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Chapter 7 - Ectopic factor VII expression is an indicator of poor survival in young women with breast cancer

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Abstract

Blood coagulation and cancer progression are engaged in a bidirectional relationship that manifests itself in malignancy-related thrombosis and coagulation factor-dependent tumor progression. Full-length tissue factor (fITF):factor(F)VIIa signaling through Protease Activated Receptor (PAR) 2 determines outcome in murine breast cancer. Whether only canonical liver-derived FVII or also tumor-expressed FVII is involved in cancer progression is unknown. The aim of the present study was to examine the association between tumor-expressed FVII and clinical outcome in breast cancer. Breast cancer specimens collected during primary surgery - at which time radiotherapy or chemotherapy had not been started - were assembled in a tissue micro array (TMA) and immuno-stained for fITF and FVII. Data on fITF expression was available from 507 patients and data on both fITF and FVII expression from 302 patients. The TMA's were independently scored by two observers, and associations with clinical and pathological parameters and survival were analyzed. fITF was abundantly expressed in cancer specimens when compared to normal healthy tissue, while FVII was detected in only 39% of the tumors. Whereas fITF expression was associated with higher tumor grade ($p < 0.001$), FVII was also associated with T status that reflects tumor size and invasiveness ($p = 0.009$), and with loss of estrogen ($p = 0.011$) and progesterone ($p < 0.001$) receptors. Patients with FVII+ tumors had a shortened relapse-free survival ($p = 0.02$), and when analyzed for double expression of fITF and FVII, we found a similar effect ($p = 0.006$). After 10 years, relapse-free rates were 58% for women younger than 55 years with fITF:FVII positive tumors compared to 82% for women > 55 years. These results confirm that fITF expression is a general phenomenon in breast cancer. This report also shows that FVII expression in breast cancer is associated with poor prognosis in young women. Ectopically produced FVII therefore may offer new perspectives in cancer staging or therapeutic options.

Introduction

For more than one century, activation of coagulation is recognized to be tightly interwoven with cancer progression¹. Patients suffering from various types of cancer may develop venous thrombosis, a condition referred to as Trousseau's syndrome. Tumoral overexpression of full-length tissue factor (fITF), a 47 kDa transmembrane protein, may cause this thrombotic tendency². Under physiological circumstances, fITF expression is limited to the surface of extravascular cells. Upon damage of the endothelium, fITF becomes exposed to the bloodstream and binds the blood-borne zymogen FVII. The fITF:FVIIa complex induces the subsequent activation of factor Xa and thrombin, thus resulting in a fibrin clot³.

Activation of coagulation also influences various stages of cancerous disease. The f1TF:FVIIa complex and thrombin have been implicated in tumor growth and metastasis⁴. f1TF expression in human colon cancer is associated with an increased risk of hepatic metastasis⁵, and in murine models, f1TF expression in colon cancer as well as in breast cancer promotes tumor growth through modulation of angiogenesis^{6;7}. Extensive research has uncovered that coagulation proteases activate protease-activated receptors (PARs). The f1TF:FVIIa complex specifically activates PAR2 resulting in altered cell behavior such as proliferation, gene expression and migration, eventually leading to enhanced tumor growth⁷. In a spontaneous murine breast cancer model, absence of PAR2 delays the angiogenic switch and as a consequence, tumor growth is delayed⁸.

FVII is normally synthesized in the liver and released into the bloodstream. When vessel damage occurs, the f1TF:FVII complex initiates coagulation in order to stem bleeding. One report claims that FVII is bound to f1TF on fibroblasts even before vascular damage occurs⁹. Whether this FVII came from the circulation or was locally produced is unknown. Furthermore, FVII can be detected in lung macrophages under inflammatory conditions¹⁰⁻¹² and in atherosclerotic plaques¹³⁻¹⁵, again suggesting a possible role for ectopic FVII production in pathophysiology.

FVII mRNA has been detected in ovarian, breast, brain, prostate, thyroid, lung and gastric cancer cell lines up to levels comparable with those found in non-malignant hepatoma cell lines, which indicates that FVII may be produced in (patho)physiologically relevant amounts by cancer cells. In ovarian and breast cancer cell lines, ectopically produced FVII is bound to f1TF, and FVII levels were sufficient to trigger coagulation and subsequent factor Xa (FXa)-dependent PAR1 activation. Consequently, FVII production in ovarian cancer cells leads to enhanced cancer cell motility and invasiveness¹⁶. In glioma cell lines, EGFRvIII-driven simultaneous expression of f1TF, FVII, PAR1 and PAR2 precedes the production of angiogenic proteins like VEGF and IL-8¹⁷. This may indicate that ectopic FVII expression facilitates the angiogenic switch in poorly vascularized tumors thus circumventing the requirement for blood-borne FVII.

