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Evaluation of the cyclin-dependent kinase inhibitor p21^{Cip1} in epithelial ovarian tumors of low malignant potential and adenocarcinomas

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Summary. Objective: In view of the controversial information on the significance of the cyclin-dependent kinase inhibitor p21^{Cip1} in ovarian cancer, we conducted a retrospective investigation to clarify the relationships of this protein to proliferation rate, clinicopathological variables and prognosis of epithelial ovarian tumors. Methods: Paraffin-embedded tissue from 43 ovarian tumors of low malignant potential (LMP) and 82 primary ovarian adenocarcinomas were stained immunohistochemically for p21^{Cip1}, p53 protein and Ki-67 antigen (a marker of cell proliferation). Results: p21^{Cip1} levels were significantly higher in LMP tumors (p<0.001) as well as in early stage adenocarcinomas (p=0.021) and those associated with minimal residual disease (p=0.008). However, no relationship existed between p21^{Cip1} expression and the proliferation rate of adenocarcinomas or LMP tumors. In the vast majority of LMP tumors p21^{Cip1} expression was not accompanied by p53 accumulation. This p21^{Cip1}-positive/p53-negative phenotype prevailed in the early stage (p=0.026), lower grade (p=0.018) adenocarcinomas as well as in those left with minimal residual disease (p=0.059). In patients with lower grade adenocarcinomas, decreased p21^{Cip1} expression was adversely related to poor overall survival on its own (p=0.0500) and when combined with p53 protein overexpression (p=0.0323). In multivariate analysis, only the stage remained as the independent predictor of survival. Conclusions: Decreased p21^{Cip1} expression is related to several indicators of aggressiveness in ovarian adenocarcinomas and seems to be differentially regulated in LMP tumors and adenocarcinomas. On the contrary, deregulation of p21^{Cip1} expression does not seem to participate in the pathogenesis of LMP tumors. Furthermore, although p21^{Cip1} alone or combined with p53 is of prognostic significance in lower grade adenocarcinomas, it does not appear to add to the information gained from traditional

prognosticators.

Key words: p21^{Cip1}, Immunohistochemistry, Ovarian carcinomas, LMP, Prognosis

Introduction

Ovarian cancer is the most common cancer in women to be diagnosed at advanced stage and is the leading cause of death from gynecological malignancy in western countries (Landis et al., 1999). Despite optimal surgical debulking and a pathologically complete response to primary chemotherapy, half of the patients will develop a recurrence and will eventually die of their disease within 2 years, denoting the inadequacy of traditional prognosticators in predicting clinical outcome in individual cases (Landis et al., 1999). Therefore, there has been a growing interest in the characterization of more reliable prognostic factors closely related to tumor cell biology, which would assist in the identification of patients who could benefit from more aggressive therapy.

Although our understanding of the molecular mechanisms intervening and conducting the biological behavior of ovarian carcinomas is certainly incomplete, the intuitive notion that ovarian cancer, like most cancers, arises as a result of perturbations in cell-cycle regulation has rapidly been appreciated (Weinberg, 1991). In particular, disruption of the mechanisms regulating transition from G1 to S phase appears to be a key issue in cancer evolution. Gaining insight into the molecular machinery coordinating orderly progression into S phase has unraveled the crucial role of various inhibitory proteins, known as cyclin-dependent kinase inhibitors (CKIs), which bind to and down-regulate the activities of cyclin/cyclin-dependent kinase (Cdk) complexes required for driving the cell through the various restriction points of the cell cycle (Granna and Reddy, 1995; Sherr, 1996). Currently, two groups of CKIs have been identified: the Cip/Kip family, with

broad cyclin-Cdk reactivity, and the INK4 family with a much more selective inhibitory effect on cyclin D-Cdk 4/6 complexes.

The WAF1/Cip1/SD1 gene product p21 is a member of the Cip/Kip family of CKIs, able to arrest the cell cycle at the G1 checkpoint, by interacting with cyclin/cdk complexes as well as with the DNA polymerase δ replicating factor proliferating cell nuclear antigen (PCNA) (Zhang et al., 1994; Luo et al., 1995). The inhibitory function of p21^{Cip1} on DNA synthesis requires the formation of quarternary complexes composed of cyclins, CDKs and PCNA (Zhang et al., 1993) and is thought to be stoichiometrically regulated, being exerted only when p21^{Cip1} is in molar excess. If the ratio of p21^{Cip1} to cdk is less than one, p21^{Cip1} serves only as an assembling factor for cdk complex and does not inhibit cdk activity. p21^{Cip1} binding prevents CDK2/cyclin E-catalysed phosphorylation of retinoblastoma gene protein (pRb) with consequent abrogation of pRb function as a negative regulator of G1/S restriction point (Cordon-Cardo, 1995). Evidence has been provided that the WAF1 gene possesses at least two p53-responsive sites in its promoter region. Binding of p53 to these elements activates the transcription of p21^{Cip1}, suggesting that the latter is a potent downstream mediator of the antiproliferative function of the wildtype p53 (El-Deiry et al., 1993). On the other hand, p21^{Cip1} induction also occurs in a p53-independent manner, having been observed in various normal tissues during development or differentiation in the absence of p53 activation and in tumor cells with mutated p53 (Johnson et al., 1994; Macleod et al., 1995; Parker et al.,

