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Laboratory markers of cardiovascular risk in pediatric SLE: the APPLE baseline cohort

SP Ardoin1, **LE Schanberg**2, **C Sandborg**3, **E Yow**2, **HX Barnhart**2, **KL Mieszkalski**2, **NT Ilowite**4, **E von Scheven**5, **A Eberhard**6, **DM Levy**7, **Y Kimura**8, **E Silverman**9, **SL Bowyer**10, **L Punaro**11, **NG Singer**12, **DD Sherry**13, **D McCurdy**14, **M Klein-Gitelman**15, **C Wallace**16, **R Silver**17, **L Wagner-Weiner**18, **GC Higgins**1, **HI Brunner**19, **LK Jung**20, **L Imundo**7, **JB Soep**21, **AM Reed**22, and **APPLE investigators**

¹Ohio State University Medical Center, Columbus, OH, USA

²Duke University Medical Center, Durham, NC, USA

³Stanford University School of Medicine, Palo Alto, CA, USA

⁴Albert Einstein College of Medicine, Bronx, NY, USA

⁵University of California at San Francisco, San Francisco, CA, USA

6Schneider Children's Hospital, New Hyde Park, NY, USA

⁷Columbia University Medical Center, New York, NY, USA

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Correspondence to: SP Ardoin, Ohio State University Medical Center, 480 Medical Center Drive, Columbus, OH 43210, USA. stacy.ardoin@osumc.edu.

Appendix The APPLE Investigators and Collaborators, in addition to the authors of this article, include the following individuals: Esi Morgan Dewitt, MD, MSCE, C. Egla Rabinovich, MD, MPH, Edward Fels, MD, Angela Robinson, MD, Rhonda Wilder, MS, Janet Ellis, RT, RDMS, RVT, Janet Wootton, RN, RSCN (Duke University Medical Center, Durham, NC); Imelda Balboni, MD, Peter Chira, MD, MS, Jennifer Frankovich, Joyce Hsu, MD, MS, Tzielan Lee, MD, Elizabeth Mellins, MD, Jane Park, MD, Vivian Saper, MD, Steven Song, MD, Jan Perea, BS, RDMS, RDCS (Stanford University School of Medicine, Palo Alto, CA); Lilliana Barillas-Arias, MD, Jude Cope, MD, Maria Espinosa, MD, Beth Gottlieb, MD, Patricia Irigoyen, MD, Jennifer Luftig, MD, Shaz Siddiqi, MD, Zhen Ni, RDMS, RVT, RCDS, Marilynn Orlando, RN, BC, Eileen Pagano, RN, MS, Marie Purugganan, MD (Schneider Children's Hospital, New Hyde Park, NY); Andrew Eichenfield, MD, Lisa F Imundo, MD, Philip Kahn, MD, Candido Batres, MD, Digna Cabral, BS (Morgan Stanley Children's Hospital of New York-Presbyterian, New York, NY); Laurie Ebner-Lyon, MD, Kathleen A Haines, MD, Suzanne C Li, MD, PhD, Jennifer Weiss, MD, Mary Ellen Riordan, RN, Beena Vaidya, RCDS (Hackensack University Medical