Hypoxia is an important upstream event that triggers ectopic FVII in tumors. Oxygen depletion leads to the activation of a hypoxia-inducible factor 2 α (HIF2) α -dependent pathway in ovarium cancer cells that eventually induces FVII expression^{16;18}. A recent study in ovarian cancer cell lines described a non-canonical interaction between hypoxia-driven and epigenetic pathways leading to ectopic FVII production¹⁹. HIF2 α and Sp1 form a complex and bind to a known Sp1-binding site in the FVII promoter region and subsequently recruit histone deacetylase 4 (HDAC4), which induces FVII expression. A different epigenetic mechanism was identified in breast cancer cell lines in which

curcumin-sensitive histone acetyltransferases (HAT) p300 and cyclic AMP-responsive element binding protein (CBP) were exclusively recruited to the promoter region of the FVII gene²⁰. Although differences may exist between tumor cells from different tissues, hypoxia alters the acetylation status of the FVII promoter region and thereby induces FVII expression.

The relation between FVII expression in human tumor specimens and patient outcome has not been investigated. In the present study, we examined a large number of human breast cancer samples and matched normal tissue from patients with a long follow-up in order to investigate how tumoral FVII and fITF expression relate to clinical outcome in human breast cancer.

Methods

Reagents- Normal goat serum and EnVision™ anti-rabbit HRP and anti-mouse HRP conjugates were purchased from DAKO (Glostrup, Denmark). FVII polyclonal antibody GTX101238 was purchased from Genetex (Irvine, CA). fITF monoclonal antibody #4509 was purchased from American Diagnostica Stamford, CT).

Study cohort- The patient population comprised patients with non-metastasized breast cancer treated with surgery in the Leiden University Medical Center between 1985 and 1994 of whom tumor material is available (N = 574) as described earlier²¹. Patients with bilateral tumors or a prior history of cancer - other than basal cell carcinoma or cervical carcinoma in situ - were excluded. The following data were recorded: age, tumor grade, histological type, TNM (TNM classification of malignant tumors, Union for International Cancer Control), stage, local and systemic therapy, locoregional or distant tumor recurrence, survival, and expression of estrogen receptor (ER), progesterone receptor (PgR) or human epidermal growth factor receptor 2 (HER2). All tumors were graded according to the Nottingham modification of the Scarff-Bloom-Richardson grading system. Of 266 patients (46%), also normal mammary tissue was available for analysis. Median follow-up was 17.9 years (range: 0.01 to 23.5). Approval was obtained from the Leiden University Medical Center Medical Ethics Committee. All samples were handled in a coded fashion, according to National ethical guidelines ("Code for Proper Secondary Use of Human Tissue", Dutch Federation of Medical Scientific Societies).

Assessment of expression of FVII and fITF- Tissue sections of 4 µm were cut from a previously constructed tissue microarray of formalin-fixed paraffin-embedded tumors of 574 patients and a tissue microarray of formalin-fixed paraffin-embedded corresponding normal mammary tissue of 266 patients. Immunohistochemical staining was performed

according to standard procedures. Briefly, sections were deparaffinized, rehydrated and endogenous peroxidase activity was blocked with 0.3% H₂O₂ in MetOH. All sections were subjected to antigen retrieval in sodium citrate buffer for 10 minutes at 100 °C. Sections were blocked for 1 hour with 10% normal goat serum in PBS/BSA 1% at room temperature. Sections were then incubated with 1 µg/ml antibody overnight at 4 °C (both anti FVII and anti fITF). Sections were washed in PBS, incubated for 30 minutes with Envision, washed and visualized in DAB solution. After counterstaining with hematoxylin, sections were dehydrated and covered. Liver and placenta sections served as a positive control for respectively FVII and fITF. The TMA's contained tumor samples from each patient in triplicate, and were randomly arranged on the array. During the immunohistochemical procedures a random loss of tissue punches occurred, finally resulting in data on fITF expression in 507 patients, on FVII expression in 331 patients and on both fITF and FVII in 302 patients.

FVII was considered to be expressed when one or more tumor cells stained positive. fITF staining was defined by percentage of positive staining tumor cells per microscopic field. Two independent observers performed all scorings in a blinded fashion. fITF staining percentages were arranged in increasing order and on basis of this, the cohort was divided into quartiles. In subsequent analyses, the first quartile was considered “negative”.

