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The RB/ p16^{INK4A} pathway but not p53 is disrupted by human papillomavirus in penile squamous cell carcinoma

Short title: HPV and protein expression in penile SCC

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Keywords: Penile squamous cell carcinoma, human papillomavirus, p53, p21, p16^{INK4A}, RB

<u>Abbreviations</u>: HPV, human papillomavirus; PCR, polymerase chain reaction; PSCC, penile squamous cell carcinoma

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Abstract

Aims: The Pathogenesis of penile squamous cell carcinoma (PSCC) is not well understood. Human papillomavirus (HPV) may be involved in carcinogenesis, but few studies have compared cell-cycle protein expression in HPV positive and negative cancers. The aim was to determine the extent of HPV infection in different histological subtypes of PSCC and its impact on the expression of key cell cycle proteins: p53, p21, p16^{INK4A} and RB.

Methods and Results: We examined 148 PSCC samples immunohistochemically for RB, $p16^{INK4A}$, p53 and p21 protein expression. 102 cases were typed for HPV by PCR. HPV DNA was detected in 56% of tumours with HPV16 present in 81%. Basaloid tumours were strongly related to HPV infection (10/13) while verrucous were not (3/13). Fifty-nine % (38/64) of usual type SCCs had HPV infection. RB protein negatively (p<0.0001) and p21 (p=0.0002) positively correlated with HPV infection. p53 did not correlate with HPV infection.

Conclusions: HPV infection is present in over half of penile cancers and it is responsible for RB pathway disruption. However, no link between HPV and p53 immunodetection was found. Only basaloid and half of usual type PSSCs correlate with HPV, confirming possible separate aetiologies for those tumours.

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Introduction

Penile carcinoma is rare in developed countries, representing 0.3-0.5% of male malignancies in Europe and the USA ¹. In the UK there are approximately 600 (rate 1 per 100 000) new cases each year, mostly after the sixth decade ^{1, 2}. The vast majority (95%) are squamous cell carcinomas ³. These may be divided into usual type (70%), followed by more aggressive basaloid (10%) and a slow growing, low grade group of 'verruciform' tumours (20%). Verruciform lesions include verrucous carcinoma, warty carcinoma and papillary squamous cell carcinomas ⁴. Mixed tumours of usual type and one or more of other subtypes of SCC also exist.

Risk factors for penile cancer include lack of circumcision during childhood, phimosis and cigarette smoking ⁵. There is some confusion on the role of infection with human papillomavirus in penile cancer ³, as contrary to cervical cancer ⁶ the incidence rate varies significantly from 15% to 77.5% ^{7, 8}, depending on detection range, population studied and tumour type ⁹. HPV16 is most prevalent infection ⁷, similar to other HPV-related ano-genital cancers ¹⁰. Basaloid and warty tumours have been shown to be strongly associated with HPV infection ¹¹. Therefore, penile cancer may resemble vulvar cancer, which has two different aetiologies, one related to HPV infection and one that is not ¹².

The carcinogenic abilities of high-risk HPV types are well known due to extensive studies on cervical cancer. Viral oncoproteins E6 and E7 can disrupt cell cycle checkpoints and apoptosis by respectively interacting with tumour suppressor proteins, RB and p53. RB regulates cell cycle progression, protecting the cell from uncontrolled proliferation and is regulated by cyclin dependent kinase phosphorylation. p16^{INK4A} can inhibit cyclin dependent kinase mediated RB phosphorylation, preventing cell cycle progression. In cells infected with high-risk HPV, viral E7 protein binds directly to RB causing its inactivation and downregulation, which prevents cell-cycle control by p16^{INK4A} ^{13, 14}. In these circumstances accumulation of p16^{INK4A} can occur, which is an indication of HPV infection ¹⁵. p53 can also

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inhibit cancer development and tumour growth through its ability to efficiently inhibit cell proliferation and promote apoptotic cell death ¹⁶. In cancers harbouring high-risk HPV, expression of viral E6 protein can inactivate p53 through its downregulation and an inverse correlation between HPV positivity and p53 overexpression has been found in some cancer sites but not others ¹⁷. HPV E7 protein can additionally overcome the inhibitory function of p21. p21 is a p53-responsive protein and arrests cell cycle in presence of DNA damage. E7 protein binds to p21 and abrogates its inhibitory functions, therefore overcoming a DNA damage-induced cell cycle arrest despite high levels of p21 ¹⁸.

