- 1 Low levels of IgG recognizing the alpha-1-antitrypsin (A1AT) <sup>50-63</sup>
- 2 peptide and its association with Taiwanese women with primary Sjögren's
- 3 syndrome

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- 40 Running Title: Low level of IgG anti-A1AT<sup>50-63</sup> in patients with pSS
- 42 Abbreviations:
- pSS, primary Sjögren's syndrome; RA, rheumatoid arthritis; SLE, systemic lupus
- erythematosus; HNE, 4-Hydroxy-2-nonenal; 1-D SDS-PAGE, one-dimensional
- sodium dodecyl sulfate polyacrylamide gel electrophoresis; nano-LC-MS/MS, nano-
- liquid chromatography-tandem mass spectrometry; HC, healthy control; Ig,
- immunoglobulin; ACR, American College of Rheumatology; IP, immunoprecipitation;
- 48 A1AG1, alpha-1-acid glycoprotein 1; A1AT, alpha-1-antitrypsin; RSD, relative
- 49 standard deviation; SD, standard deviation; ROC, receiver operating characteristic;
- 50 AUC, area under the ROC curve; CBB, Coomassie brilliant blue; RF, rheumatoid
- factor; ANA, antinuclear antibody; anti-Ro (SSA), anti-Sjögren's-syndrome-related
- antigen A; anti-La (SSB), anti-Sjögren's-syndrome-related antigen B; CRP, C-

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- Reactive protein; ESR, erythrocyte sedimentation rate; OR, odds ratio; ESSDAI,
- 54 Sjögren's syndrome disease activity index; NSAIDs, nonsteroidal anti-inflammatory
- 55 drugs; DMARDs, Disease-modifying anti-rheumatic drugs.
- 57 Key words:
- primary Sjögren's syndrome; alpha-1-antitrypsin; inhibitor; 4-hydroxy-2-nonenal;
- 59 autoantibody isotypes; serum
- 60

## Abstract

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The aim of this study was to examine oxidative stress and low level of alpha-62 1-antitrypsin (A1AT) in primary Sjögren's syndrome (pSS), and evaluate the 63 64 associated autoreactivity against unmodified and their 4-hydroxy-2-nonenal (HNE)modified peptides with pSS. Two differentially expressed proteins, alpha-1-acid 65 glycoprotein 1 (A1AG1) and A1AT, exhibited 2-fold differences, and their HNE 66 modifications were identified by depleted-albumin and immunoglobulin G (IgG) 67 68 serum protein, in-solution digestion, in-gel digestion, and nano-LC-MS/MS from pSS 69 patients and age-matched healthy controls (HCs). Furthermore, levels of proteins, confirmation of HNE modifications, HNE-protein adducts and autoreactivity against 70 unmodified and their HNE-modified peptides were further validated. Levels of the 71 72 HNE-protein adduct and A1AG1 were significantly higher in pSS patients than HCs, but levels of A1AT were significantly lower in pSS patients compared to HCs. Only 73 the HNE modification of A1AT was confirmed. Further, concentrations of anti-74 A1AT<sup>50-63</sup> IgG and anti-A1AT<sup>50-63</sup> HNE IgA were significantly lower in pSS patients 75 than HCs. Our study suggests that elevated HNE-protein adduct, oxidative stress, 76 level [odds ratio (OR) 4.877, p = 0.003], lowered A1AT level (OR 3.910, p = 0.010) 77 and a decreased level of anti-A1AT<sup>50-63</sup> IgG (OR 3.360, p = 0.010) showed an 78 79 increased risk in pSS patients compared to HCs, respectively.