Statistical analysis- Statistical analyses were performed using the statistical packages SPSS (version 16.0 for Windows, Spps Inc, Chicago, IL, USA) and Stata (version 10.0 for Windows, StataCorp, College Station, TX, USA). Cohen's kappa coefficient was used to assess the inter-observer agreement in quantification of protein expression. The χ^2 test was used to evaluate associations between protein expression and various clinicopathological parameters.

Relapse-free survival was defined as the time from date of surgery until an event (locoregional recurrence and/or a distant recurrence, whichever came first). Relapse-free survival is reported as cumulative incidence function, after accounting for death as competing risk²². The Kaplan–Meier method was used for survival plotting and log-rank test for comparison of curves. Analyses were performed for all patients and stratified for age. Relative survival was calculated by the Hakulinen method as the ratio of the survival observed among the cancer patients and the survival that would have been expected based on the corresponding (age, sex, and year) general population. National life tables were used to estimate expected survival. Relative excess risks of death were estimated using a multivariable generalized linear model with a Poisson distribution, based on collapsed relative survival data, using exact survival times. The Kaplan–Meier method was used for survival plotting and log-rank test for comparison of curves.

Results

Patterns of FVII and fITF expression in human breast cancer- A granular, cytoplasmic staining pattern for human FVII was found in 130 out of 331 patients (39.3%), whereas non-cancerous epithelium showed no FVII staining (figure 1). Inter-observer agreement was high ($\kappa > 0.8$). Tables 1 and 2 list the associations in the cohort between fITF and FVII expression and several clinical and pathological characteristics. In accordance with previous reports (reviewed in reference 1), we found that expression of fITF in breast cancer is detected in >90% of the cases. We scored the percentage of fITF positive cells, arranged the scored percentages in increasing order, and divided this range into quartiles. The 2nd to the 4th quartile comprised tumor specimens in which all cells per microscopic field were positive for fITF.

Higher histological grades were associated with fITF expression. FVII-positive tumors display a higher histological grade, advanced T-status and loss of estrogen and progesterone receptors, which are unfavorable tumor characteristics²³. Similar associations, but also a loss of Her2 receptor ($p=0.05$), was found when both fITF and FVII were positive, thus suggesting that expression of fITF and FVII coincides with loss of the estrogen, progesterone and Her2 receptor. The latter is clinically relevant as the so-called triple negative tumors are considered to have the most unfavorable phenotype in breast cancer patients.

FVII positivity is associated with impaired relapse-free survival and relative survival- The percentage of fITF expressing tumor cells was not associated with the length of the relapse-free period. Analysis of the relapse-free period in patients with FVII positive versus FVII negative tumors revealed that the length of the relapse-free survival was reduced in patients with FVII positive tumors. After 10 years of follow-up, 72% of the patients with FVII negative tumors were relapse-free compared to 60% of patients with FVII positive tumors (absolute difference 12%, log rank $P = .02$, Figure 2).

Since fITF serves as the receptor for FVII, we next determined survival in patients with fITF/FVII double positive tumors, compared to single positive or negative tumors. Different lengths of the relapse-free period were observed for these categories (either or both negative 72% relapse-free, versus 58% relapse-free when both positive, $P=.006$). The overall survival was similar to the relapse-free survival (Figure 2) as well as the differences in survival after 10 years.

Combined fITF and FVII expression is an independent risk factor for decreased RFP and relative survival- In order to investigate whether combined fITF and FVII expression is an

independent risk determinant of relapse-free survival and relative survival, we performed a Cox univariate and multivariate analysis (Table 3). As combined fITF and FVII expression were associated with grade, T-status, presence of lymph node metastasis and histological subtype, we included these determinants in our analysis. Combined fITF and FVII expression indeed independently predicted early cancer recurrence (HR 1.55; CI95% 1.095-2.205) and in a similar fashion, increased mortality (RER 1.68; CI95% 1.092-2.575).

Tumoral FVII affects relapse-free survival and relative survival in pre-menopausal women-

Next we analyzed whether age affects the association between fITF, FVII and survival. We divided the cohort, on the basis of the median age, in women aged <55 years or 55 years and older. We found an association between fITF expression and shorter relapse-free and overall survival in women aged younger than 55 years. Moreover, we found that the associations between FVII expression alone, or combined expression of fITF and FVII, with relapse-free period became more prominent for women younger than 55 years, but were absent in women aged 55 or older. After 10 years, relapse-free rates were 58% for women younger than 55 years with fITF:FVII positive tumors compared to 82% for women > 55 years (figure 3). A similar effect was observed when survival analyses were performed after dividing the cohort between women aged younger than 65 and older than 65 years (data not shown). Moreover, a Cox regression analysis that assessed the interaction between age and presence of both fITF and FVII proved significant ($p=0.006$). In summary, the effects of combined fITF and FVII expression affects breast cancer in younger age groups and this effect fades out in older aged patients.

