RHEUMATOLOGY

Original article

Anti-phosphatidylserine/prothrombin antibodies and thrombosis associate positively with HLA-DRB1*13 and negatively with HLA-DRB1*03 in SLE

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Abstract

Objectives. Emerging evidence demonstrates that aPS-PT associate with thrombotic events. Genetic predisposition, including *HLA-DRB1* alleles, is known to contribute to the occurrence of conventional aPL [anti- β_2 glycoprotein-I (anti- β_2 GPI) and aCL]. We investigated associations between aPS-PT and *HLA-DRB1** alleles and thrombosis in SLE. Conventional aPL were included for comparison.

Methods. We included 341 consecutive SLE patients, with information on general cardiovascular risk factors, including blood lipids, LA and thrombotic events. aPS/PT, anti- β_2 GPI and aCL of IgA/G/M isotypes and LA were quantified.

Results. aPS/PT antibodies associated positively with *HLA-DRB1**13 [odds ratio (OR) 2.7, P = 0.002], whereas antiβ₂GPI and aCL antibodies associated primarily with *HLA-DRB1**04 (OR 2.5, P = 0.0005). These associations remained after adjustment for age, gender and other *HLA-DRB1** alleles. *HLA-DRB1**13, but not *DRB1**04, remained as an independent risk factor for thrombosis and APS after adjustment for aPL and cardiovascular risk factors. The association between *DRB1**13 and thrombosis was mediated by aPS-PT positivity. *HLA-DRB1**03, on the other hand, associated negatively with thrombotic events as well as all aPL using both uni- and multivariate analyses. *HLA-DRB1**03 had a thrombo-protective effect in aPL-positive patients. Additionally, *HLA-DRB1**03 was associated with a favourable lipid profile regarding high-density lipoprotein and triglycerides.

Conclusions. *HLA-DRB1*13* confers risk for both aPS-PT and thrombotic events in lupus. The association between *HLA-DRB1*13* and thrombosis is largely, but not totally, mediated through aPS-PT. *HLA-DRB1*03* was negatively associated with aPL and positively with favourable lipid levels. Thus, *HLA-DRB1*03* seems to identify a subgroup of SLE patients with reduced vascular risk.

Key words: antiphosphatidylserine/prothrombin, aPL, HLA, SLE, thrombosis

Rheumatology key messages

- HLA-DRB1*13 alleles independently associate with aPS-PT and vascular events in SLE patients.
- HLA-DRB1*13 association with vascular events is, to a large extent, mediated by aPS-PT antibodies.
- *HLA-DRB1*03* associate negatively with thrombosis and aPL, and favourably with lipid profile, suggesting a thrombo-protective role.

Introduction

aPL are directed against negatively charged phospholipids and/or plasma proteins and represent the immunological hallmark of primary and secondary APS. Genetic predisposition contributes to the occurrence of aPL and to the development of APS, with the human

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leucocyte antigens (*HLA*) gene loci having a documented contributive role [1, 2].

Both family and population-based studies demonstrated that *HLA-DRB1*04* and *-DRB1*07* are more frequent in APS patients with or without SLE, as compared with healthy controls [3–9]. aCL and anti- β_2 glycoprotein I (anti- β_2 GPI) have associated with a number of *HLA-DRB1* alleles, most consistently with *HLA-DRB1*04* [2, 6, 8, 10, 11]. Among the *HLA-DRB1*04* alleles, *HLA-DRB1*0402* confers high risk for the presence of both antibodies [8]. In addition to *HLA-DRB1*04*, several studies have reported that anti- β_2 GPI antibodies also associate with *HLA-DRB1*13* alleles [9, 12, 13].

Venous and arterial vascular events associated with *HLA-DRB1*13*, and ischaemic cerebrovascular disease with *HLA-DRB1*04* also after adjustment for traditional risk factors [14]. Associations between *HLA-DRB1*13* and clinical events/aPL remained significant when excluding *HLA-DRB1*04*-positive patients, indicating an *HLA-DRB1*04*-independent risk. Other studies have also described *HLA-DRB1*07* as a risk factor for vascular events [9, 15].

Most previous immuno-genetic studies on APS only investigated aCL and anti- β_2 GPI associations, i.e. the conventional 'criteria' aPL of IgM/G isotypes. More recently, positivity for aPS-PT has emerged as a risk factor for the presence of LA and vascular events [16, 17]. In a recent publication we reported that aPS-PT antibodies associated more strongly with venous thromboembolism (VTE) and arterial thrombosis than did the conventional aPL [18]. In the current study we investigated a well-characterized Swedish SLE cohort regarding the relationship between *HLA-DRB1* alleles, aPS-PT antibodies (IgG, IgM and IgA) and vascular events. Conventional aPL were included for comparison.