## 1. Introduction

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Primary Sjögren's syndrome (pSS) is a chronic inflammatory autoimmune disease characterized by dysfunction of the exocrine glands leading to dryness of the mouth and eyes [1]. Patients with pSS feature the presence of autoantibodies mainly against the ribonucleoprotein complex SS-related antigen A (SSA, Ro) and SS-related antigen B (SSB, La) [1]. In 2000~2008, the prevalence of pSS was 16.0 (females, 28.8, males, 3.7; female: male ratio, 7.9) per 100,000 persons; the incidence rate of pSS was 10.6 (females, 18.5, males, 2.9; female: male ratio 6.3) per 100,000 personyears; and the mortality from pSS was 1304.7 (females 987.4, males 3444.2; ageadjusted female: male ratio 0.4) per 100,000 person-years in Taiwan [2]. The etiology and pathogenesis of Sjögren's syndrome are not clearly understood [3]. Jonsson and Brun proposed etiopathogenic events prior to a diagnosis of SS including a genetic predisposition, environmental triggers, autoantibodies, pathological injury, clinical disease, and clinical presentation [4]. Norheim *et al.* reported that patients with pSS have high levels of oxidative stress compared to healthy controls (HCs) [5]. Wakamatsu et al. found an increase in 4-hydroxy-2-nonenal (HNE)-protein adducts, marker of oxidative stress, in the conjunctiva of SS patients that may play a role in the pathogenesis of dry-eye disease [6,7]. HNE is one of the lipid peroxidation products that has an alkene bond and an aldehyde group which react with amino acid residues that form HNE-protein adducts via types of Michael addition and Schiff-base adducts, respectively [8]. Amino acid residues that can react with HNE include cysteine (C), histidine (H), lysine (K), arginine (R), glutamine (Q), alanine (A), and leucine (L) [8-10]. The HNE-protein adduct is an autoantigen and can elicit specific autoantibody formation [11,12]. Breit et al. indicated that several immune-mediated diseases were associated with an alpha-1-antitrypsin (A1AT) deficiency including rheumatoid arthritis (RA),

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anterior uveitis, systemic lupus erythematosus (SLE), and asthma in which A1AT 107 may play roles as an anti-inflammatory and immune regulator [13]. A1AT is a serine 108 protease inhibitor [14]. Further, two cases were reported with an A1AT deficiency in 109 patients with pSS in which the A1AT level of plasma declined by 1.28~2.10-fold 110 [15,16]. 111 In the present study, our aim was to investigate whether a low level of serum 112 113 A1AT occurs in Taiwanese women with pSS and then identify the HNE modification on A1AT using depleted-albumin and immunoglobulin G (IgG) serum, in-solution 114 digestion, one-dimensional sodium dodecylsulfate polyacrylamide gel electrophoresis 115 (1-D SDS-PAGE), in-gel digestion, and label-free nano-liquid chromatography 116 tandem mass spectrometry (nano-LC-MS/MS) from pSS patients vs. HCs. Further, we 117 also assessed associations of autoantibody isotypes against A1AT<sup>50-63</sup> and their HNE-118 modified peptides with pSS patients compared to HCs. 119 120

## 2. Results

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2.1 Identification and validation of differentially expressed serum proteins by in-122 solution digestion and LC-MS/MS 123 Enrichment of depleted-albumin and IgG serum protein samples from a single 124 pair of each of nine pooled serum samples (patients with pSS vs. HCs) was analyzed 125 in triplicate by in-solution digestion coupled to nano-LC-MS/MS (Table 1). In total, 126 127 255 proteins were detected, of which 28 differentially expressed proteins significantly varied, as shown in Table 1 and Supplementary Table 2. There were seven 128 upregulated proteins and 21 downregulated proteins; relative to the HC serum pools, 129 two of the identified proteins, alpha-1-acid glycoprotein 1 (A1AG1) and A1AT, 130 differed by a 2-fold increase or decrease in patients with pSS serum pools, and 26 131 proteins differed by a 1.7~1.9-fold increase or decrease (Table 1). 132 Next, we validated the LC-MS/MS data of A1AG1 and A1AT, and protein 133 levels of A1AG1 (~48 kDa) and A1AT (~55 kDa) were analyzed by Western blotting 134 135 (Figure 1). A1AG1 expression levels were significantly higher in pSS samples by 1.53-fold (p = 0.0001) than in HCs, but A1AT levels in pSS samples were 136 significantly lower than those in HCs by 1.84-fold (p = 0.0071, Figure 1A, B, right 137 upper panel). Equal amounts of serum proteins in these experiments were examined 138 (Figure 1A, B, right bottom panel). The AUC value, sensitivity, and specificity of 139 serum A1AG1 and A1AT in pSS samples vs. HCs were calculated based on these 140 141 results and plotted on an ROC curve. The Western blot results of A1AG1 showed that the AUC was 0.75, the sensitivity was 85.0%, and the specificity was 62.5% for pSS 142 detection at an average densitometric cutoff of 19,994.82; the AUC was 0.67, the 143 sensitivity was 77.5%, and the specificity was 60.0% for pSS detection by A1AT at 144 an average densitometric cutoff of 104,087.25 (Figure 1C). 145