Center, Hackensack, NJ); Lenny Dragone, MD, PhD, Aimee Hersh, MD, Diana Milojevic, MD, Michelle Mietus-Snyder, MD (University of California at San Francisco Medical Center); Brian McCrindle, MD, Diane Hebert, MD, Elizabeth Harvey, MD, Lawrence Ng, BSc (Hospital for Sick Children, Toronto, Ontario, Canada); Susan Ballinger, MD, Thomas Klausmeier, MD, Debra Hinchman, BS, RVT, Andrea Hudgins, CCRP, Amy Woodward, MD (Indiana University School of Medicine, Indianapolis); Maria Pascual, MD, Joele Wesley, MD, Heather Benham, MD, Dorothee Stichweh, MD, Shirley Henry, RN, PNP, Shuzen Zhang, ARDMS, RVT (Texas Scottish Rite Hospital for Children, Dallas); Elizabeth B Brooks, MD, PhD, Ira Davis, MD, Lisabeth Scalzi, MD, Stacy Miner, Nancy Szabo (University Hospitals/Case Medical Center, Cleveland, OH); Libby Dorfeld, Sarajane Wilson, BS, Tracey Wright, MD, Terri Finkel, MD, Pamela Fitch, MD, Timothy Buekelman, MD, Randy Cron, MD, Jon Burnham, MD, Eric Hanson, MD, Edward Behrens, MD, Daniel Albert, MD, Bita Arabshahi, MD, Amir Lerman, MD (Children's Hospital of Philadelphia, Philadelphia, PA); Andreas Reiff, MD, Bracha Shaham, MD, Bram Bernstein, MD, Katherine Marzan, MD, Tatiana Hernandez, BS, Jyotsna Vitale, BS, RT(M), RDMS, RVT (University of California Los Angeles Medical Center); Angela Kress, Nicole Lowe, Falguni Patel (Children's Memorial Hospital, Chicago, IL); Helen Emery, MD, Stephanie Hamilton, BSN, Chihiro Morishima, MD, Anne Stevens, MD (Seattle Children's Hospital and Regional Medical Center, Seattle, WA); Katie Caldwell, Diane Kamen, MD, MSCR (Medical University of South Carolina, Charleston); Becky Puplava, BS, Atanas Lonchev, RVT, RDCS, RDMS, Karen Onel, MD (University of Chicago, Chicago, IL); Monica Bacani, RDMS (Nationwide Children's Hospital, Columbus, OH); Thomas Griffin, MD, Thomas Graham, MD, Robert Colbert, MD, Murray Passo, MD, Minakshi Chaudhari, MD, Judith Smith, MD, David Glass, MD, Daniel Lovell, MD, Carline Wood, MD, Alexei Grom, MD, Cynthia Rutherford, BS, CRCIII, Jamie Meyers-Eaton, BS, CRCIII (Cincinnati Children's Hospital Medical Center, Cincinnati, OH); Laurie Beitz, MD, Teresa Conway, RN, MS, CCRC, Lacey Frank, Lori Kuss, ASRT, RDMS, RDCS, Troy Torgerson, MD (Creighton University Medical Center, Omaha, NE); J. Hollister, MD, Hazel Senz, RN, BSN, CCRC (Denver Children's Hospital, Denver, CO); Thomas Mason, MD, Jane Jaquith, CCRC, Diana E. Paepke-Tollefsrud, BS, RVT, RDMS, RT(R) (Mayo Clinic, Rochester, MN).