Methods

Participants and clinical characteristics

Consecutive SLE patients from the Karolinska lupus cohort (n = 341) fulfilling the 1982 revised SLE criteria [19] were included. A detailed description of the cohort was reported in our previous publications [20, 21]. All patients were subjected to a thorough clinical investigation by a rheumatologist at inclusion and medical files were reviewed. APS was diagnosed according to the revised Sydney classification criteria for APS and history of any thrombotic event (VTE: including deep venous thrombosis or pulmonary embolism. Arterial thrombosis: cerebrovascular events, myocardial infarction or ischaemic peripheral tissue loss) [22]. Arterial hypertension was defined as systolic and/or diastolic blood pressure \geq 140/90 mmHg or use of antihypertensive medications. Diagnostic methods to establish clinical diagnoses were specified in our previous publication [18]. The Global APS Score (GAPSS) was calculated for all patients [23]. BMI and lipid measurements including low-density lipoprotein (LDL), high-density lipoprotein (HDL), triglycerides

(TG) and total cholesterol levels were also included in the analyses, with hyperlipidaemia defined as LDL >3 mmol/l or treatment with lipid-lowering agents.

All participants gave written informed consent and the study conformed to the guidelines of the Declaration of Helsinki. Karolinska University Hospital approved the study [03-556 (16 December 2003)].

aPL testing

Quantification of aPS-PT, aCL and anti- β_2 GPI IgA/G/M was performed using the Aptiva system based on a particle-based multi-analyte technology (PMAT; research use only, Inova Diagnostics, San Diego, CA, USA [24]). Clinical associations for all aPL antibodies among patients were previously published in comparative studies between Sudan and Sweden, using a slightly lower number of Swedish patients (n = 332) [18, 21]. The 99th percentile values of 162 age- and sex-matched population controls were used as cut-offs [22]. Information about LA positivity determined by the modified dRVVT (Bioclot LA; Biopool, Umeå, Sweden) was available for 329 patients.

Genotyping

HLA typing was performed using sequence-specific primer PCR assay (SSP-PCR) (DR 2 digit-resolution kit; Olerup SSP, Saltsjöbaden, Sweden). The studied *HLA-DRB1* alleles were *01, *03, *04, *07, *08, *09, *10, *11, *12, *13, *14, *15 and *16.

Statistical analyses

For comparisons between categorical variables, Chi²-test was used with Fisher's exact test applied when appropriate, and for quantitative differences between groups, Mann–Whitney's *U* test was used. To estimate the individual impact of independent variables we conducted multivariable logistic regression including *HLADRB1* alleles with significant associations to the specific outcomes in univariate analyses. Linear regression was also used to adjust for confounders. We used stepwise forward regression analyses when logistic regression models were unfeasible due to number of independent variables. We performed a series of regression analyses to estimate whether aPS-PT positivity acts as a mediator between the relationship of *HLA-DR*13* and thrombosis.

All statistical analyses were conducted using JMP statistical software (SAS institute, Cary, NC, USA), except for the mediation analysis where the packages tidy-verse [25], lavaan [26] and mediation [27] for R software (version 4.1.1 Kick Things) were used. *P*-values <0.05 were considered significant.

Results

Seventy-eight SLE patients (23.0%) had presented with at least one vascular event, and 31 (9.6%) were confirmed as secondary APS [22]. Among all SLE patients, 50 (15.0%) tested positive for aPS-PT antibodies of any isotype, 100 (30.2%) were positive for anti- β_2 GPI and/or aCL antibodies and LA was positive in 60 patients (18.2%). Patient demographics, history of thrombotic events, cardiovascular risk factors and antibody prevalence are shown in supplementary Table S1, available at *Rheumatology* online.

HLA-DRB1 genotype associations with aPS-PT antibodies

HLA-DRB1 allele frequencies among all patients are presented in Table 1. aPS/PT-positive patients had higher frequency of HLA-DRB1*13 allele compared with aPS-PT-negative patients [odds ratio (OR) 2.7 (95% CI 1.4, 5.2), P = 0.002]. This association remained significant after exclusion of patients positive for any anti-B2GPI and aCL isotypes (IgG/M/A): 8.7% vs 1.2% in DRB1*13 carriers vs non-carriers [OR 7.8 (95% CI 1.4, 44.3), P=0.02]. No other positive association was observed with aPS-PT. Occurrence of anti-B2GPI and/or aCL associated also with HLA-DRB*13 [OR 2.0 (95% CI 1.2, 3.4). P = 0.011 but even more strongly with HLA-DRB1*04 [OR 2.5 (95% CI 1.5, 4.1), P=0.0005]. Negative associations were found between HLA-DRB1*03 and all aPL including LA, and also between aPS-PT and anti- β_2 GI/aCL with HLA-DRB1*15. Using multiple logistic regression, where the HLA-DRB1*04, *13, *03 and *15 were independent variables and status for aPS-PT, anti- β_2 GPI/aCL and LA as the outcome, similar results were obtained except that the negative associations with HLA-DRB1*15 were lost (Fig. 1A-C). When we used age, gender, HCQ treatment and the HLA-DRB1*04, *13, *03 and *15 in a forward step-wise regression model, associations remained significant as shown in Fig. 1A-C, except for the negative association between anti-β2GPI/aCL and HLA-DRB1*03 that was lost.