Recently, a number of investigators have looked into $p21^{Cip1}$ expression in ovarian cancer with contradictory findings (Anttila et al., 1999; Costa et al., 1999; Shimizu et al., 1999; Werness et al., 1999; Ferrandina et al., 2000 Schmider et al., 2000; Sengupta et al., 2000). This controversial information regarding the clinical significance of $p21^{Cip1}$ in ovarian carcinomas has prompted us to undertake the present study. The aim was to determine the possible associations of $p21^{Cip1}$ expression with clinicopathological parameters, p53 expression and patients' outcome in a large series of epithelial ovarian tumors.

Materials and methods

Patients

A total of 125 consecutive cases of primary epithelial ovarian tumors of low malignant potential (LMP) and adenocarcinomas, diagnosed and treated at Alexandra General Hospital and Iasso Gynecological Institute of Athens between 1989 and 1999, were enrolled in the present study. Eighty-two specimens were defined as ovarian adenocarcinomas, whereas 43 specimens fulfilled the criteria of LMP tumors (Scully and Sobin, 1987; Russel, 1994). Histological classifica-

tion of tumors was carried out according to the World Health Organization System (Scully and Sobin, 1987). Poorly differentiated adenocarcinoma was the diagnosis for those adenocarcinomas that did not show evident cellular differentiation. Adenocarcinomas were graded as well, moderately and poorly differentiated (Scully and Sobin, 1987) and patients were assigned a clinical stage according to the International Federation of Gynecology and Obstetrics (FIGO, 1989) standards. Surgical and pathological findings and postoperative abdomino-pelvic computerized tomography (CT) scans were used to determine the FIGO stage for the ovarian adenocarcinomas and the residual disease after the initial surgery. Persistence of tumor masses of <2 cm was defined as minimal residual disease, whereas the presence of masses of >2 cm in diameter was defined as bulk residual disease (Anttila et al., 1999; Sengupta et al.,

All patients with carcinomas had undergone total abdominal hysterectomy with bilateral salpingo-oophorectomy followed by chemotherapy. None had received preoperative chemotherapy or radiothrerapy. The clinicopathological characteristics of these patients are summarized in Table 1. Follow-up information was available in 71 patients. By the time this study was undertaken, 22 patients had died of their disease after a median survival of 24 (1-60) months whereas the median follow-up period for the remaining 49 patients was 52.5

Table 1. Clinical data of patients with ovarian adenocarcinomas.

Age (years) Median (range)	53 (20-78)
, ,	33 (20-70)
Histological type	00 (050/)
Serous	29 (35%)
Mucinous Endometrioid	10 (12%)
Clear cell	17 (21%) 19 (23%)
Adenocarcinoma poorly differentiated	7 (9%)
• •	7 (970)
Histological grade	(= (10+i)
1	15 (18%)
2	39 (48%
3	28 (34%)
FIGO stage	
I	31 (38%)
II	4 (5%)
III	36 (44%)
IV	7 (9%)
Not specified	4 (5%)
Primary residual tumor	
None	35 (43%)
<2 cm	10 (12%)
>2 cm	37 (45%)
Adjuvant chemotherapy	
Platinum-containing	57 (70%)
Non platinum-containing	21 (26%)
No data	4 (5%)
Clinical outcome	, ,
Died of ovarian cancer	22 (27%)
Alive	49 (60%)
Lost to follow-up	11 (13%)
	(,-,

(14-126) months.

Immunohistochemistry

Tissues were fixed immediately after removal in 10% formalin and embedded in paraffin. Sections for immunohistochemistry were cut at 3 µm, mounted on Superfrost/Plus glass slides and left to dry overnight at 37 °C. All cases were stained for p21^{Cip1}, p53 and Ki-67 using the following monoclonal antibodies: 4D10 for p21^{Cip1} (Novocastra, Burlingame, CA) diluted 1:20; D0-1 for p53 (Dako, Carpinteria, CA) diluted 1:100; and MIB-1 for Ki-67 (YLEM s.r.l., Rome, Italy) purchased prediluted by the manufacturer. The incubation time was 1 hour at room temperature for p53 and Ki-67 and 18 hours at 4 °C for p21^{Cip1}. Before applying the primary antibody, sections were treated in a microwave oven at 750 W for a total of 30 minutes (three cycles of 10 minutes each) in 10 mmol/L sodium citrate buffer pH 6.0. Slides were left to cool down in the antigen retrieval solution for 20 minutes. Then, the standard streptavidinbiotin complex immunoperoxidase technique (Histostain-SP Kit, Zymed Laboratories, San Francisco, CA) was used. Peroxidase activity was visualized with chromogen 3,3'-diaminobenzidine tetrachloride in 0.5 mmol/L Tris Buffer. The slides were counterstained with hematoxylin. Known positive controls (a colorectal carcinoma for DO-1, normal tonsillar tissue for MIB-1 and a previously irradiated piece of skin for 4D10) as well as negative controls (sections in which the primary antibody was substituted with non-immune mouse serum) were also stained in each run.

