86.8	5.3	0.0	7.9	13.2	
	Hepatocellular carcinoma	50	46	100.0	0.0	0.0	0.0	0.0	
	Ductal adenocarcinoma of the pancreas	612	489	84.7	5.3	3.7	6.3	15.3	
	Pancreatic/Ampullary adenocarcinoma	89	64	92.2	1.6	6.3	0.0	7.8	
	Acinar cell carcinoma of the pancreas	13	12	100.0	0.0	0.0	0.0	0.0	
Gastrointestinal stromal tumor (GIST)	50	42	100.0	0.0	0.0	0.0	0.0		
Tumors of the urinary system	Non-invasive papillary urothelial carcinoma, pTa G2 low grade	177	116	0.0	0.0	2.6	97.4	100.0	
	Non-invasive papillary urothelial carcinoma, pTa G2 high grade	141	106	0.0	0.9	8.5	90.6	100.0	
	Non-invasive papillary urothelial carcinoma, pTa G3	187	132	9.8	6.1	15.2	68.9	90.2	
	Urothelial carcinoma, pT2-4 G3	1117	732	18.9	7.4	9.0	64.8	81.1	
	Small cell neuroendocrine carcinoma of the bladder	18	18	88.9	5.6	5.6	0.0	11.1	
	Sarcomatoid urothelial carcinoma	25	18	44.4	11.1	5.6	38.9	55.6	
	Clear cell renal cell carcinoma	858	509	100.0	0.0	0.0	0.0	0.0	
	Papillary renal cell carcinoma	255	155	100.0	0.0	0.0	0.0	0.0	
	Clear cell (tubulo) papillary renal cell carcinoma	21	10	100.0	0.0	0.0	0.0	0.0	
	Chromophobe renal cell carcinoma	131	96	100.0	0.0	0.0	0.0	0.0	
Tumors of the male genital organs	Oncocytoma	177	110	100.0	0.0	0.0	0.0	0.0	
	Adenocarcinoma of the prostate, Gleason 3+3	83	78	100.0	0.0	0.0	0.0	0.0	
	Adenocarcinoma of the prostate, Gleason 4+4	80	71	100.0	0.0	0.0	0.0	0.0	
	Adenocarcinoma of the prostate, Gleason 5+5	85	76	100.0	0.0	0.0	0.0	0.0	
	Adenocarcinoma of the prostate (recurrence)	330	181	100.0	0.0	0.0	0.0	0.0	
	Small cell neuroendocrine carcinoma of the prostate	17	15	100.0	0.0	0.0	0.0	0.0	
	Seminoma	50	46	100.0	0.0	0.0	0.0	0.0	
	Embryonal carcinoma of the testis	50	42	97.6	0.0	2.4	0.0	2.4	
	Yolk sack tumor	50	38	97.4	0.0	2.6	0.0	2.6	
	Teratoma	50	33	36.4	9.1	39.4	15.2	63.6	
Tumors of endocrine organs	Adenoma of the thyroid gland	114	105	100.0	0.0	0.0	0.0	0.0	
	Papillary thyroid carcinoma	392	363	89.0	6.1	5.0	0.0	11.0	
	Follicular thyroid carcinoma	158	150	98.7	1.3	0.0	0.0	1.3	

Table 1 p63 immunostaining in human cancers (Continued)