The mechanisms of oncogenesis in penile cancer are not fully understood. There are no data on RB expression in penile cancer and results on p53 in relation to HPV are inconclusive. Limited data is available on p16^{INK4A} immunodetection ^{19, 20} and p21 expression ²¹ in penile tumours. Therefore, we investigated HPV infection type in one of the largest series of penile SCC to test the hypothesis that HPV type and its association with key cell cycle proteins had differential effects on the tumour subtypes, in order to elucidate their role in tumour pathogenesis.

P P

Materials and Methods

The study was conducted upon approval from East London and The City Research Ethics Committee. We retrospectively reviewed the Cellular Pathology Department Registry of St George's Hospital to identify patients treated for penile SCC between 2001 - 2007. We retrieved 148 penile SCCs. 97 samples were usual type SCCs, 17 basaloid, 15 pure verrucous carcinomas, 7 mixed verrucous/usual type, 7 mixed verrucous/warty, 2 warty and 3 warty/usual types. 21 cases were obtained from excision biopsies/circumcisions, 82 from glansectomies and 45 from partial/total penectomies. All cases were re-reviewed by an expert uropathologist (C.C.) including subtyping, grading and staging by standard methodologies ⁴, ²².

Polymerase chain reaction (PCR).

102 wax blocks from penile SCC cases were available for DNA extraction with a QIAamp DNA Mini kit (51304; Qiagen, Crawley, U.K.). Beta-Globin polymerase chain reaction was performed using primers B1 and B19 to confirm the adequacy of the extracted DNA. Validated samples were tested for the presence of HPV DNA by a broad-spectrum HPV PCR method using SPF10 primers which amplify a 65-bp fragment of the L1 open reading frame and HPV genotypes identified by the INNO-LiPA line probe assay (Innogenetics NV, Ghent, Belgium)¹⁹.

Immunohistochemistry.

Tissue microarray blocks were prepared using a manual microarrayer. Three x 1mm tissue cores were taken from each tumour. Four um sections were cut and immunostained using standard heat-induced antigen retrieval methods and the ABC kit (Vector Laboratories, PK-6200), according to manufacturer instructions ²³. Primary antibodies dilutions were: 1:50 for RB (Novocastra, NCL-RB-358), 1:100 for p16^{INK4A} (Neomarkers, MS-1064-PO), 1:1000 for p53 (Dako, M7001) and p21 (Dako, M7202). The antibody used against RB (clone 13A10) binds to the N-terminal region of the protein and detects RB regardless of phosphorylation status. Positive controls included CIN III for p16^{INK4A}, placenta for p21, anaplastic thyroid

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cancer for p53 and tonsil for RB. The staining pattern of RB, p53 and p21 was nuclear. p16^{INK4A} showed both nuclear and cytoplasmic staining.

Sections were scored semiguantitatively by a consultant genitourinary pathologist (D.B.). For nuclear positivity each core was given an estimated visual score between 0 - 100%, representing the percentage of positively stained neoplastic nuclei. The intensity of staining was also measured as: 1 (weak), 2 (medium) and 3 (strong). The final score was deduced by multiplying the percentage of staining by intensity to give an expression score from 0-300. p53 expression was always strong; therefore nuclear score alone was applied. Cytoplasmic expression of p16^{INK4A} was determined by intensity of staining alone. The core with highest score was selected for analysis. Statistical analysis was performed using StatsDirect software, version 2.60.6000. The correlations between antibodies were evaluated using Spearman's rank correlation test and the tumour type or HPV infection was evaluated by Chi-Square test or Fisher's exact probability test. Comparisons between antibodies expression in different histological subtypes of SCCs were restricted to usual type, verrucous and basaloid only. The warty group of tumours was too heterogeneous and included only 2 pure warty samples. Similar to previous papers, the cut-off points selected for antibody positivity were: >0 for p16^{INK4A 24}, \geq 5 for p53 and >5 for p21^{25, 26}. As RB is normally highly expressed in tissue we chose the median value of 240 and above (cut off \geq 240) as a high expression of RB in penile SCC. All analyses were 2-sided, p<0.05 was considered to be significant.

