Expression of Apocrine Differentiation Markers in Neuroendocrine Breast Carcinomas of Aged Women

Anna Sapino, M.D., Luisella Righi, M.D., Paola Cassoni, M.D., Mauro Papotti, M.D., Patrizia Gugliotta, Bsc, Gianni Bussolati, M.D., Frc Path Department of Biomedical Sciences and Human Oncology, University of Torino, Torino, Italy

Neuroendocrine (NE) breast carcinomas are a rare entity in young women; however, their frequency increases in aged patients. The present work demonstrates that NE breast carcinomas in elderly women can also express an apocrine immunophenotype and analyzes the histological and clinical aspects of such differentiation. A selected series of 50 NE tumors (positive for NE markers in \geq 50% of the cells) was tested for the immunocytochemical expression of gross cystic disease fluid protein-15 (GCDFP-15). The results demonstrated that about 50% of moderately (G2) and well-differentiated (G1) NE breast carcinomas (mucinous, solid papillary, and solid cohesive histotypes) coexpressed the apocrine marker. In these cases, specific mRNA for GCDFP-15 (PIP) and for chromogranin A (ChA) was demonstrated using in situ hybridization (ISH). Carcinomas of the alveolar subtype (G2) and poorly differentiated carcinomas (G3), including one case of atypical carcinoid, were pure NE carcinomas, devoid of apocrine differentiation. The steroid receptor status of these lesions was evaluated to test a possible involvement of androgen receptors in apocrine differentiation. We demonstrated that the level of AR and the mean age of patients at diagnosis were significantly higher in apocrine than in nonapocrine differentiated tumors. The histological grade and the expression of estrogen receptor (ER) significantly influenced the prognosis of these NE carcinomas, either pure or NE-apocrine differentiated. The most original result of our study is therefore the demonstration of a possible divergent apocrine differentiation of NE breast carcinomas that might be regulated by the activation of androgen receptors in elder patients. In addition, the possibil-

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ity for using Chs or GCDFP-15 serum values in the follow-up of these patients, as demonstrated in two cases of the present series, can justify the immunophenotyping of the tumors.

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Carcinomas of the breast are a multifaceted entity that may show various types of differentiation. In breast carcinomas of usual type, focal neuroendocrine (NE) and apocrine differentiation is occasionally detected in scattered cells by the immunocytochemical expression of NE and apocrine markers (1–4). In addition, Nesland *et al.* (5) described scattered hormone-producing cells in two cases of apocrine carcinoma. Expression of chromogranin-A (ChA) has also been demonstrated in 3 to 5% of cells of a particular variant of invasive lobular carcinoma, the so-called histiocytoid carcinoma, which is diffusely immunoreactive for the apocrine protein GCDFP-15 (6, 7).

Apocrine differentiation in breast carcinomas is consistently linked to overexpression of androgen receptors (AR; 8) and, on the other hand, a series of "carcinoid" of the breast was shown to express AR in >50% of cases (9).

Taken together, these data suggest a link between NE and apocrine differentiation in breast carcinomas. To test this hypothesis, we investigated the possible production of the apocrine marker GCDFP-15 by NE-differentiated breast carcinomas. In addition, we analyzed the expression of steroid receptors and, in particular, of AR, as mediators of NE-apocrine differentiation.

In other nonendocrine organs, *pure NE tumors* are considered lesions homogeneously positive for NE markers (10, 11) and correlated to a specific clinical evolution. As a consequence, we recently defined as *pure NE-differentiated* breast carcinomas those tumors having a relevant percentage (\geq 50%) of cells expressing NE markers (2, 12). In the present study, we followed the same immuno-histochemical approach to define tumors as pure

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Address reprint requests to: Anna Sapino, M.D., Department of Biomedical Sciences and Human Oncology, University of Turin, Via Santena, 7, 10126 Torino, Italy; e-mail: anna.sapino@unito.it; fax: 0039-011-663-5267.

NE or as NE–apocrine differentiated, and we evaluated whether the different immunophenotype could be related to specific clinical features.

MATERIALS AND METHODS

Case Series

A series of 50 NE breast carcinomas that expressed at least one NE marker (namely chromogranin A [ChA], chromogranin B [ChB], or synaptophysin [Syn]) in \geq 50% of their cells was studied for the presence of the apocrine marker GCDFP-15. Forty-five tumors were part of a previously reported series of NE breast carcinomas (2). Five other cases, fitting the above reported immunocytochemical features, were obtained from a series of 153 breast carcinomas collected over a 9-month period in our department. The whole series of 50 NE carcinomas was subdivided according to our proposed classification (2) and the Elston and Ellis histological grading system (13).