Hackensack University Medical Center, Hackensack, NJ, USA 9Toronto Hospital for Sick Children, Toronto, Ontario, Canada ¹⁰Indiana University School of Medicine, Indianapolis, IN, USA Texas Scottish Rite Hospital for Children, Dallas, TX, USA University Hospitals/Case Medical Center, Cleveland, OH, USA Children's Hospital of Philadelphia, Philadelphia, PA, USA University of California Los Angeles School of Medicine, Los Angeles, CA, USA Northewstern University Feinberg School of Medicine, Chicago, IL, USA Seattle Children's Hospital, Seattle, WA, USA 17 Medical University of South Carolina, Charleston, SC, USA 18University of Chicago, Chicago, IL, USA Cincinnati Children's Hospital Medical Center, Cincinnati, OH, USA ²⁰Children's National Medical Center, Washington, DC, USA The Children's Hospital, Denver, CO, USA Mayo Clinic, Rochester, MN, USA

Abstract

As part of the Atherosclerosis Prevention in Pediatric Lupus Erythematosus (APPLE) Trial, a prospective multicenter cohort of 221 children and adolescents with systemic lupus erythematosus (SLE) (mean age 15.7 years, 83% female) underwent baseline measurement of markers of cardiovascular risk, including fasting levels of high-density lipoprotein (HDL), low-density lipoprotein (LDL), triglycerides (TG), lipoprotein A (Lpa), homocysteine and high-sensitivity Creactive protein (hs-CRP). A cross-sectional analysis of the baseline laboratory values and clinical characteristics of this cohort was performed. Univariable relationships between the cardiovascular markers of interest and clinical variables were assessed, followed by multivariable linear regression modeling. Mean levels of LDL, HDL, Lpa, TG, hs-CRP and homo-cysteine were in the normal or borderline ranges. In multivariable analysis, increased Systemic Lupus Erythematosus Disease Activity Index (SLEDAI), prednisone dose, and hypertension (HTN) were independently associated with higher LDL levels. Higher hs-CRP and creatinine clearance were independently related to lower HDL levels. Higher body mass index (BMI), prednisone dose, and homocysteine levels were independently associated with higher TG levels. Only Hispanic or non-White status predicted higher Lpa levels. Proteinuria, higher TG and lower creatinine clearance were independently associated with higher homocysteine levels, while use of multivitamin with folate predicted lower homocysteine levels. Higher BMI, lower HDL, and longer SLE disease duration, but not SLEDAI, were independently associated with higher hs-CRP levels. The R^2 for these models ranged from 7% to 23%. SLE disease activity as measured by the SLEDAI was associated only with higher LDL levels and not with hs-CRP. Markers of renal injury (HTN, proteinuria, and creatinine clearance) were independently associated with levels of LDL, HDL, and homocysteine, highlighting the importance of renal status in the cardiovascular health of children and adolescents with SLE. Future longitudinal analysis of the APPLE cohort is needed to further examine these relationships.

Keywords

atherosclerosis; cardiovascular; lipid; pediatric; SLE; systemic lupus erythematosus

Introduction

Premature atherosclerosis is a well-recognized complication of systemic lupus erythematosus (SLE). Although a focus of intense research, the pathophysiology underlying this association remains incompletely understood. Medication side effects, chronic inflammation, autoantibodies, hypertension, renal disease, and abnormal lipid profiles likely contribute to the enhanced cardiovascular risk evident in SLE.¹ Given their lifelong exposure to atherogenic risk factors, children and adolescents with SLE are at particularly high risk for developing premature atherosclerosis and are therefore ideal candidates for primary prevention. However, optimal primary prevention strategies for children and adolescents with SLE are not yet evidence-based. Several markers of atherosclerotic risk have been extensively investigated as targets for primary and secondary prevention in the general population, and efforts are underway to define optimal prevention strategies for SLE patients.

In large-scale observational studies of the general population, abnormal levels of lowdensity lipoprotein (LDL), high-density lipoprotein (HDL), and triglycerides (TG) are clearly linked with heightened risk of atherosclerotic events, and well established guidelines for primary and secondary prevention, including those published by the National Cholesterol Education Program (NCEP), are widely accepted.² High rates of abnormal HDL and LDL levels have also been described in adult and pediatric SLE cohorts.3,4 Across the age spectrum, active SLE has been associated with low HDL levels.^{4,5} Although lowering LDL and increasing HDL decrease atherosclerotic events in the general population, at least 50% of vascular events occur in the absence of overt dyslipidemia.⁶ Thus, there is great interest in the development of novel markers of atherosclerotic risk to complement the prognostic capabilities of LDL, HDL, and TG.

Among newer candidate markers for athero-sclerotic risk, high-sensitivity C-reactive protein (hs-CRP) has stimulated the most interest. Elevated levels of hs-CRP independently predict myocardial infarction, peripheral arterial disease, and sudden cardiac death in large-scale adult epidemiologic studies, adding prognostic benefit beyond LDL and the traditional Framingham risk score.⁷ Potential pro-atherogenic mechanisms of hs-CRP include enhanced expression of vascular adhesion molecules, impaired fibrinolysis, and endothelial dysfunction.⁷ Among adults with SLE in a large epidemiologic study, hs-CRP was associated with cardiovascular mortality.⁸ One case-control study showed no difference in hs-CRP levels in children with SLE compared with healthy controls.⁹ Normal, borderline, and elevated levels of hs-CRP have been defined for the general adult population, but not for children and adolescents or for individuals with SLE.10 Despite the chronic inflammation