2.2 Novel HNE modification identification of serum proteins by in-gel digestion and 147 LC-MS/MS 148 In addition to serum protein levels, we further identified HNE modifications of 149 A1AG1 and A1AT. The average coverage of amino acid sequences in A1AG1 and 150 A1AT were estimated to be 52% and 70%, respectively. No HNE modification was 151 identified on A1AG1 (data not shown). Novel HNE modifications of A1AT were 152 153 identified by manual examination of the modified spectra using the PeaksPTM module in PEAKS 7 software. Further, HNE modifications of A1AT were confirmed 154 in the two pooled serum samples (patients with pSS vs. HCs) through IP-Western 155 blotting, which detected signals of approximately 55 kDa (Figure 2). Because low 156 coverage of A1AG1 was identified, we also confirmed HNE modifications of A1AG1 157 using IP-Western blotting, but no signal was detected (data not shown). 158 MS/MS spectrum data of HNE-modified peptides on A1AT are presented in 159 Supplementary Figure 1B and Supplementary Table 3. The peptide <sup>50</sup>-160 ITPNLAEFAFSLYR-<sup>63</sup> was used to identify A1AT as pSS-specific and was found to 161 have an HNE modification at A58. Identification of the peptide moieties was based on 162 the presence of b- and y-series ions derived from the peptide, and HNE-modified 163 residues were confirmed by an unmodified b8 ion followed by a modified y6 ion that 164 corresponded to a mass increase of 156.11504 Da (Supplementary Figure 1C, upper 165 panel). The peptide <sup>360</sup>-AVLTIDEK-<sup>367</sup> was used to identify A1AT as HC-specific 166 and was found to have an HNE modification at A360. Identification of the peptide 167 moieties was based on the presence of b- and y-series ions derived from the peptide, 168 and HNE-modified residues were confirmed by an unmodified y7 ion followed by a 169 modified b1 ion that corresponded to a mass increase of 138.10446 Da 170 (Supplementary Figure 1C, bottom panel). 171