The correlation of immunophenotype (pure NE or NE–apocrine) with the expression of the different NE markers (ChA, ChB, Syn) and with the hormonal receptor types was analyzed by Yates corrected χ^2 .

Age was obtained from all patients and follow-up, from 35 patients. Correlation of these clinical data with the immunophenotype, the histological grade, and the steroid receptor expression was analyzed by one-way ANOVA. Univariate analysis was based on Kaplan-Meier product-limit estimate of survival distribution using the Wilcoxon test. All data were analyzed by using a selected computer program for statistics.

Immunohistochemistry

Cases were either formalin fixed or Bouin fixed and were paraffin embedded. Parallel sections of each tumor were immunostained using the antibodies listed in Table 1. Immunohistochemical analysis of steroid receptors was possible on formalin fixed tissues only; thus, in 13 Bouin's-fixed cases, the results of the biochemical estrogen re-

TABLE 1. Panel of Antibodies Applied

Marker	Reagent	Source	Dilution
Chromogranin A	LK2H10	BioGenex, San Ramon,	1:100
	mAb	CA	
Chromogranin B	PE-11	Dr R. Fischer-Colbrie,	1:1000
	Polyclonal	Innsbruck, Austria	
Synapthophysin	Polyclonal	Biomeda, Foster City,	1:50
		CA	
GCDFP-15	Polyclonal	Dr D. Haagensen,	1:10000
		Boston, MA	
Androgen receptor	F39.41	BioGenex	Prediluted
	mAb		
Estrogen receptor	1D5	Zymed, San Francisco,	1:50
	mAb	CA	
Progesterone receptor	PgR-ICA	Dakopatts,	1:5
	mAb	Copenhagen, Denmark	

ceptor (ER) and progesterone receptor (PR) assay were used (cutoff value for positive cases was >20 fmol/mg).

Endogenous peroxidase activity was inhibited with 6% H₂O2 for 5 minutes before the incubation with the primary antibodies. When antigen retrieval was necessary, the sections were incubated with specific avidin- and biotin-blocking reagents (avidin/biotin blocking kit, Biogenex, San Ramon, CA), as described elsewhere (14). Heat-induced antigen retrieval was performed by pretreating the sections in a microwave oven (three 3-minute passages at 750 W in 1 mM EDTA buffer, pH 8.0, or in 10 mM citrate buffer, pH 6.0). In selected cases, the immunoreactivity for ChA was enhanced using biotinyl-tyramide to increase the sensitivity of the detection system.

In all cases, the sections were then washed in phosphate buffered saline and the reaction was revealed using the LSAB2 kit peroxidase streptavidin universal system (DAKO, Glostrüp, Denmark) for 30 minutes and developed in a solution containing 15 mg/100 mL phosphate buffered saline of 3,3'-diaminobenzidine (3,3'-diaminobenzidine) and 0.3% H₂O₂. Nuclei were counterstained using hemalum.

To evaluate the coexpression of ChA and GCDFP-15 in selected cases, a two-step double immunostaining was performed. In the first step, ChA was revealed with 3,3'-diaminobenzidine, whereas the second step involved the detection of GCDFP-15 immunoreactivity by β -galactosidase–conjugated secondary antibody. The reaction was developed with 5'bromo-4chloro-3 indoxyl β -galactosidase (15). Sections were then mounted in Canada balsam.

In Situ Hybridization

In 21 cases positive for GCDFP-15, ISH for GCDFP-15 mRNA (PIP) and for ChA mRNA was performed as described elsewhere (4–16).

A 600-base pair cDNA encoding human PIP cloned into the pVZI bluescript plasmid vector (pPIP-8–3-cDNA clone, kindly supplied by Dr. R. Shiu and Y. Myal, Winnipeg, Manitoba, Canada) was used for ISH. The probe was cleaved with the restriction enzyme *Xba*I. Antisense RNA molecules were obtained using a T7 RNA polymerase and digoxigenin RNA-labeled nucleotide mixture containing DIG-11dUTP (DIG RNA labeling kit, Boehringer, Mannheim, Germany).