Assessment of staining

Staining for all antibodies was assessed by two observers without any knowledge of the clinical data. Whenever a difference of greater than 5% was observed between the two readings, slides were reviewed jointly and a consensus was reached. To this end, nuclei from about 2000 tumor cells from 20 systematically randomized fields (x400) throughout the whole section were counted and the labeling index (LI) was calculated as the percentage of labeled nuclei out of the total number of tumor cells counted. All clearly identifiable nuclear staining beyong background was recorded as positive for Ki-67 (Anttila et al., 1999; Palazzo et al., 2000), but for p53 and p21^{Cip1} the threshold of positivity was raised to 10% and 1% respectively, according to published convention (Palazzo et al., 2000; Schmider et al., 2000; Sengupta et al., 2000). Given that the immunostaining intensity did not significantly vary from case to case, the expression of various markers was assessed quantitatively.

Statistical analysis

p21^{Cip1} LI, Ki-67 LI and p53 LI were considered as continuous variables. The relationships between the

expression of the three markrers and clinicopathological parameters were examined by the Mann-Whitney U-test and Kruskal-Wallis analysis of variance. The associations of the categorized form of p21^{Cip1} (<1% vs \geq 1%) and p53 (<10% vs \geq 10%) were tested with Pearson's chi-square test with continuity correction and Fisher's exact test. The interrelations among p53, p21^{Cip1} and Ki-67 levels were analysed with Spearman's correlation coefficient.

Survival analysis was restricted to adenocarcinomas, given that LMP tumors follow a substantially more favorable clinical course. The prognostic effect of the various parameters (age, histological type, grade, FIGO stage, volume of residual disease, type of chemotherapy, Ki-67 LI, p53 expression, p21^{Cip1} LI and the combined p21^{Cip1}/p53 phenotype) was tested by univariate and multivariate analysis. Univariate analysis was based on life tables and differences among groups were tested for significance using the log-rank test. Age was categorized on the basis of the median value. p53 and p21^{Cip1} expression was categorized as positive versus negative using 10% and 1% as cut-off points respectively. Ki-67 LI was categorized on the basis of 10% (Palazzo et al., 2000; Sengupta et al., 2000). Multivariate analysis was performed using the stepwise Cox's regression model (forward selection of variables) to evaluate the predictive power of each variable independently of the others. In order to avoid any "data-driven" categorization, numerical variables were entered in continuous form only. Statistical calculations were performed using the SPSS for Windows Software (SPSS Inc., Chicago IL). A p-value of less than or equal to 0.05 was considered indicative of a statistically significant difference.

Results

p21^{Cip1} expression in LMP tumors and ovarian adenocarcinomas

Using 1% as the cut-off point, positive staining for p21^{Cip1} was observed in 35/43 (81.4%) LMP cases and in 39/82 (47.6%) adenocarcinoma cases (χ^2 , p=0.001). p21^{Cip1} LIs ranged from 1% to 90% in the former group and from 1% to 60% in the latter. Positivity was confined to the nucleus and had a granular quality (Fig. 1) The distribution of positive nuclei was uniform throughout the neoplastic tissue. In a few (10%) adenocarcinoma cases a faint cytoplasmic reactivity was also noted which was disregarded as nonspecific. Exclusively cytoplasmic localization of the protein was not observed in our series. Non-neoplastic elements such as ovarian surface epithelium, ovarian cortical cells, lymphocytes and endothelial cells were not labeled. The median p21^{Cip1} LI was significantly higher in LMP tumors (8%) compared to carcinomas (3%) (Mann-Whitney U-test, p<0.001, Table 2). When comparisons between LMP cases and carcinomas were made for

serous and mucinous types separately, a difference in the incidence of $p21^{Cip1}$ positivity between carcinomas and LMP tumors was recorded for serous (Fisher's exact test, p<0.001; Mann-Whitney U-test, p<0.001), but not for mucinous type (Fisher's exact test, p=0.240; Mann-Whitney U-test, p=0.248) (Table 2). However, mucinous LMP tumors were less often (57.1%) $p21^{Cip1}$ -positive than serous ones (93.1%) (Fisher's exact test, p=0.009).