Associations between *HLA-DRB1* alleles and the different aPL isotypes are shown in Fig. 2. IgA and IgM, but not IgG, aPS-PT associated with *HLA-DRB1*13*, and all aPS-PT isotypes associated negatively with *HLA-DRB1*03*, whereas no isotype showed association with *HLA-DRB1*04*. On the contrary, all isotypes of anti- β_2 GPI and aCL associated with *HLA-DRB1*04*, with the highest ORs observed for IgM. Only aCL IgG among the conventional aPL associated with *HLA-DRB1*13*.

HLA-DRB1 genotype associations with vascular events

In univariate analyses, *HLA-DRB1*13* was more frequent in patients with a history of any thrombosis, VTE or APS diagnosis, compared with patients without (Table 2). No clinical association with *HLA-DRB1*04* was observed, except for higher GAPSS. In multivariate analysis including *HLA-DRB1*03*, **04*, **13* and **15* alleles as independent factors and thrombosis as outcome, only *HLA-DRB1*13* and *HLA-DRB1*03* showed significant

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| HLA-DRB1 | HLA-DRB1, n (%) total patients | Anti-PS/PT positive (any isotype) <i>n</i> = 48, <i>n</i> (%) | OR (95% CI); P | Anti-l}₂GPI or aCL positive (any isotype) n = 96, n (%) | OR (95% Cl); <i>P</i> | LA <i>n</i> = 57, <i>n</i> (%) | UR (95% CI); H |
|-------------|-----------------------------------|--|-----------------------|---|------------------------|--------------------------------|-------------------------|
| *01 | 41 (12.9) | 4 (8.3) | 0.6 (0.2, 1.7); 0.3 | 11 (11.4) | 0.8 (0.4, 1.7); 0.6 | 6 (10.5) | 0.8 (0.3, 1.9); 0.6 |
| *03 | 147 (46.5) | 13 (27.1) | 0.4 (0.2, 0.7); 0.004 | 33 (34.4) | 0.5 (0.3, 0.8); 0.006 | 11 (19.2) | 0.2 (0.1, 0.4); <0.0001 |
| *04 | 94 (29.7) | 18 (37.5) | 1.6 (0.8, 2.9); 0.2 | 41 (42.7) | 2.5 (1.5, 4.1); 0.0005 | 27 (47.4) | 2.4 (1.4, 4.4); 0.002 |
| <i>20</i> * | 28 (8.9) | 6 (12.5) | 1.5 (0.6, 4); 0.4 | 9 (9.4) | 1 (0.4, 2.4); 0.9 | 7 (12.2) | 1.6 (0.6, 4.0); 0.3 |
| *08 | 28 (8.9) | 6 (12.5) | 1.6 (0.6, 4.3); 0.3 | 9 (9.4) | 1.2 (0.5, 2.7); 0.7 | 5 (8.8) | 0.9 (0.3, 2.6); 0.9 |
| <i>60</i> * | 9 (2.8) | 1 (2.1) | 0.7 (0.1, 5.5); 0.7 | 2 (2.1) | 0.6 (0.1, 3.0); 0.5 | 1 (1.7) | 0.7 (0.1, 5.1); 0.6 |
| 01* | 7 (2.2) | 0 0 | NA | 2 (2.1) | 0.9 (0.2, 4.6); 0.9 | 00 | NA |
| +11 | 27 (8.5) | 6 (12.5) | 1.6 (0.6, 4.3); 0.3 | 8 (8.3) | 0.9 (0.4, 2.2); 0.8 | 6 (10.5) | 1.3 (0.5, 3.3); 0.6 |
| *12 | 7 (2.2) | 0.0 | NA | 1 (1.0) | 0.4 (0.05, 3.7); 0.4 | 1 (1.7) | 0.7 (0.1, 6.1); 0.8 |
| *13 | 79 (25) | 21 (43.7) | 2.7 (1.4, 5.2); 0.002 | 33(34.3) | 2 (1.2, 3.4); 0.01 | 23 (40.3) | 2.4 (1.3, 4.5); 0.003 |
| *14 | 6 (1.9) | 2 (4.2) | 3.7 (0.6, 23); 0.1 | 2 (2.1) | 1.4 (0.2, 8.9); 0.8 | 2 (3.5) | 2.2 (0.4, 12.5); 0.3 |
| *15 | 118 (37.3) | 12 (25.0) | 0.5 (0.2, 0.9); 0.045 | 27 (28.1) | 0.5 (0.3, 0.9); 0.01 | 18 (31.6) | 0.7 (0.4, 1.4); 0.4 |
| *16 | 8 (2.5) | 1 (2.1) | 0.8 (0.1, 6.4); 0.8 | 4 (4.2) | 2.2 (0.5, 9.2); 0.2 | 2 (3.5) | 1.5 (0.3, 7.5); 0.6 |