Briefly, after proteinase K digestion, the slides were treated with the hybridization mixture containing the antisense digoxigenin–labeled RNA probe. After incubation, the nonspecifically bound single-strand RNA probe was removed by RNAse digestion. The specifically bound hybridized probe was revealed using anti-digoxigenin antibodies conjugated with alkaline phosphatase. Color was developed using a nitroblue tetrazolium salt substrate. Nuclei were counterstained with methyl green. Slides were then dehydrated and mounted.

For ChA ISH, 10 oligonucleotide probes were synthesized with an automated DNA synthesizer from published nucleotide sequences (17), and the probes were digoxigenin labeled. Briefly, the sections were treated in 10 mM of citrate buffer, pH 6.0, in a microwave oven at 800 W for 5 minutes, then they were incubated in proteinase K and, finally, in the hybridization mixture containing the oligonucleotide-labeled probes. The specifically bound hybridized probes were then revealed using peroxidase-conjugated anti-digoxigenin antibodies and developed in a 3,3'-diaminobenzidine solution. Nuclei were counterstained using hemalum.

RESULTS

Histotypes of NE Breast Carcinomas

Thirteen cases were NE mucinous carcinomas (Type B; 18, 19). The cells (plasmacytoid, spindle, or with signet ring appearance) formed cribriform or solid islands floating within the mucin lakes (Fig. 1, A–C). Mitoses ranged from 4 to 10 per 10 high power fields (HPF). Nine cases were well differentiated (G1), and four were moderately differentiated (G2) carcinomas.

Nine cases corresponded to those described by Maluf and Koerner (20) as "solid papillary carcinomas" and by Tsang and Chan (21) as "endocrine ductal carcinoma in situ" of the breast. These tumors produced expansive lesions (Fig. 1, D-F) formed by solid sheets of cells. In six cases, both invasive and in situ lesions were present. Benign intraductal papillomas were present in three cases, and a "pagetoid spread" of cells within the papillomas, as described by Tsang and Chan (21), was observed in one case. These cases presented a variable amount of extracellular mucin that ranged from small pools to large lakes, as seen in mucinous carcinomas (three cases). Mitoses ranged from 2 to 6 per 10 HPF. Six of these tumors were G1, and three were G2. One of the patients developed lung metastases with a mucinous component 3 years after diagnosis. The recurrence of the disease was suspected because of the increase in ChA serum level up to 143 ng/mL (cutoff value: 20 to 100 ng/ mL) observed during patient follow-up.

Fifteen cases defined as solid-cohesive carcinoma (2) or low-grade insular carcinoma (22) showed, at variance with solid papillary carcinomas, an infiltrating appearance because of a dense collagen core surrounded by highly cellular nests, or trabeculae of cells, reminding one of carcinoid tumors (Fig. 1, G–I). To these histological patterns corresponded polygonal, or plasmacytoid, or spindle cells with eosinophilic granular cytoplasm. Mitoses never exceeded 4 per 10 HPF. Seven of these tumors were G1, and eight were G2.

The NE alveolar carcinomas (five cases) featured round alveolarlike structures, separated by scanty dense stroma that reproduced an infiltrative growth pattern (Fig. 2, C–D), similar to the alveolar variant of lobular carcinoma (23). A homogeneous population of large clear cells, with faintly granular periodic acid-Schiff–negative cytoplasm, formed these NE alveolar carcinomas. Mitoses ranged from 6 to 12 (mean, 10) per 10 HPF. All cases were G2.

Eight cases were classified as poorly differentiated (G3) NE carcinomas. The small-cell variety was excluded from this group (24). Poorly differentiated NE carcinoma showed crowded clusters of cells, with moderate to abundant cytoplasm; nuclei with vesicular to finely granular chromatin; and high numbers of mitoses (ranging from 18 to 65 per 10 HPF; Fig. 2, E–F). Focal areas of necrosis were present. Within this group, one case similar to the atypical carcinoid showed whirls of spindle cells and focal necrosis (19, 10). No patients of this group were known to have nonmammary carcinomas. One patient with a poorly differentiated NE carcinomas and axillary lymph node metastases had high ChA serum level at diagnosis, which decreased to basal level after surgery.

Expression of NE Markers and GCDFP-15 and Correlation with Histological Grade

The expression of the NE markers and GCDFP-15 varied in the different histotypes (Table 2). In analogy to the quantitative approach followed to define NE differentiation, cases were considered NE–apocrine differentiated when \geq 50% of the cells were positive for GCDFP-15.