Correlation of p21^{Cip1} expression in adenocarcinomas with clinicopathological parameters

The rate of p21^{Cip1} positivity significantly decreased with advancing stage (I-II versus III versus IV) ($\chi^2,$ p=0.044) as well as with bulk residual disease ($\chi^2,$ p=0.023) (Table 3) but was not related to tumor grade ($\chi^2,$ p=0.423). The same applied to p21^{Cip1} LI (Mann-Whitney U-test, p=0.021, p=0.008, p=0.294 respectively). There was an almost significant relationship between p21^{Cip1} expression and histological type ($\chi^2,$ p=0.059), in the sense that serous and

mucinous types tended to be less often positive than clear-cell and endometrioid types. $p21^{Cip1}$ levels were also lower in the former two types (Kruskal-Wallis ANOVA, p=0.042)

p53 immunostaining

p53 immunoreactivity was confined to tumor cell nuclei. Normal surface epithelium as well as ovarian cortical cells and inflammatory cells did not express p53. According to the 10% cut-off point, 39 out of 82 (47.6%) adenocarcinomas and only one out of 43 (2.3%) LMP tumors were considered p53-positive (x^2 , p<0.001). Serous adenocarcinomas were more often p53-positive than their LMP counterparts (χ^2 , p<0.001), but mucinous adenocarcinomas were not (χ^2 , p=0.355, Table 2). Within the adenocarcinoma group, the rate of p53-positivity significantly increased with high grade (χ^2 , p=0.001) and marginally with advancing stage category (χ^2 , p=0.075), but remained unaffected through histological types (χ^2 , p=0.250) and residual disease (χ^2 ,

Table 2. p21^{Cip1}, p53 and Ki-67 expression in LMP tumors and ovarian adenocarcinomas.

HISTOLOGY	p21 ^{Cip1}	p21 ^{Cip1} EXPRESSION		p53 EXPRESSION	
	Pos/neg	p21 ^{Cip1} LI (%) (median,range)	Pos/neg	p53 LI (%) (median, range)	(median, range)
LMP tumors (n=43)	35/8	8.0, 1.0-90.0	1/42	40.0, -	5.0, 0.5-50.1
Serous (n=29)	27/2	8.0, 1.0-90.0	0/29	-, -	3.0, 0.5-12.1
Mucinous (n=14)	8/6	22.5, 1.0-90.0	1/13	40.0, -	10.0, 1.1-50.2
Carcinomas (n=82)	39/43	3.0, 1.0-60.0	39/43	65.0, 10.0-99.2	15.0, 0.8-95.1
Serous (n=29)	9/20	2.0, 1.0-5.0	18/11	67.5, 25.0-99.1	20.0, 0.8-55.1
Mucinous (n=10)	3/7	10.0, 5.0-10.2	3/7	30.0, 20.1-50.2	11.0, 4.0-60.2

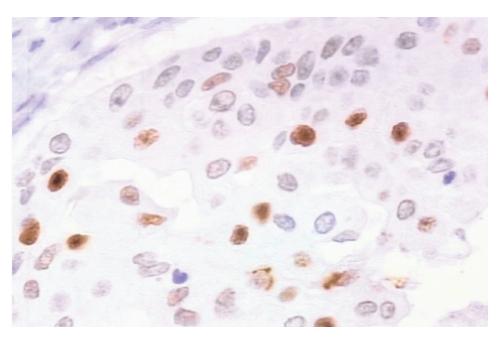


Fig. 1. p21^{Cip1} immunostaining in an endometrioid grade 2 adenocarcinoma. Many neoplastic nuclei are stained. p21^{Cip1}LI in this case was 25.5%.x 250

p=0.254). There was also no relationship between p53 LI and histological type (p=0.263), histological grade (p=0.229), FIGO stage (p=0.984) or the volume of residual disease (p=0.572) (Table 3).

Ki-67 immunostaining

Ki-67 immunoreractive cells were readily identified in all cases but Ki-67 LI was much higher in

adenocarcinomas compared with LMP tumors (Mann-Whitney U-test, p<0.001) and in mucinous LMP compared to serous LMP tumors (p=0.001) (Table 2). High grade adenocarcinomas displayed higher percentages of Ki-67-positive cells (Kruskal- Wallis test, p=0.002) but Ki-67 LI was unrelated to stage (p=0.082) or histological type (p=0.598). In addition, higher proliferation rate correlated with bulk residual disease (Mann-Whitney U-test, p=0.002) (Table 3).

Table 3. p21^{Cip1}, p53 and Ki-67 expression as related to clinicopathological variables in ovarian adenocarcinomas.