Several epidemiologic studies have established lipoprotein A (Lpa) as an independent cardiovascular risk factor.^{11,12} Lpa is a lipid moiety consisting of an LDL-like particle, linked by its apolipoprotein B 100 component to a molecule of apolipoprotein a. Among lipid-lowering therapies, only niacin appears to lower Lpa levels significantly; however, no randomized trial data show that lowering Lpa levels prevents cardiovascular morbidity.¹³ No guidelines for routine Lpa screening or intervention have been established. While this has not been studied extensively, adult SLE cohorts have shown elevated Lpa levels compared with healthy controls and higher Lpa levels with increased disease activity.¹⁴ However, in a small pediatric SLE cohort study, Lpa levels did not differ from controls.¹⁵ Published data from the Atherosclerosis Prevention in Pediatric Lupus Erythematosus (APPLE) trial showed that higher Lpa levels at baseline were associated with thicker carotid intima media.¹⁶

Elevated homocysteine levels are associated with endothelial dysfunction, platelet activation, and accelerated LDL oxidation.¹⁷ In the general population, elevated homocysteine levels are associated with increased risk of cardiovascular disease (CVD).¹⁸ While it is known that supplementation with vitamins B6, B12, and folic acid reduces homocysteine levels, clinical trials have failed to show that lowering homocysteine levels improves cardiovascular outcomes in the general population.¹⁹ Among adults with SLE, higher homocysteine levels are associated with atherosclerosis progression, arterial stiffness, left ventricular hypertrophy, and coronary artery calcification.^{20–23} Among children, small studies have suggested that homocysteine levels are higher in pediatric SLE than in healthy controls. One study in pediatric SLE patients found no correlation with endothelial dysfunction.¹⁵

In this study, we investigate the levels and clinical associations of laboratory markers of cardiovascular risk – LDL, HDL, TG, Lpa, homocysteine, and hs-CRP – in a large and diverse pediatric SLE cohort enrolled in the APPLE trial. In approaching this exploratory analysis, our hypothesis was that children and adolescents with SLE would have abnormal levels of LDL, HDL, TG, Lpa, homocysteine, and hs-CRP. We anticipated that abnormalities of LDL, HDL, Lpa, and hs-CRP would be influenced by SLE disease activity and that prednisone dose would be related to levels of LDL and TG.

Patients and methods

Patient population: baseline APPLE cohort

A prospective multicenter cohort of 221 children and adolescents with SLE from 21 sites in North America underwent measurement of fasting LDL, HDL, TG, Lpa, homocysteine, and hs-CRP as part of the APPLE Trial. The APPLE Trial is a double-blind randomized placebo-controlled study designed to determine the efficacy and safety of atorvastatin in preventing the progression of atherosclerosis in children and adolescents with SLE. The primary endpoint is the rate of progression of mean carotid intimal thickness (CIMT) in the

common carotid artery over 3 years. Subjects are randomized to receive either placebo or atorvastatin (10 or 20 mg daily depending upon weight). Prescription of multivitamin (MVI) with folate, hydroxychloroquine, and low-dose aspirin was recommended for all subjects, as were dietary recommendations and risk factor counseling. Enrollment was completed in 2006, with study results expected in 2010.

All subjects met the ACR 1997 revised diagnostic criteria for $SLE²⁴$ Full inclusion and exclusion criteria for the APPLE trial are as previously published.16 Inclusion criteria included weight 25 kg, outpatient status, and age between 10 and 21 years, and patients were excluded for active nephrotic syndrome, myositis, liver disease, renal insufficiency, or hypercholesterolemia warranting treatment (total cholesterol > 350 mg/dl) at enrollment.

Clinical variables of interest

The APPLE baseline assessment included demographics, a history and physical examination performed by a pediatric rheumatologist, chart review, and subject and parent questionnaires. Variables of interest for this analysis were SLE clinical and laboratory parameters, non-lupus and lupus-related medical history, family history, current and prior medication use, traditional risk factors for atherosclerosis, and SLE disease activity. Race and ethnicity were self-reported.