Tumor type	TMA (n)	p63 immunohistochemistry						positive (%)
		analyzable (n)	negative (%)	weak (%)	moderate (%)	strong (%)		
Medullary thyroid carcinoma	107	92	100.0	0.0	0.0	0.0	0.0	
Anaplastic thyroid carcinoma	45	38	78.9	2.6	2.6	15.8	21.1	
Adrenal cortical adenoma	50	47	100.0	0.0	0.0	0.0	0.0	
Adrenal cortical carcinoma	26	23	100.0	0.0	0.0	0.0	0.0	
Phaeochromocytoma	50	44	100.0	0.0	0.0	0.0	0.0	
Appendix, neuroendocrine tumor (NET)	22	10	100.0	0.0	0.0	0.0	0.0	
Colorectal, neuroendocrine tumor (NET)	10	7	100.0	0.0	0.0	0.0	0.0	
Ileum, neuroendocrine tumor (NET)	49	41	100.0	0.0	0.0	0.0	0.0	
Lung, neuroendocrine tumor (NET)	19	12	100.0	0.0	0.0	0.0	0.0	
Pancreas, neuroendocrine tumor (NET)	100	83	98.8	1.2	0.0	0.0	1.2	
Colorectal, neuroendocrine carcinoma (NEC)	11	7	85.7	14.3	0.0	0.0	14.3	
Gallbladder, neuroendocrine carcinoma (NEC)	4	2	100.0	0.0	0.0	0.0	0.0	
Pancreas, neuroendocrine carcinoma (NEC)	13	10	100.0	0.0	0.0	0.0	0.0	
Tumors of haematopoietic and lymphoid tissues								
Hodgkin Lymphoma	45	38	100.0	0.0	0.0	0.0	0.0	
Non-Hodgkin Lymphoma	48	45	60.0	22.2	11.1	6.7	40.0	
Tumors of soft tissue and bone								
Tenosynovial giant cell tumor	45	45	100.0	0.0	0.0	0.0	0.0	
Granular cell tumor	53	50	100.0	0.0	0.0	0.0	0.0	
Leiomyoma	50	47	100.0	0.0	0.0	0.0	0.0	
Angiomyolipoma	91	91	100.0	0.0	0.0	0.0	0.0	
Angiosarcoma	73	66	97.0	0.0	0.0	3.0	3.0	
Dermatofibrosarcoma protuberans	21	21	100.0	0.0	0.0	0.0	0.0	
Ganglioneuroma	14	14	100.0	0.0	0.0	0.0	0.0	
Kaposi sarcoma	8	8	100.0	0.0	0.0	0.0	0.0	
Leiomyosarcoma	87	84	100.0	0.0	0.0	0.0	0.0	
Liposarcoma	132	124	100.0	0.0	0.0	0.0	0.0	
Malignant peripheral nerve sheath tumor (MPNST)	13	13	100.0	0.0	0.0	0.0	0.0	
Myofibrosarcoma	26	26	88.5	11.5	0.0	0.0	11.5	
Neurofibroma	117	117	100.0	0.0	0.0	0.0	0.0	
Sarcoma, not otherwise specified (NOS)	75	75	92.0	8.0	0.0	0.0	8.0	
Paraganglioma	41	41	100.0	0.0	0.0	0.0	0.0	
Primitive neuroectodermal tumor (PNET)	23	23	95.7	0.0	0.0	4.3	4.3	
Rhabdomyosarcoma	7	7	100.0	0.0	0.0	0.0	0.0	
Schwannoma	121	121	100.0	0.0	0.0	0.0	0.0	
Synovial sarcoma	12	12	100.0	0.0	0.0	0.0	0.0	
Osteosarcoma	43	39	94.9	5.1	0.0	0.0	5.1	
Chondrosarcoma	39	27	96.3	3.7	0.0	0.0	3.7	



cancers with inappropriate fixation or other preanalytical issues leading to tissue damage [80]. Our findings in 1038 analyzed urothelial cancers further indicate that p63 expression loss can occur as a result of cellular dedifferentiation during cancer progression. The progressive loss of p63 expression from pTa G2 low grade (0%) small cell urothelial cancer (89%) represents an example of progression associated p63 loss. The fact that p63 loss represents an ominous feature in urothelial carcinomas is also demonstrated by the worse prognosis in p63 negative as compared to p63 positive pT2–4 urothelial carcinomas. These data are in line with reports from other investigators also describing associations of p63 loss with advanced stage and poor prognosis in bladder cancer [41]. Other authors have also described a link between low p63 expression and poor prognosis in squamous cell carcinomas of the esophagus [47] and the larynx [48].

Only nine of 115 cancer types (7.8% of analyzed tumor categories) had a prevalence of p63 positivity between 25 and 90%. These included teratoma and Non-Hodgkin lymphoma. In teratoma the “p63 immunostaining result” of a TMA spot was obviously driven by whether or not epithelial components represented in the spot physiologically expressed p63. In malignant lymphoma, the staining intensity was often moderate. This is reflective of the role of p63 in normal lymphocytes where few cells regularly showed weak to moderate expression. Occasional p63 expression in B-lymphocytes, mainly in germinal centers have been described in previous studies [2, 60]. The high rate of p63 positive B-cell Non-Hodgkin lymphoma is in line with reports from other authors describing common p63 expression, mainly in large cell Non-Hodgkin lymphomas but also in chronic lymphocytic leukemia (CLL), and follicular lymphoma [2, 60,