VARIABLE	p21 ^{Cip1} EXPRESSION		p53 E	p53 EXPRESSION	
	Pos/neg	p21 ^{Cip1} LI (%) (median,range)	Pos/neg	p53 LI (%) (median, range)	(median, range)
Histological type					
Serous (n=29)	9/20	2.0, 1.0-5.0	18/11	67.5, 25.0-99.1	20.0, 0.8-55.1
Mucinous (n=10)	3/7	10.0, 5.0-10.0	3/7	30.0, 20.1-50.2	11.0, 4.0-60.2
Endometrioid (n=17)	12/5	2.5, 1.0-35.0	8/9	65.0, 10.1-95.2	15.0, 8.2-80.3
Clear cell (n=19)	11/8	4.0, 1.0-60.0	6/13	45.0, 10.0-90.1	12.0, 4.1-60.3
Adenocarcinoma poorly differentiated (n=7)	4/3	2.5, 2.0-20.0	4/3	67.5, 60.0-90.3	20.0, 8.2-95.1
Histological grade					
1 (n=15)	9/6	3.0, 1.0-10.0	2/13	52.5, 30.2-75.1	12.0, 0.8-70.1
2 (n=39)	19/20	5.0, 1.0-60.0	17/22	50.0, 10.0-99.2	12.0, 3.1-65.2
3 (n=28)	11/17	2.0, 2.0-20.0	20/8	70.0, 10.0-99.1	37.5, 5.1-95.2
FIGO stage					
I/II (n=35)	22/13	3.0, 1.0-60.0	12/23	60.0, 10.0-99.2	12.0, 0.8-70.3
III (n=36)	12/24	2.5, 1.0-20.0	22/14	62.5, 10.0-99.0	20.0, 3.0-95.2
IV (n=7)	3/4	2.0, 1.0-10.0	3/4	50.0, 30.1-95.2	35.0, 3.0-75.1
Primary residual tumor					
Absent or minimal (n=45)	27/18	3.0, 1.0-60.0	18/27	70.0, 10.0-99.1	12.0, 0.8-70.1
Bulk (n=37)	12/25	3.0, 1.0-20.0	21/17	60.0, 10.0-95.2	30.0, 3.0-95.2

Table 4. Associations of the combined p21^{Cip1}/p53 expression with clinicopathological variables of ovarian tumors.

VARIABLE	p21 ^{Cip1} -/p53+	p21 ^{Cip1} -/p53-	p21 ^{Cip1} +/p53-	p21 ^{Cip1} +/p53+	
Histology					
LMP tumors	2	6	35	0	
Adenocarcinomas	24	19	24	15	
	p<0.001	(df=3)			
Histological type					
Serous	15	5	6	3	
Mucinous	1	6	1	2	
Endometrioid	2	3	6	6	
Clear cell	4	4	9	2	
Adenocarcinoma poorly differentiated	2	1	2	2	
, ,	p=0.018	(df=12)			
Histological grade					
1/2	11	15	20	8	
3	13	4	4	7	
	p=0.018	(df=3)			
FIGO stage					
1/11	5	8	15	7	
III/IV	17	11	7	8	
	p=0.026	(df=3)			
Primary residual tumor					
Absent or minimal	9	9	18	9	
Bulk	15	10	6	6	
	p=0.059				

Correlations among p21Cip1, p53 and proliferation rate

A moderately strong negative correlation emerged between p21^{Cip1} expression and proliferation rate (rho=0.337, p<0.001) as well as between p21^{Cip1} and p53 (rho=-0.347, p<0.001). When, however, LMP tumors and adenocarcinomas were examined separately, the relationship between p21^{Cip1} and proliferation rate disappeared. p53 expression, on the other hand, exerted a positive, though less strong, effect on proliferation rate (rho=0.496, p<0.001).

Combined p21^{Cip1} /p53 expression

According to the combined p21Cip1/p53 expression, our cases were divided into four subsets: p21^{Cip1}positive/p53-negative, p21^{Cip1}-negative/p53-positive, p21^{Cip1}-positive/p53-positive and p21^{Cip1}-negative/p53negative. The vast majority (81.4%) of LMP tumors belonged to the p21^{Cip1}-positive/p53-negative group whereas in adenocarcinomas p21^{Cip1}-positive/p53negative and p21^{Cip1}-negative/p53-positive phenotypes were equally common (28%) (χ^2 , p<0.001) (Table 4). The distribution of these subsets was also significantly different across the various grade categories (χ^2 , p=0.018), in the sense that the p21^{Cip1}-positive/p53negative phenotype prevailed in grades 1 and 2, whereas the p21^{Cip1}-negative/p53-positive phenotype was most often encountered in grade 3 cases. The combined $p21^{Cip1}/p53$ phenotype was also related to stage (χ^2 , p=0.026) and marginally to the presence of residual tumor (χ^2 , p=0.059), with the majority of p21^{Cip1}positive/p53-negative cases presenting with early (I-II) stages and being left with minimal residual tumor after