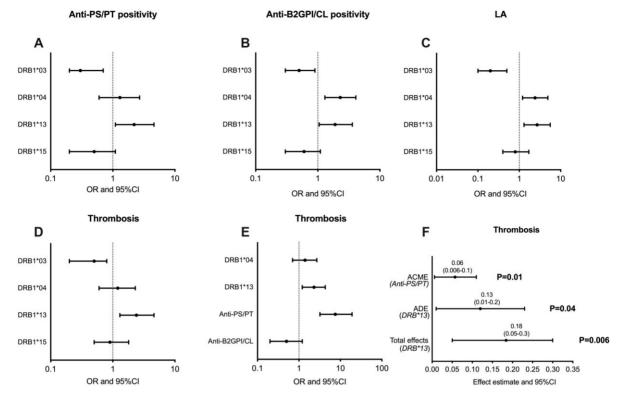


Fig. 1 Associations of HLA-DRB1 alleles with anti-phospholipid phenotypes and thrombosis in 341 SLE patients

Odds ratios (OR) with CIs of the independent variables (*DRB1*03*, *04, *13, *15) are illustrated for the outcomes (**A**) occurrence of any isotype of anti-phosphatidylserine/prothrombin, (**B**) any isotype of anti- β_2 GPI or aCL, (**C**) LA and (**D**) thrombosis. In (**E**), thrombosis as outcome is related to aPL, *HLA-DRB1*04* and *13. Multiple logistic regression models were used to calculate OR (central bullets) and CI (brackets). In (**F**), using mediation analysis, *HLA-DRB*13* and aPS-PT effect estimates with CI and *P*-values are shown for the outcome thrombosis. Anti- β_2 GPI/aCL: anti- β_2 glycoprotein I and/or anti-cardiolipin; ACME: average causal mediation effects; ADE: average direct effect of *DRB*13*.

associations (Fig. 1D). After adjustment for HCQ/prednisolone use and cardiovascular thrombotic risk factors (i.e. age, gender, hypertension, smoking, BMI and hyperlipidaemia) using one full forward stepwise regression model as well as several follow-up step-wise models excluding, one at a time, the independent variables that had significant associations with each other (supplementary Table S2, available at *Rheumatology* online), the associations to *HLA-DRB1*13 and HLA-DRB1*03* remained significant, with age, hyperlipidemia and steroids use also showing positive associations.

In a separate logistic regression model including anti- β_2 GPI/aCL, aPS-PT, *HLA-DRB*13* and *04 as predictors, occurrence of thrombosis associated only with *HLA-DRB*13* and aPS-PT positivity (Fig. 1E). The effect of *HLA-DRB1*13* on thrombosis [β (95% CI): 0.18 (0.05, 0.3)] was partially mediated by aPS-PT positivity [β (95% CI): 0.06 (0.006, 0.1)] (Fig. 1F), implying that aPS-PT positivity accounts for about one-third (31.4%) of the total effect of *HLA-DRB1*13* on thrombosis. In SLE patients without secondary APS, only *HLA-DRB*13*, among HLA-DRB* alleles, associated with thrombosis [OR 1.2 (CI 1.1–3.9), P = 0.04] and cerebrovascular events [OR 3.1 (CI 1.1–9.6), P = 0.03].

HLA-DRB1*03 negative associations

*HLA-DRB1*03* associated negatively with any thrombosis, VTE or APS diagnosis (Table 2). Associations between aPL positivity and VTE were significant, but after stratification for *HLA-DRB1*03* status, these associations were lost in the *HLA-DRB1*03* carrier patient group, for both aPS-PT and conventional aPL (Table 3). In addition, *HLA-DRB1*03* carriers had significantly lower levels of aPL compared with non-carriers (supplementary Table S3, available at *Rheumatology* online).

*HLA-DRB1*03* associated with higher levels of HDL (median 1.4 vs 1.2 mmol/l, P = 0.02) and lower levels of TG (median 0.9 vs 1.1 mmol/l, P = 0.04), Fig. 3. When correcting for age, gender, BMI and treatment with lipid-lowering agents, these associations remained, for HDL (standardized $\beta = 0.11$, P = 0.04) as well as TG (standardized $\beta = 0.12$, P = 0.03; data not shown). No other *HLA-DRB1* alleles associated with TG, HDL, LDL or total cholesterol (data not shown).

