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Anti-Fibrillar Antibody in African American Patients with Systemic Sclerosis: Immunogenetics, Clinical Features, and Survival Analysis

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Abstract

Background—Anti-U3-RNP or anti-fibrillarin antibodies (AFA) are detected more frequently among African American (AA) patients with systemic sclerosis (SSc) compared to other ethnic groups and are associated with distinct clinical features. The current study examines the immunogenetic, clinical, and survival correlates of AFA in a large group of AA patients with SSc.

Methods—Overall, 278 AA SSc patients and 328 unaffected AA controls were enrolled from three North American cohorts. Clinical features, autoantibody profile, and HLA-class-II genotyping were captured. To compare the clinical manifestations, relevant clinical features were adjusted for disease duration. The Cox proportional hazards regression was used to determine the effect of AFA on survival.

Results—Fifty (18.5%) AA patients had AFA. After Bonferroni correction, HLA-*DRB1**08:04 was associated with AFA, compared to unaffected AA controls (OR=11.5, $p<0.0001$) and AFA negative SSc patients (OR=5.2, $p=0.0002$). AFA positive AA patients had younger age of disease onset, higher frequency of digital ulcers, diarrhea, pericarditis, higher Medsger Perivascular and lower Lung Severity Indices ($p=0.004$, $p=0.014$, $p=0.019$, $p=0.092$, $p=0.006$, and $p=0.016$, respectively). After adjustment for age at enrollment, AFA positive patients did not have different survival compared with patients without AFA ($p=0.493$).

Conclusion—These findings demonstrate strong association between AFA and HLA-*DRB1**08:04 allele in AA patients with SSc. Moreover, AA SSc patients with AFA had younger age of onset, higher frequency of digital ulcers, pericarditis, and severe lower gastrointestinal involvement, but less severe lung involvement compared to AA patients without AFA. However, presence of AFA did not change survival.

Keywords

Scleroderma; GENISOS; anti-U3-RNP; digital ulcer; HLA DRB1; and Scleroderma Family Registry

INTRODUCTION

African American (AA) patients with systemic sclerosis (SSc; scleroderma) are reported to have a worse overall prognosis than Caucasians which may be reflected by a younger age of disease onset, higher frequency of diffuse cutaneous involvement, digital ulcers and pits, more severe lung involvement and younger age at onset of pulmonary artery hypertension (PAH) (1–5).

Anti-U3-RNP or anti-fibrillarin antibody (AFA) is directed against a 35 kD protein component of a nucleolar ribonucleoprotein called fibrillarin, which is an early marker for the site of formation of nucleolus in dividing cells (6). The frequency of AFA differs across ethnic groups, ranging from zero in a large cohort of Italian patients with SSc (7) to 50% in an African American SSc population (8). The higher prevalence of AFA in the sera of African American (AA) patients with SSc has been noted in several studies (9–13).

Previous studies have shown that *HLA-DRB1*08* and *DQB1*03:01* are associated with AFA in African Americans (10;14). Clinically, SSc patients with AFA have been reported to have younger ages of disease onset, higher frequency of diffuse cutaneous involvement, pulmonary artery hypertension (PAH), SSc-associated musculoskeletal and cardiac involvement, and lower frequency of arthritis (9–11;15–17). However, there is a lack of large and robust studies on the immunogenetic associations, clinical manifestations, and survival effect of AFA in African American (AA) patients with SSc.

This study compared the HLA class-II alleles in AA SSc patients with AFA with ethnic-gender matched unaffected controls and with SSc patients without AFA. In addition, we investigated the clinical features and survival effect of AFA in AA patients with SSc.