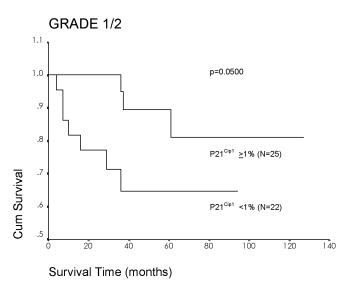


Fig. 2. Overall survival of patients with grade 1 and 2 adenocarcinomas in relation to p21^{Cip1} status. Patients with p21^{Cip1}-negativity tended to fare substantially worse than those with p21^{Cip1}-positive tumors. (Kaplan Meier curves, p=0.0500).

surgery. Most clear-cell adenocarcinomas expressed the $p21^{Cip1}$ -positive/p53-negative phenotype in contrast to serous adenocarcinomas, most of which belonged to the $p21^{Cip1}$ -negative/p53-positive group (χ^2 , p=0.018) (Table 4).

Survival analysis

In univariate survival analysis, high grade (p=0.0387), advanced stage (p=0.0003), the presence of bulk residual tumor (p=0.0010) and increased (>10%) Ki-67 LI (p=0.0127) portended a decreased overall survival in patients with adenocarcinomas (Table 5). p21^{Cip1} failed to reach significance in this regard. However, when patients were stratified by grade, the absence of p21Cip1expression exerted an unfavorable effect on the survival of patients with grades 1 and 2 (p=0.0500). The mean survival rate was 66 months for the p21^{Cip1}-negative group as opposed to 112 months for the p21^{Cip1}-positive one (Fig. 2) The combined p21^{Cip1}/p53 expression also affected survival in patients with lower grade adenocarcinomas only (p=0.0323), with p21^{Cip1}-negative/p53-positive cases displaying the shorter mean survival time (43 months) among the four p21^{Cip1}/p53 phenotypes (Fig 3) All the examined parameters were entered into Cox's analysis. However, stage emerged as the single independent predictor of overall survival (Table 5).

Discussion

The molecular basis of ovarian cancer has been a subject of intense scrutiny in recent years. However, histological and biological heterogeneity, as well as

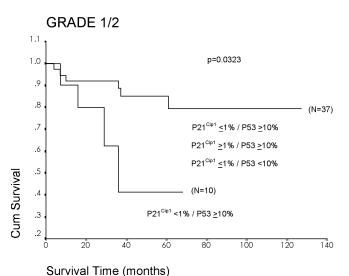


Fig. 3. Overall survival of patients with grade 1 and 2 adenocarcinomas in relation to the dichotomized p21^{Cip1}/p53 phenotype. Survival rates were lower in the p21^{Cip1}-negative/p53-positive group than in the remaining three phenotypes. (Kaplan Meier curves, p=0.0323).

uncertainties about the cell of origin, have impeded progress in our understanding of the underlying molecular mechanisms in these neoplasms. In search of the key events intervening in the development and progression of ovarian carcinomas, considerable attention has been directed toward the aberrations of cell-cycle regulatory proteins.

In the present study, we selected an immunohistochemical approach to evaluate the expression of p21^{Cip1} in ovarian cancer. Given that mutations in the coding region of WAF1 gene are extremely rare in a wide variety of human tumors (Shiohara et al., 1994), including those of the ovary (Wan et al., 1996), immunohistochemistry provides valuable information on the WAF1 gene functional status. The frequency of p21^{Cip1} expression in ovarian carcinomas (analysed by immunohistochemical and molecular methods) reported in the literature fluctuates between 42% to 78% (Barboule et al., 1995; Anttila et al., 1999; Costa et al., 1999; Werness et al., 1999; Palazzo et al., 2000; Schmider et al., 2000; Sengupta et al., 2000), although in two studies this percentage is much lower (Lukas et al., 1997; Shimizu et al., 1999). In our series the rate of p21^{Cip1} immunopositivity rises to 47.6%, comparing well with most published studies. This finding, in association with the fact that nonneoplastic ovarian tissue, including celomic epithelium, revealed no p21^{Cip1} expression (Werness et al., 1999; Ferrandina et al., 2000), has led some investigators to hypothesize that p21^{Cip1} activation may be associated with ovarian tumorigenesis (Ferrandina et al., 2000), in alignment with data reported by other investigators on lung, bladder, breast and brain tumors (Jung et al., 1995; Korkolopoulou et al., 1998, 2000; Rey et al., 1998; Takeshima et al., 1998).