Table 2 p63 immunostaining and tumor phenotype

		n	p63 IHC result (%)				p
			negative	weak	moderate	strong	
Urinary bladder cancer	all cancers	1038	13.4	5.9	9.1	71.7	
	pTa G2 low	116	0.0	0.0	2.6	97.4	<0.0001
	pTa G2 high	106	0.0	0.9	8.5	90.6	
	pTaG3	132	9.8	6.1	15.2	68.9	
	pT \geq 2 G3	672	18.6	7.0	9.2	65.2	
	pT \geq 2 G3 sarcomatoid	18	44.4	11.1	5.6	38.9	0.0052 (vs >pT2G3)
	pT \geq 2 G3 small cell ca.	18	88.9	5.6	5.6	0.0	<0.0001 (vs \geq pT2G3)
	pN0	111	13.5	8.1	9.0	69.4	0.9138
	pN+	71	15.5	5.6	8.5	70.4	
Pancreatic adenocarcinoma	all cancers	422	84.4	5.7	3.1	6.9	
	pT1	12	91.7	0	0	8.3	0.2753
	pT2	63	92.1	3.2	1.6	3.2	
	pT3	318	82.4	6.9	3.1	7.5	
	pT4	27	85.2	0	7.4	7.4	
	G1	12	100	0	0	0	0.3483
	G2	295	82.7	7.1	3.1	7.1	
	G3	92	85.9	3.3	2.2	8.7	
	pN0	91	85.7	3.3	4.4	6.6	0.5777
	pN+	329	83.9	6.4	2.7	7	
	pM0	334	83.2	6.3	3	7.5	0.515
	pM1	87	88.5	3.4	3.4	4.6	
	R0	212	83.5	7.1	2.4	7.1	0.5384
R1	174	85.6	4.6	4	5.7		
Endometrial cancer, endometrioid	all cancers	160	83.1	4.4	12.5	0.0	
	pT1	101	83.2	3.0	13.9	0.0	0.6733
	pT2	23	87.0	4.3	8.7	0.0	
	pT3-4	33	78.8	9.1	12.1	0.0	
	pN0	43	83.7	4.7	11.6	0.0	0.3704
	pN+	28	75.0	14.3	10.7	0.0	
Ovarian cancers, serous	all cancers	374	91.7	5.1	2.4	0.8	
	pT1	29	93.1	3.4	3.4	0.0	0.2274
	pT2	39	84.6	5.1	10.3	0.0	
	pT3	257	91.8	5.4	1.6	1.2	
	pN0	81	88.9	4.9	4.9	1.2	0.3673
	pN+	166	91.0	6.6	1.2	1.2	
Stomach cancer	all cancers	355	96.1	1.1	0.6	2.3	
	diffuse	85	97.6	2.4	0.0	0.0	0.1478
	intestinal	79	97.5	0.0	0.0	2.5	
	mixed	56	94.6	1.8	0.0	3.6	
	pT1-2	57	96.5	0.0	1.8	1.8	0.3119
	pT3	117	94.9	0.9	0.9	3.4	
	pT4	118	96.6	2.5	0.0	0.8	
	pN0	74	100.0	0.0	0.0	0.0	0.0208

Table 2 p63 immunostaining and tumor phenotype (Continued)

	n	p63 IHC result (%)				p
		negative	weak	moderate	strong	
pN1	62	93.5	1.6	3.2	1.6	
pN2	55	92.7	0.0	0.0	7.3	
pN3	101	96.0	3.0	0.0	1.0	

81–84]. Due to the fact that both the TP63 gene and the Bcl-6 gene are located on chromosome 3q27(–29) and Bcl-6 gene rearrangements are often seen in diffuse large B-cell lymphomas (DLBCL) [85–88], it has been speculated that the close vicinity of TP63 to Bcl-6 may contribute to its potential involvement in DLBCL tumor progression [60]. However, associations between p63 and BCL-6 expression was not found in another study [84]. Moreover, it remains unclear whether the common translocation at 3q27 affects the expression or structure of the more distal p63 gene [60, 87].

A total of 29 (25%) cancer types and subtypes showed p63 positivity in 3 to 25% of analyzed cases. Very often, p63 staining did not involve the entire tumor mass in these cancers. Almost all of these entities are derived from tissues not normally expressing p63 suggesting neo-expression of p63 occurring during cancer development and progression. At least in a fraction of these tumors p63 neo-expression was obviously linked to focal squamous cell differentiation. Tumor types that are particularly known for occasionally containing squamous elements were common in the category of tumors with p63 positivity between 10 and 25% and included endometroid cancer and malignant mixed Mullerian

tumors of the uterus as well as ovarian, pancreatic and cholangiocellular carcinomas. In other tumors with occasional occurrence of p63 positive cells, the phenomenon may reflect stemness properties as earlier shown for p63 expressing normal and cancerous cells [89–91] or be caused by incidental and possibly biologically irrelevant p63 neo-expression in dysregulated cancer cells. The fact that p63 neo-expression was unrelated to features of cancer aggressiveness in cohorts of 422 pancreatic, 374 ovarian, and 160 endometrium cancers argues against a major biologic impact of p63 expression in these tumors. The significantly higher rate of nodal metastasis in p63 positive gastric cancers may be explained by the known poor prognosis of adenosquamous gastric cancers [92].

