



Dermatomyositis autoantibodies: how can we maximize utility?

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Abstract: The past 15 years has seen significant advances in the characterization of myositis-specific autoantibodies (MSAs) and their associated phenotypes in patients with dermatomyositis (DM). As more careful studies are performed, it is clear that unique combinations of clinical and pathological phenotypes are associated with each MSA, despite the fact that there is considerable heterogeneity within antibody classes as well as overlap across the groups. Because risk for interstitial lung disease (ILD), internal malignancy, adverse disease trajectory, and, potentially response to therapy differ by DM MSA group, a deeper understanding of MSAs and validation and standardization of assays used for detection are critical for optimizing diagnosis and treatment. Like any test, the diagnostic sensitivity and specificity of assays for various MSAs is not perfect. Currently tests for MSAs are helpful at minimum for a clinician to assess relative risk or contribute to diagnosis and perhaps counsel the appropriate patient about what to expect. With international standardization and larger studies it is likely that more antibody tests will make their way into formal schemata for diagnosis and actionable risk assessment in DM. In this review, we summarize key considerations for interpreting the clinical and pathologic associations with MSA in DM and identify critical gaps in knowledge and practice that will maximize their clinical utility and utility for understanding disease pathogenesis.

Keywords: Dermatomyositis (DM); autoantibodies; immunoassay

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Dermatomyositis (DM) is one of the idiopathic inflammatory myopathies (IIMs) whose classification has continued to evolve over time (1-4). A widely adopted classification system separates the IIMs into DM, overlap myositis including mainly antisynthetase syndrome, immune-mediated necrotizing myopathy, and inclusion body myositis (1,5). Other classification systems include polymyositis as a separate entity, pure polymyositis becoming rarer as more becomes known about the IIMs (2).

For the clinician, one of the challenging aspects about

DM is its heterogeneity. This creates difficulty in diagnosis as well as assessing patient risk for organ involvement and associated, often occult, internal malignancy. Diagnostically, DM can be confused with other rheumatic disorders (e.g., systemic lupus erythematosus, mixed connective tissue disease, rheumatoid arthritis, cutaneous lupus erythematosus), inflammatory myopathies, and cutaneous eruptions (e.g., drug eruptions, photosensitive dermatoses, psoriasis, and others). In terms of end-organ disease, patients can have variable skin morbidity, musculoskeletal

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involvement, and lung disease, and, more rarely, severe gastrointestinal (GI) involvement. In some cases, such as patients with rapidly progressive lung disease or GI vasculopathy, early diagnosis and aggressive therapy may be required to avert morbidity or death. Because 7–32% of DM patients have an associated internal malignancy which is often occult at DM onset, the clinician is left with a quandary regarding how aggressively to screen DM patients for cancer (6–9). Having markers that enhance diagnosis and early detection of adverse prognostic features would theoretically allow earlier actionable therapy and improved patient outcomes.

It is now recognized that DM is associated with an intriguing diversity of myositis-specific autoantibodies (MSAs) that often have characteristic associated systemic and cutaneous manifestations (10–17). These MSAs define subclasses of disease with each autoantibody having characteristic clinical associations, organ pathology, HLA associations, and microRNA profiles (18–22). The comprehensive nature of these associations has been the subject of several excellent reviews and will not be reviewed here (3,10–12,14,15,17,23,24). Instead, we focus on defining the current limitations and contradictions that limit their clinical utility for optimizing patient care and performing translational research.

Studies often differ in the reported prevalence of MSAs and their association with particular clinical features. Possible explanations for this include variable patient inclusion criteria, demographics, and MSA detection assays. The only MSA included as part of the EULAR/ACR classification criteria is anti-Jo-1, mostly due to the fact that during development of these guidelines, and even in many places presently, testing for the other DM-associated MSA was not widely available (25–27). The lack of validated, widely available, and affordable assays for MSA detection is one of the major factors that has hampered the use of MSAs in devising classification criteria and in clinical decision-making. However, it is very likely that other MSAs will be useful in the diagnosis and classification of DM even in light of current limitations (28).

DM is heterogenous, and there is a great need for better tools for diagnosis and prognosis. MSAs may be good biologic classifiers and risk management tools, but there are shortcomings with MSAs that presently limit this. Developing evidence-based screening guidelines for patients with DM is a priority, as leading causes of death in these patients are interstitial lung disease (ILD) and cancer (29,30). In one study, for example, 5-year survival among

DM patients was only 65%, with cause of death segregating according to MSA type: 37% of anti-MDA-5 patients died of ILD and 28% of anti-TIF-1 γ /anti-TIF-1 α patients died of cancer (16).