More than 50% of mucinous (Fig. 1, B–C), solid papillary (Fig. 1, E–F), and solid-cohesive (Fig. 1, H–I) histotypes were NE–apocrine carcinomas. NE markers and GCDFP-15 could be expressed either in different cells or in the same cell, as demonstrated by double immunostaining (Fig 2, A–B).

The entire alveolar (Fig. 2D) and the poorly differentiated NE carcinomas (Fig. 2F) expressed ChA and were negative or focally positive (two of eight cases of poorly differentiated NE carcinoma) for GCDFP-15. ChA expression was, in fact, significantly correlated with the pure NE phenotype (χ^2 = 8.26; *P* = .004), whereas ChB expression was correlated with the NE-apocrine phenotype (χ^2 = 11.52; *P* = .0007). Syn expression was not correlated with any particular phenotype.

When the two immunophenotypes were correlated with the grade of differentiation, the great majority of pure-NE carcinomas were found to be high-grade tumors (G2, 66.7%; G3, 100%), whereas NE–apocrine carcinomas were predominantly lowgrade tumors (G1, 68.4%; P = .006).

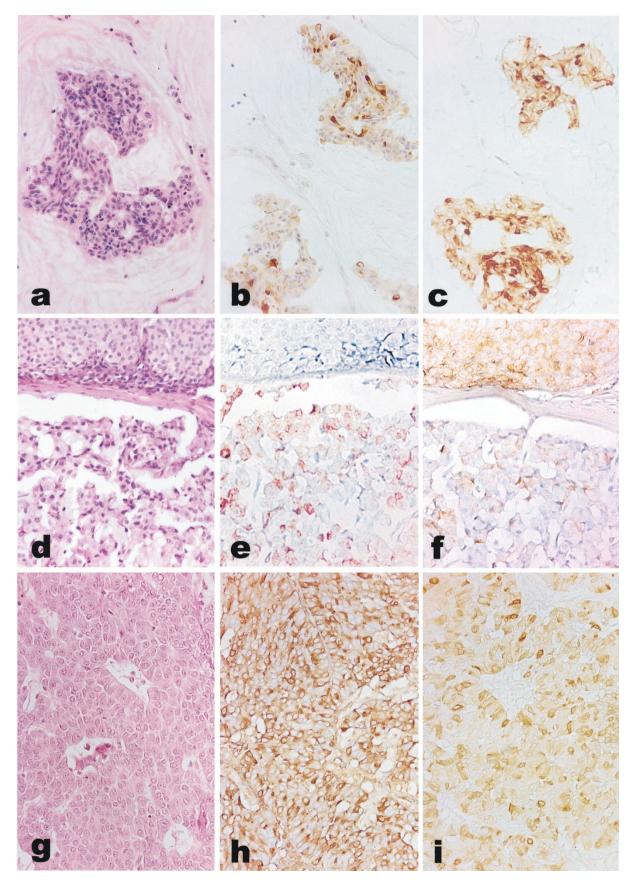


FIGURE 1. Expression of ChA and GCDFP-15 in different histotypes of NE breast cancers (mucinous, solid papillary, and solid cohesive types). A cribriform sheet of cells is floating within the mucin lakes in a mucinous NE–apocrine carcinoma (**A**). The same tumor is positive for ChA (**B**) and GCDFP-15 (**C**) in >50% of cells. In a solid papillary carcinoma, a solid nest of cells (*upper*) is separated from a mucin-producing area by sclerotic stroma (**D**). ChA is expressed only in the mucinous component (**E**), whereas GCDFP-15 is expressed in both counterparts (**F**). Solid cohesive NE–apocrine carcinoma showing rosettelike structures and peripheral cell palisading are reminiscent of carcinoid tumors (**G**). Both ChA (**H**) and GCDFP-15 (**I**) are expressed.