Discussion

The notion that fITF mediates signaling events in breast cancer, and thereby influences cancer progression, either via direct PAR2 signaling or indirectly via coagulation activation, is supported by several *in vivo* and clinical studies¹. Both pathways depend on the presence of blood-borne coagulation factors that are produced in the liver, but it is unclear how tumors have access to coagulation factors when they are still poorly vascularized. Hypoxia-induced expression of ectopic FVII expression in cancer may at least partially answer this conceptual question. In order to examine whether local FVII expression in tumors is related to cancer progression in humans, we investigated the association between FVII or fITF expression and clinical and pathological outcome in breast cancer patients.

We observed that FVII expression is present in only 39% of the investigated tumor specimens, whereas fITF expression is virtually present in all specimens. This suggests that fITF expression is a general phenomenon in breast cancer and that expression of FVII is

associated with cancer progression in a subgroup of breast cancer patients. We found associations between tumoral FVII expression and unfavorable tumor characteristics, such as high histological grade and loss of hormone receptors. Both relapse-free and relative survival were associated with expression of FVII. Moreover, we showed that combined expression of both fITF and FVII further increased the strength of our survival analyses, possibly through an association with triple negative breast cancer, a condition that is associated with aggressive breast cancer at young age.

We found that the effects of combined fITF and FVII expression on outcome were only apparent in patients under 55 years of age. This may suggest that the onset of menopause and related changes in hormonal status is associated with FVII expression in breast cancer. However, a similar association was found when a cut-off of 65 years was applied, which makes menopause a less likely determinant. Future studies in other solid tumors, e.g. colon cancer, will show whether the results that we here report are breast cancer- and gender-specific or rather a general feature of FVII expression in solid tumors.

There are two potential limitations to our study. First, this study is immunohistochemistry-based and potentially blood-borne FVII may have become internalized after binding to fITF. Therefore, a RNA-based follow-up study may take away uncertainties about the origin of FVII that we detected in the tumor samples. The second drawback is a lack of data that confirm that FVII expression in tumor cells indeed triggers fITF:FVII:PAR2 signaling. An earlier study, investigated the role of phosphorylated fITF in relation to clinical outcome and survival²⁴. Upon activation of PAR2, the cytoplasmic domain rapidly becomes phosphorylated and detection of phosphorylated fITF can serve as a read-out of PAR2 signaling in these tumor specimens. Indeed, phosphorylation of fITF was associated with poor survival and in future studies it would be interesting to investigate whether fITF phosphorylation is associated with ectopic FVII production.

Regarding other avenues for future studies, it is warranted to assess whether other coagulation proteins that are canonically expressed in the liver become expressed in tumors during oncogenesis as well. Since FXa, thrombin and activated protein C also have effects on cellular processes via either PAR-dependent pathways¹ studies on ectopic expression of these proteins in tumors may prove informative in further understanding the relation between coagulation proteins and cancer.

In conclusion, our study describes a strong association between ectopic FVII expression in breast tumors and poor survival in women younger than 55 years. Whether these effects are partially caused by hormonal effects at pre-menopausal age is unclear and remains to be investigated. Future research may also unravel the role of tumor-expressed FVII in fITF-

mediated PAR2 signaling, and potentially also of other ectopically, tumor-expressed coagulation factors.