SLE disease activity and damage were assessed using the modified SELENA Systemic Lupus Erythematosus Disease Activity Index (SLEDAI)25 and Systemic Lupus International Collaborating Clinics/ACR Damage Index (SDI)26 For these analyses, the total SLEDAI score was treated as a continuous variable. SDI scores were analyzed as a dichotomous variable with subjects grouped as a score of 0 (no damage) or a score of $\frac{1}{2}$ (damage) present). Proteinuria was defined as spot urine protein/creatinine ratio 0.5 or timed urine with protein $>$ 500 mg/24 h. A history of glomerulonephritis referred to current or prior history of nephritis or prior history of nephrotic syndrome. Similarly, a history of hypertension referred to the treating clinician's report of current or historical hypertension. Creatinine clearance was calculated using the Schwartz formula.27 Leukopenia was defined as a white blood cell count $<$ 3000/mm³. A family history of CVD was defined as subjectreported cerebrovascular accident, myocardial infarction, angina, or atherosclerosis in a parent or grandparent. A family history of hyperlipidemia was defined as subject-reported hyperlipidemia in a parent or grandparent. Current oral glucocorticoid doses were weightadjusted and recorded as prednisone equivalent in mg/kg/d. For the purposes of this analysis, use of immunosuppressant medications and non-steroidal anti-inflammatory drugs (NSAIDs) referred to current usage.

Laboratory assessment

The following tests were obtained following a 12-h fast and were measured in a central laboratory: total cholesterol, TG, HDL, apolipoprotein A1 (ApoA1), apolipoprotein B (ApoB), Lpa, homocysteine, and hs-CRP. All other laboratory assessments were performed locally. Total cholesterol and TG were measured by Roche Enzymatic assay. HDL was measured by Roche Enzymatic with MnCl precipitation. LDL was calculated using the Friedewald equation.²⁸ ApoA1 and ApoB were measured using Dade-Behring

Immunoturbimetric Nephelometry. Lpa and hs-CRP were measured using Polymedco Immunoturbimetric Assay. Homocysteine was measured using high performance liquid chromatography.

Statistical methods

The characteristics of the study sample were summarized using descriptive statistics, with dichotomous or ordinal data presented as percentages and continuous data as medians with interquartile ranges or as means with standard deviations (SD). Differences between groups were assessed using either the χ^2 test for dichotomous data or the non-parametric Wilcoxon test for continuous data.

Using cross-sectional baseline data from the APPLE trial, we first investigated potential risk factors and their univariable relationships with each of the six outcomes of interest (baseline levels of LDL, HDL, TG, Lpa, homocysteine, and hs-CRP). The following potential predictors were considered: age, gender, Hispanic or non-White status, BMI, duration of SLE, SDI, current immunosuppressive use, current NSAID use, current prednisone dose, creatinine clearance, proteinuria, history of glomerulonephritis, history of hypertension, family history of CVD, and family history of hyperlipidemia. Lipid values (total cholesterol, HDL, LDL, TG, ApoA1, ApoB, and Lpa) were also studied for potential relationships with hs-CRP and homocysteine models. Given the anticipated high correlation, relationships between HDL, LDL, TG, ApoA1, and ApoB were not assessed. Homocysteine and hs-CRP were considered potential predictors for HDL, LDL, TG, and Lpa in the multivariable models. Hydroxychloroquine use was not included as a potential predictor in these models because 96% of subjects were taking this medication. Information on duration of hydroxychloroquine use was not available. Use of MVI was considered in the homocysteine analyses, as vitamins B6, B12, and folate are known to impact homocysteine levels.

Separate univariable analyses using linear regression models were performed, examining the relationship between each predictor and each of the six outcomes. Residual plots of the regression models were used to assess the linear models. Scatter plots with overlay of locally weighted scatterplot smoothing (LOESS) were created to examine the relationships between continuous variables and outcome variables.29 Histograms of each of the outcome variables according to presence or absence of categorical predictors were produced to assess the relationship between categorical predictors and outcome variables. When the relationship between the continuous predictor and the outcome variable was not linear, appropriate transformations or piecewise linear regression models were pursued to better understand the relationship between variables. Upon examination of the distributions of the continuous variables, hs-CRP, TG, and Lpa and hs-CRP were log-transformed to achieve a more normal distribution before proceeding to multivariable linear regression.

As this was an exploratory analysis, potential predictors with *p* values <0.15 in univariable analysis were included in the multivariable linear regression models. Forward, backward, and stepwise regression model building procedures were used to derive the final regression model for each of the six outcomes. We assessed each model with co-linearity diagnostics. Bonferroni adjustments for multiple comparisons were not performed due to the exploratory nature of our analyses. Statistical associations were considered significant in the final

models if *p* values were <0.05. All statistical calculations were performed using SAS 8.2 and SAS Enterprise Guide 4.0 (SAS Enterprise Institute, Cary, NC).