2.3 Autoreactivity against A1AT<sup>50-63</sup> and A1AT<sup>50-63</sup> HNE peptides

Serum samples were assessed with autoantibody isotypes against A1AT<sup>50-63</sup> 174 and A1AT<sup>50-63</sup> HNE peptides by an ELISA. The level of the anti-A1AT<sup>50-63</sup> IgG 175 antibody in RA was significantly higher than that of HCs by 1.80-fold (p = 0.0002), 176 that of SLE vs. HC was 2.41-fold higher (p < 0.0001), that of RA vs. pSS was 2.52-177 fold higher (p < 0.0001), that of SLE vs. pSS was 3.38-fold higher (p < 0.0001), and 178 that of SLE vs. RA was 1.34-fold higher (p = 0.0321); however, that of pSS was 179 significantly lower than that of the HCs by 1.40-fold (p = 0.0488, Figure 3A, left 180 panel). Anti-A1AT<sup>50-63</sup> IgM expression levels did not significantly differ among 181 patients with pSS, RA, SLE, and HCs, except that of RA was significantly higher than 182 that of pSS by 1.57-fold (p = 0.0143, Figure 3A, middle panel). Levels of the anti-183 A1AT<sup>50-63</sup> IgA antibody did not significantly differ among patients with pSS, RA, 184 SLE, and HCs (Figure 3A, right panel). 185 The level of the anti-A1AT<sup>50-63</sup> HNE IgG antibody in RA was significantly 186 higher than that of HCs by 2.10-fold (p < 0.0001), that of SLE vs. HC was 2.70-fold 187 higher (p < 0.0001), that of RA vs. pSS was 2.69-fold higher (p < 0.0001), that of SLE 188 vs. pSS was 3.48-fold higher (p < 0.0001), and that of SLE vs. RA was 1.29-fold 189 higher (p = 0.0469); however, that of pSS did not significantly differ from that of the 190 HCs (Figure 3B, left panel). Anti-A1AT<sup>50-63</sup> HNE IgM expression levels did not 191 significantly differ among patients with pSS, RA, SLE, and HCs, except that of SLE 192 was significantly higher than that of HCs by 1.36-fold (p = 0.0433) and that of SLE vs. 193 pSS was 1.47-fold higher (p = 0.0088) (Figure 3B, middle panel). Levels of the anti-194 A1AT<sup>50-63</sup> HNE IgA antibody did not significantly differ among patients with pSS, 195 RA, SLE, and HCs, except that of pSS was significantly lower than that of the HCs by 196 1.15-fold (p = 0.0484) and that of RA vs. pSS was 1.17-fold higher (p = 0.0346) 197 (Figure 3B, right panel). 198

2.4 Determination of HNE-protein adducts

of autoantibodies (Table 2).

The level of the HNE-protein adduct can present the oxidative stress status and plays important pathogenic roles in several diseases including cancer, and neurodegenerative, chronic inflammatory, and autoimmune diseases [28]. As shown in Supplementary Table 1, serum levels of the HNE-protein adduct in pSS patients were significantly higher compared to those of the HCs (1.27-fold, p=0.0004).

2.5 Association of elevated HNE-protein adduct, lowed A1AT level or decreased autoreactivity against A1AT<sup>50-63</sup> and A1AT<sup>50-63</sup> HNE peptides with pSS patients

In Table 2, HNE-protein adduct, serum A1AT and anti-A1AT<sup>50-63</sup> IgG of pSS patients carried a 4.887-fold risk (p=0.003, power = 0.708), 3.910-fold risk (p=0.010, power = 0.802) showed a significant difference compared to HCs after adjusting for age, in the logistic regression analyses, respectively. Risks did not significantly differ after the age-adjusted logistic regression, suggesting that they were associated with other low levels

## 3. Discussion

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This is the first study to investigate the association between decreased serum levels of autoantibody isotypes against A1AT<sup>50-63</sup> and A1AT<sup>50-63</sup> HNE peptides and the risk of low A1AT levels in pSS patients. In the present study, two differentially expressed proteins, A1AG1 and A1AT, had 2-fold differences in depleted-albumin and IgG serum protein pools of nine pSS patients vs. nine HCs, identified in triplicate from in-solution digestion coupled to LC-MS/MS (Table 1). However, A1AG1 (1.53fold increase, AUC = 0.75) and A1AT (1.84-fold decline, AUC = 0.67) showed acceptable diagnostic values for discriminating between pSS patients and HCs according to a Western blot analysis (Figure 1C). In this study, significantly higher serum levels of HNE-protein adducts indicated increments in the oxidative stress status of pSS patients (Supplementary Table 1); these results are consistent with those of previous studies [5, 6]. A1AG is an acute-phase protein, and its serum levels are elevated in response to a local inflammatory stimulus in several diseases including depression, cancer, and acquired autoimmune deficiency syndrome [25]. A1AG may have anti-inflammatory and immunomodulatory properties [26]. In the macrophage deactivation process, A1AG1 may act as a signaling molecule in the maintenance of tissue homeostasis and remodeling [27]. Rantapaa-Dahlqvist et al. reported that pSS patients with pericarditis had significantly higher levels of A1AG than did pSS patients without pericarditis; however, no information on A1AG1's involvement in the development of pSS has been reported. In this study, serum protein levels of A1AG1 in pSS were significantly higher than those of HCs by 1.53-fold (Figure 1A). Saroha et al. indicated that altered glycosylation and expression of plasma A1AG may play a role in RA progression [28].