Overall, these data demonstrate a broad diagnostic utility of p63 IHC for the categorization of cancers, all of which have previously been suggested. For example, p63 expression in a kidney tumor argues for urothelial carcinomas and against poorly differentiated or sarcomatoid renal cell carcinoma in kidney masses [93]. Although several sarcomas showed limited p63 expression, the striking positivity in many sarcomatoid urothelial carcinomas suggests that p63 positivity - in organs where p63 positive epithelial cells are common - argues for sarcomatoid carcinoma and against sarcoma [93, 94]. p63 positivity in a poorly differentiated urinary bladder tumor is literally ruling out infiltration by a prostate cancer even though - considering the general likelihood of these conditions - most p63 negative cancers in the bladder are still representing urothelial carcinomas. In addition, p63 IHC is well established for facilitating the sometimes difficult and clinically highly relevant distinction of adenocarcinoma and squamous cell carcinoma of the lung [22, 52] as well as ruling out invasive cancer by demonstrating a basal cell or myoepithelial cell layer in prostate, breast and salivary gland tumors [24, 67, 95]. It is of note, that in case of a p63 positive solid tumor, Non-Hodgkin lymphoma always remains a diagnostic option.

Importantly, all prevalences described in this study are specific to the reagents and the protocol used in our laboratory. The sum of data that had been earlier collected on p63 expression in cancers -