MATERIALS AND METHODS

Study population

Between 1985 and 2010, 3033 patients with SSc were enrolled in the following cohorts: (a) the Genetics versus ENvironment In Scleroderma Outcomes Study (GENISOS) (3;5;18), (b) the NIH/NIAMS Scleroderma Family Registry and DNA Repository (19), and (c) Division of Rheumatology at University of Texas Health Science Center at Houston (UTHSC-H) (10). Patients were included if they met the American College of Rheumatology (ACR; formerly, the American Rheumatism Association) classification criteria for SSc (20) or had at least three of the five CREST (calcinosis, Raynaud's phenomenon, esophageal dysmotility, sclerodactyly, and telangiectasias) features (21). We included all African American patients from these cohorts ($n=278$). Patients enrolled in more than one of the above-mentioned cohorts were identified and duplicate entries were omitted. Furthermore, we enrolled 328 unaffected AA controls to determine any HLA class II allele associations with AFA. The unaffected AA individuals were volunteers with no personal or family history of SSc or other autoimmune disease by screening questionnaire. All enrolled study subjects (SSc patients and unaffected controls) provided written informed consent and the institutional review board of all participating institutions approved the study.

Autoantibody profile and HLA class II allele genotyping

All autoantibody determinations and HLA class II allele typing were conducted in the Division of Rheumatology at UTHSC-H Houston, TX and the Mitogen Advanced Diagnostics Laboratory, University of Calgary, Calgary, Canada. Anti-nuclear antibodies (ANA) and anti-centromere antibodies (ACA) were determined using indirect immunofluorescence with HEp-2 cells as the antigen substrate (Antibodies Inc., Davis, CA, USA). Passive immunodiffusion gels against calf thymus extract were used to examine sera for anti-topoisomerase-I (ATA; Scl-70), anti-Ro/SS-A, anti-La/SS-B, and anti-U1-RNP autoantibodies (INOVA Diagnostics, San Diego, California, USA). Anti-RNA polymerase III (RNAP III) was detected by enzyme-linked immunosorbent immunoassay (ELISA) kits (MBL Co. Ltd, Nagoya, Japan) and AFA were determined by a line immunoassay at a serum dilution of 1:1000 using purified recombinant fibrillarin protein (Euroline-WB: Euroimmun, Lubeck, Germany) in patients who had a positive ANA in anti-nucleolar pattern on the indirect immunofluorescence.

As previously described (5;22), we genotyped HLA class II alleles (*DRB1*, *DQA1*, *DQB1*, and *DPB1*) on extracted and purified genomic DNA. Furthermore, we examined the HLA class-II allele binding peptide by ProPred MHC Class II Binding Peptide Prediction Server in order to predict binding peptides of human fibrillar protein. This prediction is based on quantitative matrices derived from the literature (23;24).

Clinical manifestation

Age, gender, disease type (categorized as limited or diffuse cutaneous involvement at time of enrollment (21)), disease duration (calculated from the onset of the first non-Raynaud's phenomenon symptom attributable to SSc), and modified Rodnan Skin Score (MRSS (25)) were recorded.

To assess the severity of the individual organ system involvement, the Medsger severity indices (13;26) of eight organ systems were captured: peripheral vessels, skin, joints/tendons, skeletal muscle, gastrointestinal (GI) tract, lung, heart, and kidney. However, these data were available only for the patients enrolled in the GENISOS cohort ($n=78$). The presence of digital ulcers was determined based on the participating rheumatologist's clinical assessment. Arthritis was defined as presence of joint swelling and tenderness on physical examination not attributable to osteoarthritis, crystalline arthropathy, or trauma. A decrease in range of motion $> 25\%$ in at least one joint axis was defined as joint contracture. Dysphagia, diarrhea attributable to SSc, and history of SSc renal crisis were recorded. Electrocardiography and 2-dimensional echocardiography findings and/or presence of an auscultatory friction rub determined the presence of pericarditis or clinically significant pericardial effusion.

As previously described (18), pulmonary function tests were obtained at enrollment. Interstitial lung fibrosis was defined as chest radiographs showing fibrosis and/or a forced vital capacity (FVC) of less than 75% predicted value was recorded.

For the purpose of our review, pulmonary artery hypertension (PAH) was defined if the patient had (a) mean pulmonary artery pressure (mPAP) equal or higher than 25 mm Hg on right heart catheterization, (b) right ventricular systolic pressure ≥ 40 mmHg on 2-dimensional echocardiography, or (c) the ratio of FVC % predicted to diffusion capacity of carbon monoxide (DLCO) % predicted ≥ 1.6 . Serum creatinine kinase (CK) levels were recorded and myositis was diagnosed if the patient had proximal muscle weakness with at least one of the following: elevated levels of CK, features of myositis on electromyography, and/or a characteristic muscle biopsy.