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A1AT is also an acute-phase protein that has anti-inflammatory and tissueprotective properties and is an immune regulator [12, 29]. Human A1AT protein levels can increase to inhibit elastase and serine-type proteinase during inflammation [13]. Serum protein levels of A1AT in patients with pSS were significantly lower than those of HCs by 1.84-fold (Figure 1B); these results are consistent with previous studies [14, 15]. In this study, patients with pSS showed a feature of low A1AT level (Figure 1B). Thus, low serum level of A1AT is a risk factor for the development of pSS (Table 2). Further, serum levels of anti-A1AT<sup>50-63</sup> IgG and anti-A1AT<sup>50-63</sup> HNE IgA were significantly lower in pSS patients (Figure 3). However, Elshikha et al. demonstrated that the human A1AT protein has protective effects through inhibition of dendritic cell (DC) activation and function to attenuate autoimmunity in RA mouse models [30]. Ciobanu et al. indicated that significantly lower levels of A1AT in rheumatoid synovial fluid can decrease the anti-protease activity in RA [31]. Stefanescu et al. showed that levels of anti-A1AT antibodies were significantly elevated in RA [32]. In several previous studies, elevated IgA-A1AT complex levels were reported in RA, SLE, mixed connective tissue disease, and ankylosing spondylitis compared to HCs [33-35]. Further, Lacki et al. suggested that a high level of the IgA-A1AT complex may cause worsening of bone erosion in RA [35]. In this study, levels of IgA-A1AT<sup>50-63</sup> and their HNE-modified peptide complexes did not significantly differ among patients with RA and SLE compared to HCs (Figure 3A, B, right panel). Levels of IgG-A1AT<sup>50-63</sup> and their HNE-modified peptide complexes were significantly higher among patients with RA and SLE compared to HCs (Figure 3A, B, left panel). Further, we observed that low levels of the anti-A1AT<sup>50-63</sup> IgG antibody obviously increased the risk against pSS, but the anti-A1AT<sup>50-63</sup> HNE IgG antibody reduced the risk against pSS (Table 2). The presence of self-reactive IgG autoantibodies in human serum is thought to represent as pathogenic antibodies in

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patients with pSS [29]. Further, the HNE-modified epitope belong to oxidation-specific epitopes (OSEs) [30]. OSEs are present on damaged proteins and induce specific autoantibodies formation [11,12]. Anti-OSEs autoantibodies have conveyed protection from autoimmune pathogenesis [29,31,32]. Importantly, oxidative stress remained in patients with pSS (Supplementary Table 1).

#### 4. Materials and methods

#### 4.1 Patient samples

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This study was approved by the institutional review board of the study hospital, and all volunteers provided informed consent before being allowed to participate. Serum samples from 168 female patients [49 with pSS (55.50  $\pm$  12.85 years old), 40 with RA (54.30  $\pm$  11.30 years old), and 30 with SLE (40.60  $\pm$ 11.18 years old)] and 49 age-matched female HCs (55.40  $\pm$  11.67 years old) were obtained from the Division of Allergy, Immunology and Rheumatology, Department of Internal Medicine and Department of Laboratory Medicine, Shuang-Ho Hospital (New Taipei City, Taiwan). Patients with pSS, RA, or SLE were diagnosed by a rheumatologist and had satisfied appropriate classification criteria. RA patients had received a diagnosis from a rheumatologist and had fulfilled appropriate classification criteria—either the 2010 American College of Rheumatology (ACR)/European League Against Rheumatism classification criteria [33] or the 1987 ACR classification criteria [34]. pSS patients were diagnosed according to the American-European Consensus Group (AECG) classification criteria [35]. SLE patients fulfilled the 1997 ACR SLE classification criteria [36]. Differentially expressed serum proteins were identified through in-solution digestion and nano-LC-MS/MS using pooled depleted-albumin and IgG serum protein samples randomly selected from nine RA patients and nine age-matched HCs. Two differentially expressed proteins, A1AG1 and A1AT, exhibited 2-fold differences in patients with pSS compared to HCs, and these were selected to examine protein levels through Western blotting using individual serum samples randomly selected from another 40 pSS patients and 40 age-matched HCs. HNE modifications of A1AT and A1AG1 were identified by ingel digestion and nano-LC-MS/MS. HNE modifications of proteins were evaluated through immunoprecipitation (IP) and Western blotting using the aforementioned 40