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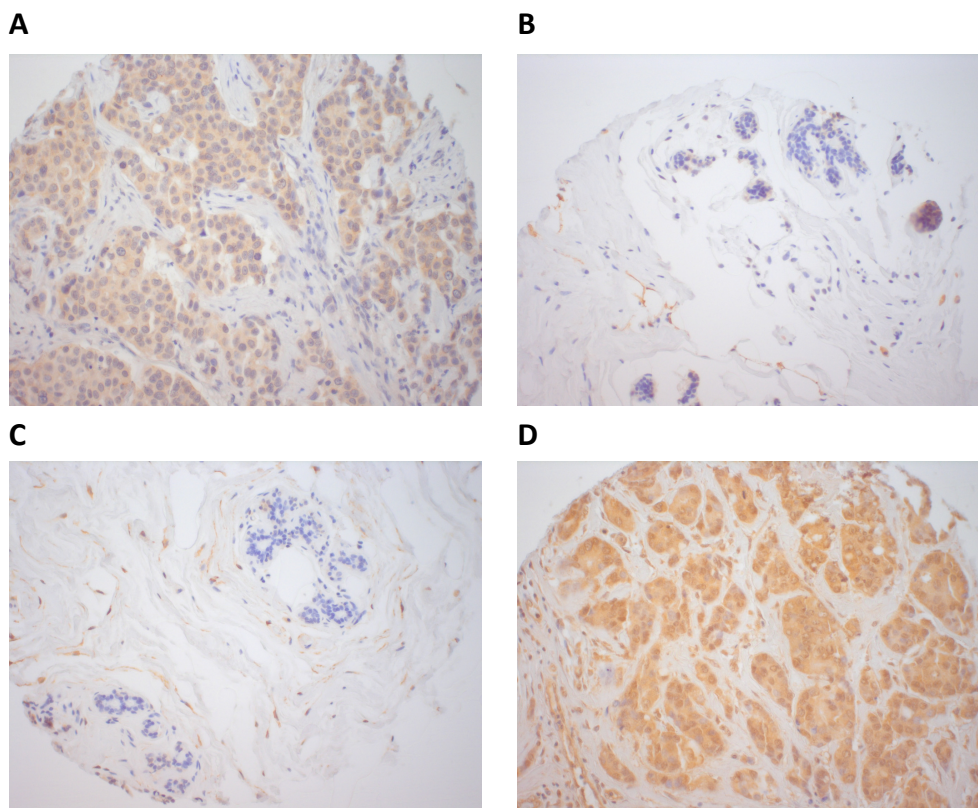


Figure 1. Expression patterns of FVII and fITF in human breast cancer. Breast cancer and matched normal breast specimens were assembled in a TMA fashion and stained with antibodies for either FVII or fITF. Panel A: non-malignant breast tissue, stained for FVII; panel B: FVII positive breast cancer specimen; panel C: non-malignant breast tissue, stained for fITF; panel D: fITF positive breast cancer specimen.