Current understanding of MSAs

Myositis-specific antibodies can be detected in more than 60% of patients with myositis and it is likely for DM patients that number approaches 80–90% using appropriately sensitive assays (our unpublished data) (18,31). The major MSAs associated with DM are anti-Mi-2, anti-MDA-5, anti-NXP-2, anti-TIF-1 γ , and anti-SAE-1/2 (*Table 1*) (32). In addition, eight anti-synthetase autoantibodies (ASAs) have been defined that, in some cases, can mimic features of DM. These antibodies are: anti-Jo-1 (anti-histidyl-tRNA synthetase), anti-PL-12 (anti-alanyl-tRNA synthetase), anti-PL-7 (anti-threonyl-tRNA synthetase), anti-EJ (anti-glycyl-tRNA synthetase), anti-OJ (anti-isoleucyl-tRNA synthetase), anti-KS (anti-asparaginy-tRNA synthetase), anti-Zo (anti-phenylalanyl-tRNA synthetase), and anti-Ha/YRS (anti-tyrosyl-tRNA synthetase) (*Table 1*) (33). There is controversy about whether all patients with anti-synthetase antibodies have their own syndrome, given that some have features of DM (1,2). Anti-synthetase syndrome and DM may not be completely separate and evaluation of criteria is ongoing (1,2,4). There are also a number of myositis-associated antibodies (MAAs) including anti-PM/Scl, anti-Ro52, and anti-U1RNP, antibodies which are not specific to IIMs (5,34).

Discovery of the MSAs dates back to over 40 years ago, and there has been slow evolution of the concept that they define biologically relevant subgroups of the IIMs, including subgroups within DM. Anti-Jo-1 antibodies were discovered in 1980 and subsequently identified in 15–25% of polymyositis and DM patients (35,36). Anti-Jo-1 antibodies are highly specific for the anti-synthetase syndrome and, although preferentially seen in myositis, can rarely also be found in patients with other autoimmune disorders such as systemic lupus erythematosus, systemic sclerosis, or other diseases (33,36). Anti-Mi-2 autoantibodies were found to be specific for DM in the 1980s after anti-Mi-2 autoantibodies were initially described in polymyositis and DM (37,38). Anti-NXP-2 antibodies were first described in 1997 and were initially denoted anti-MJ antibodies (39–43). Anti-MDA-5 antibodies were discovered in 2005 and were originally called anti-CADM-140 antibodies given their association with clinically amyopathic dermatomyositis (CADM) (44). Anti-155/140 autoantibodies were discovered

Table 1 Myositis-specific antibodies and associated clinical phenotypes

Antigen	Antibody prevalence among patients with DM (%)	Clinical features
Transcription intermediary factor 1 γ (TIF-1 γ)	8–41 (USA, Europe), 7–14 (Japan)	Malignancy, severe rash, “red on white” poikiloderma, ovoid palatal patch
Melanoma differentiation-associated gene 5 (MDA-5)	0–13 (USA, Europe), 11–37 (Asia)	ILD, clinically amyopathic, arthritis, cutaneous ulcers, alopecia, hyperferritinemia, calcinosis
Nuclear matrix protein (NXP-2)	3–30 (USA, Europe), 2–4 (Japan)	Calcinosis, malignancy severe dysphagia, myalgia, distal weakness, intestinal vasculopathy
Small ubiquitin-like modifier activating enzyme (SAE)	5–10 (USA, Europe), 2–3 (Asia)	Dysphagia, “angel wing” rash on back, skin predominant precedes variable myositis
Nucleosome-remodeling deacetylase complex (Mi-2)	10–21 (USA, Europe), 2–30 (Asia), ~8 (Brazil)	Photo-distributed rash, good prognosis with common relapse, high CK with nearly universal muscle involvement
Aminoacyl tRNA synthetases (ASAs)	<5–10	Myositis, polyarthritis, ILD, mechanic’s hands, Raynaud phenomenon

DM, dermatomyositis; ILD, interstitial lung disease; ASAs, anti-synthetase autoantibodies.

in 2007 and later shown to recognize TIF-1 γ /anti-TIF-1 α proteins, respectively, with TIF-1 β being a target of antibodies much less frequently (45–47). In 2007, anti-SAE antibodies were reported, and appear to delineate another class of patients that otherwise would be seronegative for the known MSA (48).

The MSAs recognize diverse antigens and it has been challenging to identify a single theme that unites them with regards to cellular localization or function (*Table 1*). Mi-2 is part of the nucleosome-remodeling deacetylase complex which is involved in regulation of chromatin structure with effects on transcription and DNA repair as well as end-organ function, such as regulation of myoblast differentiation during muscle regeneration (49,50). MDA-5 encodes a retinoic acid-inducible gene I (RIG-I)-like receptor that is a cytosolic double-stranded RNA sensor recognizing viral RNA for innate immunity (51). The nuclear matrix protein (NXP-2) has unclear function, but is known to have a role in p53 regulation (39,52). TIF-1 γ , a member of the tripartite motif (TRIM) family of proteins that also includes TIF-1 α and TIF-1 β , has been shown to regulate transcription, tumor growth, DNA damage repair, and TGF- β signaling (45,47,53–56). SAE-1/2 are the A and B subunits of small ubiquitin-like modifier 1 (SUMO-1) activating enzyme, forming the heterodimer SAE that is involved in sumoylation, a post-translational process that regulates protein localization, stability and function (48). The anti-synthetase antibodies target tRNA synthetases,

cytoplasmic enzymes that generate aminoacyl tRNAs for protein translation, but may have other roles in the cell as well (36).