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pairs of pooled serum samples. Next, autoantibody isotypes against unmodified and their HNE-modified peptides were assessed among 49 pSS, 40 RA, and 30 SLE patients, and 49 HCs. Serum was stored at -20 °C until being analyzed. Clinical and demographic characteristics of pSS, RA, and SLE patients, and HCs are presented in Supplementary Table 1. However, the age of patients with SLE was significantly lower compared to those of the pSS, RA, and HC cohorts (Supplementary Table 1). 4.2 Depleted-albumin and IgG serum proteins, in-solution digestion, and protein identification by LC-MS/MS Protein concentrations of serum were determined using a Coomassie Plus (Bradford) Assay Kit according to the manufacturer's protocol. Albumin and IgG of serum samples were removed using an Albumin and IgG Depletion SpinTrap column according to the protocol of Uen et al. [37]. Three micrograms of depleted-albumin and IgG serum proteins was used to perform in-solution digestion using an In-Solution Tryptic Digestion and Guanidination Kit according to the manufacturer's instructions. Tryptic peptide mixtures were analyzed in triplicate using NanoLCnanoESi-MS/MS that was performed on a nanoAcquity system (Waters, Milford, MA, USA) connected to an LTQ-Orbitrap XL<sup>TM</sup> hybrid mass spectrometer (Thermo Fisher Scientific, Bremen, Germany) equipped with a nanospray interface (Proxeon, Odense, Denmark). Differentially expressed proteins were quantified using label-free peptide quantification by the Peaks Q module of the PEAKS 7 software (Bioinformatics Solutions, Waterloo, Canada) [38]. Details are provided in "Supplementary information".

325 4.3 Western blotting

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Serum protein levels of differentially expressed proteins showing 2-fold differences in pSS patients vs. HCs were examined using a Western blot analysis. A1AG1 (2 µg of protein in 10% SDS-PAGE) or A1AT (2 µg of protein in 8% SDS-PAGE) was evaluated using a mouse anti-A1AG1 monoclonal antibody (sc-69753, Santa Cruz Biotechnology, Dallas, TX, USA) or a mouse anti-A1AT monoclonal antibody (sc-69752, Santa Cruz Biotechnology). Details are provided in "Supplementary information". 4.4 1-D SDS-PAGE, in-gel digestion, and HNE identification by LC-MS/MS Fifty-microgram protein samples (pooled serum proteins of A1AG1 or A1AT) were run on 10% SDS-PAGE with in-gel digestion according to a previously described method with minor modifications (Supplementary Figure 1A) [39]. HNE modifications were identified in triplicate using tryptic peptide mixtures of gel slices by the aforementioned nano-LC-MS/MS (nanoAcquity system and LTQ-Orbitrap XL<sup>TM</sup> hybrid mass spectrometer). HNE-modified peptide sequences and sites of serum A1AG1 and A1AT were identified using the PeaksPTM module of the PEAKS 7 software (Bioinformatics Solutions). Details are provided in "Supplementary information". 4.5 Immunoprecipitation (IP) An IP experiment for A1AG1 or A1AT was performed using a mouse anti-A1AG1 monoclonal antibody (sc-69753, Santa Cruz Biotechnology) or a mouse monoclonal antibody (sc-69752, Santa Cruz Biotechnology). HNE modifications of A1AG1 or A1AT were evaluated through a Western blot analysis with a goat polyclonal anti-HNE antibody (MBS536107, MyBioSource, San Diego, CA, USA). Details are provided in "Supplementary information".