Death search

The vital status of the patients was determined through the National Death Index (NDI), at Centers for Disease Control and Prevention (CDCP), which provided data up until 2007. We then reviewed the Social Security Death Index (SSDI) to update our results as of August 2010. SSDI is an online death search tool that provides fatality reports based on observing death certificates and family confirmation. Those patients who were alive based on NDI report and no further records were found on SSDI were assumed to be alive.

Statistical analysis

Homozygosity for alleles at each of the tested HLA loci was not suggestive of recessive inheritance, regardless of whether the referent comparison group was disease-free controls or AFA-negative cases. Besides, there were too few homozygous subjects to distinguish between additive *vs.* dominant modes of inheritance, regardless of the referent. Therefore, a dominant mode of inheritance approach was used to compare the HLA association with

AFA. Heterozygosity and homozygosity for a particular allele were both re-coded as '1' in a binary (zero or one) variable created for each specific HLA gene of interest. In other words, subjects negative for the gene on both of their alleles for the particular HLA locus were coded "0" for the gene on the new binary variable. Bonferroni correction for multiple comparisons was performed for HLA allelic analyses.

Moreover, age, gender, disease type, and duration between AFA positive and AFA negative patients was evaluated utilizing χ^2 and student's t-test accordingly. SSc clinical manifestations might change over the course of disease. Therefore, logistic regression was used to adjust for disease duration as a possible confounding factor in clinical features and to examine the independent effect of AFA.

We utilized a Cox proportional hazards regression analysis to examine the association of AFA with survival. Besides, we investigated the potential association of relevant HLA class-II with survival of the AA patients with SSc. The survival analysis was corrected for age at enrollment. Survival was calculated from the date of enrollment.

ATA and AFA are the two most common anti-nuclear antibodies among AA patients with SSc. We also compared the clinical features and survival of AA scleroderma patients with AFA ($n=50$) to those with ATA ($n=61$) in order to conduct our comparative analysis between more homogenous groups.

All the statistical analyses were performed with SAS Version 9.2 (SAS Institute Inc., Cary, NC) and STATA 11 (StataCorp, College Station, TX). The hypothesis testing was 2-sided with a $p \leq 0.05$ significance level.

RESULTS

Study population, disease, and autoantibody characteristics

All 278 AA scleroderma patients from the three cohorts were included in the study. The mean age (\pm SD) of patients at enrollment was 46.9 (13.9) years and 237 (85.3%) were female. At enrollment, 171 (61.5%) AA patients with SSc were diagnosed with diffuse cutaneous involvement. Average disease duration (\pm SD) was 6.0 (6.5) years.

Anti-nuclear antibodies (ANA) on HEp-2 substrate were detected in 93.1% of AA SSc patients. ATA, RNAP-III, and AFA were present in 21.8%, 15.4%, and 18.5% of patients, respectively (table 1).

HLA class II allelic frequencies

As illustrated in table 2, comparing the HLA class II allelic frequencies of AFA positive patients with 329 ethnically matched unaffected controls demonstrated that the HLA-*DRB1*08:04* allele was seen more frequently in AFA positive patients (47.6% vs. 6.4%; odds ratio (OR): 11.52; 95% confidence interval (CI): 5.43, 24.40; *corrected p* < 0.0001). Two other alleles, which are located on the same haplotype *DQA1*04:01* and *DQB1*03:01*, had similar patterns. However, the increased frequency of *DQA1*04:01* was not statistically significant.

Moreover, the frequency of HLA-*DRB1*08:04* in AFA positive patients also was higher in comparison to AA patients without AFA, even after correction for multiple comparisons (47.6% vs. 14.9%; OR: 5.21; CI: 2.44, 11.09; *corrected p*= 0.0002). Both HLA *DQA1*04:01* and *DQB1*03:01* showed similar trends. However, neither of them remained significant after correction for multiple comparisons.