MSAs tend to occur with different frequencies in various patient populations and several MSAs are known to be genetically associated with HLA haplotypes. Anti-Mi-2 autoantibodies are associated with HLA DRB1*0302 and DRB1*0701/DQA1*0201, differing in that DRB1*0302 is often seen in African Americans and DRB1*0701/DQA1*0201 is often seen in European Americans (57–61). HLA-DRB1*04 and HLA-DQA1*03 are risk factors for the development of anti-Mi-2 autoantibodies in Native Americans (61). While associated with development of anti-Mi-2 antibodies, the linked alleles DRB1*0701/DQA1*0201 may be protective against the development of anti-Jo-1 antibodies (58). Anti-MDA-5 autoantibodies are associated with DRB1*0401, DRB1*1202, DRB1*1201, and DRB1*0901 in Chinese (62,63). Anti-TIF-1 γ antibodies are associated with DQB1*02:02 in Caucasian adults and DQB1*02:01 in Caucasian children (21).

Beyond genetic associations, certain MSAs may be associated with environmental exposures. In particular, anti-Mi-2 and anti-TIF-1 γ antibodies may be associated with UV exposure (64–68).

MSAs are associated with a variety of clinical phenotypes that potentially could impact clinical care decision making (*Table 1*). Two of these clinically important outcomes are the risk of internal malignancy and ILD. In a large study in

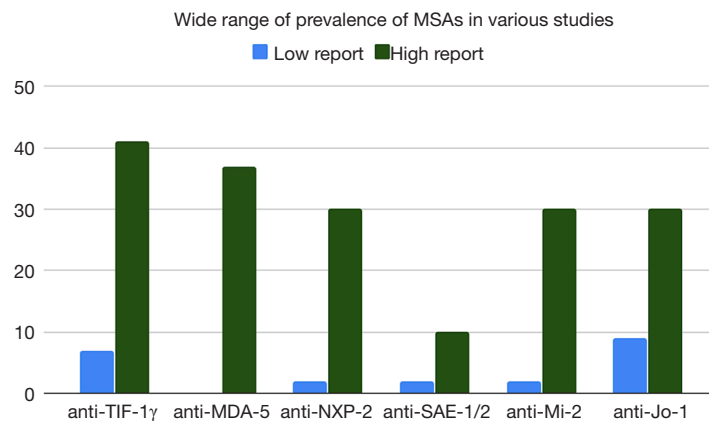


Figure 1 Wide range of prevalence of major MSAs in various studies. Anti-TIF-1 γ : low report: 7% (Japan, anti-TIF-1 γ /anti-TIF-1 α , IP) (16); high report: 41% (USA, IP) (86). Anti-MDA-5: low report: 0% (Hungary, IP) (87); high report: 37% (China, ELISA) (88). Anti-NXP-2: low report: 2% (Japan, IP) (42); high report: 30% (Italy, IP) (89). Anti-SAE-1/2: low report: 2% (Japan, IP) (90); high report: 10% (UK, IP) (48). Anti-Mi-2: low report: 2% (Japan, IP) (16); high report: 30% (India, LIA) (91). Anti-Jo-1: low report: 9% (Italy, LIA) (92); high report: 30% (USA, African Americans, IP) (93). MSAs, myositis-specific autoantibodies; IP, immunoprecipitation; ELISA, enzyme-linked immunosorbent assay.

USA, 83% of DM patients who developed cancer had anti-TIF-1 γ or anti-NXP-2 antibodies, and numerous studies have associated anti-TIF-1 γ antibodies with diagnosis of malignancy (6,9). However, these data are limited by the fact that they are only relative risks and say nothing about the general risk for cancer in each antibody group compared to control patients. It is quite possible that, compared to a control population, the other DM-specific antibodies are also associated with an increased risk of internal malignancy. Similarly, anti-MDA-5 antibodies are associated with ILD, but both the prevalence as well as severity of the anti-MDA-5-associated ILD vary greatly between studies of patients from very different populations (16,51,69-79).

MSAs may be associated with differential response to treatments. As examples, anti-Mi-2 patients experience greater and more rapid benefit from rituximab, patients with anti-MDA-5 antibodies have lower chance of achieving cutaneous clinical remission even after aggressive systemic therapies, and patients with anti-SAE-1/2 antibodies are more likely to have hydroxychloroquine-related skin eruptions whereas anti-MDA-5 antibodies appear to be protective for this (16,80-85).

Uncertainty regarding prevalence of MSAs

Because there is high variability in reports of MSA prevalence and strength of association with various DM-

associated morbidities (Figure 1, Table 1), the utility of a test result to guide clinical decision-making is presently unclear.

