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## Close Temporal Relationship Between Onset of Cancer and Scleroderma in Patients with RNA Polymerase I/III Antibodies

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### Abstract

**Objective**—We examined the temporal relationship between scleroderma development and malignancy, and evaluated whether this differed by autoantibody status among affected patients.

**Methods**—Participants had a diagnosis of scleroderma, cancer, an available serum sample, and a cancer pathology specimen. Sera were tested for autoantibodies against topoisomerase I, centromere, and RNA polymerase I/III by immunoprecipitation and/or ELISA. Clinical and demographic characteristics were compared across autoantibody categories. Expression of RNA polymerases I and III was evaluated by immunohistochemistry using cancerous tissue from patients with anti-RNA polymerase antibodies.

**Results**—Twenty three subjects were enrolled. Six subjects tested positive for anti-RNA polymerase I/III (Pol), 5 for anti-topoisomerase I (Topo), 8 for anti-centromere (CENP), and 4 recognized none of these antigens (Negative). The median duration of scleroderma at cancer diagnosis differed significantly between groups:  $-1.2$  years (Pol),  $+13.4$  years (Topo),  $+11.1$  years (CENP), and  $+2.3$  years (Negative) ( $p=0.027$ ). RNA polymerase III demonstrated a robust nucleolar staining pattern in 4 of 5 available tumors from patients with antibodies to RNA polymerase I/III. In contrast, nucleolar RNA polymerase III staining was not detected in any of 4 examined tumors in the RNA polymerase antibody-negative group ( $p=0.048$ ).

**Conclusions**—There is a close temporal relationship between onset of cancer and scleroderma in patients with antibodies to RNA polymerase I/III, which is distinct from scleroderma patients with other autoantibody specificities. In this study, autoantibody response and tumor antigen expression are associated. We propose that malignancy may initiate the scleroderma-specific immune response and drive disease in a subset of scleroderma patients.

### Introduction

Patients with scleroderma may have an increased risk of malignancy compared to the general population (1–6). A wide array of cancers has been reported in scleroderma, although lung and breast cancers are thought to be the most common (3,4,6,7). Although it is controversial whether malignancy risk is truly increased in scleroderma patients, reports detailing a close, at times concurrent onset of scleroderma and malignancy raise the possibility of malignancy triggering an autoimmune disease process in a subset of scleroderma patients (8–10). Among scleroderma patients, this tight temporal association is most striking in breast cancer, with the majority of cases developing scleroderma within 18 months of cancer diagnosis (11–14). In 2 case series reviewing scleroderma patients with

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breast cancer, it has been estimated that up to 50% of breast cancer cases closely preceded or were diagnosed simultaneously with scleroderma (12,14). Additionally, it is reported that prompt therapy of a malignancy can abrogate the scleroderma disease process (8,9,15), suggesting that in these unique cases the biological response to the malignancy or the malignant process itself may be driving the expression of scleroderma.

Despite this reported association between malignancy and scleroderma onset, few studies have evaluated scleroderma disease characteristics that associate with the presence or risk of malignancy, and little is known about potential mechanisms underlying this connection. We hypothesize that scleroderma-specific autoantibody production in a subset of patients with scleroderma is a manifestation of the immune response to tumor antigens that may associate with or induce the scleroderma disease process. In this study, we evaluated whether clinical characteristics, including the temporal relationship between scleroderma and malignancy onset, differed by autoantibody status among patients with scleroderma and cancer. After demonstrating a temporal clustering between cancer onset and scleroderma in the RNA polymerase antibody-positive group, we investigated the expression of RNA polymerases I and III in cancerous tissue of these scleroderma patients compared to cancers from RNA polymerase antibody-negative patients, as well as noncancerous tissue from controls.