HLA *DPB1*01:01* was also seen more frequently among AFA positive AA patients compared to unaffected controls and SSc patients without AFA while HLA *DRB1*11:01* seemed to be protective. HLA *DPB1*01:01* and HLA *DRB1*11:01* are not in linkage disequilibrium with HLA *DRB1*08:04*. However, the association of these two alleles with AFA did not withstand correction for multiple comparisons. The frequencies of all relevant HLA class II alleles in AA SSc patients and unaffected individuals are illustrated in supplement table 1.

HLA *DRB1*08:04* binding peptides—Using virtual matrix for HLA *DRB1*08:04*, at a threshold of 1% (the percentage of best scoring natural peptides), we identified four binding peptides (FRSKLAAAI, FRGRGRGGG, IHKPGAKV and FVISIKANC) from the human fibrillar protein that could serve as potential binding sites within the antigen binding groove.

Clinical features

AA SSc patients with AFA were younger at disease onset ($p=0.004$) but gender, disease type, and duration were not significantly different between the two SSc groups. Table 3 illustrates the comparison of clinical manifestations between AFA positive and AFA negative AA patients with SSc.

After adjusting for disease duration, AA SSc patients with AFA had were 3.31 times more likely to have digital ulcers ($p=0.014$). Diarrhea and pericarditis occurred more frequently in AFA positive AA SSc patients (OR 4.84; $p=0.019$ and OR: 2.45; $p=0.092$, respectively) than AA patients without AFA. However, there was no difference between AFA positive and negative AA SSc patients in MRSS, dysphagia, pulmonary artery hypertension (PAH), SSc-associated interstitial lung fibrosis, FVC and DLCO % predicted values, SSc renal crisis, myositis or muscle weakness, serum CK, joint contracture, or sicca symptoms.

Moreover, AFA positive patients had higher Medsger peripheral vascular severity index (regression coefficient (b): 0.79; CI: 0.27, 1.30; $p=0.003$), indicating more severe peripheral vascular involvement and lower Medsger lung severity index (b: -0.82 ; CI: -1.50 , -0.14 ; $p=0.019$), indicating less severe lung involvement. The other Medsger severity indices were not significantly different (table 3).

Survival analysis

At the time of analysis, 30 % of AFA positive patients and 29.5% of AFA negative patients were deceased (table 3). After correction for age at enrollment, AFA positive patients did not have different survival compared to AFA negative AA SSc patients (Hazard ratio=0.79, $p=0.493$). In addition, none of the relevant HLA class-II was predictor of mortality in AA patients with SSc (table 4).

AFA Versus ATA among AA patients with SSc

Although, the age at the onset of first non-Raynaud phenomenon symptom was not statistically different between two groups, the AA scleroderma patients with AFA had higher frequency of digital ulcer, lower GI tract involvement, pericarditis, and Medsger peripheral vascular severity index (Supplement 2). While AFA positive patients had lower Medsger lung severity index, higher FVC and DLCO % predicted values, less cases of PAH. Despite, less severe lung disease, after adjusting for age of disease onset, AFA positive patients did not have better or worse survival compared to ATA positive (table 4).

DISCUSSION

At a frequency of 18.5%, AFA is the second most common anti-nuclear antibody among AA patients with SSc (second to ATA). The present report is the first study investigating the genetic associations, clinical manifestations, and survival impact of AFA in a large population of AA SSc patients.

Distinct HLA class-II allelic associations of SSc-specific autoantibodies in different ethnic groups have been demonstrated in several studies (5;10;14;27;28). In a large sample of Caucasian patients, we have previously reported that the HLA *DRB1*13:02*, *DQB1*06:04/06:05* haplotype correlated with AFA (14). In the current study, we did not observe a similar pattern among AA patients with AFA. Our results indicated that HLA *DRB1*08:04* is strongly associated with AFA in AA patients with SSc, compared to either unaffected individuals or AFA negative AA patients with SSc.