In LMP tumors, p21^{Cip1} expression was present in higher levels than in adenocarcinomas providing evidence for a different pathogenetic basis of these two major categories of epithelial ovarian tumors. Analogous findings have been recently published by us and other authors (Masciullo et al., 1999; Korkolopoulou et al., 2002) regarding the differential expression of p27^{Kip1}, a

CKI structurally related to p21^{Cip1}, in LMP and adenocarcinomas. This picture is in harmony with the well-known growth inhibitory role of p21^{Cip1} suggested by in vitro data (El-Deiry et al., 1993) and, seemingly, in contrast with the proposed role of p21^{Cip1} in ovarian tumorigenesis as mentioned above (Ferrandina et al., 2000). To explain this disparity between theory and practice, it has been postulated that tumor cells may modify p21^{Cip1} expression attenuating its inhibitory activity (Jung et al., 1995). Alternatively, Ferrandina et al. (2000) have proposed that elevation of p21^{Cip1} might represent a biochemical adaptation aiming to overcome the rapid proliferation rate of tumorous tissue. Additional experiments (i.e. gene transfection, transgenic and knock out experiments) are required to elucidate the role of p21^{Cip1} in ovarian cancer initiation.

Opinions differ as to the association between p21^{Cip1} expression and clinicopathological indicators of aggressiveness in ovarian adenocarcinomas. In most studies, an inverse relationship between p21^{Cip1} levels and stage, grade or volume of residual disease (Anttila et al., 1999; Costa et al., 1999; Ferrandina et al., 2000; Schmider et al., 2000) has been established, but other authors (Werness et al., 1999; Sengupta et al., 2000) have disputed such a relationship. This variation largely results from the different positivity criteria imposed in these studies. In the present investigation, in order to circumvent the danger of using cut-off points, p21Cip1 exression was considered a continuous variable and all the associations between loss of p21^{Cip1} and clinicopathological parameters of aggressiveness were also obtained when p21^{Cip1} was tested as a continuous variable. Our findings are consonant with those of the former group of investigations and seem to advocate that p21^{Cip1} loss of function might be a late event in the progression of these tumors. Within LMP tumors, p21^{Cip1} positivity was more often associated with serous histology, strengthening the observations of Palazzo et al. (2000). However, within adenocarcinomas, p21^{Cip1} overexpression prevailed in clear cell and endometrioid categories, in agreement with the data reported by Werness et al. (1997), Anttila et al. (1999), Costa et al. (1999) and Shimizu et al. (1999). What emerges from

Table 5. Univariate and multivariate analysis of overall survival.

VARIABLE	UNIVARIATE ANALYSIS		MULTIVARIATE ANALYSIS (Cox's model)		
	(log-rank test)	p-value	Hazard ratio (HR)	95% confidence limits	
Age (<53/>53)	0.0527	0.134			
Histological type (serous/mucinous/endometrioid/clear cell/adenocarinoma)	0.4488	0.159			
Histological grade (1/2/3)	0.0387	0.136			
FIGO stage (I-II/III/IV)	0.0003	0.002	3.423	1.599-7.327	
Residual tumor (absent-minimal/bulk)	0.0010	0.494			
Chemotherapy	0.9951	0.127			
$p21^{Cip1} LI (<1\%/\ge 1\%)$	0.2751	0.332			
<i>p53 LI</i> (<10%/≥ 10%)	0.1674	0.580			
$p21^{Cip1}/p53$ (p21 ^{Cip1} +/p53- vs p21 ^{Cip1} +/p53 + or p21 ^{Cip1} -/p53- vs p21 ^{Cip1} -/p	53+) 0.3050	0.128			
Ki-67 LI (<10%/>10%)	0.0127	0.239			

these findings is that the role of p21^{Cip1} in the progression of ovarian tumors may be histology-dependent.

Consistent with the wild p53-dependent activation of p21^{Cip1} (El-Deiry et al., 1993), we observed a significant negative correlation between p53 and p21^{Cip1} expression, lending support to the results by Anttila et al. (1999), Costa et al. (1999) and Sengupta et al. (2000). However, the correlation was not very strong, implying that in a minority of ovarian tumors, p21^{Cip1} is p53independently regulated. Relevant to this issue are the differences in the distribution of the various p21^{Cip1}/p53 phenotypes between LMP and adenocarcinomas as well as among the various grade, stage and residual disease categories. Thus, the p21^{Cip1}-positive/p53-negative phenotype characterized the vast majority of LMP tumors as well as adenocarcinomas belonging to early stage, low grade and minimal residual disease categories. The opposite applied to the double-positive p21^{Cip1}/p53 phenotype, this being commoner in adenocarcinomas compared to LMP and to grade 3 cases. Given that p53 immunoreactivity in ovarian adenocarcinomas is a surrogate marker for p53 gene mutation (Skilling et al., 1996), and taking into account the limitations due to the lack of p53 gene analysis in our study, we may presume that the concomitant overexpression of p53 and p21^{Cip1} speaks in favor of p53-independent induction of p21Cip1, while the expression of p21^{Cip1} in the absence of p53 might be indicative of its p53 dependence (Anttila et al., 1999). Following this line of argument, it sounds logical to hypothesize that the p53-dependent mechanisms of p21 induction operate mainly in LMP tumors and adenocarcinomas of a favorable phenotype, in contrast to the p53-independent mechanisms which, not unexpectedly, are associated with features of aggressive