## Patients and Methods

### Patients

Participants were scleroderma patients followed at the Johns Hopkins Scleroderma Center who had (i) a new or past diagnosis of malignancy, (ii) an available serum sample, and (iii) an existing cancer pathology specimen available for histologic confirmation of cancer diagnosis. Among established patients, subjects were identified as having had a prior diagnosis of malignancy from the Center's research database. Eligibility included informed consent and meeting either American College of Rheumatology criteria for scleroderma (16), having at least 3 of 5 features of the CREST syndrome (calcinosis, Raynaud's, esophageal dysmotility, sclerodactyly, telangiectasias), or having definite Raynaud's phenomenon, abnormal nailfold capillaries and the presence of a scleroderma-specific autoantibody. For all patients, the closest available serum sample to cancer diagnosis was studied.

Demographic data, scleroderma subtype (limited versus diffuse skin disease), disease duration, date of cancer diagnosis, smoking status (never, former, or current), most recent Medsger disease severity scores (17), peak modified Rodnan skin scores (18), medication use prior to cancer diagnosis, autoantibody status, pulmonary function tests, and echocardiogram data were obtained from the Center's database and, when necessary, medical chart review. Patients were classified according to LeRoy et al (19) as having limited cutaneous disease if scleroderma skin changes were noted only on the face and/or distal to the knees and elbows; diffuse cutaneous disease involved the trunk and/or proximal extremities. Disease duration was defined as the period of time from the first non-Raynaud's phenomenon symptom to the date of cancer diagnosis. Cancer onset was defined by the date of cancer diagnosis. All forced vital capacity (FVC) and diffusion capacity (DLCO) results were standardized by age, gender and height according to NHANES criteria and Knudson et al., respectively (20,21). Interstitial lung disease (ILD) was defined by an FVC < 70% predicted, and pulmonary hypertension (PH) was defined by a right ventricular systolic pressure (RVSP)  $\geq$  45mmHg on resting echocardiography (22) or by right heart catheterization evidence of pulmonary arterial hypertension.

All studies on human materials were performed on samples provided in compliance with Johns Hopkins IRB and HIPAA regulations. Surgical procedures were performed for patient

management; the research tissue used in our studies was in excess of the biopsied tissue required for routine diagnostic purposes. Serum samples were collected under an IRB approved protocol. Each serum was tested for autoantibodies against topoisomerase I by ELISA using commercially available kits (Inova Diagnostics). The presence of antibodies against RNA polymerase I/III in each patient serum was assessed using two different assays. Sera were tested by immunoprecipitation using radiolabeled Hela cell extracts. The presence of anti-RNA polymerase I/III antibodies was determined based on co-migration of immunoprecipitated bands with those detected using an RNA polymerase I/III scleroderma reference serum, which was included in each precipitation set (data not shown). The findings for RNA polymerase III were also validated using a commercially available RNA polymerase III ELISA kit (Inova Diagnostics); in all cases, the identified sera had antibodies against RNA polymerase using both assays. The presence of anti-centromere antibodies was determined by immunoprecipitation using *in vitro* transcription translated <sup>35</sup>S-methionine labeled centromere protein B as described (23).

### Immunohistochemistry

Paraffin sections from affected cancerous tissue in 6 patients with RNA polymerase antibodies were initially available for study. Since the patient tissues were obtained and paraffin embedded by various pathology units at different times, we first assessed fixation variation (and subsequent loss of antigenicity) by staining sections of each tissue with a monoclonal antibody to CD31 (Dako). This evaluation confirmed excellent tissue preservation and comparable antigenicity in tissues from 4 different patients (subjects 1 and 35 (both breast cancer), subject 2 (lung cancer) and subject 4 (ovarian cancer)). As 2 of the original 6 patients with RNA polymerase antibodies did not have adequate tissue preservation, tissue from 1 additional patient (subject 42) with anti-RNA polymerase antibodies and breast cancer was obtained for further validation of our preliminary findings. Normal breast and ovary paraffin sections were purchased from US Biomax (Rockville, MD). Cancerous and normal tissue sections were stained with a monoclonal antibody against RNA polymerase I (polypeptide C) (Abnova, Taipei, Taiwan) or a polyclonal antibody against RNA polymerase III (POLR3A, Santa Cruz Biotechnology, CA). Staining was visualized with diaminobenzidine per the manufacturer's directions (Dako, Carpinteria, CA), and all sections were counterstained with Mayer's hematoxylin.

