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INTRODUCTION

Oropharyngeal cancer has increasing incidence in several continents. Oropharyngeal squamous cell carcinoma (OSCC) accounts for 90-95% of the tumor. In addition to the consumption of tobacco and alcohol, some Human Papillomavirus types have been associated with OSCC and in these cases are related to presentation and biological behavior. Hypomethylation of the Long Interspersed Nucleotide Element-1 (*LINE-1*) repetitive elements, a widely accepted surrogate of overall genomic DNA methylation content, was found to be associated with a poor prognosis in several cancers.

OBJECTIVE

To evaluate the profile of *LINE-1* methylation status in patients with OSCC diagnosed between 1999-2012.

METHODS

Records from 346 patients were collected, immunoexpression of p16 protein, qPCR for HPV16 *E6* and *LINE-1* methylation were performed, analyzed and submitted to bivariate (χ^2 and Mann Whitney) and survival (Kaplan-Meier and Log rank) analysis.

RESULTS

White (48%) males (89.8%), aged between 41-60 years (59%), alcohol-drinkers (88.2%) and smokers (94.5%) were most affected. Predominated moderately-differentiated tumors (79.8%), clinical stage IV (54.9%) and treated with radiotherapy (56.9%). 84 were identified cases of disease progression, 106 cases of recurrence, mostly local (72.6%). 271 came to death, of these 256 patients had confirmed death from cancer. Negative cases were predominant for p16 immunoexpression (90.8%) and for *E6* HPV16 qPCR (88.1%). Regarding HPV status, only 21 cases (6.1%) were classified as HPV positive. *LINE-1* methylation levels were largely heterogeneous (range, 16.39-73.95% and median, 54.33%) (Table 1).

Table 1: Sociodemographic, clinical-pathological and molecular features of patients (No=346).