TABLE 2. Apocrine Differentiation in 50 NE Breast Carcinomas

Total Cases	Morphological Type	ChA, <i>n</i> (%)	ChB, <i>n</i> (%)	Syn, <i>n</i> (%)	GCDFP-15 (%)
#13	Mucinous	8 (61)	6 (46)	3 (23)	7 (54)
#9	Solid papillary	4 (44)	6 (67)	1 (11)	6 (66)
#15	Solid cohesive	8 (53)	9 (60)	2 (13)	8 (53)
#5	Alveolar	5 (100)	neg	neg	neg
#8	Poorly differentiated	8 (100)	4 (50)	2 (25)	2 foc (0)

ChA, chromogranin A; ChB, chromogranin B; Syn, synaptophysin; GCDFP-15, gross cystic disease fluid protein-15; %, percentage of positive cases in each histological type; neg, negative; foc, focal.

Specific mRNA for ChA and for GCDFP-15 was demonstrated in all cases expressing the related protein immunocytochemically (Fig. 2H).

Expression of Steroid Receptors

The expression of steroid receptors on formalinfixed tissues is summarized in Table 3. AR were expressed by 80% of cells of the NE-apocrine–differentiated tumors tested (11/11 cases; Fig. 2I). In these cases, ER and PGR were expressed in about 50 to 60% of the cells. Noteworthy was the diffuse expression of AR in NE-apocrine mucinous carcinomas (4/4), whereas NE-mucinous (nonapocrine) carcinomas (three cases) were weakly positive in <5% of cells. The pure NE carcinomas rarely expressed high nuclear levels of AR (3/13). On the contrary, ER and PGR were widely expressed.

The analysis of steroid receptor expression within the different tumor immunophenotypes revealed a significant correlation only between AR distribution and apocrine differentiation ($\chi^2 = 13.3$; P = .0003).

Clinical Features

The immunophenotype of tumors was correlated with the age of patients at diagnosis. The mean age of patients with pure-NE-differentiated tumors (62.3 years) was significantly younger than that of patients with NE-apocrine tumors (72.4 years; P < .002).

Survival data were available for 35 patients. The mucinous differentiation was a favorable prognostic factor. The patients with mucinous and solid papillary carcinomas (producing some degree of mucin) had a significantly longer survival time than did patients with tumors of the other histological types (P = .05; Fig. 3A). In short-term follow-up, the survival was not significantly modified by the immunophenotype (pure-NE *versus* NE-apocrine tumors, P = .1), but 5 years after surgery, the association with apocrine differentiation improved long-term survival of patients (Fig. 3B). The histological grade was significantly correlated with prognosis. All G1 cases survived at >13-year follow-up, whereas the G3 NE tumors were very

aggressive, and all those patients died within 6 years (P = .007; Fig. 3C).

The expression of ER (P < .0001) was correlated with a favorable prognosis (Fig 3D), followed by PR expression (P = .02; Fig. 3E). AR expression did not significantly change the prognosis (P = .19).

DISCUSSION

The present study demonstrates that NE and apocrine differentiation may coexist in the same breast tumor. Cells producing NE markers or GCDFP-15 in the absence of specific NE or apocrine morphology are present in the normal breast (1, 21, 25). Similarly, in breast carcinomas, the immunocytochemical expression of apocrine markers, such as GCDFP-15, is wider than expected by morphology alone (6). As a consequence, it has been suggested that immunohistochemical techniques should be used to identify apocrine and NE differentiation in breast carcinomas, as hematoxylin and eosin identification is highly subjective and unreliable (26, 2).

In the present study, we considered as NE and apocrine differentiated those tumors expressing specific markers in \geq 50% of cells. ChB expression was typically correlated with the NE-apocrine immunophenotype; on the other hand, ChA was the prevalent or the only marker produced by pure-NE carcinomas but was also present in NE-apocrinedifferentiated tumors. When the serum levels of ChA were measured in two patients affected respectively by a pure NE and an NE-apocrine tumor, they were highly above the cutoff value. This result is in agreement with the observation of high ChA serum level in small-sized (<2 cm) but intensely ChAimmunoreactive NE tumors (27). In addition, a potential role of GCDFP-15 as a marker of breast cancer micrometastasis has been recently proposed (28). Thus, the possibility of using Chs or GCDFP-15 serum values in the follow-up of these patients can justify the immunophenotyping of the tumors. In addition, in previous studies, the apocrine differentiation of tumors was correlated with a better prognosis (4, 29). In the present work, the clinical evolution was highly correlated with the histological grade, but at long-term follow-up, there was a tendency for a longer survival in patients with NEapocrine-differentiated tumors. In agreement with other studies (30), we here demonstrated that the mucin production of tumors was correlated to a better prognosis.