### Statistical analyses

Clinical and demographic characteristics were compared across autoantibody categories. Statistical significance testing included the Kruskal Wallis test for continuous variables and the Fisher's exact test for binomial and categorical variables. Comparison of tissue nucleolar RNA polymerase III antigen expression by serum anti-RNA polymerase III antibody status was performed by the Fisher's exact test. Statistical analyses were performed using Stata 10.0 (Stata Corporation, College Station, TX, USA). *P* values are 2-sided and were considered significant at  $\alpha = 0.05$ .

### Results

Among 2367 patients seen at the Johns Hopkins Scleroderma Center, 210 have a known history of malignancy. Thirty seven of these patients were seen between February and August of 2008 and were screened for entry into our study. Both serum and pathology samples could be obtained in 23 of these individuals. These 23 cases therefore comprised our study population. The mean ages at scleroderma and cancer diagnosis were 50.1 years (SD 12.1 years) and 57.3 years (SD 11.0 years), respectively. The mean duration of scleroderma at cancer diagnosis was +7.2 years (SD 10.4), and the average duration of Raynaud's phenomenon (RP) at cancer diagnosis was +10.1 years (SD 14.1). The majority

of subjects were female (95.7%) and white (95.7%). Nineteen subjects (82.6%) met ACR criteria for diagnosis of systemic sclerosis, and the remaining 4 met CREST criteria. Fourteen subjects (60.9%) had limited cutaneous disease, and 9 subjects (39.1%) had diffuse skin involvement. The mean modified Rodnan skin score was 15.0 (SD 14.6), and 5 individuals had ILD, 2 had scleroderma renal crisis, and 3 had a myopathy suggestive of inflammatory myositis. The types of malignancies were varied, but the majority (91.3%) were epithelial cell tumors with breast being the primary site in 13 cases (Table 1).

Among these 23 individuals, 6 tested positive for anti-RNA polymerase I/III, 5 for anti-topoisomerase I, 8 for anti-centromere, and 4 for none of these 3 antibodies (Table 2). No individual produced antibodies to more than one of these tested autoantigens. Age, gender, race, smoking status, disease severity indices, medication use prior to cancer diagnosis, and the frequency of ILD or evidence for PH did not differ statistically between groups. In contrast, the median duration of scleroderma at cancer diagnosis differed significantly between groups:  $-1.2$  years in the anti-RNA polymerase I/III group (range  $-2$  to  $+1.3$  years),  $+13.4$  years in the anti-topoisomerase I group (range  $+0.25$  to  $+29$  years),  $+11.1$  years in the anti-centromere group (range  $-2.0$  to  $+36.9$  years), and  $+2.3$  years in the group negative for all of these antibodies (hereafter referred to as the “antibody negative group”) (range  $-1.2$  to  $+5.0$  years) ( $p=0.027$ ) (Figure 1). The median duration of RP at cancer diagnosis followed a similar trend with a duration of  $+0.25$  years in the anti-RNA polymerase I/III group (range  $-2.4$  to  $+1$  years),  $+13.2$  years in the anti-topoisomerase I group (range  $+0.25$  to  $+34$  years),  $+23.8$  years in the anti-centromere group (range  $-5.0$  to  $+36.9$  years), and  $+4.0$  years in the antibody negative group (range  $-1.2$  to  $+7.9$  years) ( $p=0.113$ ). Patients in the anti-RNA polymerase I/III group exclusively had diffuse disease, whereas 40% of the anti-topoisomerase I, none of the anti-centromere, and 25% of subjects in the antibody negative group had diffuse cutaneous disease ( $p<0.001$ ). Correspondingly, the median modified Rodnan skin score was significantly higher (36) in the anti-RNA polymerase I/III group than in the anti-topoisomerase I (9), anti-centromere (4) or antibody negative (6) groups ( $p=0.012$ ).