352 4.6 Detection of autoreactivity against A1AT<sup>50-63</sup> and their HNE-modified peptides 353 Polypeptides corresponding to the 50~63 amino acid sequence of human 354 A1AT, i.e., ITPNLAEFAFSLYR (named A1AT<sup>50-63</sup>) were synthesized (Yao-Hong 355 Biotechnology, New Taipei City, Taiwan) and their HNE-modified A1AT<sup>50-63</sup> (named 356 A1AT<sup>50-63</sup> HNE) used in an enzyme-linked immunosorbent assay (ELISA). In total, 357 168 serum samples were assessed for the presence of IgG, IgM, and IgA isotypes of 358 anti-A1AT<sup>50-63</sup> and anti-A1AT<sup>50-63</sup> HNE peptide antibodies. The absorbance was 359 measured at 450 nm with the reference filter set to 620 nm. All samples were treated 360 in duplicate. Details are provided in "Supplementary information". 361 362 4.7 Detection of serum HNE-protein adducts 363 Levels of HNE-protein adducts were quantified using 168 serum samples for 364 the ELISA protocol of Weber et al. [40]. All samples were analyzed in duplicate. 365 366 Details are provided in "Supplementary Information". 367 4.8 Statistical analyses 368 Student's *t*-test was used to determine the significance of differences in blot 369 densitometry, levels of serum proteins, and levels of HNE-protein adducts, and levels 370 of autoantibody isotypes against A1AT<sup>50-63</sup> and A1AT<sup>50-63</sup> HNE peptides. GraphPad 371 Prism (vers. 5.0; Graphpad Software, San Diego, CA, USA) was used to assess 372 differences in Student's t-tests between groups, and a dot plot was drawn. Ages and 373 clinical test results are presented as the mean  $\pm$  standard deviation (SD). Spectral 374 count data are presented as the mean  $\pm$  relative SD (RSD). The RSD is a coefficient of 375 variation (CV) and is calculated as a percentage. Multiples of change were defined as 376

(mean of pSS-normalized spectral counts) / (mean of HC-normalized spectral counts).

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The threshold for up- or downregulated proteins was a 1.0-fold change in expression. Comparisons of pSS vs. HC serum samples were performed. Proteins that had a 2-fold difference were selected for validation by a Western blot analysis. Univariate and multiple logistic regression models were further used to estimate the adjusted odds ratios (ORs) and their 95% confidence intervals (CIs) for the pSS risk. Power estimations were determined using SAS (vers. 9.3; SAS Institute, Cary, NC, USA). Receiver operating characteristic (ROC) curves were generated to evaluate the diagnostic performance of differentially expressed proteins using MedCalc Statistical Software (vers. 15.4; MedCalc Software, Ostend, Belgium). The area under the ROC curve (AUC), sensitivity, and specificity were estimated at a 95% confidence level. For all statistical tests, the significance level was set to p < 0.05.

5. Conclusions

We identified HNE modifications on the human serum A1AT protein *in vivo* to investigate autoantibody isotypes against A1AT<sup>50-63</sup> and A1AT<sup>50-63</sup> HNE peptides associated with pSS patients. Our results showed that low levels of the anti-A1AT<sup>50-63</sup> IgG antibody had an increased risk in pSS patients. However, this possibility needs to be confirmed in larger studies.

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#### **Author Contributions:**

Ching-Yu Lin, Che-Chang Chang, Yu-Sheng Chang, Kai-Leun Tsai and Sheng-Hong Lin conceived and designed the experiments; Han-Wen Chou, Chih-Chun Tai and Yi-Fang Lin performed the experiments; Jin-Hua Chen, Chih-Hong Pan, Yi-Ying Lu and Han-Wen Chou analyzed the data; Ching-Yu Lin and Che-Chang Chang wrote the paper.