The steroid receptor level was also evaluated, as apocrine carcinomas consistently overexpress AR (8), and secretion of GCDFP-15 in breast carcinoma cell lines is stimulated by androgen treatment *in vitro* (31). Our work confirms that the AR level is higher in apocrine differentiated tumors. The apocrine tumors of our series are typical of elderly women; thus, it can

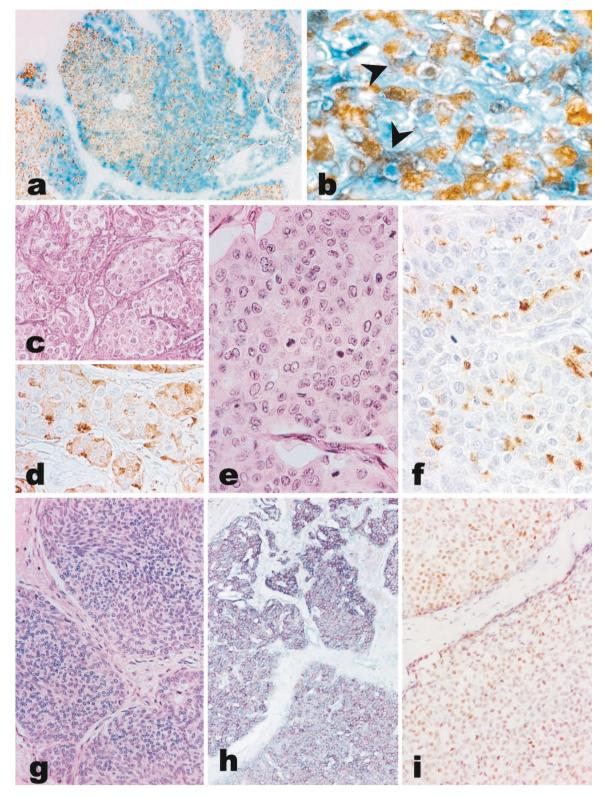


FIGURE 2. Expression of ChA, ChB, GCDFP-15, and AR in different histotypes of NE breast cancers (solid cohesive, alveolar, and poorly differentiated types). Double immunostaining of a case of solid-cohesive carcinoma (immunoperoxidase and beta-galactosidase). ChA-positive NE cells are in brown, whereas apocrine-differentiated cells expressing GCDFP-15 are in blue (A). At higher magnification, the coexpression of ChA and GCDFP-15 in the same cells (**arrows**) is evident (**B**). Pure NE alveolar carcinoma featuring round alveolarlike structures, separated by scanty dense stroma, is formed by a homogeneous population of large clear cells, with faintly granular cytoplasm (**C**), positive for ChA (**D**). Poorly differentiated NE carcinoma shows clusters of cells, with moderate to abundant cytoplasm, nuclei with vesicular to finely granular chromatin, and high number of mitoses (**E**). Immunostaining for ChB shows an intense cytoplasmic granular staining (**F**). In a case of solid-cohesive spindle cell NE-apocrine carcinoma (**G**), the presence of specific mRNAs for ChA GCDFP-15 (nitroblue tetrazolium salt substrate) is demonstrated with *in situ* hybridization (**H**). The nuclei of the same tumor are intensely and diffusely positive for AR (**I**).

TABLE 3. Analysis of Steroid Receptors Related to Tumor Immunophenotype

Tumor Immunophenotype	Steroid Receptors			
	AR	ER	PGR	
NE	3/16	15/22	9/20	
Apocrine-NE	9/9	15/16	13/15	

All data are listed as immunoreactive tumors/examined tumors. AR, androgen receptors; ER, estrogen receptors; PR, progesterone receptors.

be speculated that the relative prevalence of androgenic hormones in the menopausal period may induce the production of apocrine proteins through the activation of specific AR receptors. Finally, this study demonstrates that the morphological features traditionally considered typical of NE breast carcinomas may be consistent also with an apocrine differentiation. Immunoelectron microscopy studies demonstrated the presence of ChA-negative dense core granules in breast carcinomas reactive for ChA in light microscopy (32), and ChA-negative dense core granules were found on the cell apex of solid papillary NE carcinomas (33). The authors presumed that these granules could be positive for other types of Chs. However, the possibility that they could correspond to those clustered towards the apex of cells in apocrine carcinomas (34) has to be considered.