Characteristics of these 6 patients with anti-RNA polymerase I/III and a tight temporal clustering of scleroderma onset and malignancy diagnosis are provided in Table 3. Five of the malignancies were epithelial cell tumors, 3 of which were breast in origin. Cancer diagnosis closely preceded scleroderma onset in 4 individuals, and the range of scleroderma duration at cancer diagnosis was  $-2$  to  $+1.3$  years. Similarly, cancer preceded RP onset in 2 individuals and occurred concurrently in one subject. These subjects had aggressive skin disease with skin scores ranging from 14 to 48. Scleroderma renal crisis developed in 2 patients.

As adequate tissue preservation permitted only 4 of these 6 subjects' tissues to be studied further, one additional patient with scleroderma, cancer, an available cancer pathology specimen, and anti-RNA polymerase I/III autoantibodies was recruited (subject 42). Subject 42 had a breast ductal carcinoma that was detected 1.5 years after scleroderma onset.

To evaluate levels of RNA polymerase I and RNA polymerase III expression *in vivo*, paraffin sections from cancerous breast ( $n=3$ ; subjects # 1, #35, and #42), cancerous lung ( $n=1$ ; subject # 2) and cancerous ovarian ( $n=1$ ; subject #4) tissue were analyzed by immunohistochemistry (selection of these tissues is detailed in the Methods section). Tumors from 4 RNA polymerase antibody-negative patients were also evaluated (cancerous breast,  $n=3$  and cancerous lung,  $n=1$ ; 3 of these 4 subjects had anti-topoisomerase I antibodies). Breast, ovarian and lung paraffin sections were also obtained from normal individuals for the purposes of comparison. Robust and extensive nuclear staining was detected in all of the cancerous tissue sections stained with anti-RNA polymerase I antibody,

irrespective of the subject's RNA polymerase antibody status (Figure 2, panels A, B, E and F and data not shown). When staining was performed under identical conditions using an isotype matched IgG1 antibody instead of the RNA polymerase I monoclonal antibody, no staining was detected (data not shown). In contrast to the prominent staining noted in cancerous tissues, normal breast, ovarian and lung paraffin sections showed minimal/limited RNA polymerase I staining (Figure 2, panels C, D, G and H and data not shown). Of note, in normal breast, RNA polymerase I staining was restricted to ductal cells (Figure 2, panels G and H). The pattern of staining of RNA polymerase III in cancers was strikingly different. An exclusively nucleolar staining pattern with the anti-RNA polymerase III antibody was detected in 4 of the 5 cancerous tissues from RNA polymerase antibody-positive patients (Figure 3, panels A, B, E and F, and data not shown). In contrast, this nucleolar pattern was absent in all 4 of the tumors from RNA polymerase antibody-negative patients ( $p=0.048$ ). Additionally, it was also not detected in the normal tissue sections, nor in the cancerous sections when staining was performed under identical conditions except that normal goat serum was substituted for the anti-RNA polymerase III goat polyclonal antibody (Figure 3, panels C, D, G and H and data not shown). Tumor RNA polymerase III staining and not tumor RNA polymerase I staining is therefore strikingly associated with the scleroderma patient's RNA polymerase antibody status.

## Discussion

In this pilot study, we evaluated whether clinical characteristics, including the temporal relationship between scleroderma and malignancy onset, differed by autoantibody status among patients with scleroderma and cancer. After identifying all patients with a history of cancer seen in our center, we evaluated the first 23 patients in whom histologic confirmation of cancer diagnosis was possible. Within this group, we found that RNA polymerase I/III autoantibodies strongly associate with malignancy that occurs contemporaneously with scleroderma onset. In all patients who produced anti-RNA polymerase I/III antibodies, scleroderma developed within 2 years of cancer diagnosis. Because of this association, we evaluated expression of RNA polymerases I and III in the tumors of patients with anti-RNA polymerase I/III antibodies compared to those patients without these antibodies. Interestingly, we found that nucleolar RNA polymerase III expression was enhanced exclusively in patients with RNA polymerase antibodies. Although RNA polymerase I was expressed at high levels in tumors, this was not restricted to tumors from patients with the anti-RNA polymerase immune response. This association of tumor RNA polymerase III expression with autoantibodies in those patients suggests that RNA polymerase III could be driving the immune response in these individuals. These preliminary findings require confirmation in a larger patient sample with an expanded control population. It is of interest that another subset of patients, the autoantibody "negative" group, also had a similar close relationship between scleroderma and cancer onset. This group may elaborate unique autoantibodies and express novel tumor antigens that remain to be identified.