# **Competing interests**

The authors declare that there are no competing interests.

**Legends of Figures and Tables** 418 Figure 1. Protein levels of A1AG1 and A1AT in serum were examined using anti-419 A1AG1 (A), and anti-A1AT (B) antibodies through Western blotting. Average blot 420 densitometric values were calculated from duplicate data. Percentages of SDS-PAGE 421 gel and loading amounts of serum proteins used in Western blotting were 10% and 2 422 423 ug for A1AG1, and 8% and 2 ug for A1AT, respectively. A duplicate gel was stained with Coomassie brilliant blue (CBB) as a loading control (right, bottom panel). The 424 red arrow indicates the A1AG1 or A1AT protein. Receiver operating characteristic 425 (ROC) curves were generated according to blot densitometry of A1AG1 and A1AT. 426 The area under the ROC curve (AUC), sensitivity, and specificity were further 427 estimated (C). 428 429 Figure 2. 4-Hydroxy-2-nonenal (HNE) modification of the serum A1AT protein was 430 431 validated using IP and Western blotting. A1AT was immunoprecipitated from pooled serum samples [40 patients with primary Sjögren's syndrome (pSS) and 40 healthy 432 controls (HCs)] using anti-A1AT antibodies and then subjected to Western blotting 433 with anti-HNE antibodies (upper panel). Individually selected random serum samples 434 (patient with pSS and HC) were used as controls; these were simultaneously used for 435 Western blotting with anti-HNE antibodies. A duplicate gel was stained with 436 437 Coomassie brilliant blue as a loading control (bottom panel). The red arrow indicates 438 the A1AT protein. 439 Figure 3. Dot plot of serum concentrations (absorbance units at 450/620 nm) of IgG, 440 IgM, and IgA autoantibody isotypes recognizing A1AT<sup>50-63</sup> (A) and A1AT<sup>50-63</sup> 4-441 hydroxy-2-nonenal (HNE) (B) in healthy controls (HCs), patients with primary 442

Sjögren's syndrome (pSS), rheumatoid arthritis (RA), and systemic lupus 443 erythematosus (SLE) with an ELISA. OD<sub>450/620</sub>, optical density at 450/620 nm. 444 445 Table 1. Differentially expressed serum proteins identified by in-solution digestion 446 and LC-MS/MS analysis in patients with primary Sjögren's syndrome (pSS) and 447 448 healthy controls (HCs). 449 Table 2. Association among HNE-protein adduct, A1AT, anti-A1AT<sup>50-63</sup> and their 450 HNE-modified peptides antibodies and pSS patients, in patients with pSS vs. healthy 451 controls. 452 453 Supplementary Figure 1. The gel was rapidly stained with Coomassie brilliant blue, 454 and gel bands were cut into slices according to the molecular weight of A1AG1 (48 455 kDa) and A1AT (55 kDa), respectively (A). Identification of 4-hydroxy-2-nonenal 456 (HNE) modifications of A1AT (B). A representative MS/MS spectrum of the peptide 457 sequence of <sup>50</sup>-ITPNLAEFAFSLYR-<sup>63</sup> and the modified peptide bearing the HNE 458 modification at alanine 58 in primary Sjögren's syndrome (pSS) (C, upper panel). The 459 MS/MS spectrum of <sup>360</sup>-AVLTIDEK-<sup>367</sup> and the modified peptide bearing the HNE 460 modification at alanine 360 in healthy controls (HCs) (C, bottom panel). 461 462 Supplementary Table 1. Demographic and clinical characteristics of individual 463 subjects contributing to serum for the healthy controls (HCs), and patients with 464 primary Sjögren's syndrome (pSS), rheumatoid arthritis (RA), and systemic lupus 465 erythematosus (SLE). 466 467

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Supplementary Table 2. A list of 255 identified proteins and peptides of depletedalbumin and IgG serum protein.

Supplementary Table 3. Post-translational modifications (PTMs), identified proteins,
and peptides of serum A1AT.

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