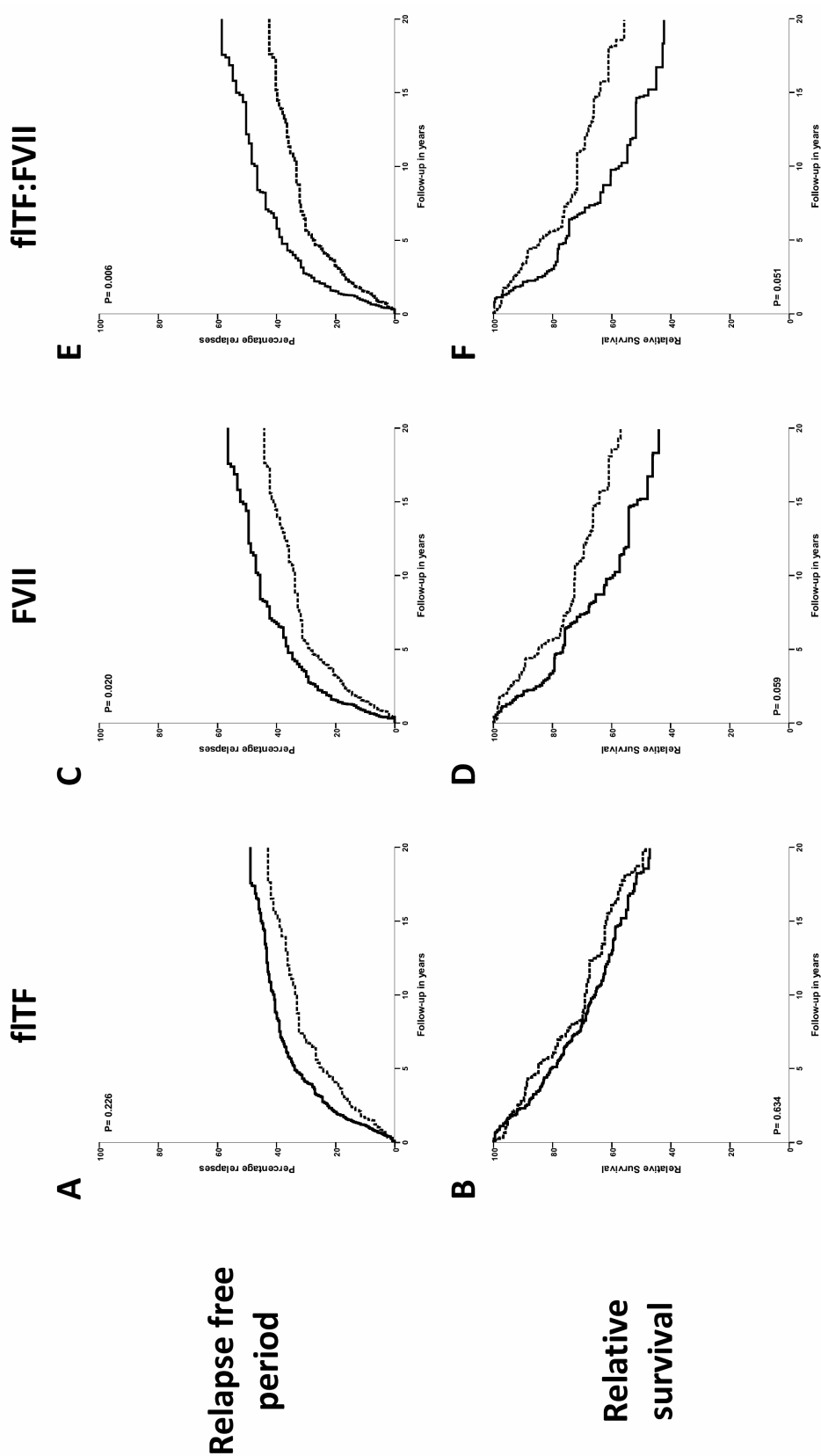


Figure 2. Relapse-free period and relative survival Kaplan-Meier curves. For fITF respectively panel A and B; for FVII respectively panel C and D; for FVII and fITF double expression respectively panel E and F. Continuous lines indicate high expression, dotted lines indicate low or absent expression.