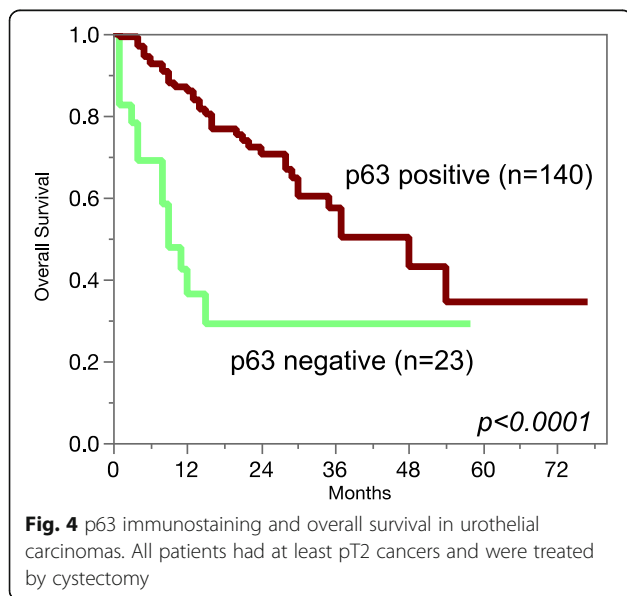


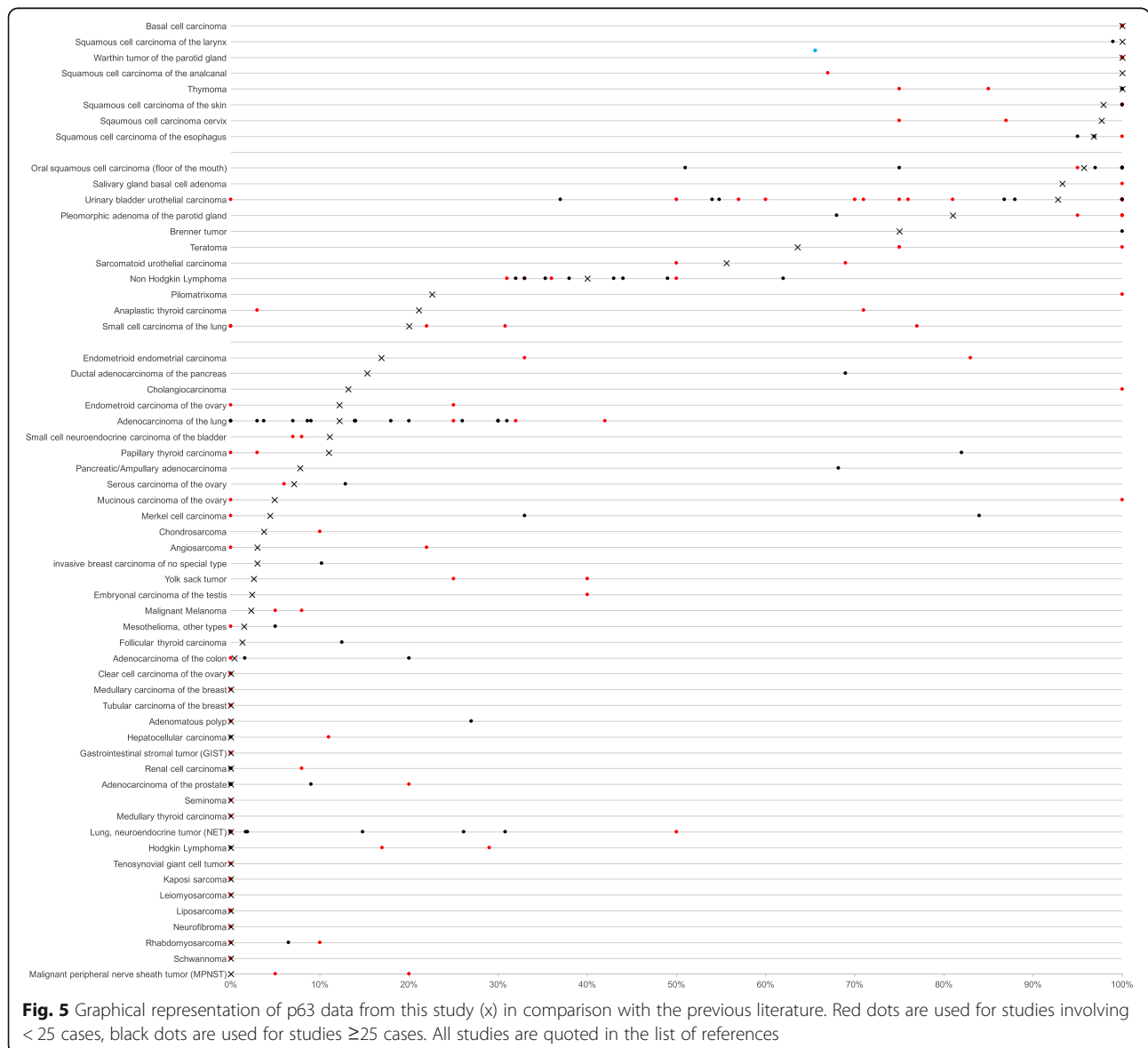
Fig. 4 p63 immunostaining and overall survival in urothelial carcinomas. All patients had at least pT2 cancers and were treated by cystectomy

summarized in Fig. 5 - would not necessarily have led to the same conclusions as drawn from this study. The use of different antibodies, protocols and interpretation criteria have jointly caused highly diverse literature data on p63 expression in cancer. It is well known, that different antibodies designed for the same target protein can vary to a large extent in their binding properties and that protocol modifications greatly impact the rate of cases considered “positive” for a certain protein [19, 96–102]. A further limitation of the study could be the missing evaluability of about 20% of the tumor samples. However, a statistical bias, which could potentially result from exclusion of non-interpretable samples, is highly unlikely in our study as non-interpretable

samples were evenly distributed across all pathological diagnoses.

Conclusion

Strong and abundant p63 expression is seen in only few cancer types primarily including squamous and urothelial cancers as well as tumors derived from the thymus. Occasional strong p63 expression can, however, occur in a large number of other neoplasias including cancer subtypes with a propensity for focal squamous differentiation. The study also demonstrates the high value of extensive tissue validation using large-scale TMAs for determining the diagnostic utility of antibodies.



Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s40364-021-00260-5>.

Additional file 1.

Abbreviations

DLBCL: Diffuse Large B Cell Lymphoma; IHC: Immunohistochemistry; TMA: Tissue Micro Array; p63: Tumor Protein 63

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Authors' contributions

SS, AHM, RS, and GS designed the study. SS, FB, CR, AML, AH, DH, SW, CF, KM, AM, CB, PL, TSC, RU, WW, DD, SM, EB, RK, and TK performed the immunohistochemical analyses and/or contributed to the pathological validation of the tumors, the tissue microarray construction, and data collection. MF and MR contributed to data collection. MK, CHM, and RS carried out the data analyses. GS, RS, AHM, SS and MK wrote the first draft of the manuscript. All authors contributed toward data analysis, drafting and critically revising the paper, gave final approval of the version to the published, and agree to be accountable for all aspects of the work.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Ethics approval and consent to participate

The use of remnants of archived diagnostic tissues for manufacturing of TMAs and their analysis for research purposes as well as patient data analysis has been approved by local laws (HmbKHG, §12) and by the local ethics committee (Ethics commission Hamburg, WF-049/09). All work has been carried out in compliance with the Helsinki Declaration.

Consent for publication

Not applicable.

Competing interests

RS developed the antibody 7B4.

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