Our study suggests that cases of paraneoplastic scleroderma may demonstrate hallmark scleroderma specific reactivity. Prior studies have not investigated or detected this association between contemporaneous onset of scleroderma and malignancy and anti-RNA polymerase I/III antibodies in scleroderma patients with cancer. This relationship may have been missed because: (i) RNA polymerase I/III antibody testing was not commercially available until recently and (ii) prior investigations have focused on whether the relationship between scleroderma and cancer differed by scleroderma subtype or tumor origin and histology. By seeking whether the relationship between scleroderma and malignancy onset differed by autoantibody status among patients with scleroderma and cancer, we were able to detect this association even in a relatively small group of patients.



This striking temporal relationship between scleroderma and malignancy onset among anti-RNA polymerase I/III subjects with cancer is similar to that observed in dermatomyositis (24–26) and systemic lupus erythematosus (27) and suggests that cancer and autoimmunity onset might be mechanistically related. There is strong evidence that anti-cancer immunity and autoimmunity are related. For example, effective initiation of anti-cancer immunity during immunotherapy is often accompanied by autoimmunity (28–33). Multiple immune effector pathways are likely involved in causing tissue damage, with prominent involvement of cytotoxic killing pathways. We therefore hypothesize that tumors expressing high concentrations of RNA polymerase III initiate an immune response to these autoantigens. In the appropriate setting, possibly involving enhanced expression of the same autoantigens in damaged or perturbed blood vessels, this anti-tumor immune response may also be directed against specific host tissues, with consequent tissue damage that generates the ongoing rheumatic phenotype. Direct visualization of specific autoantigen expression in tissues targeted in scleroderma is an important future priority.

Although the groups of patients with cancer and autoantibodies to centromere and topoisomerase-1 had a prolonged interval between scleroderma onset and cancer diagnosis, there were outliers in each group (2 anti-centromere and 1 anti-topoisomerase I) in whom cancer and scleroderma onset occurred close together in time. It is of interest that increased topoisomerase I expression has been detected in a variety of cancers (34–36), and in two cases of patients with pre-existing scleroderma, anti-topoisomerase I titers markedly increased in patients at the time of lung cancer diagnosis, recognizing distinct epitopes (37). The data suggest that in some cases, anti-topoisomerase I antibody production might also be driven by malignancy (37). In our series of patients, this group appears to be a minority.

A variety of other mechanisms could explain the relationship between malignancy and scleroderma. Immunosuppressive therapy for autoimmune disease could account for the increase in malignancy risk in a subset of patients. Additionally, treatment of malignancy could result in the development of scleroderma. For example, multiple chemotherapeutic agents have been implicated as potential causes of scleroderma, scleroderma-like disease, or severe Raynaud's phenomenon (38–42). Radiation therapy may also result in severe skin thickening in patients with scleroderma (43) or cause localized scleroderma in patients without a prior history of connective tissue disease (44). If cancer therapy were the inciting agent that triggered the development of scleroderma, we would not expect our findings to segregate by autoantibody status. Chronic inflammation and repair due to the scleroderma disease process may predispose cells to malignant transformation; this may especially be true of late lung cancers and esophageal adenocarcinomas in the setting of pulmonary fibrosis and longstanding gastroesophageal reflux disease, respectively. Other possible explanations for the relationship between cancer and scleroderma include genetic susceptibility to both malignancy and the development of autoimmune disease, or a common inciting exposure.