Anti-TIF-1 γ , anti-NXP-2, and anti-SAE antibodies tend to be more prevalent in studies conducted in USA and Europe than in studies conducted in Asia. For example, anti-TIF-1 γ antibody prevalence in DM ranges from 8–41% in USA and Europe compared to only 7–14% in Japan (6,16,46,86,87,94-96). Similarly, anti-NXP-2 antibody prevalence in DM ranges from 3–30% in USA and Europe and only 2–4% in Japan (6,42,43,87,89,97-99). Less pronounced is the anti-SAE antibody prevalence difference in DM which ranges from 5–10% in USA and Europe compared to 2–3% in Asia (48,87,90,100-104). In contrast, anti-MDA-5 autoantibodies are found in a much larger percentage of Asian DM patients, 11–37%, compared to only 0–13% in patients from Europe and USA (16,72,77,87,88,105). Anti-Mi-2 antibodies are found in similar percentages in DM patients around the world: in Asia ranging from 2–30%, in Brazil around 8%, and in Europe and USA ranging from 10–21% (16,44,64,91-93,106-112). Meta-analysis of prevalence of anti-Mi-2 antibodies across studies found 9% prevalence in DM patients with 95% confidence interval of 9–14% (113). Among ASAs, anti-Jo-1 is most common with prevalence in DM patients from Asia ranging from 10–14% compared to 9–22% in USA and Europe, though African American DM patients have a higher prevalence of around 30% (44,91-93,109-111).

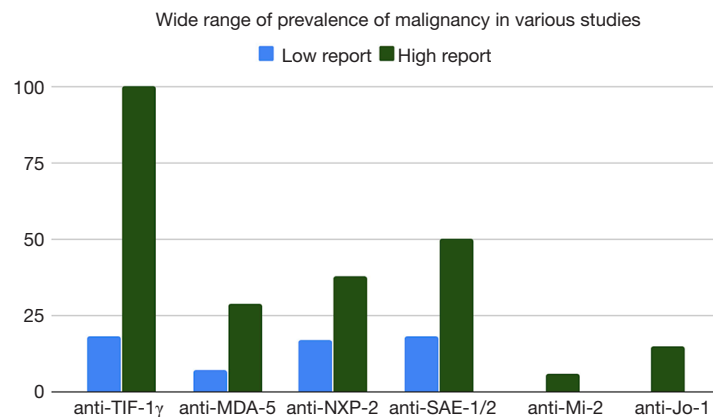


Figure 2 Wide range of prevalence of malignancy in various studies. Anti-TIF-1 γ : low report: 18% (USA, IP) (6); high report: 100% (Japan, anti-TIF-1 γ /anti-TIF-1 α , IP) (70). Anti-MDA-5: low report: 7% (China, ELISA) (77); high report: 29% (ELISA and immunoblot) (78). Anti-NXP-2: low report: 17% (Europe, IP) (18); high report: 38% (Japan, IP) (42). Anti-SAE-1/2: low report: 18% (UK, IP) (100); high report: 50% (Japan, ELISA and IP) (115). Anti-Mi-2: low report: 0% (Japan, IP) (116); high report: 6% (USA, ELISA) (37). Anti-Jo-1: low report: 0% (Hungary, immune serology otherwise unspecified) (117); high report: 15% (Japan, IP) (114). IP, immunoprecipitation; ELISA, enzyme-linked immunosorbent assay.

Meta-analysis found 11% prevalence of anti-Jo-1 in DM patients with 95% confidence interval of 9–14% (113). Also in this meta-analysis, anti-PL-7 antibodies were detected in 2% of DM patients, anti-PL-12 in 3%, anti-KS in 1%, anti-OJ in 1%, and anti-EJ in 1% (113). Restricting to DM-like patients with antisynthetase antibodies, a study in Japan found the prevalences to be anti-Jo-1 (36%), anti-EJ (23%), anti-PL-7 (18%), anti-PL-12 (11%), anti-KS (8%), and anti-OJ (5%) (114).

Antibodies and clinical associations with malignancy and ILD: uncertain risks

Anti-TIF-1 γ antibodies have a very strong association with cancer, but there is a large range in internal malignancy prevalence in the various reported studies (*Figure 2*) (6,16,18,46,70,86,96,118-121). Though the prevalence of cancer in anti-TIF-1 γ patients tends to be higher in Japan (38–68%) versus the USA (18–46%), the ranges vary between studies and are overlapping (6,16,46,86,96). In patients with anti-NXP-2 antibodies, around 38% in Japan and around 24% in the USA develop internal malignancy (6,42). Anti-NXP-2 autoantibodies were found to have a 3.68 increased risk of cancer relative to the general population in a study of 56 anti-NXP-2 patients compared with 179 other DM patients (97). However, in a study of 20 anti-NXP-2 patients compared

with 158 other DM patients, the increase in prevalence of internal malignancy was not statistically significant, and a recent large study out of Europe that included 1,483 IIM patients did not find an association between anti-NXP-2 antibodies and cancer (18,43). Anti-aminoacyl-tRNA synthetase (anti-ARS) patients have up to 12% malignancy with rates for anti-Jo-1 (15%), anti-OJ (25%), anti-PL-12 (16%), and anti-KS (15%) being somewhat higher, and rates for anti-EJ (3%) and anti-PL-7 (7%) being somewhat lower (114). However, an earlier study of 103 DM patients found 16% had anti-Jo-1 antibodies none of whom had associated malignancy (117). Anti-Mi-2 patients have generally been considered to be associated with relative lower rates of malignancy, for example 0% in Japan and 6% in USA (37,81,116). However, this is not uniform, and recent studies have found significant association between anti-Mi-2 antibodies and malignancy (18,122). A recent large study suggested an anti-Mi-2 cancer association with odds ratio 2.50 for developing cancer in anti-Mi-2 patients compared to odds ratio of 4.67 for anti-TIF-1 γ patients (18). Reports of malignancy in anti-MDA-5 patients are rare likely due to the major concern of ILD in these patients, however one study in China reports that 7% of anti-MDA-5 patients had malignancy and another study in Spain reported 29% (77,78).