Discussion

Risk factors, such as HLA alleles as well as autoantibody profiles, have the potential to predict thrombotic events

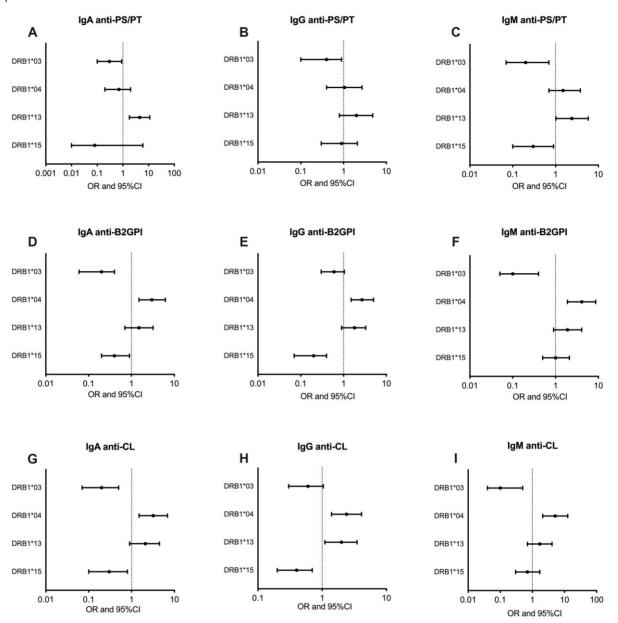


Fig. 2 Associations of HLA-DRB1*03, *04, *13, *15 with the occurrence of individual anti-phospholipid isotypes in SLE patients

Odds ratios (OR) with CIs of the illustrated *HLA-DRB1* alleles for the occurrence of IgA, IgG, IgM antibodies against phosphatidylserine/prothrombin (A, B, C), β_2 glycoprotein I (D, E, F), cardiolipin (G, H, I) are demonstrated. Chi2 tests were used to calculate OR (central bullets) and CI (brackets). Anti- β_2 GPI: anti- β_2 glycoprotein I.

in patients with APS. Therefore, understanding the association between these groups of risk factors and clinical variables is instrumental in the understanding and management of APS. Here we demonstrate that *HLA-DRB1*13* is a risk factor for expression of aPS-PT autoantibodies in SLE patients, while in contrast to conventional aPL, no association between aPS-PT and *HLA-DRB1*04* was observed. In addition, we demonstrate that aPS-PT antibodies mediate the association between *HLA-DRB1*13* and vascular events. Interestingly, *HLA-* *DRB1*03* associated negatively with aPL and thrombotic events, and positively with a favourable lipid profile.

To our best knowledge, only Bertolaccini *et al.* have previously investigated the associations between *HLA* and aPS-PT antibodies [28]. Also, they reported a higher prevalence of *HLA-DRB1*13* alleles among 82 patients from the UK with primary and secondary APS and positivity for aPS-PT as compared with healthy controls. However, this association was lost when anti- β_2 GPI-positive patients were excluded [28]. We here confirm

TABLE 2 Associations of the various HLA DRB1 genotypes with APS-associated features