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Results

148 tumours were analysed which comprised 97 usual type SCCs, 17 basaloid, 15 verrucous, 7 mixed verrucous/usual type and 12 mixed warty and other SCC subtypes. The histopathological features of the tumours are listed in Table 1. Of these, 102 cases were also suitable for HPV analysis.

HPV infection

HPV DNA was detected in 57/102 (56%) penile SCCs. Of these HPV positive tumours, 39/57 (68%) were single, 17/57 (30%) multiple HPV type infections containing up to 6 low and high risk HPV types, and one sample (2%) contained unidentified HPV (Table 2). High-risk type 16 was the most prevalent type, present in 46/57 (81%) of HPV positive tumours. HPV18 was not detected. In the majority of HPV positive tumours, 33/57 (58%) HPV16 was the only HPV type detected (Table 2).

Differences in the HPV infections were observed between the histological subtypes of PSCC. For the usual type, HPV DNA was detected in 38/64 (59%) tumours, with high risk HPV16 present in 33/38 (87%) cases. Mixed warty subtypes were positive for HPV DNA in 6/11 (55%) cases, with HPV16 present in 3/6 (50%) cases. Basaloid tumours showed higher positivity, 10/13 (77%) for HPV DNA, with HPV type 16 present in 100% of these cases. In contrast, HPV was detected in only 3/13 (23%) verrucous tumours and HPV16 was not found (0%).

Immunohistochemistry

The positive expression of proteins and mean values are listed in Table 3. High RB expression (Figure 1A) was detected in 85/147 (58%) of penile SCC and significant differences were observed between histological groups (p<0.0001). A high percentage of verrucous cases (87%), an intermediate number of usual type (60%) and few basaloid cancers (12%) expressed high RB levels. The mean RB expression was two-fold lower in basaloid than in verrucous and usual subtypes.

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p16^{INK4A} demonstrated both nuclear and cytoplasmic staining (Figure 1B). Overall 47% of PSCCs are positive for cytoplasmic p16^{INK4A} expression, with significant differences between histological groups: 13% of verrucous, 52% usual and 94% of basaloid cases were p16^{INK4A} positive. Basaloid samples had very high mean expression of nuclear and cytoplasmic p16^{INK4A}, while mean expression of p16^{INK4A} in verrucous samples was very low and usual type showed intermediate values. There was a significant inverse correlation between RB and p16^{INK4A} expression (p<0.0001) in penile SCCs.

p53 immunodetection (Figure 1C) showed no significant difference between histological subtypes of SCC and was present in 79% of cases overall. Intensity of p21 staining was weak (Figure 1D) and present in 62%, with no difference in the expression between different SCC subtypes. No relationship was detected between p53 and p21 expression. There was a positive correlation between p21 and p16^{INK4A} expression (Spearman's $\rho = 0.658793$, p<0.0001) and negative correlation with RB (Spearman's $\rho = -0.499952$, p<0.0001).

There was also a strong positive correlation between HPV infection and p21 (p=0.0002) and p16^{INK4A} (p<0.0001) immunodetection and negative correlation with RB expression (p<0.0001) in penile SCC. p53 did not show any correlation with HPV infection (p=0.5682).

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Discussion

The rate of HPV infection in penile cancer varies widely, depending on the population studied and sensitivity and specificity of the method used. Our results suggest that in a developed country, unlike cervical cancer, penile cancer has at least two aetiologies: one HPV related and one unrelated (similar to vulvar cancer) as we detected HPV DNA in 56% (57/102) of PSCC cases. This is consistent with our previous report of HPV prevalence in PSCC of 54% ¹⁹ and the recent review that found 48% of 1,266 cases from 30 studies of invasive penile cancer were HPV positive ²⁷. We confirm existence of differences in HPV infection between histological subtypes ¹¹. HPV prevalence in usual type SCC varies between 11% and 71% ^{7, 8, 28}, and 59% (38/64) reported by us falls well within this range. As previously reported by our group, verrucous tumours were mostly HPV negative confirming the lack of HPV involvement in this neoplasm ²⁹, while basaloid carcinomas showed strong correlation to HPV infection ^{11, 28}.