The lack of agreement on predictive values of antibody tests for malignancy can create uncertainty on the part

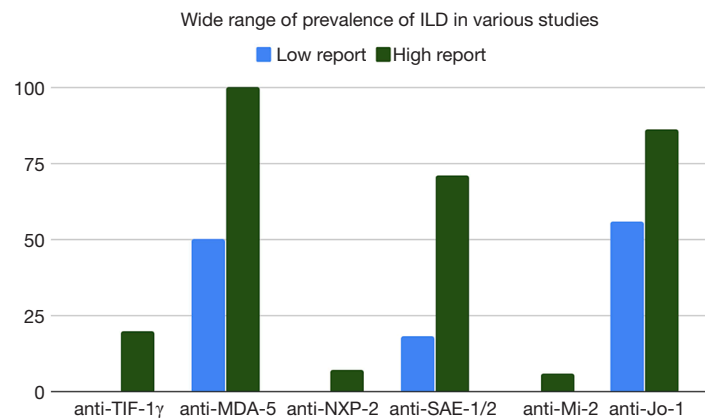


Figure 3 Wide range of prevalence of ILD in various studies. Anti-TIF-1 γ : low report: 0% (Spain, IP, method of evaluating ILD unspecified) (118); high report: 20% (Japan, IP, chest radiography and high-resolution CT) (70). Anti-MDA-5: low report: 50% (USA, ELISA, pulmonary fibrosis seen on chest radiography or high-resolution CT) (74); high report: 100% (Japan, IP, chest radiography and high-resolution CT) (70). Anti-NXP-2: low report: 0% (Japan, IP, standard clinical criteria) (42); high report: 7% (USA, IP, percentage of FVC) (97). Anti-SAE-1/2: low report: 18% (UK, IP, high resolution CT scan) (100); high report: 71% (Japan, IP, method of evaluating ILD unspecified) (90). Anti-Mi-2: low report: 0% (India, LIA, high-resolution CT) (91); high report: 6% (Brazil, LIA, high-resolution CT) (127). Anti-Jo-1: low report: 56% (Japan, IP, chest radiograph and high-resolution CT) (114); high report: 86% (USA, ELISA, abnormalities on chest radiograph or high-resolution CT or biopsy) (128). ILD, interstitial lung disease; IP, immunoprecipitation; CT, computed tomography; LIA, line immunoblot assay.

of both physician and patient. For example, a patient in USA with a new diagnosis of DM and a positive test for anti-TIF-1 γ antibodies may get the impression that this is a cancer-associated antibody, although, in USA for example, 75–80% of patients in this population will still be cancer-free (6). This may lead to unnecessarily aggressive cancer screening, cost, patient concern, and anxiety over ultimately benign lesions from comprehensive imaging procedures. In addition, internal malignancy can occur in all MSA subgroups, albeit at varying frequencies, and so, understandably, with the current data a physician may be tempted to screen all patients equivalently for cancer. Indeed, because of this uncertainty, malignancy and ILD screening are still currently recommended for all patients with DM regardless of autoantibody type (7,123). Data from two large USA cohorts suggest that blind screening for malignancy consisting largely of computed tomography (CT) scans will indeed detect a significant number of cancers that would otherwise have been missed (124). Other data derived from insurance claims data in USA similarly suggest that age and sex-appropriate cancer screening likely does not detect a significant proportion of occult malignancies in young patients with DM (125). A greater understanding of the role of antibodies in the context of

other clinical factors to stratify malignancy risk would help alleviate cost considerations as well as adverse effects of unnecessary screening procedures.

Variability also exists with regards to the risk of ILD, and especially rapidly-progressive ILD, in patients with anti-MDA-5 antibodies. In an early study of 82 DM patients, 95% of anti-MDA-5 patients identified by immunoprecipitation (IP) had ILD (126). However, a study of adult patients using enzyme-linked immunosorbent assay (ELISA) at the University of Pittsburgh found that only 50% of anti-MDA-5 patients had ILD compared with 26% of DM patients who were not anti-MDA-5 positive (*Figure 3*) (74). Though in pooled studies for anti-MDA-5-positive patients, 5-year survival was only 56%, mainly due to ILD, this may be due to results from East Asia where anti-MDA-5-associated ILD is associated with a rapidly progressive course and high mortality (16,51,69-79). While patients with anti-MDA-5 antibodies may have ILD with rapidly progressive course in other populations, the incidence of this outcome appears to be lower (72,74).