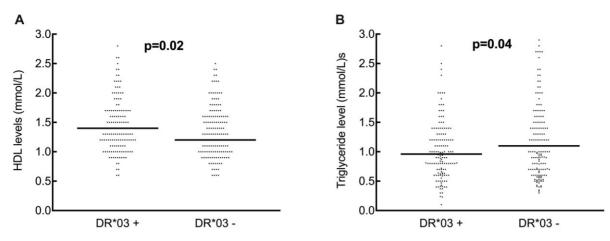
| HLA- DRB1 | Any thrombosis <i>n</i> = 73, <i>n</i> (%) | OR (95% CI); P | VTE n = 48, n (%) | , OR (95% CI); <i>P</i> | AT n = 36, n (%) | OR (95% CI); P | APS diagnosis <i>n</i> = 30, <i>n</i> (%) | OR (95% CI); <i>P</i> | GAPSS median (mean) in DRB1 (pos/neg) | Р |
|--------------|---|-----------------------|----------------------|-------------------------|---------------------|----------------------|--|------------------------|--|-------|
| *01 | 8 (10.9) | 0.9 (0.3, 1.8); 0.6 | 3 (6.2) | 0.4 (0.1, 1.3); 0.1 | 6 (17.1) | 1.4 (0.5, 3.7); 0.5 | 5 (16.7) | 1.5 (0.5, 4.2) 0.4 | 3 (5.8)/4 (5.8) | 0.6 |
| *03 | 23 (31.5) | 0.4 (0.2, 0.8); 0.004 | 14 (29.2) | 0.4 (0.2, 0.8); 0.01 | 11 (31.4) | 0.5 (0.2, 1.1); 0.06 | 6 (20.0) | 0.2 (0.09, 0.6); 0.002 | 3 (4.3)/4 (7.1) | 0.001 |
| *04 | 24 (32.9) | 1.2 (0.7, 2.1); 0.5 | 18 (37.5) | 1.5 (0.8, 2.8); 0.2 | 11 (31.4) | 1.1 (0.5, 2.3); 0.8 | 12 (40.0) | 1.6 (0.7, 3.4); 0.2 | 4 (7.7)/3 (4.9) | 0.003 |
| *07 | 6 (8.2) | 0.9 (0.3, 2.2); 0.8 | 2 (4.2) | 0.4 (0.09, 1.7); 0.2 | 4 (11.4) | 1.3 (0.4, 4.2); 0.6 | 2 (6.7) | 0.7 (0.1, 3.2); 0.7 | 4 (7.3)/4 (5.7) | 0.2 |
| *08 | 6 (8.2) | 0.9 (0.3, 2.2); 0.8 | 4 (8.3) | 0.9 (0.3, 2.7); 0.9 | 3 (8.6) | 0.9 (0.3, 3.3); 0.9 | 3 (10.0) | 1.1 (0.3, 4.0); 0.8 | 3 (5.7)/4 (5.8) | 0.6 |
| *09 | 2 (2.7) | 1.3 (0.2, 6.9); 0.7 | 2 (4.2) | 2.2 (0.4, 11.9); 0.3 | 2 (5.7) | 3.3 (0.6, 17.6); 0.1 | 1 (3.3) | 1.3 (0.1, 10.9); 0.8 | 3 (3.7)/4 (5.9) | 0.5 |
| *10 | 1 (1.3) | 0.5 (0.1, 4.5); 0.5 | 0 0 | NA | 1 (2.9) | 1.3 (0.1, 11.3); 0.8 | 0 0 | NA | 4 (3.2)/4 (5.9) | 0.7 |
| *11 | 5 (6.8) | 0.7 (0.3-2.1); 0.6 | 5 (10.4) | 1.3 (0.5, 3.7); 0.6 | 2 (5.7) | 0.6 (0.1, 2.8); 0.5 | 1 (3.3) | 0.3 (0.04, 2.6); 0.3 | 4 (5.1)/4 (5.9) | 0.9 |
| *12 | 3 (4.1) | 2.5 (0.5, 11.4); 0.2 | 2 (4.2) | 2.2 (0.4, 11.9); 0.3 | 1 (2.9) | 1.3 (0.1, 11.3); 0.8 | 1 (3.3) | 1.8 (0.2, 16.2); 0.6 | 2 (4.2)/4 (5.8) | 0.3 |
| *13 | 29 (39.7) | 2.5 (1.4, 4.4); 0.001 | 19 (39.6) | 2.3 (1.2, 4.3); 0.01 | 12 (34.3) | 1.7 (0.8, 3.5); 0.2 | 14 (46.7) | 2.9 (1.4, 6.4); 0.004 | 4.5 (7.7)/3 (5.1) | 0.007 |
| *14 | 3 (4.1) | 3.3 (0.6, 16.9); 0.1 | 3 (6.2) | 5.8 (1.1, 29.5); 0.049 | 00 | NA | 00 | NA | 9.5 (9.5)/4 (5.7) | 0.6 |
| *15 | 25 (34.3) | 0.8 (0.5, 1.4); 0.5 | 15 (31.2) | 0.7 (0.4, 1.4); 0.3 | 13 (37.1) | 0.9 (0.5, 2.0); 0.9 | 10 (33.3) | 0.8 (0.4, 1.8); 0.6 | 3 (4.8)/4(6.4) | 0.1 |
| *16 | 1 (1.3) | 0.4 (0.05, 3.8); 0.4 | 00 | NA | 1 (2.9) | 1.1 (0.1, 9.5); 0.9 | 1 (3.3) | 1.5 (0.2, 13.1); 0.7 | 6.5 (7.7)/4 (5.8) | 0.2 |

TABLE 3 Associations of aPL with thrombotic events among patients, stratified for the occurrence of HLA-DRB1*03 alleles