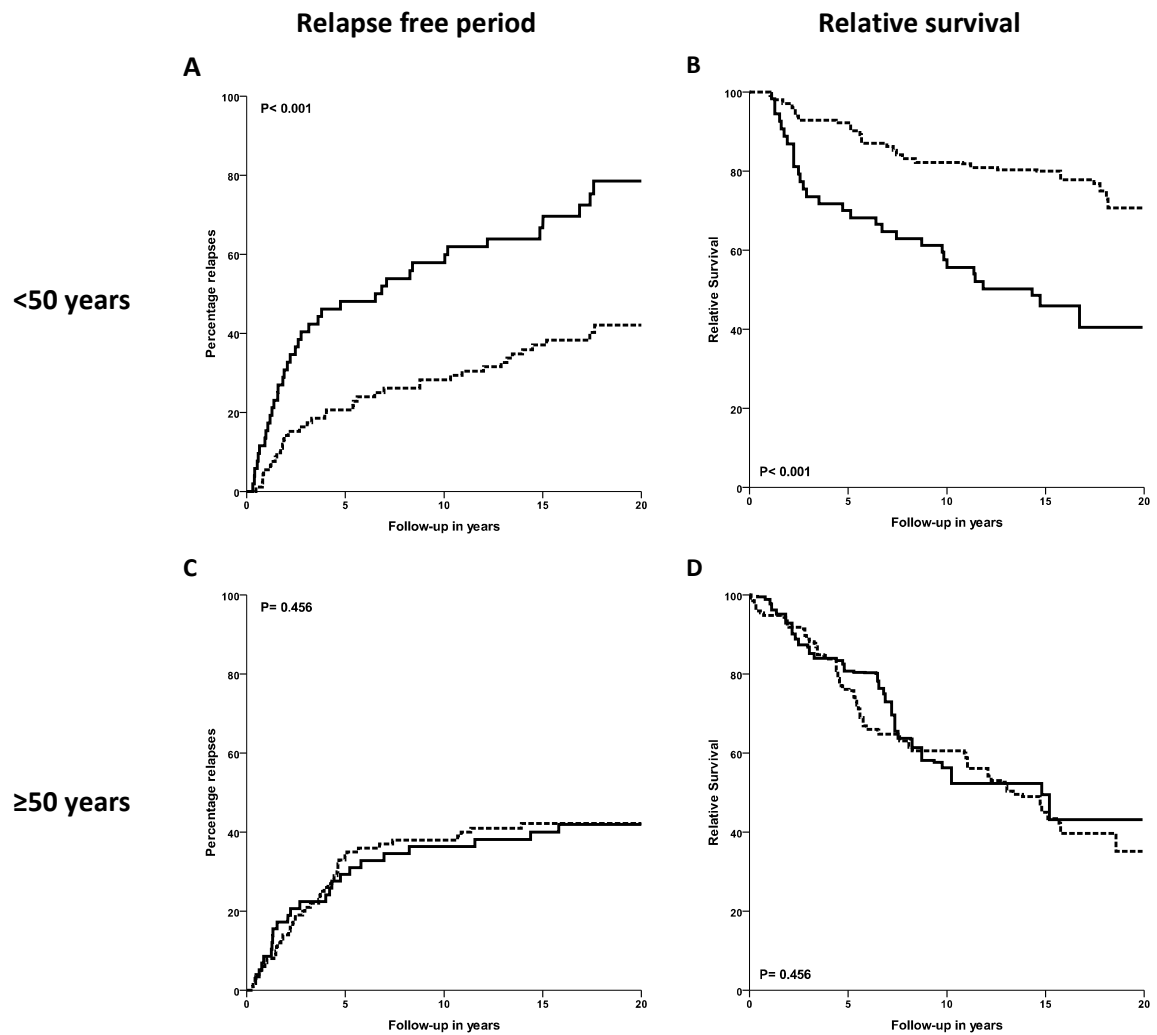


Figure 3. Relapse-free period and relative survival Kaplan-Meier curves for FVII and fITF in patients aged younger than 55 years (repectively panel A and C) or older than 55 years (repectively panel B and D).

	Factor VII				p-value	fITF				p-value
	Negative		Positive			Negative		Positive		
	N	%	N	%		N	%	N	%	
Total	201	100	130	100		156	100	351	100	
Age										
<40	20	9.9	14	10.8		15	9.6	26	7.4	
40-55	80	39.8	45	34.6		57	36.5	132	37.6	
55-65	36	17.9	32	24.6		30	19.2	77	21.9	
>65	65	32.3	39	30	0.479	54	34.6	116	33.0	0.768
Missing	0	0.0	0	0.0		0	0.0	0	0.0	
Grade										
I	33	16.4	9	6.9		30	19.2	42	12.0	
II	110	54.7	55	42.3		88	56.4	157	44.7	
III	55	27.4	63	48.5	<0.001	36	23.1	148	42.2	<0.001
Missing	3	1.5	3	2.3		2	1.3	4	1.1	
Histological type										
Ductal	179	89.1	121	93.1		134	85.9	320	91.2	
Lobular	20	9.9	6	4.6	0.057	20	12.8	27	7.7	0.065
Missing	2	1.0	3	2.3		2	1.3	4	1.1	
T-status										
T1	87	43.3	39	30.0		63	40.4	120	34.1	
T2	81	40.3	76	58.5		71	45.5	171	48.7	
T3/4	27	13.4	14	10.8	0.009	17	10.9	51	14.5	0.294
Missing	6	3.0	1	0.8		5	3.2	9	2.6	
N-status										
N0	114	56.7	66	50.8		92	59.0	182	51.9	
N1-3	84	41.8	60	46.2	0.359	62	39.7	158	45.0	0.198
Missing	3	1.5	4	3.0		2	1.3	11	3.1	
ER-status										
Negative	63	31.3	60	46.2		57	36.5	124	35.3	
Positive	128	63.7	67	51.5	0.011	96	61.5	217	61.8	0.849
Missing	10	5.0	3	2.3		3	1.9	10	2.9	
PgR-status										
Negative	60	29.9	65	50.0		54	34.6	149	42.5	
Positive	127	63.2	63	48.5	0.001	96	61.5	190	54.1	0.1
Missing	14	6.9	2	1.5		6	3.9	12	3.4	
Her2-status										
Overexpression-	129	64.2	92	70.8		112	71.8	260	74.1	
Overexpression+	13	6.5	16	12.3	0.166	13	8.3	34	9.7	0.730
Missing	59	29.3	22	16.9		31	19.9	57	16.2	
fITF										
low expression	87	43.3	14	10.8						
high expression	91	45.3	110	84.6	<0.001					
Missing	23	11.4	6	4.6						
Factor VII										
Low expression						87	55.8	91	25.9	
High expression						14	9.0	110	31.3	
Missing						55	35.2	150	42.7	<0.001

Table 1. Associations between clinical and pathological parameters, and expression of FVII or fITF. Abbreviations N number of patients; ER estrogen receptor; PR progesterone receptor; HER2 human epidermal growth factor receptor 2.