We propose that in scleroderma patients who produce anti-RNA polymerase antibodies but have not been diagnosed with cancer that the full expression of an underlying malignancy was aborted by the now scleroderma-specific (originally anti-tumor) immune response. In the paraneoplastic neurological diseases, available data suggest that a patient's immune response recognizes antigens expressed in the tumors and the target tissue, and that often patients have very small or undetectable tumors at disease diagnosis (45–47). Further investigation is needed to determine whether anti-RNA polymerase I/III antibodies are a marker for increased malignancy risk in scleroderma and whether more aggressive cancer screening should be performed in this patient population.

It is important to note that our small sample size limits the generalizability of these conclusions, and these results need validation in a larger patient sample with nonscleroderma cancer controls. The association of cancer RNA polymerase III expression, RNA polymerase III autoantibodies and interval between cancer and scleroderma diagnosis observed in scleroderma patients does not in any way predict that such antigen expression patterns are restricted to scleroderma-associated cancers. Indeed, it is likely that similar RNA polymerase III expression patterns occur in tumors from patients who do not have scleroderma, and that additional pathways and events are required to generate both the scleroderma-specific immune response and clinical phenotype. We acknowledge that we cannot establish a causal relationship between cancer and scleroderma with our retrospective study design that focused on scleroderma patients with a history of malignancy. Another limitation of our study was that in some cases, we lacked serum samples that were concurrent with malignancy onset. To address this, we evaluated the closest available serum sample to cancer diagnosis; the median duration between cancer diagnosis and serum sample studied was 2.3 years. There are many issues that can only be addressed in a prospective study including changes in autoantibody profiles in response to cancer therapy (range of autoantibodies targeted and titers).

We have demonstrated a tight temporal relationship between scleroderma onset and malignancy diagnosis in scleroderma patients with cancer who produce anti-RNA polymerase autoantibodies. Expression of RNA polymerase III is enhanced exclusively in the tumors from patients with RNA polymerase antibodies, demonstrating that tumor antigen expression and scleroderma autoantibodies are strongly associated, and highlighting RNA polymerase III as the tumor-associated antigen target. These findings argue for a mechanistic relationship between malignancy, the immune response and development of scleroderma, and raise the possibility that RNA polymerase I/III autoantibodies are markers of malignancy in newly diagnosed scleroderma patients. These findings may have both important diagnostic and therapeutic implications.

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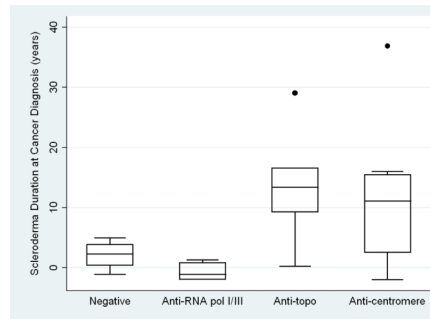
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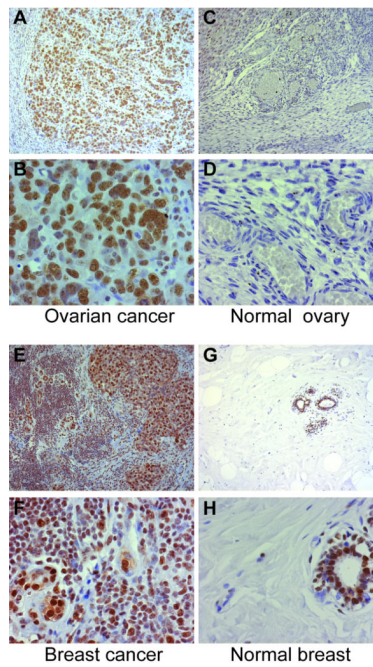


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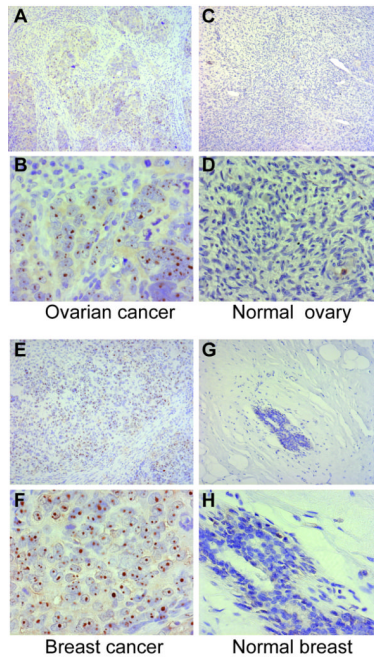