| | Any thrombosis antibody pos, <i>n/N</i> (%) | No thrombosis antibody pos, <i>n/N</i> (%) | OR (95% CI); <i>P</i> | VTE antibody pos, <i>n/N</i> (%) | No VTE antibody pos, <i>n/N</i> (%) | OR (95% CI); <i>P</i> |
|---------------------------------------|---|---|--------------------------|-------------------------------------|--|--------------------------|
| aPS/PT (any isotype) | | | | | | |
| All patients | 28/76 (36.8) | 22/248 (8.9) | 6 (3.1, 11.3); <0.0001 | 23/50 (46) | 27/275 (9.8) | 7.8 (3.9, 15.5); <0.0001 |
| HLA-DRB1*03 carriers | 4/23 (17.4) | 9/115 (7.8) | 2.5 (0.7, 8.9); 0.1 | 3/14 (21.4) | 10/124 (8.1) | 3.1 (0.7, 13); 0.1 |
| HLA-DRB1*03 non-carriers | 23/49 (46.9) | 12/116 (10.3) | 7.7 (3.4, 17.4); <0.0001 | 19/33 (57.7) | 16/133 (12) | 9.9 (4.2, 23.6); <0.0001 |
| Anti- β_2 GPI/aCL (any isotype) | | | | | | |
| All patients | 30/76 (39.5) | 70/246 (28.4) | 1.6 (0.1, 2.8); 0.07 | 25/50 (50) | 75/273 (27.4) | 2.6 (1.4, 4.9); 0.001 |
| HLA-DRB1*03 carriers | 3/23 (13) | 30/114 (26.3) | 0.4 (0.1, 1.5); 0.2 | 3/14 (21.4) | 30/123 (24.3) | 0.8 (0.2, 3.2); 0.8 |
| HLA-DRB1*03 non-carriers | 27/49 (55.1) | 36/115 (31.3) | 2.7 (1.3, 5.3); 0.004 | 22/33 (66.7) | 41/132 (31.1) | 4.4 (1.9, 10); 0.0002 |

Odds ratios (OR) and CIs were calculated using Chi^2 tests and data are expressed as fractions of antibody positive individuals among patients with and without thrombotic events, respectively. Significant associations are underlined. Anti- β_2 GPI/aCL: anti- β_2 glycoprotein I or aCL; VTE: venous thromboembolism.

Fig. 3 HLA-DRB1*03 allele association with HDL and LDL levels



Distribution of (A) HDL levels and (B) triglyceride levels among SLE patients stratified by carriage of *HLA-DRB1*03* alleles. HDL and TG data were available from 307 patients. HDL were outside the y axis limits for one patient and TG data for 13 patients; statistics and median bars represent data from all patients. HDL: high-density lipoprotein; LDL: low-density lipoprotein; TG: triglycerides.

associations between *HLA-DRB1*13* and aPS-PT, but in our larger cohort this association remain significant after exclusion of patients with anti- β_2 GPI/aCL.

Among the conventional aPL markers, only aCL IgG associated with HLA-DRB1*13. Instead, there were strong associations between all isotypes of anti-B2GPI and aCL and HLA-DRB1*04, in agreement with previous studies [12-14, 29]. In the study conducted by Lundström et al. on 665 Caucasian SLE patients, where parts of the current cohort were included, both HLA-DRB1*13 and *04 associated with conventional aPL, although the DRB1*13 allele associated only with anti-β₂GPI/aCL IgG, while HLA-DRB1*04 associated with all conventional aPL measures [14]. Arnett et al. studied HLA risk haplotypes in a multiethnic North American cohort with primary and secondary APS [12]. They also documented a significant association between HLA-DRB1*04 and anti-B2GPI in patients with Caucasian ancestry, without association with HLA-DRB1*13. In that study, HLA-DRB1*13 was associated with anti-β2GP1 among patients with African genetic ancestry. We observed, in a previous publication, positive associations between HLA-DRB1*13 and antiphospholipid-associated microvascular damage in kidney biopsies from patients with SLE [30]. None of these studies analysed associations for aPS-PT, which, as shown in the present study, differ from a main HLA-DRB1*04 association present for conventional aPL in SLE. We could not confirm the previous association between aPL and HLA-DRB1*07 [6, 7].

In our study, *HLA-DRB1*13* but not *04 increased the risk for thrombosis using both univariate and multivariate analyses. Lundström *et al.* reported that *HLA-DRB1*13* associated with any vascular events, while *HLA-DRB1*04* associated with cerebrovascular events. This *HLA-DRB1*04* association disappeared when excluding *HLA-DRB1*13* carrier patients. On the other hand, associations between *HLA-DRB1*13* and thrombosis remained significant after exclusion of *HLA-DRB1*04*

carrier patients, indicating that HLA-DRB1*13 confers an independent contribution to thrombosis [14]. In a study by Caliz et al., both HLA-DRB1*04 and DRB1*13 were more frequent in primary and secondary APS as compared with controls, but only HLA-DRB1*13 associated with arterial and venous thromboses [9]. Similarly, Tkachenko et al. showed that HLA-DRB1*04 associated with anti-β₂GPI carriage but not with APS diagnosis, whereas HLA-DRB1*13 associated with APS [13]. Collectively these data and our results indicate that HLA-DRB1*13 is a stronger risk factor for thrombosis than HLA-DRB1*04 in SLE patients. It is noteworthy that HLA-DRB1*13 alleles were shown previously to be more frequent in SLE patients with aPL-associated nephritis compared with LN, without HLA-DRB1*04 showing such associations [30].