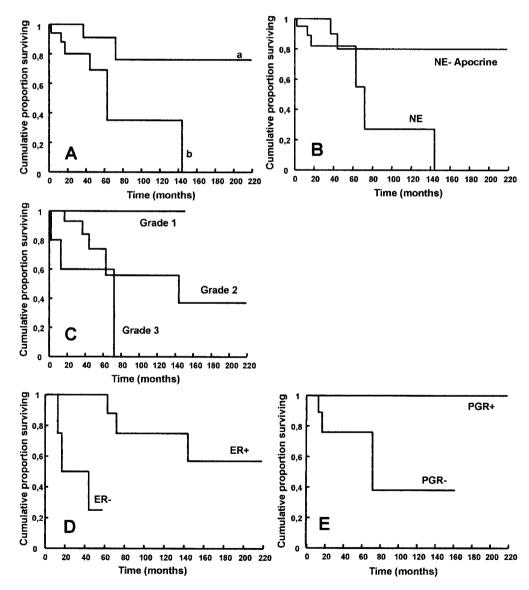


FIGURE 3. Kaplan-Meier curves of 35 patients. (A) Patients' survival is influenced by the histological type of tumors. The patients with mucinous and solid papillary carcinomas (producing some degree of mucin; a) have a significantly longer survival than patients with tumors of the other histological types (P = .05; b). (B) In short-term follow-up, the survival is not significantly modified by the immunophenotype (pure NE *versus* NE-apocrine tumors, P = .1), but 5 years after surgery, there is a tendency to a better prognosis for patients with NE-apocrine tumors. (C) The histological grade influenced prognosis; in fact, patients with G1 tumors are all alive at 13 years, those with G2 tumors have an intermediate prognosis; and all patients with G3 tumors had died by the 6-year follow-up. Patients with ER- (D) and PR- (E) tumors have significantly poorer prognosis than patients with ER+ and PR+ tumors.

In conclusion, the present cases may be a good example of multidirectional differentiation of breast tumors. Mixed or divergent differentiation with production of NE and exocrine protein is rather a frequent phenomenon in tumors of endocrine and nonendocrine organs (11, 35, 36). The origin from different cell lines as colliding NE and apocrine tumors is possible. However, the immunocytochemical and in situ hybridization coexpression of NE and apocrine markers in tumors with identical morphological substrate may lead to the hypothesis of an uncommitted stem cell capable of differentiating toward both NE and apocrine lineage. Cells of welldifferentiated tumors maintain this potential, whereas those of poorly differentiated carcinomas do not show this multidifferentiation capacity and express the NE phenotype only. Our preliminary in vitro data demonstrate the presence of low levels of specific ChA mRNA in breast carcinoma cell lines producing apocrine proteins and expressing AR.

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Book Review

Hamilton SR, Aaltonen LA, editors: Pathology and Genetics of Tumours of the Digestive System 314 pp, Lyon, France, IARC Press 2000 (\$75.00).

This is the second volume in the new WHO series on Histologic and Genetic Typing of Human Tumors. In this volume, the authors provide a review of the neoplasms of the alimentary tract including the esophagus, gastrointestinal tract, liver, biliary system, and exocrine pancreas. Each chapter is dedicated to a discussion of a specific organ and is divided into sections. The sections discuss the specific tumors of the organ or site and are authored by distinguished investigators in the respective areas. The chapters begin with the WHO histologic classification of tumors, the TMN classification, and stage grouping. Each section within the chapter begins with the WHO definition of the tumor(s), the ICD-O codes, followed by sections on the epidemiology, etiology, clinical features, including imaging (both endoscopic and radiographic), microscopy, staging,

and grading. What is so beneficial in this monograph is that in each section the authors present an excellent summary of the genetics of the tumors. There are outstanding photographs throughout the textbook, demonstrating the macroscopic and microscopic features of the tumor, as well as excellent figures. The textbook, by its complete discussions of the clinical, pathologic, and genetic information, provides a unique opportunity for clinicians, pathologists, and geneticists to have a reference text (with more than 2000 citations) pertaining to alimentary system tumors suitable for all. This should be present in all departments of Medicine and Pathology/resident's libraries. In summary, this is a superb effort on the part of the editors and the contributors and is highly recommended.

Peter S. Amenta, M.D., Ph.D.

UMDNJ—Robert Wood Johnson Medical School Piscataway, New Jersey