Previous studies investigated potential association of HLA *DRB1*08:04* with other rheumatic conditions like SLE (29) and RA (30). Reveille *et al.* detected no difference in frequency of HLA *DRB1*08:04* between 88 AA patients with SLE or 88 unaffected AA controls. In another study by Hughes LB *et al.* no difference was reported in frequency of HLA *DRB1*08:04* between 321 AA patients with RA and 564 unaffected individuals (30). Previously, we showed that HLA *DRB1*08:04* might be a susceptibility gene for SSc among AA (14); while the results of the current study demonstrated that the higher frequency of HLA *DRB1*08:04* with SSc in AA patients is mainly driven by its strong association with AFA in this ethnic group. Furthermore, through the Binding Peptide Prediction Server for HLA *DRB1*0804*, we identified four potential binding peptides from the human fibrillarin protein that could serve as potential binding sites within the antigen binding groove. The large effect sizes (table 2 and supplement table 1) and predicted binding peptides should prompt more studies to investigate their potential causal and/or environmental relationship of these autoantibodies.

In this regard, an animal model for induction of AFA has been extensively studied and may provide clues to an environmental trigger in humans with AFA-positive SSc. Certain mouse strains possessing specific H2 (the murine counterpart for HLA) haplotypes develop a non-SSc autoimmune disease and high titer AFA following administration of mercuric chloride or silver nitrate (31–34). Of note, one study of urinary mercury levels in SSc patients noted higher levels in those with AFA. However, this observation did not maintain statistical significance following corrections (35). Interestingly, heavy metals have previously been noted to be highly concentrated in the nucleolus (36). It was noted by Pollard *et al.* (37) that most if not all of the SSc-specific autoantigens were at sometime during their life cycle localized to the nucleolus. Clearly, larger and more targeted studies of heavy metal and other environmental exposures in AFA-positive SSc patients, perhaps selected for the associated HLA-class II alleles (*DRB1*13:02* in Caucasians and *DRB1*08:04* in AAs) and AFA negative SSc patients, as well as well-matched normal controls are warranted.

Confirming our previous findings (10), we showed that HLA *DQB1*03:01* has higher frequency among AFA positive AA patients compared to AA unaffected individuals. However, there is no difference between AFA positive and negative AA patients with SSc.

AA SSc patients are younger at SSc onset compared to other ethnic groups (1;2;5;28). Moreover, other studies have shown that SSc patients with AFA have younger age of onset (2;12;13;16;17). In support of these findings, we further demonstrated that AA SSc patients with AFA had younger age of onset in comparison to AA patients without AFA.

Higher Medsger peripheral vascular severity index and prevalence of digital ulcers in AA patients with AFA compared to those without AFA are novel findings. These findings were also present when we compared AFA vs. ATA positive AA patients with SSc. Previous studies have shown higher rate of digital ulcers among AA patients with SSc compared to Caucasians (2;4). Higher frequency of AFA in AAs might contribute to this finding. In another study, Steen *et al* noted higher frequency of digital ulcers in AFA positive patients (12). However, these findings were not stratified for ethnic background.

In agreement with previous studies reporting more severe GI involvement in AFA positive patients (regardless of ethnicity) (10;12), we observed a higher frequency of SSc-associated diarrhea in AFA positive AA patients. The higher frequency of lower GI tract involvement was more significant when AFA positive AA patients were compared to ATA positive patients. It is possible that AFA positive patients have more severe lower GI tract hypomotility and bacterial overgrowth that contribute to diarrhea.

Our results imply a less severe lung involvement among AFA positive AA patients with SSc, as assessed by lower Medsger lung severity index. The comparison of AFA vs. ATA positive AA scleroderma patients further demonstrated less severe lung involvement (higher FVC and DLCO % predicted values and lower Medsger lung severity index). In agreement with our findings, in an ethnically homogenous cohort of Japanese patients with SSc, AFA positive patients had less severe lung involvement (17). While data from several multi-ethnic cohorts suggested a higher frequency of isolated PAH and/or pulmonary fibrosis in the SSc patients with AFA (9;10;12;13;38), these comparisons were adjusted neither for ethnicity nor for other antibodies i.e. ATA, as potential confounders. Therefore, the higher frequency of lung fibrosis and PAH might be due to a sizeable AA population in AFA positive group and large number of Caucasian SSc patients in AFA negative group. More severe SSc-associated lung involvement in AA patients with SSc compared to other ethnic groups has been reported in several studies (2;18;28;39;40).

Based on the current findings, AFA positive AA patients with SSc have a higher prevalence of pericarditis, compared to AFA negative as well as and ATA positive patients. This is in agreement with former studies indicating higher frequency of cardiac involvement in AFA positive (10;12) and AA patients with SSc (39).