**Figure 1. Distribution of Scleroderma Duration at Cancer Diagnosis by Autoantibody Status**  
 There is a tight temporal relationship between malignancy diagnosis and scleroderma onset in patients producing anti-RNA polymerase I/III antibodies (Anti-RNA pol I/III) compared to patients with anti-topoisomerase I antibodies (Anti-topo) and patients with antibodies to centromere (Anti-centromere). The antibody negative group (Negative) has a similar close relationship between scleroderma and cancer onset.



**Figure 2. RNA polymerase I staining is prominent in cancerous ovary and breast tissue from scleroderma patients compared to normal ovary and breast. Paraffin sections from cancerous ovary (panels A and B) and breast (panels E and F) from scleroderma patients with cancer, as well as normal ovary (panels C and D) and normal breast (panels G and H) were stained with antibodies against RNA polymerase I as described in the methods section**

In all panels, the brown color represents RNA polymerase I staining, with nuclei in blue (Mayers' hematoxylin counterstain). Magnifications are 10 $\times$  (upper panels of each set - panels A, C, E and G) and 40 $\times$  (lower panels of each set - panels B, D, F and H). In each set, the 40 $\times$  panel is a magnification of part of the field shown at 10 $\times$ . The cancer sections shown were from subjects # 4 (ovarian cancer) and # 42 (breast cancer).



**Figure 3. RNA polymerase III staining is prominent in cancerous ovary and breast tissue from scleroderma patients compared to normal ovary and breast. Paraffin sections from cancerous ovary (panels A and B) and breast (panels E and F) from scleroderma patients with cancer, as well as normal ovary (panels C and D) and normal breast (panels G and H) were stained with antibodies against RNA polymerase III as described in the methods section**

In all panels, the brown color represents RNA polymerase III staining, with nuclei in blue (Mayers' hematoxylin counterstain). Magnifications are 10 $\times$  (upper panels of each set - panels A, C, E and G) and 40 $\times$  (lower panels of each set - panels B, D, F and H). In each set, the 40 $\times$  panel is a magnification of part of the field shown at 10 $\times$ . The cancer sections shown were from subjects # 4 (ovarian cancer) and # 42 (breast cancer).



**Table 1**

Cancer Site and Histology of Study Participants, N=23

Cancer site and histology	Number (%)
Breast, no. (%)	13 (56.5)
Ductal carcinoma in situ, no.	4
Invasive ductal carcinoma, no.	5
High grade adenocarcinoma, no.	1
Invasive lobular carcinoma, no.	3
Lung, no. (%)	2 (8.7)
Small cell carcinoma, no.	1
Adenocarcinoma, no.	1
Lymphomas, no. (%)	2 (8.7)
Skin – squamous cell carcinoma, no. (%)	1 (4.3)
Ovary – poorly differentiated carcinoma, no. (%)	1 (4.3)
Tongue – squamous cell carcinoma, no. (%)	1 (4.3)
Uterus – endometrial adenocarcinoma, no. (%)	1 (4.3)
Anus – squamous cell carcinoma, no. (%)	1 (4.3)
Vagina – squamous cell carcinoma, no. (%)	1 (4.3)

