The Minor Protective Allele at rs1876453 is Associated with Increased Age of Onset of Systemic Lupus Erythematosus

Ani Oganesyan¹, Jennifer A Kelly², Stuart B Glenn², Adam J Adler², Adrienne H Williams³, Marta E Alarcón-Riquelme^{4,5}, Graciela S Alarcón⁶, Juan-Manuel Anaya⁷, Sang-Cheol Bae⁸, Dam Kim⁸, Hye-Soon Lee⁸, Lindsey A Criswell⁹, Barry I Freedman¹⁰, Gary S Gilkeson¹¹, Joel M Guthridge², Chaim O Jacob¹², Judith A James^{2,13,14}, Diane L Kamen¹¹, Joan T Merrill¹⁵, Kathy Moser Sivils^{2,13}, Timothy B Niewold¹⁶, Michelle A Petri¹⁷, Rosalind Ramsey-Goldman¹⁸, John D Reveille¹⁹, R Hal Scofield^{2,14,20}, Anne M Stevens²¹, Luis M Vilá²², Timothy J Vyse²³, Kenneth M Kaufman^{24,25}, John B Harley²⁴, Carl D Langefeld³, Patrick M Gaffney², Elizabeth E Brown²⁶, Jeffrey C Edberg⁶, Robert P Kimberly⁶, Betty P Tsao¹¹, Daniela Ulgiati²⁷, Kenneth L. Jones²⁸, Susan A Boackle^{1,29}

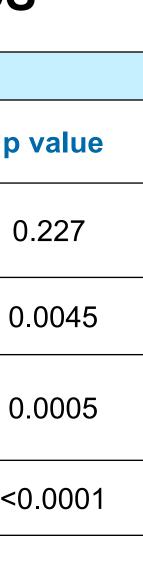
¹Division of Rheumatology, University of Colorado School of Medicine, Aurora, Colorado, USA; ³Department of Biostatistical Sciences and Center for Public Health Genomics, Wake Forest School of Medicine, Winston-Salem, North Carolina, USA; ⁴Medical Genomics, GENYO. Centre for Genomics and Oncological Research: Pfizer/University of Granada, Spain; ⁵Unit of Inflammatory Chronic Diseases, Institute of Environmental Medicine, Karolinska Institutet, Solna, Sweden; ⁶Department of Medicine, University of Alabama at Birmingham, ⁹Rosalind Russell/Ephraim P. Engleman Rheumatology Research Center, University of California, USA; ¹⁰Department of Internal Medicine, Wake Forest School of Medicine, Winston-Salem, North Carolina, USA; ¹¹Division of Rheumatology, Medical University of South Carolina, Charleston, South Carolina, USA; ¹²Department of Nedicine, University of Oklahoma City, Oklahoma City, Oklahoma, USA; ¹⁴Department of Medicine, University of Oklahoma Health Sciences Center, Oklahoma City, Oklahoma, USA; ¹⁵Department of Clinical Pharmacology, Oklahoma Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland, USA; ¹⁶Colton Center for Autoimmunity, NYU School of Medicine; ¹⁷Department of Medicine; ¹⁷Department of Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland, USA; ¹⁸Division of Rheumatology, Northwestern University Feinberg School of Medicine, Chicago, Illinois, USA; ²⁰US Department of Veterans Affairs Medical Center, Oklahoma City, Oklahoma, USA; ²¹Janssen Pharmaceuticals, Spring House, Pennsylvania, USA; ²²Division of Genetics and Molecular Medicine and Immunology, King's College London, London, UK; ²⁴Cincinnati Children's Hospital Medical Center, Cincinnati, Ohio, USA; ²⁵US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁷US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁷US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁷US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁷US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁷US Department of Veterans Affairs Medical Center, Cincinnati, Ohio, USA; ²⁶US Department of Veterans Affairs Medical Cent of Health and Disease, The University of Western Australia, Crawley, Western Australia, Crawley, Western Australia, Crawley, Western Australia, Crawley, Western Australia, Colorado, USA

BACKGROUND

- rs1876453 in CR2 is associated with decreased susceptibility to lupus (p^{meta}=4.2 × 10⁻⁴, OR 0.85) (Zhao et al, Ann Rheum Dis 75:242, 2016)
- This association was particularly striking when subjects with lupus were stratified based on the presence of anti-dsDNA autoantibodies $(p^{meta}=7.6 \times 10^{-7}, OR 0.71)$ (Zhao et al, Ann Rheum Dis 75:242, 2016)
- Anti-dsDNA antibodies are detected in patients with lupus a mean of 1.2 years before their initial symptom and a mean of 2.2 years before their diagnosis (Arbuckle et al, N Engl J Med 349:1526, 2003)

LLAS2 Demographics				
Characteristics	GG n=4703	AG n=648	AA n=36	р
Females, n (%)	4320 (92)	586 (90)	35 (97)	(
European American, n (%)	2643 (56)	370 (57)	30 (83)	C
African American, n (%)	1164 (25)	204 (31)	6 (17)	C
Hispanic, n (%)	896 (19)	74 (11)	0	<
n, number in sample				