Anti-SAE-positive patients appear to have a high percentage of ILD (64–71%) in Asia, but a much lower percentage (around 18%) in Europe (90,100,104). Other MSAs have very low levels of ILD: anti-Mi-2 (0% in Asia and

6% in Brazil), anti-NXP-2 (0% in Japan and 7% in USA), and anti-TIF-1 γ (0–20% in Japan and 0–5% in Europe and USA) (42,46,70,86,91,96,97,99,116,118,126,127). Although many groups have a relatively lower risk of ILD, it is still possible that their risk of ILD is much higher than healthy comparators. To provide patients with early treatment, which is presumably associated with better outcomes, it is important to accurately identify groups that are susceptible. Overly aggressive treatment in a patient who is not at risk has its costs including sometimes fatal consequences (129).

The current state of MSA testing in the clinic

Depending on the geographic area of clinical practice, MSA assays are not always readily available to the practicing clinician. Even in situations in which they are available, many physicians do not have a clear idea which laboratory is the preferred vendor for accurate testing. For example, in USA, many laboratories have their own proprietary assay for testing that can vary widely. This means that the results of a given MSA assay need to be interpreted very differently depending on the laboratory, since assay platform is critical for interpreting results. In addition, many busy clinicians are unaware of these important nuances, and may interpret a false positive or negative result with misguided concern or comfort with regards to patient risk. Also, many of these assays have turnaround times of several weeks to even months which has obvious potentially adverse clinical implications as in the situation in which a patient is suffering from rapidly progressive lung disease whose etiology is unclear.

The importance of population characteristics in interpreting autoantibody significance

There is a critical need to consider population characteristics in order to interpret significance of any test or study. Population characteristics include not only geography, environmental exposures, ethnicity, gender, and age, but also the underlying disease of the patient. Autoantibody associations may depend on the underlying disease in which the study takes place. For example, anti-Ro52 antibodies are associated with Sjogren's disease in one population, with neonatal lupus and congenital heart block in another, with certain mucocutaneous features in lupus patients, with increased risk of ILD in systemic sclerosis, and with more severe muscle disease, and possibly ILD, in myositis

(130,131). For each MSA, varying numbers of patients may have only skin disease, only muscle disease (myositis), or varying states of overlap or DM-like disease—thus one might expect different associations depending on the clinical phenotype being studied. For example, originally, DM classification criteria required muscle involvement, and amyopathic DM was not included until recently (132–134). Recent criteria are still not very sensitive to detect amyopathic DM which may represent approximately 20% of DM cases with similar morbidities and mortalities as myopathic DM including cancer rates (7,25,135–138). Thus, the classification of DM continues to be a moving target and remains a real challenge for standardizing phenotypes associated with specific autoantibodies.

In addition, antibody associations for the MSA appear, at least in some instances, to depend on underlying demographics of the patients. For example, rapidly progressive ILD in the anti-MDA-5 population appears to be more likely in Asian than non-Asian populations (16,51,69–79). Another notable population difference is seen in the association of anti-SAE antibodies and malignancy which is around 50% in Japan, but only 13–20% in USA and Europe (87,100,101,115). In Japan, the standardized incidence ratio for cancer in anti-SAE patients was 13 relative to a matched normal population and may be enriched for cancers of the GI tract (139). The sensitivity and specificity of anti-TIF-1 γ for cancer in adult patients with DM is approximately 70% and 89% respectively (140). In addition, younger adults and children with DM with anti-TIF-1 γ antibodies do not appear to have a significantly increased risk of malignancy, pointing to the importance of age in assessing antibody phenotype (6,47,141,142). Thus, antibody-associated risk for cancer, and its specific types, might be influenced by the demographics of the patient.

The critical importance of autoantibody assay platform in interpreting autoantibody significance

Though differences in MSA prevalence may be partially due to genetic and environmental factors, autoantibody prevalence also differs significantly between studies conducted in similar populations, raising questions about the consistency of various MSA detection methods (65,143).

It is well known that there are many methodologies currently employed to detect autoantibodies. The gold standard assay is IP using native protein as the antigen, as it is clear that the immune system overwhelmingly

Table 2 Autoantibody detection assays and their characteristics

Assay	Benefits	Limitations
IP	Gold standard—mimics native antigen structure	Labor intensive, less reproducible, lack of precise quantitation
ELISA, bead-based assays	Reproducible, quantitative, affordable, scalable	Need validation against IP for each specificity
LIA and immunodot	Reproducible, semi-quantitative, scalable, widely available as commercial kits	Antigen is denatured (potential for false positive/negative), requires validation against IP for each specificity. LIA poorly sensitive for detecting anti-TIF1- γ antibodies

IP, immunoprecipitation; ELISA, enzyme-linked immunosorbent assay; LIA, line immunoblot assay.

sees antigen in its native conformation, but this is highly labor intensive and may be influenced by protein-protein interactions and post-translational modifications of the target antigen (*Table 2*) (144). It should be noted that this doesn't mean IP is always "correct", just that studies should always be comparing data from their platform with that from IP, and preferentially with some phenotypic characterization of the groups to delineate how different platforms might be detecting different subsets of patients. ELISA is more affordable with advantages that include standardization, large-scale reproducibility, and quantitative results, but does not necessarily always provide the same set of antigenic epitopes as do IP assays (36). Similarly, bead-based assays have been employed to detect certain MSAs (145). Finally, assays to detect non-native denatured forms of the antigens are in wide use, in the form of line-blot or dot-blot assays (146). In clinical laboratories, it is common to use a combination of these methodologies—a popular example would be screening with the use of indirect immunofluorescence followed by verification with automated monospecific immunoassays or multi-specific immunoassays, often commercial line/dot immunoassays (5,27,147).