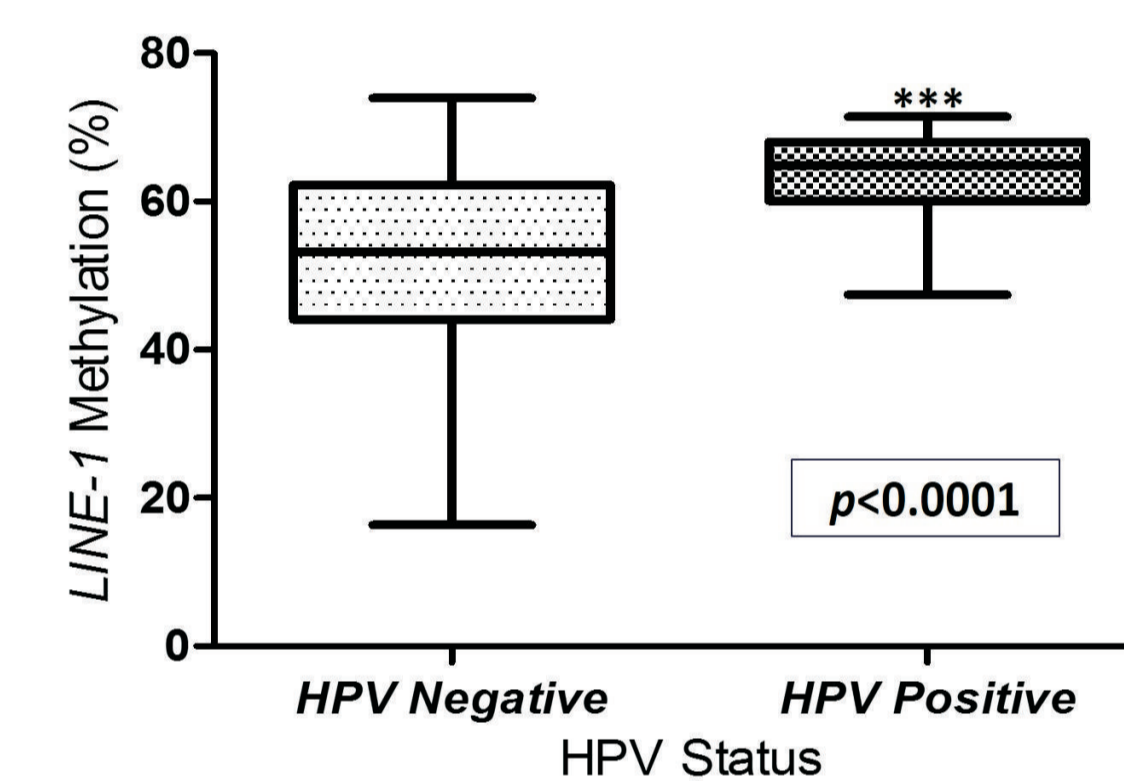
VARIABLE	CATEGORY	No. (%)
Gender	Male	308 (89.8%)
	Female	38 (11.2%)
Race	White	166 (48.0%)
	Brown	114 (32.9%)
	Black	66 (19.1%)
Age	≤40 years	8 (2.3%)
	41-60 years	204 (59.0%)
	>60 years	134 (38.7%)
Smokers	Yes and ex	327 (94.5%)
	No	19 (5.5%)
Alcohol-drinkers	Yes and ex	305 (88.2%)
	No	41 (11.8%)
Multiple tumor sites	One site	108 (31.2%)
	Two or more sites	238 (68.8%)
	Surgery	30 (8.7%)
Initial treatment	Radiotherapy	197 (56.9%)
	Radiotherapy and chemotherapy	68 (19.7%)
	Support	51 (14.7%)
Adjuvant treatment	Yes	27 (7.8%)
	No	319 (92.2%)
Clinical staging	I	26 (7.5%)
	II	43 (12.4%)
	III	87 (25.1%)
	IV	190 (54.9%)
Disease progression	Yes	84 (24.3%)
	No	262 (75.7%)
Recurrence	Yes	106 (30.6%)
	No	240 (69.4%)
Type of recurrence	Local	77 (72.6%)
	Regional	18 (17.0%)
	Distance	11 (10.4%)
Second primary tumor	Yes	17 (4.9%)
	No	329 (95.1%)
Death	Yes	271 (78.3%)
	No	75 (21.7%)
Death from cancer	Yes	256 (97.0%)
	No	8 (3.0%)
WHO grading	Well differentiated	12 (3.5%)
	Moderately differentiated	276 (79.8%)
	Poorly differentiated	58 (16.8%)
p16 immunoexpression	Positive	32 (9.2%)
	Negative	314 (90.8%)
E6 HPV16 qPCR	Positive	40 (11.9%)
	Negative	295 (88.1%)
HPV Status	Positive	21 (6.1%)
	Negative	325 (93.9%)
LINE-1 Methylation	Hypermethylation (63.41%)	20 (8.0%)
	Hypomethylation (52.64%)	231 (92.0%)

Association between advanced clinical stage and treated with radiotherapy ($p<0.0001$), smokers ($p=0.013$), disease progression ($p=0.022$), recurrence ($p=0.008$) and died ($p=0.001$) (Table 2).

Table 2: Distribution of socio-demographic and clinical-pathological variables according to clinical staging (No=346).

VARIABLE	CATEGORY	CLINICAL STAGING		
		Early (I + II) No. (%)	Advanced (III + IV) No. (%)	p value
Smokers	Yes and ex	69 (19.9%)	258 (74.6%)	0.013
	No	0 (0.0%)	19 (5.5%)	
Initial treatment	Surgery	18 (5.2%)	12 (3.5%)	<0.0001
	Radiotherapy	42 (12.1%)	155 (44.8%)	
	Radiotherapy and chemotherapy	7 (2.0%)	61 (17.6%)	
Disease progression	Support	2 (0.6%)	49 (14.2%)	0.002
	Yes	10 (2.9%)	74 (21.4%)	
Recurrence	No	59 (17.1%)	203 (58.7%)	0.008
	Yes	30 (8.7%)	76 (22.0%)	
Death	No	39 (11.3%)	201 (58.1%)	0.001
	Yes	44 (12.7%)	277 (65.6%)	
Death	No	25 (7.2%)	50 (14.5%)	0.001
	Yes	44 (12.7%)	277 (65.6%)	

LINE-1 methylation level was significantly higher in HPV-positive OSCC patients than in HPV-negative ones (median, 63.41% and 52.64%, respectively; ($p<0.0001$)) (Graphic 1).