PARTICIPANT CHARACTERISTICS



• Lupus patients with the minor protective allele at rs1876453 will have delayed onset of lupus compared to patients with the major

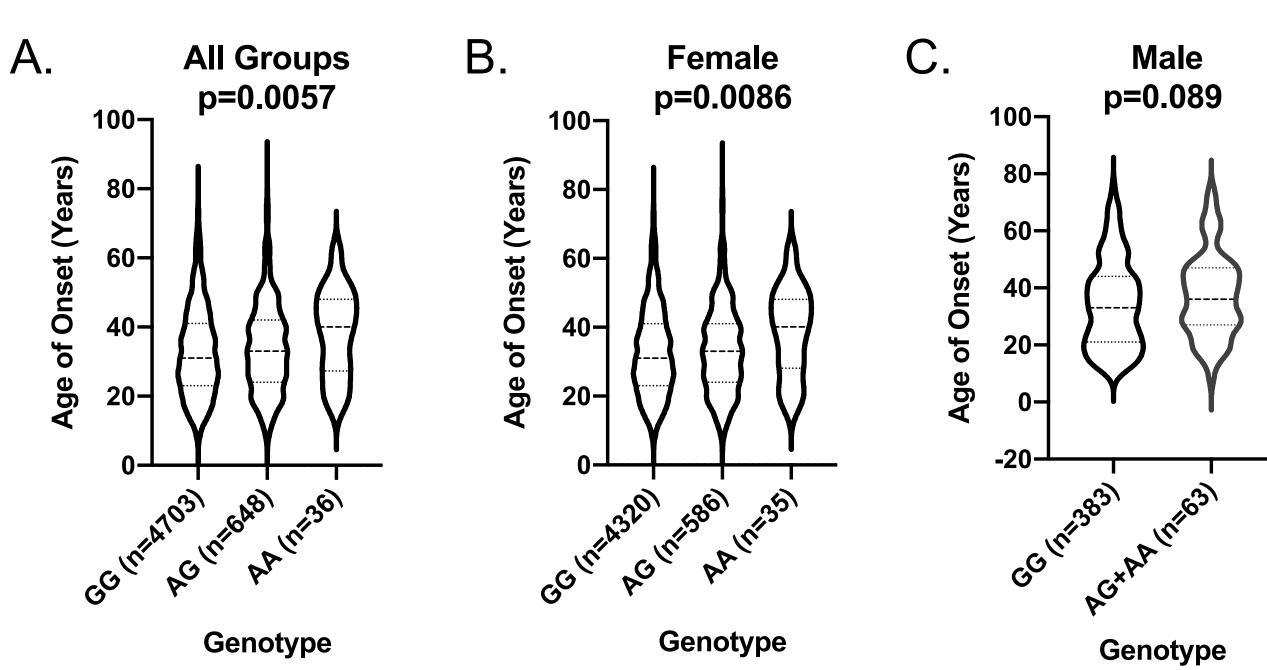


Figure 1. Lupus onset is delayed in subjects with the minor allele at rs1876453. The median age of lupus onset for subjects with AG or AA at rs1876453 was significantly higher than subjects with GG at the SNP [median (IQR) 40 (27.25-48) for AA, 33 (24-42) for AG, and 31 (23-41) for GG] (A). When stratified based on sex, females with the protective allele had significantly delayed disease onset (B) [median (IQR) 40 (28-48) for AA and 33 (24-41) for AG versus 31 (23-31) for GG], and males showed a similar difference (C) [median (IQR) 36 (27-47) for AG+AA versus 33 (21-44) for GG], though this was not statistically significant because the male cohort only had 40% power to detect a difference of 3.09 between the means with a significance level of 0.05 (two tailed).

HYPOTHESIS

RESULTS

- ancestry informative markers
- GraphPad Prism software.

- allele
- lupus pathogenesis

ACKNOWLEDGEMENTS

- NIAMS T32 AR007534 34

METHODS

• DNA from 5382 patients with SLE recruited from multiple sites was genotyped for rs1876453 on the Oklahoma Medical Research Foundation (OMRF) Illumina iSelect platform as part of the Large Lupus Association Study 2 (LLAS2)

Global ancestry was estimated based on the genotype of

• Age of onset was collected by chart review

• Kruskal-Wallis and Mann-Whitney tests were used to detect differences between groups. A p value of <0.05 was considered significant. Statistics and graphs were generated using

 Institutional review board approval was obtained for LLAS2 at OMRF and at each of the contributing sites.

CONCLUSIONS

• The minor protective allele at rs1876453 delays onset of lupus • Subjects with one copy of the protective allele develop lupus a median of 2 years later than subjects without the protective

• Subjects with two copies of the protective allele develop lupus a median of 9 years later than subjects without the protective

• These data confirm the relevance of this polymorphism in

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