Early data using the MSAs suggested that assay platform is critical in interpreting results. For example, though multiple studies have shown that anti-Mi-2 antibodies are specific to DM, a large group of patients with polymyositis were detected by ELISA using fragments of the full-length Mi-2 protein, raising doubt as to whether these patients were the same population that would be considered classic anti-Mi-2 patients (37,81).

In light of this, several recent studies have cross-validated some of these platforms by testing the same sera using multiple assays (148-150). This discussion will be limited to studies that compared results to IP as the latter is considered

an important reference. One early study compared commercial line immunoblot assay (LIA) with IP for 208 patients, finding 100% specificity and comparable sensitivity for anti-Jo-1 (92). Several recent studies compared the Euroline line immunoassay with IP (145,151,152). Most of these studies find good test performance for anti-Jo-1 antibodies (145). These studies highlight a major issue that arises in these validation studies—that is, they include a large majority of seronegative patients, which can give falsely high concordance rates driven by the negative samples. Thus, one study found "good" agreement between line blot and IP for detecting antibodies against TIF-1 γ and MDA-5, despite the fact that only 4 patients tested positive for each antibody (151). Later studies have confirmed that the Euroline blot misses a significant number of patients that are positive for anti-TIF-1 γ antibodies by IP (149,153). For example, our recent data revealed that the Euroline line blot was shown to miss 8/26 anti-TIF-1 γ positives by IP (149). Early studies also suggested that, the Euroline line blot assay, which detects anti-Mi-2 α and anti-Mi-2 β separately, is fraught with false positives. It has more recently been confirmed that, using the Euroline, anti-Mi-2 α is sensitive and specific, but anti-Mi-2 β suffers from lower sensitivity and much lower specificity (154,155). It was suggested that more stringent cutoff values would improve the performance of the line assay, but this was not helpful for the performance for anti-TIF-1 γ , and obviously would not ameliorate instances where sensitivity is already a problem (156,157). For example, for anti-NXP2 and anti-SAE-1, there are significant numbers of "false" positives on the Euroline (e.g., not positive on IP), while the line blot appears to have an issue with lower sensitivity in detecting anti-NXP2 antibodies (much as the case for anti-TIF-1 γ) (149,151-153).

Other platforms have been less frequently validated

against IP for the DM-specific MSA. In a study of 157 IIM patients from the United Kingdom, a fully automated particle-based multi-analyte technology (PMAT) test was compared with IP and demonstrated high agreement between the methods for all of the DM-specific antibodies, though this will need to be validated in other studies (153).

ELISA assays are being commonly employed for some specificities in detecting MSA in DM. Three ELISA assays have been developed in Japan for detecting antibodies against MDA-5, TIF-1 γ , and Mi-2 β and were shown to be highly specific and sensitive in cohorts of Japanese patients (158,159). Using the ELISA to detect anti-TIF-1 γ antibodies, a proportion of patients had low titers and it was suggested that these represent false positive patients that are actually anti-Mi-2-positive but have cross-reacting antibodies that can detect TIF-1 γ (159). Our studies in USA patients also suggest that the anti-MDA-5 ELISAs are highly sensitive and specific, but that the ELISA detects another 25% positive for antibodies against Mi-2 and TIF-1 γ , whose significance is unclear, but these do not appear to be due to cross-reactivity (149).

One question arises as to if the IP assay is always the “correct” gold standard. In a recent study, LIA, IP, and ELISA detected anti-TIF-1 γ antibodies in increasingly larger nested sets of patients (149). The question then arises which is the “correct” assay to be considering. If one considers association with internal malignancy as a critical phenotype that “validates” anti-TIF-1 γ positivity, this study found decreasing prevalence of malignancy using the LIA, IP, and ELISA assays, respectively (149). However, the “lowest” rate of malignancy found in the “ELISA-positive only” group still was greater than that found in patients without anti-TIF-1 γ antibodies, leaving open the question if this is still a unique subset of patients that is not equivalent to the anti-TIF-1 γ -negative group. Further studies in large numbers of patients with careful phenotypic characterization will be required to answer these questions.

