Table 1. Intensity of p53 immunostaining

Melanocytic lesions	Staining intensity 0/(+)	+	++/+++	Total
Regular nevi (n = 10)				
PAb-1801	0	7	3	10
PAb-240	0	8	1	9
Dysplastic nevi (n = 11)				
PAb-1801	2	7	2	11
PAb-240	4	6	1	11
Superficial spreading Melanoma (n = 12)				
PAb-1801	2	6	4	12
PAb-240	4	7	1	12
Nodular melanoma (n = 10)				
PAb-1801	1	6	3	10
PAb-240	0	6	3	9

histochemical detection of a reliable tumour marker would therefore be a great advantage.

Immunostaining of p53 protein was examined in 43 benign and malignant melanocytic lesions using two common antibodies (Oncogene Science, NY): PAb-1801 against human wild-type and mutant p53 and PAb-240 against mutant p53 protein. Formalin-fixed and paraffin-embedded routine specimens were used to examine the practical value of p53 immunostaining in diagnostic evaluation of these lesions.

Table 1 shows that there was no significant difference between benign, premalignant and malignant melanocytic lesions with respect to intensity of p53 immunostaining (χ^2 test), although some of the malignant melanomas showed a strong, dot-like staining in the cytoplasm of the tumour cells. As a rule, both nuclear and cytoplasmic positivity were present, corresponding to the results with fresh tissue [4]. These findings indicate that immunopositivity does not discriminate between benign and malignant melanocytic proliferations.

The positive staining in a large proportion of regular nevi was surprising. However, recent reports stress that deterioration of the p53 protein may be introduced by formalin fixation [5], and positivity may therefore be artificial and without biological significance in some cases. For these reasons, p53 immunostaining using the present antibodies on formalin-fixed, paraffin-embedded specimens should not be used in the routine evaluation of melanocytic lesions.

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Unexpected *Prad*-1 Amplification in Multiple Simultaneous Localisations of Squamous Cell Carcinoma of the Head and Neck

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PRAD-1 IS the most recent addition to the list of genes localised in 11q13 region. This gene may be equivalent to the cyclin D1 and it has been proposed that *Prad*-1 plays a key role in the regulation of cell growth [1]. Its clinical significance is not yet defined, but its amplification has been described in parathyroid adenomas, centrocytic lymphomas, breast adenocarcinomas, as well as in squamous cell carcinoma cell lines [2–5]. Analysed by polymerase chain reaction (PCR) techniques, *Prad*-1 is often coamplified with *hst*1 and *int*-2 genes, and in squamous cell carcinomas, these amplifications show a trend to be associated with poor prognosis [6]. In our previous multiparametric and prospective study, we found *Prad*-1 amplifications (with dopamine receptor gene as a control) in 27 out of 51 (53%) squamous cell carcinomas, and amplifications were related to small T volume and tumour vascularisation [7]. 7 other patients, presenting multiple simultaneous head and neck localisations, and biopsied at the principal tumour site, were analysed by PCR. Five tumours were amplified (three non-amplified normal

Table 1.

Patient number	Localisation	TNM (AJC-UICC 86)	Number of *Prad*-1 copies
5	Oral cavity and piriform sinus	T4N1	4
45	Hypopharynx and oesophagus	T1N0	2 (not amplified)
76	Oral cavity and oropharynx	T4N0	7
93	Oral cavity and oropharynx	T4N2c	4
114	Hypopharynx and oesophagus	T4N0	2 (not amplified)
120	Oral cavity and oropharynx	T3N0	12
141	Hypopharynx and oropharynx	T3N2c	7

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mucosas being control). Interestingly, amplifications were detected in primary tumours except in association with oesophageal localisations (Table 1). Our previous results suggest that *Prad*-1 may play an important role in the early steps of carcinogenesis in squamous cell carcinoma and may also be implicated in the pathogenesis of multilocalisations.

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