Our study adds an understanding that the *HLA-DRB1*13* contribution to thrombosis risk is, to a large extent, mediated via aPS-PT. It is intriguing to note that besides being associated with aPS-PT, we find that *HLA-DRB1*13* is also independently associated with vascular events. This might point to a separate contributive role in vascular disease, perhaps linked to another not yet defined group of aPL, or to other factors.

The lower prevalence of *HLA-DRB1*03* in SLE patients with aPL and/or thrombotic events was significant also after adjustment for the presence of *HLA-DRB*04*, **13* and **15*, and other possible confounding factors. Asherson *et al.* previously reported total absence of *HLA-DRB1*03* among 13 patients with primary APS [29]. In another study, *HLA-DRB1*03* was rare in APS patients compared with non-APS SLE patients and healthy controls [31]. Murali *et al.* also described lower prevalence of *HLA-DRB1*03* in patients with ischaemic stroke compared with healthy controls [15]. In contrast, other studies reported increased *HLA-DRB1*03* in APS, when comparing APS secondary to lupus with healthy controls [13, 32]. We suspect that their findings are

due to the previously well-known association between *DRB1*03* and SLE, and not to an association with aPL [33]. Therefore, the interpretation of results is complicated not only by the fact that *HLA-DRB1*03* is common in SLE, but also that this genotype is specifically associated with anti-SSA/anti-SSB autoantibodies [34–36]. We have recently demonstrated that an unsupervised cluster analysis based on 13 SLE-associated autoantibodies, not including aPS-PT, identified four SLE subgroups that differ regarding *HLA-DRB1* associations, clinical and immunological presentations. The *HLA-DRB1*03* genotype is accumulated in a separate subgroup of SLE patients, which is commonly positive for anti-SSA/anti-SSB autoantibodies [37].

When we stratified SLE patients into HLA-DRB1*03 carriers and non-carriers, the association between aPS-PT and thrombotic events among carriers was lost among carriers, but became stronger among non-carriers; this was also evident for conventional aPL. Likewise, HLA-DRB1*03 carriers had lower levels of aPL compared with non-carriers, perhaps explaining the less likely risk of thrombosis observed. In line with these favourable effects, our results also demonstrate that only HLA-DRB1*03 carrier state, and no other studied HLA-DRB1 allele, had a beneficial effect on lipid levels. To our knowledge there is only one paper describing a thrombo-protective effect of DRB1*03, but no study other than the current one has related a protective role of DRB1*03 to lower aPL levels or the favourable lipid levels [15]. We have only found one study relating HLA-DRB alleles to lipid levels. A study by Giger et al. on postmenopausal African American women reported a strong association between DRB1*07 and low HDL levels, and a weaker association between HLA-DRB1*13 and high HDL levels, but no association with HLA-DRB1*03 [38]. An older study by Kovalev et al. evaluated HLA class I alleles in relation to atherosclerotic disease and also reported relation to HDL levels. No association was found to HLA-B8, a finding that could imply that there is also no association with HLA-DRB1*03 as both belong to the common ancestral HLA-A1/ B8/DR3 haplotype [39].

In univariate analysis we found a negative association between *HLA-DRB1*15* and aPL, which has also been reported by others [12, 13]. These associations were lost in multivariate analyses, and were not further analysed.

Our findings that individual *HLA-DRB1* alleles not only associate with specific aPL but also constitute independent risk factors for thrombosis imply that occurrence of *HLA-DRB1*03*, *04 and *13 might be evaluated as components in future APS risk scores, as today's scoring systems do not take HLA into account [23].

The fact that *HLA-DRB1*04* and *HLA-DRB1*13* associate with distinct groups of aPL suggests that the emergence of these groups of autoantibodies is driven by different initiating T lymphocytes. There is currently a vast literature on anti- β_2 GPI-reactive T cells in APS [40, 41]. These T lymphocytes recognize β_2 GPI peptides in the context of DR53 [42], an HLA-DR paralogue in linkage disequilibrium with *HLA-DRB1*04* but not with *HLA-DRB1*13* [43], in agreement with our findings. The literature concerning PS/PT-reactive T cells is essentially

lacking; we found only one short communication dealing with prothrombin-reactive T cells in APS [44].

Limitations to our current investigation are the retrospective inclusion of thrombotic events and that the aPL assay we used was a prototype test not yet in clinical use at the time of analysis.

To conclude, our study is the first to confirm that *HLA-DRB1*13* is a risk factor for aPS-PT, and that *HLA-DRB1*13* is a stronger risk factor than *HLA-DRB1*04* for thrombosis in SLE patients. In contrast, *HLA-DRB1*03* seems to confer a protection for vascular events, being associated with lower levels of aPL and favourable lipid profiles. Our data give further support to the notion that different *HLA-DRB1* genotypes associate with different SLE subsets. Prospective large studies are required to examine this narrative.

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Data availability statement

The data underlying this article will be shared at reasonable request to the corresponding author.

Supplementary data

Supplementary data are available at *Rheumatology* online.

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