As implied above, preferred assays for MSA detection may vary by antibody type. For example, ELISA assays for anti-MDA-5 may be superior to IP, given their comparable sensitivity and specificity, rapid turnaround, high-throughput, and availability for quantitation (71,149). Line blot assays appear sufficient for detecting anti-Jo-1 antibodies, and, if one uses anti-Mi-2 α as the readout, also for Mi-2 antibodies (154). The data on using the line blot for detecting anti-SAE-1 antibodies are conflicting, with some studies suggesting that there are a significant number of false positives by line blot (145,149). Current

data suggest that line blot assay for anti-NXP-2 antibodies is neither adequately sensitive nor specific when compared to IP (145,149).

While there is a need to standardize the more modern scalable immunoassays for DM autoantibodies to the classic IP assay, there might also be a need to standardize the classic IP assay as well. Some workers have used K562 cell extract as autoantigen source in their IP assays while other workers have used Hela cells (16,42,87,118). It is conceivable that DM autoantigen configurations could vary between different cultured cell lines used as cell extracts in IP assays. It is also conceivable that Mycoplasma contamination of cultured cell line extracts could alter DM autoantigen configurations. Thus, it would seem reasonable that any cooperative effort to standardize modern DM autoantibody assays should also include efforts to standardize cell culture extracts in IP assays.

Increasing the utility of the MSAs: define the patient population

For the reasons mentioned previously, when reporting studies and results, the patient population needs to be well-defined. The recent ACR/EULAR classification criteria have gone a long way to help define myositis and its subgroups (25). However, it is still clear that these criteria do not capture up to a third of DM patients with skin-predominant disease (135). Further work is needed to define the population of DM that presents with mostly cutaneous manifestations, as there is currently lack of validation and diagnostic utility of the diverse cutaneous manifestations of DM (*Table 1*) (43,72,76,86,130,136,160-162). A project is ongoing to validate skin criteria for DM classification, including morphology, distribution, symptoms, pathology, and contextual factors (135). This may assist to verify that patients have DM and not another related condition before being included in studies.

In many instances broad statements about autoantibody associations need to be nuanced with regards to the context of the particular population of included patients. More homogenous populations in terms of demographics need to be used in studies, and, although there is often power in large numbers, there may be great utility in resisting the urge to pool data from multiple diverse populations.

More detailed definition of outcomes is needed to highlight the nuances of autoantibody associations. For example, ovarian cancer seems more likely in anti-TIF-1 γ patients and GI cancers seem more likely in anti-SAE

patients holding potential to direct screening procedures to correct populations (139,163).

Increasing the utility of the MSAs: optimize and standardize testing

In addition to stratifying patient populations, more data are needed with respect to implications of antibody titer and isotype. Some data suggest that anti-MDA-5 and anti-TIF-1 γ titers tend to go down with decreased activity and remission and go up with potential flares (47,75,164-166). The IgG2 isotype of anti-TIF-1 γ may be predictive of malignancy (167).

There is a need to continue to validate various assay platforms for each autoantibody against the “reference standard” of IP, and this needs to be done in multiple populations. IP has shortcomings as it is slow, not quantitative, difficult to reproduce, and not scalable, but other platforms must be validated before they can be adopted. Where differences exist between a given platform and IP, other data should be used to characterize specifically populations that are discordant in order to better understand what these tests might be detecting. In addition, many of the published validation studies do not have high numbers of positives for many of the DM-specific MSAs, and so apparently acceptable agreement between the assays is driven largely by all of the true negatives, with significant numbers of false positives and false negatives remaining (145,149-151). In order to test and validate assays, large populations of patients likely positive for the antibody in question are needed in addition to negative controls (153). Carefully defined cutoffs for assigning positive antibody status are also needed for each assay (155,168).

To be useful in helping understand differences between population, standardizing testing is a priority that will require assembling international groups of experts in order to look at all data to decide which platforms should be considered acceptable both clinically and for translational research (31,149). There is currently an effort lead by the International Myositis Assessment and Clinical Studies (IMACS) group that has developed a Myositis Autoantibody Scientific Interest Group, established in 2018. One of the goals of this group is to define current practice and knowledge gaps with the ultimate goal of identifying valid testing platforms and harmonizing international testing practices (31). At present, most of the choices around selection of assay platform, at least in USA, is driven by

economics and not data. Ultimately, these newer kinds of efforts should lead to endorsement of particular MSA assay platforms and best practices by national and international clinical societies that will drive changes in standard of practice and put the field in order. Despite concerns regarding reliability of results, commercial immunoassays are already being used globally to inform clinical decision making and there is no putting the genie back in the bottle (31). It is therefore critical to have reliable information about these assays and well-defined standards.

Closing

It is a challenging fact of life that DM and related disorders have overlapping and variably penetrant phenotypes. Like any test, the diagnostic sensitivity and specificity of assays for various MSAs will not be perfect. However, this does not mean that we should not continue to optimize methods of using the tests as effectively as possible. Currently tests for MSAs are helpful at minimum for a clinician to assess relative risk or contribute to diagnosis and perhaps counsel the appropriate patient about what to expect (e.g., NXP-2 patient to watch for calcinosis) (169,170). With international standardization and larger studies, it is likely that more antibody tests will make their way into formal schemata for diagnosis and actionable risk assessment in DM.

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