Our study did not confirm previous reports of worse (13) or better (11) survival in AFA positive patients with SSc. The worse survival of AFA positive patients in a previous report (13) might be attributable to the confounding or modifying effects of ethnicity in studies that are not stratified by ethnicity. As AA ethnicity is associated with AFA positivity as well as poorer survival (1;10;12;13).

This study has some limitations. Although potentially important, the data on heavy metal exposure were not collected in the current study. The Medsger severity indices were only available in the patients from the longitudinal GENISOS cohort. High-resolution computed tomography scan (HRCT) and echocardiography were not performed on all patients, which might have lead to underreporting of pulmonary involvement, despite being the largest genetic study ever reported in AA patients with SSc, we might be underpowered to detect more subtle HLA associations with AFA in AA population.

In conclusion, AFA was the second most common anti-nuclear antibody in African Americans with SSc. Presence of AFA was strongly associated with the HLA *DRB1**08:04 in the AA SSc patients. In addition, AA SSc patients with AFA had a younger age of disease onset, higher frequency of digital ulcer and pericarditis, more severe lower GI involvement, and less severe pulmonary involvement. Studies in the future should focus on environmental

factors, such as heavy metal exposure, that may influence the B cell response and the immunopathology of the disease.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Table 1

Study population ($n=278$)

Age, mean (\pm SD), yrs	46.9 (13.6)
Gender, female, n (%)	237 (85.3)
Cutaneous involvement, diffuse, n (%)	171 (61.5)
Disease duration, mean (\pm SD), yrs	6.0 (6.5)
MRSS, mean (\pm SD)	17.3 (12.3)
Deceased patients, n (%)	83 (29.7)
Survival time (from time of enrollment), mean (\pm SD), yrs	5.8 (5.0)
Autoantibody profile, %	
ANA *	93.1
ACA **	6.2
ATA †	21.8
ANoA ‡	44.5
AFA ^{β}	18.5
Pol III ^{γ}	15.4
U1-RNP ^{Δ}	12.2
PM/Scl ^{\S}	3.2
Ro ^{Φ}	9.3

* ANA, anti-nuclear antibodies;

** ACA, anti-centromere antibodies;

† ATA, anti-topoisomerase I antibodies;

‡ ANoA, anti-nucleolar antibodies;

^{β} AFA, anti-fibrillarin antibodies;

^{γ} Pol III, RNA polymerase III;

^{Δ} U1-RNP, U1-ribonucleoprotein;

^{\S} PM/Scl, polymyositis-scleroderma antigen;

^{Φ} Ro, Ro-SS-A/60.

Table 2

Frequency of HLA-class II alleles in AFA positive AA patients with SSC compared to ethnically matched AFA negative patients and unaffected controls

HLA class-II alleles	AFA positive (n=50)	AFA negative (n=221)	Unaffected controls (n=329)	AFA positive AA patients vs. unaffected AA controls		AFA positive vs. negative AA patients	
				Odds ratio (95%CI)	p-value	Odds ratio (95%CI)	p-value*
<i>DRB1</i> *08:04	47.6%	14.9%	6.4%	13.20 (6.24,27.94)	<0.001	5.21 (2.44,11.09)	<0.001
<i>DQA1</i> *04:01	33.3%	17.5%	20.6%	1.95 (0.97,3.90)	0.060	2.37 (1.10,5.09)	0.026
<i>DQB1</i> *03:01	69.1%	56.5%	39.0%	3.49 (1.75,6.95)	<0.001	1.72 (0.83, 3.56)	0.153
<i>DPB1</i> *01:01	80%	50.8%	47.8%	4.38 (1.08,25.21)	0.019	4.12 (1.07,15.89)	0.041
<i>DRB1</i> *11:01	0%	16.9%	11.8%	N/A [†]	0.019	N/A [†]	0.004

* corrected p-value,

** NS: not significant,

[†] N/A: Not applicable.