We also observed a significant negative correlation between p21^{Cip1} expression and proliferation rate in the entire cohort, which, however, disappeared when our cases were stratified into LMP and carcinoma categories. This fact implies that this correlation actually reflected the differences in both proliferation rate and p21^{Cip1} expression between these two categories of tumors. Palazzo et al. (2000) claim that such a lack of correlation between p21^{Cip1} expression and proliferation rate is to be expected in tumors with a low proliferation index, indicating that most cells are still able to exit from the cell-cycle into G0. Interestingly, we obtained similar findings when studying the effect of p27Kip1 on the proliferation rate of LMP tumors (Korkolopoulou et al., 2002). Explaining the abrogation of growth-regulatory effect of p21^{Cip1} in tumors with a high content of cycling cells, such as ovarian carcinomas, is more difficult. Probably, in these cases cells are allowed to bypass the G1/S restriction point despite the presence of p21^{Cip1}, because of the inability of the latter to form active complexes with cyclin/cdks when its stoichiometry is

altered (Barboule et al., 1995). Alternatively, it could be hypothesized that the antiproliferative action of p21^{Cip1} is neutralized by the positive effect exerted by p53 on proliferation rate.

Admittedly, the rather small number of deaths from disease-related causes is a drawback of the present series. However, the proportion of advanced to early stages compares well with previous consecutive series (Anttila et al., 1999; Shimizu et al., 2000) and the results of survival analysis recapitulate many of the factors that have been advanced as important determinants of clinical outcome in ovarian cancer, namely FIGO stage, the volume of residual disease and, to a lesser extent, histological type and grade (Hakes, 1993) which supports that our cohort is representative and that statistical analysis is valid. The prognostic utility of p21^{Cip1} in ovarian carcinomas is a contentious issue and no consensus has been reached thus far. Anttila et al. (1999), Costa et al. (1999), Ferrandina et al. (2000) and Schmider et al. (2000) agreed that loss of p21^{Cip1} alone and/or combined with p53 accumulation predicted poor overall (Anttila et al., 1999; Costa et al., 1999; Schmider et al., 2000) and progression-free survival (Anttila et al., 1999; Ferrandina et al., 2000), but Sengupta et al. (2000), Shimizu et al. (1999) and Werness et al. (1999) insisted that p21Cip1 failed to provide any significant information in this regard. Our data concur with those of the latter group of authors (Sengupta et al., 2000; Shimizu et al., 1999; Werness et al., 1999), although in the present series a lower survival probability was attributed to the loss of p21^{Cip1} expression and to the p21^{Cip1}-negative/p53-positive phenotype in lower grade adenocarcinomas. The mechanisms by which loss of p21^{Cip1} adversely affects survival are not known in detail. Perhaps the most obvious explanation is an abrogated cell-cycle block leading to less strict G1/S surveillance with facilitated transit into S phase. The absence, however, of a relationship between p21^{Cip1} and proliferation does not quite support this scenario. p21^{Cip1} may also mediate growth arrest by inhibiting further DNA synthesis, thus allowing the cells to continue their differentiation. Furthermore, the unfavorable prognostic effect of p21^{Cip1} loss of expression (as well as that of p21^{Cip1}-negative/p53-positive phenotype) may reside in its relationship with stage and the volume of residual disease. The latter explanation seems to be entirely consistent with the loss of p21^{Cip1} and of p21^{Cip1}/p53 predictive value once stage has entered the Cox's model. It is also conceivable that since p21^{Cip1} may confer sensitivity to apoptosis, its overexpression will increase the susceptibility of ovarian cancer cells to platinumbased chemotherapy (Costa et al., 1999).

In conclusion, we present evidence that p21^{Cip1} levels are related to various clinicopathological indicators of aggressiveness in ovarian adenocarcinomas and that different mechanisms may regulate its expression in LMP and adenocarcinomas. On the contrary, p21^{Cip1} deregulation does not seem to participate in the pathogenesis of LMP tumors nor is it a

major player in cell proliferation of ovarian carcinomas. More importantly, although p21^{Cip1}, alone or combined with p53, is of prognostic significance in patients with lower grade adenocarcinomas, it does not appear to add to the information gained from traditional prognosticators. Prospective studies with a larger number of patients are warranted to validate this assumption.

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