HPV16 is the most prevalent type in our study and was detected in 81% (46/57) of positive samples and in over half of these as a single infection suggesting that this HPV genotype is more likely to contribute to the carcinogenic process. However, 17/57 (30%) patients had multiple HPV type infections which is similar to other penile studies ^{30, 31}. The significance of multiple HPV infections in cancer development is not clear and reports on increased risk for carcinoma in women with multiple HPV infections, as compared to those infected with a single HPV type are contradictory ^{32, 33}.

Comparing these results with protein expression reveals that basaloid tumours have an aetiology related to high-risk HPV infection, which manifests itself in high p16^{INK4A} and decreased RB expression, as has been shown in cervical cancer ³⁴. The aetiology of usual type SCC can only be attributed to HPV infection in approximately half of the tumours, showing corresponding loss of RB and gain of p16^{INK4A} protein expression. This is analogous

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to the involvement reported for HPV in carcinogenesis of anal SCC ³⁵ and tonsillar carcinoma ³⁶.

Detection of p53 protein by immunohistochemistry in penile cancer varies between 41.5% and 89% ^{21, 37, 38} and there is a lack of reports comparing different histological types. We detected p53 immunostaining in 79% (103/143) of penile SCCs and the value was very similar regardless of histology (Table 3). There was no correlation between high-risk HPV infection and p53 immunostaining, which is in agreement with previous reports on penile SCC ^{37, 38}. We used the DO-7 clone antibody which mainly detects mutated p53 but is also able to recognise wild type (wt) p53. Therefore, it may detect wt p53 when it is highly expressed in cells. High risk HPV infection also causes oncogenic stress to the cell which induces normal cellular responses such as increased p53 levels in order to arrest the cell cycle. Keratinocytes or fibroblasts expressing high risk HPV E7 protein are well known to overexpress wt p53, however, there is evidence that in E7 expressing cells wt p53 is transcriptionally inactive³⁹. Additional studies on p53 mutations are necessary to confirm its mutational status in these tumours.

p21 was expressed in 62% (88/143) of penile SCC with no significant difference between tumour subtypes. Lam and Chan ²¹ showed lower p21 expression in penile cancer but on much smaller cohort. Interestingly p21 did not correlate with p53 and some cases expressed high levels of p21 despite low or absent p53 protein, suggesting p53-independent activation of p21. Surprisingly, similar to tonsilar SCC ³⁶, we found positive correlation of p21 with HPV infection (p=0.0002). Additionally, p21 positively correlated with p16^{INK4A} (p<0.0001) and negatively with RB expression (p<0.0001). Funk at al ¹⁸ reported that high-risk HPV16 E7 protein can directly bind to p21 and abrogate a DNA damage-induced cell cycle arrest, despite high levels of p21. He suggested it is possible that the release of E2F from RB and inactivation of p16^{INK4A} and p21 are all necessary for the ability of E7 to bypass cell cycle

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arrest signals. On the other hand, there is emerging evidence that p21 in certain cancers may itself act as an oncogene and actually promote proliferation ⁴⁰.

To our knowledge this is the largest study to examine penile cancer pathogenesis by comparing HPV type with proteins commonly affected by HPV infection. We demonstrated, for the first time, that HPV infection in penile SCC disrupts the RB/ p16^{INK4A} pathway through downregulation of RB and elimination of cell-cycle control from p16^{INK4A}, manifesting itself in accumulation of p16^{INK4A}, which fails to block cell cycle progression. p21 was widely expressed, consistent with abnormal cell cycle regulation. However, p21 seems to be regulated independently from p53 and may be involved in oncogenic process. We confirm that penile tumours seem to have two different aetiologies: one related to HPV and one unrelated. These data suggest that use of the bivalent HPV16/18 prophylactic vaccine in men could reduce occurrence of penile SCC by about 45%.