	Both fitF+/FVII+		fitF-/FVII+		fitF+/FVII-		Both fitF-/FVII-		p-value
	N	%	N	%	N	%	N	%	
Total	110	100	14	100	91	100	87	100	
Age									
<40	13	11.8	0	0.0	3	3.3	11	12.6	0.264
40-55	39	35.5	5	35.7	37	40.7	36	41.4	
55-65	25	22.7	5	35.7	17	18.7	15	17.2	
>=65	33	30.0	4	28.6	34	37.4	25	28.7	
Missing	0	0.0	0	0.0	0	0.0	0	0.0	
Grade									
I	8	7.3	1	7.1	12	13.2	17	19.5	0.001
II	43	39.0	9	64.3	51	56.0	49	56.3	
III	56	51.0	4	28.6	28	30.8	19	21.8	
Missing	3	2.7	0	0.0	0	0.0	2	2.3	
Histological type									
Ductal	103	93.6	12	85.7	81	89.0	75	86.2	0.142
Lobular	4	3.6	2	14.3	10	11.0	10	11.5	
Missing	3	2.7	0	0.0	0	0.0	2	2.3	
T-status									
T1	34	30.9	3	21.4	35	38.5	40	46.0	0.04 1
T2	62	56.4	10	71.4	35	38.5	37	42.5	
T3/4	13	11.8	1	7.1	17	18.7	8	9.2	
Missing	1	0.9	0	0.0	4	4.4	2	2.3	
N-status									
N0	57	51.8	8	57.1	49	53.8	55	63.2	0.741
N1-3	49	44.5	6	42.9	41	45.1	31	35.6	
Missing	4	3.6	0	0.0	1	1.1	1	1.2	
ER-status									
Negative	52	47.2	8	57.1	27	29.7	28	32.2	0.018
Positive	57	51.8	6	42.9	64	70.3	57	65.5	
Missing	1	0.9	0	0.0	0	0.0	2	2.3	
PgR-status									
Negative	54	49.1	9	64.3	32	35.2	25	28.7	0.010
Positive	55	50.0	5	35.7	57	62.6	58	66.7	
Missing	1	0.9	0	0.0	2	2.2	4	4.6	
Her2-status									
expression-	81	73.6	9	64.3	69	75.8	65	74.7	0.050
expression+	14	12.7	4	28.6	8	8.8	4	4.6	
Missing	15	13.6	1	7.1	14	15.4	18	20.7	

Table2. Associations between clinico-pathological parameters and combined expression of FVII and fitF.
Abbreviations N number of patients; ER estrogen receptor; PR progesterone receptor; HER2 human epidermal growth factor receptor 2.

Characteristic	Relapse Free Survival						Relative Survival						
	Univariate analysis			Multivariable analysis			Univariate analysis			Multivariable analysis			
	N	HR	95% CI	P	HR	95% CI	P	RER	95% CI	P	RER	95% CI	P
Age													
<40	48	1.00						1.00			1.00		
40-50	145	0.97	0.612-1.539	0.422				0.95	0.593-1.527	0.252	1.14	0.460-2.811	0.071
50-60	132	1.17	0.734-1.853					1.32	0.825-2.104		1.91	0.778-4.677	
>60	249	0.90	0.574-1.408					1.28	0.779-2.095		1.00		
Grade													
I	80	1.00			1.00		0.184	1.00			1.00		
II	282	1.43	0.945-2.172	0.001	1.49	0.728-3.058	0.184	1.57	0.865-2.851	0.011	1.14	0.460-2.811	0.071
III	203	2.02	1.326-3.078		1.88	0.896-3.938		2.22	1.221-4.043		1.91	0.778-4.677	
Histological type													
Ductal	513	1.00		0.291				1.00			1.00		0.158
Other	53	1.24	0.832-1.846					1.48	0.933-2.348	0.095	1.77	0.800-3.936	
Tumor stage													
pT1	211	1.00			1.00		0.290	1.00			1.00		
pT2	272	1.59	1.205-2.093	<0.001	1.14	0.763-1.715	0.290	2.46	1.670-3.632	<0.001	1.15	0.680-1.932	0.200
pT3/4	72	2.49	1.706-3.635		1.55	0.896-2.675		4.41	2.782-6.988		1.77	0.923-3.402	
Nodal stage													
Negative	307	1.00		<0.001	1.00		<0.001	1.00			1.00		0.001
Positive	250	3.06	2.379-3.945		2.53	1.759-3.629		3.18	2.285-4.416		2.29	1.431-3.663	
ER status													
Negative	203	1.00		0.725				1.00		0.105			
Positive	337	1.05	0.808-1.359					0.77	0.599-1.056				
PgR status													
Negative	223	1.00		0.744				1.00		0.152			
Positive	313	0.96	0.743-1.236					0.79	0.577-1.089				
HER2 status													
Negative	378	1.00		0.401				1.00		0.152			
Positive	44	1.21	0.776-1.883					1.43	0.877-2.324				
TF:FVII													
Either or both	192	1.00		0.006	1.00		0.014	1.00		0.051	1.00		0.018
low	110	1.59	1.141-2.218		1.55	1.095-2.205		1.55	0.998-2.409		1.68	1.092-2.575	
Both high													

Table 3. Cox univariate and multivariate analysis for recurrence free period and relative survival for TF:FVII. Abbreviations N number of patients; HR hazard ratio; RER relative excess risk; 95%CI 95% Confidence Interval; ER estrogen receptor; PR progesterone receptor; HER2 human epidermal growth factor receptor 2.