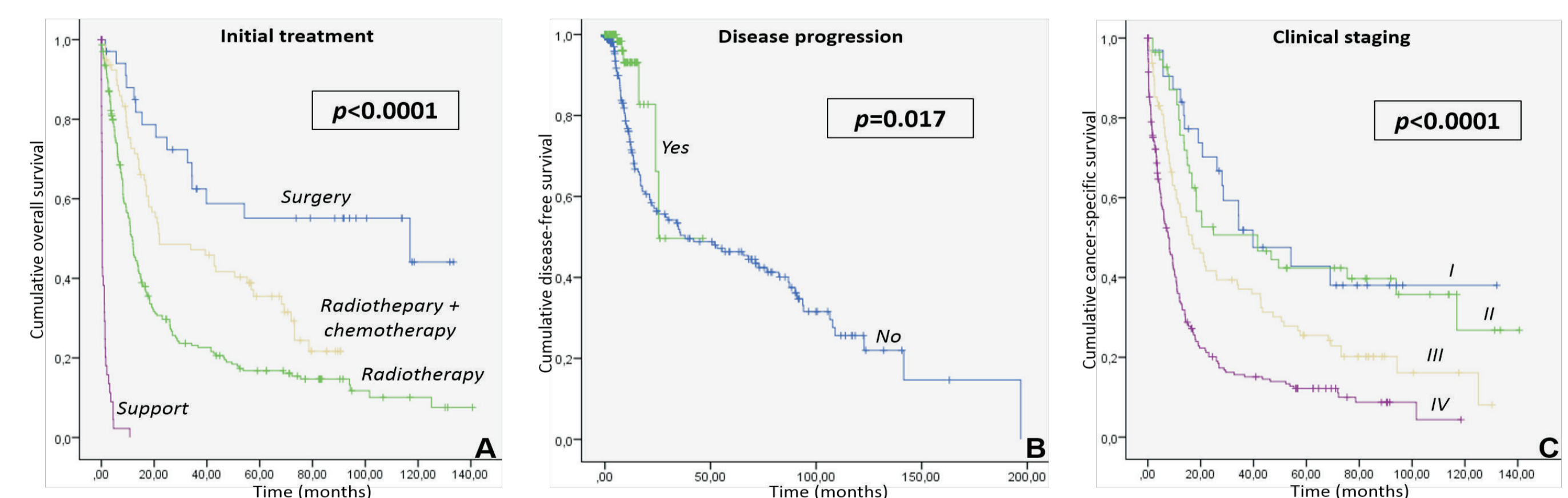


Graphic 1: Cumulative *LINE-1* methylation index of HPV-negative OSCC (52.64%; n=231) and HPV-positive OSCC (63.41%; n=20) depicted in box plots (No=251).

Overall survival (OS) median was 28.39 months. Patients in clinical stage IV ($p<0.0001$), in supportive treatment ($p<0.0001$) (Graphic 2A) and with disease progression ($p<0.0001$) had a lower OS.

Already disease-free survival (DFS) was 22.96 months. Group with disease progression ($p=0.017$) (Graphic 2B), death ($p<0.0001$) and death from cancer ($p=0.027$) had a lower DFS.

Regarding the cancer-specific survival (CSS), clinical stage IV ($p<0.0001$) (Graphic 2C), supportive treatment ($p<0.0001$) and disease progression ($p<0.0001$) had a lower CSS.



Graphic 2: Kaplan-Meier curves. A - Overall survival; B - Disease-free survival; C - Cancer-specific survival.

CONCLUSIONS

- The profile of patients with OSCC analyzed doesn't differ from literature for developing countries.
- HPV-negative status associated with *LINE-1* hypomethylation in OSCC.
- Habits, treatment, staging and death are important prognostic factors in this population, showing the aggressiveness of OSCC.

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