**Table 2**

Characteristics of Study Participants by Autoantibody Status

Variables	Pol I/III (N=6)	Topo (N=5)	CENP (N=8)	Negative (N=4)	P-value
Age at Scleroderma Diagnosis (years), median	51.8	45.1	48.9	54.6	0.511
Age at Cancer Diagnosis (years), median	51.0	64.9	60.9	56.9	0.337
Female sex, no. (%)	6 (100)	5 (100)	7 (87.5)	4 (100)	1
Race, no. (%)					0.391
White	6 (100)	4 (80)	8 (100)	4 (100)	
African-American	0 (0)	1 (20)	0 (0)	0 (0)	
ACR Criteria Met, no. (%)*	6 (100)	5 (100)	5 (62.5)	3 (75)	0.176
Scleroderma Classification, no. (%)					<0.001
Limited	0 (0)	3 (60)	8 (100)	3 (75)	
Diffuse	6 (100)	2 (40)	0 (0)	1 (25)	
Scleroderma Duration at Cancer Diagnosis (years), median	-1.2	13.4	11.1	2.3	0.027
Raynaud's Duration at Cancer Diagnosis (years), median	0.25	13.2	23.8	4.0	0.113
Smoking, no. (%)					0.832
Never	4 (66.7)	2 (40)	5 (62.5)	3 (75)	
Former	2 (33.3)	3 (60)	3 (37.5)	1 (25)	
Disease Severity, no. (%)					
Severe Raynaud's Phenomenon**	1 (16.7)	3 (60)	5 (62.5)	0 (0)	0.100
Abnormal General Severity Score §	4 (66.7)	2 (40)	3 (37.5)	1 (25)	0.681
History of renal crisis	2 (33.3)	0 (0)	0 (0)	0 (0)	0.123
History of myopathy	2 (33.3)	0 (0)	0 (0)	1 (25)	0.206
Modified Rodnan Skin Score, median	36	9	4	6	0.012
Interstitial lung disease, † no. (%)	2 (33.3)	2 (40)	0 (0)	1 (25)	0.231
Pulmonary arterial hypertension, ‡ no. (%)	1 (16.7)	1 (20)	2 (25)	1 (25)	1
Medications Prior to Cancer Diagnosis, no. (%)					
Hormone replacement therapy	1 (16.7)	1 (20)	2 (25)	0 (0)	0.892
Prednisone	1 (16.7)	1 (20)	1 (12.5)	0 (0)	1

Variables	Pol I/III (N=6)	Topo (N=5)	CENP (N=8)	Negative (N=4)	P-value
Methotrexate	1 (16.7)	0 (0)	1 (12.5)	0 (0)	1
Azathioprine	0 (0)	1 (20)	0 (0)	0 (0)	0.391
Cyclophosphamide	0 (0)	2 (40)	0 (0)	0 (0)	0.063
Mycophenolate mofetil	0 (0)	1 (20)	0 (0)	1 (25)	0.202

Pol I/III: anti-RNA polymerase I/III, topo: anti-topoisomerase I, CENP: anti-centromere, negative: no autoantibody

\* Remaining patients met CREST criteria.

\*\* Defined as Medsger Raynaud's Severity Score  $\geq 2$ .

§ Defined as Medsger General Severity Score  $\geq 1$ ; based on presence of anemia or weight loss.

‡ Defined by FVC  $< 70\%$  predicted

¥ Defined by RVSP  $\geq 45$ mmHg or right heart catheterization evidence of PAH.

**Table 3**

Characteristics of Initial 6 Anti-RNA polymerase I/III positive subjects

Subject #	Gender	Malignancy	Scleroderma Duration at Cancer Diagnosis (years)	RP Duration at Cancer Diagnosis (years)	mRSS*	Scleroderma Complications**
1	Female	Breast invasive ductal carcinoma	-2.0	0.7	42	Renal crisis ILD
2	Female	Lung small cell carcinoma	-1.0	0	47	ILD
4	Female	Ovarian poorly differentiated metastatic carcinoma	-1.3	-0.3	21	Renal crisis Myopathy
9	Female	Non-Hodgkin lymphoma	0.8	0.5	14	Myopathy PAH
13	Female	Breast invasive ductal carcinoma	1.3	1.0	30	None
35	Female	Breast ductal carcinoma in situ	-2.0	-2.4	48	None

\* Maximal mRSS during disease course

\*\* Complications evaluated include renal crisis, myopathy, ILD, and PAH.

Abbreviations: RP – Raynaud’s phenomenon, mRSS – modified Rodnan skin score, ILD – interstitial lung disease, PAH – pulmonary arterial hypertension