Table 3

Clinical manifestations of African American patients with AFA compared to those without AFA (adjusted for disease duration)

	AFA positive (n=50)	AFA negative (n=221)	Odds ratio (95% CI)	p-value
Age [*] , mean (±SD), yr	41.7 (13.31)	47.9 (13.25)	-6.29 (-10.59, -1.99)**	0.004
Gender [*] , Female, %	86.0	84.6	0.89 (0.31, 2.24)	0.806
Cutaneous involvement [*] , diffuse, %	36.0	39.5	0.86 (0.43, 1.69)	0.642
Disease duration at enrollment [*]	5.34 (5.14)	6.28 (6.85)	-0.93 (-3.51, 1.64)**	0.475
Deceased, %	30.0	29.5	0.99 (0.47, 2.03)	0.892
MRSS [†] , mean (±SD)	14.31 (7.62)	17.83 (13.50)	-2.55 (-9.66, 4.54)**	0.476
Digital ulcer, %	79.3	53.8	3.31 (1.27, 8.62)	0.014
Dysphagia, %	60.0	54.8	1.24 (0.49, 3.19)	0.619
Diarrhea, %	53.9	19.2	4.84 (1.29, 18.13)	0.019
Pericarditis, %	28.2	12.8	2.45 (0.86, 6.93)	0.092
PAH [‡] , %	10.0	22.4	0.38 (0.09, 1.19)	0.081
SSc interstitial lung fibrosis, %	28.2	47.3	0.65 (0.28, 1.52)	0.322
FVC % predicted, mean (±SD)	77.8 (17.7)	72.9 (23.2)	6.08 (-4.54, 16.69)**	0.259
DLCO % predicted, mean (±SD)	67.7 (17.2)	57.9 (24.6)	9.45 (-2.24, 21.14)**	0.112
SSc renal crisis, %	10.5	8.4	1.28 (0.28, 4.54)	0.921
Myositis or muscle weakness, %	30.0	40.6	1.05 (0.21, 5.28)	0.953
Elevated serum CK	28.6	30.9	0.88 (0.31, 2.54)	0.817
Arthritis, %	21.9	31.0	0.53 (0.13, 2.13)	0.370
Joint contracture	22.9	21.4	1.44 (0.56, 3.72)	0.447
Sicca [§] symptoms	20.0	28.1	0.64 (0.11, 3.69)	0.621
Medsker Severity Index, mean(±SD)				
<i>General</i>	0.6	0.5	0.06 (-0.53, 0.64)**	0.838
<i>Peripheral vascular</i>	2.2	1.4	0.79 (0.27, 1.30)**	0.003
<i>Skin</i>	1.5	1.7	-0.27 (-0.84, 0.31)**	0.361
<i>Joint</i>	0.8	0.9	-0.09 (-0.92, 0.73)**	0.813
<i>Muscle</i>	0.2	0.3	-0.10 (-0.41, 0.21)**	0.521
<i>GI Tract</i>	0.6	0.6	0.02 (-0.43, 0.39)**	0.935
<i>Lung</i>	1.1	1.9	-0.82 (-1.50, -0.14)**	0.019
<i>Heart</i>	0.5	0.3	0.16 (-0.27, 0.58)**	0.457
<i>Kidney</i>	0.1	0.3	0.19 (-0.68, 0.29)**	0.441

* These comparisons were not adjusted for disease duration. Student's t-test and χ^2 were utilized for comparisons, accordingly.

** numbers indicate the mean differences.

[†] modified Rodnan Skin Score,

[‡] PAH: pulmonary artery hypertension,

§ having two of three symptoms of dry mouth, dry eye and/or enlarged parotid.

Table 4

Cox proportional hazards regression model analysis of AA patients with systemic sclerosis (SSc)

	Hazard Ratio (95% Confidence interval)	p-value
AFA	0.80 (0.41, 1.53)	0.493
AFA Vs. ATA	0.84 (0.42, 1.69)	0.623
HLA-DRB1*08:04	1.00 (0.55, 1.83)	0.996
HLA-DQA1*04:01	1.50 (0.84, 2.68)	0.170
HLA-DQB1*03:01	0.85 (0.51, 1.41)	0.520
HLA-DQB1*01:01	1.13 (0.49, 2.60)	0.766
HLA-DRB1*11:01	1.17 (0.57, 2.39)	0.671