Acknowledgements

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Titles and legends to Figures

Figure 1. Immunostaining results for penile squamous cell carcinoma. (A) Strong RB expression throughout the tumour. (B) p16^{INK4A} expression showing both cytoplasmic and nuclear positivity. (C) Strong positivity of p53 in basal areas of the tumour. (D) p21 expression, showing scattered staining only

Titles and legends to Tables

Table 1. Histological grade and stage of different subtypes of penile squamous cell carcinoma

Table 2. Human papillomavirus DNA detection by polymerase chain reaction method in different histological subtypes of penile squamous cell carcinoma

Table 3. Positive expression of RB, nuclear and cytoplasmic p16^{INK4A}, p53 and p21 in all penile squamous cell carcinomas and in regard to histological subtypes

P P P

SCC subtype		C	arade	(1-3)	S	tage (⁻	1-4)		
		1	2	3	1	2	3	4	No data
All SCCs	n=148	35	59	54	49	70	16	7	6
Usual type	n=97	13	49	35	28	46	12	6	5
Verrucous	n=15	12	3	0	8	7	0	0	0
Basaloid	n=17	0	2	15	5	7	3	1	1
Mixed warty	n=12	9	2		7	4	1	0	0

Table 1. Histological grade and stage of different subtypes of penile squamous cell carcinoma

SCC, squamous cell carcinoma; No data, there was no tumour stage available for six patients, which underwent penile circumcision or excision biopsy.

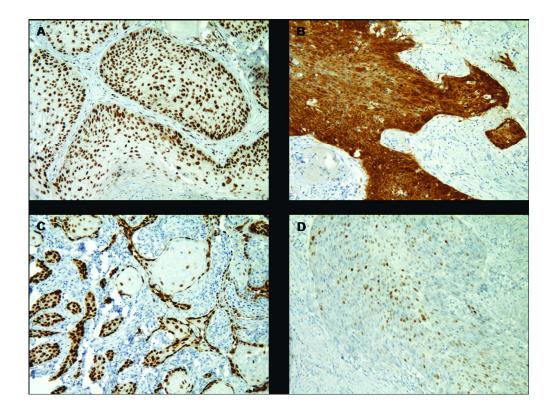
Table 2. Human papillomavirus DNA detection by polymerase chain reaction method in different histological subtypes of penile squamous cell carcinoma

			HF	PV g	enoty	ype c	distri	buti	on												
Histological subtype of SCC	n	HPV DNA positive	6	11	16	33	6 16	11 16	11 45	16 31	16 35	16 45	16 52	31 33	51 52	6 11 16	11 16 31	6 16 33 45	31 33 44 45	6 11 16 31	Х
																		-10	-0	33	
Usual type	64	38 (59%)	2	1	24	0	1	3	1	1	0	2	0	0	0	1	0	1	0	45 0	1
Basaloid	13	10 (77%)	0	0	9	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
Verrucous	13	3 (23%)	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	0	0
Mixed verrucous/usual type	1	0 (0%)	0	0	0	0	0	0	0	0	•0	0	0	0	0	0	0	0	0	0	0
Mixed warty	11	6 (55%)	0	1	0	1	0	0	0	0	1	0	0	1	0	0	1	0	0	1	0

SCC, squamous cell carcinoma; HPV, human papillomavirus; X, unclassified HPV genotype.

Table 3. Positive expression of RB, nuclear and cytoplasmic p16^{INK4A}, p53 and p21 in all penile squamous cell carcinomas and in regard to histological subtypes

SCC subtype	RB (≥24	10)	nuc p16 ^{INK4A} (>0)		0) cyt p16 ^{INK4A} (>0)			p53 (≥ 59	%)	p21 (>5%)		
	n (%)	mean	n (%)	mean	n (%)	mea	n	n (%)	mean	n (%)	mean	
All SCCs	85/147 (58)	205.6	65/144 (45)	54.2	67/144 (47)	1	.1	113/143 (79)	19.4	88/143 (62)	38.1	
Usual type	58/97 (60)	212	48/93 (52)	62.3	48/93 (52)	1	.2	75/93 (81)	20.9	58/94 (62)	41.7	
Verrucous	13/15 (87)	248	1/15 (7)	2.7	2/15 (13)	0	.1	13/15 (87)	15.4	8/13 (62)	17.8	
Basaloid	2/17 (12)	118.2	16/17 (94)	102.4	16/17 (94)	2	.2	14/17 (82)	17.5	15/17 (88)	47.4	
Mixed warty	6/11 (55)	199	2/12 (17)	16.7	3/12 (25)	0	.4	7/12 (58)	7.8	5/12 (42)	30.4	
IN U.Z.A.A			112.4.4	IN II Z A				0.				
nuc p16 ^{INK4A} ,	nuclear p16 ^{INK4A}	; cyt p16 [⊪]	, cytoplasm	ic p16 ^{™ĸ₄}	A							



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