Results

A summary of baseline characteristics of the APPLE cohort is depicted in Table 1. Of the 221 enrolled subjects, mean age was 15.7 years (range 10.1–21.7 years), 83% were female, and 65% were Hispanic or non-White. Subjects had mean disease duration of 31 months (median 24 months, interquartile range 8 to 46 months) and had mild disease activity with a mean SLEDAI score of 4.6 (range 0 to 20, median 4.0, interquartile range 2 to 6). As summarized in Table 1, mean LDL was 86.4 mg/dl (normal < 110 mg/dl); mean HDL was 46.3 mg/dl (normal > 45 mg/dl); mean TG levels were 114 mg/dl (normal < 90, borderline 90–129 mg/dl); and mean Lpa levels were 23.1 mg/dl (normal $<$ 30 mg/dl).^{30,31} Mean levels of homocysteine (7.5 μmol/l) were between the 75th and 95th percentiles for US children aged 3–19 years, and mean hs-CRP level (3.6 mg/l) in this cohort represented the 90th percentile for US children aged $3-19$ years.³² As shown in Figure 1, the most commonly detected abnormalities were elevated TG and Lpa levels.

LDL

As shown in Table 2, univariable analysis suggested that the following factors were associated with higher LDL levels: Hispanic or non-White ethnicity, history of glomerulonephritis, history of hypertension, higher SLEDAI score, proteinuria, higher prednisone dose, higher homocysteine levels, and current use of cyclophosphamide, mycophenolate mofetil, or an ACE-inhibitor. Methotrexate use and family history of CVD were associated with lower LDL levels. Complete univariable modeling results are shown in Table 3. Multivariable modeling showed independent relationships of increased LDL levels with increased SLEDAI, increased prednisone dose, and history of hypertension.

HDL

Univariable analysis revealed that male gender, higher BMI, hs-CRP and creatinine clearance all predicted lower HDL levels, while higher prednisone dose was associated with higher HDL. The final multivariable model showed that only higher hs-CRP and higher creatinine clearance were independently associated with lower HDL levels.

TG

Univariable analysis suggested that the following risk factors were independently associated with higher TG levels: proteinuria, history of glomerulonephritis, leukopenia, history of hypertension, current use of cyclophosphamide or mycophenolate mofetil, higher prednisone dose, homocysteine, SLEDAI score, and BMI. Current use of methotrexate predicted lower TG levels. The final multivariable indicated that higher BMI, prednisone dose, and homocysteine levels were independently associated with higher TG levels.

Lpa

For Lpa, univariable analysis showed that Hispanic or non-White status, higher hs-CRP and SDI score were risk factors for higher Lpa; however, multivariable modeling showed a significant relationship only between Hispanic or non-White status and Lpa.

hs-CRP

Hispanic or non-White status, BMI, duration of SLE, history of glomerulonephritis, SDI score, higher creatinine clearance, lower HDL or apoA1, NSAID use, and higher Lpa levels were independently related to higher hs-CRP levels. A family history of CVD was associated with lower levels of hs-CRP. In multivariable modeling, current NSAID use, higher BMI, lower HDL, and longer SLE disease duration predicted higher hs-CRP levels.

Homocysteine

Age, male gender, higher BMI, SLEDAI, proteinuria, leukopenia, lower creatinine clearance, higher prednisone dose, current use of cyclophosphamide, and higher TG, LDL, and apoB levels were independently related to higher homocysteine levels in univariable analysis. Use of MVI with folate was associated with lower homocysteine levels. In multivariable analysis, proteinuria, higher TG and lower creatinine clearance were independently associated with higher homocysteine levels, while use of MVI with folate predicted lower homocysteine levels.

Additional analyses

To further explore the relationship between hs-CRP and NSAIDs, additional analysis was performed, which revealed that the relationship between NSAID use and hs-CRP was strongest at two sites. When these two sites were removed from the multivariable model, the relationship between NSAID use and hs-CRP was no longer significant, but none of the other observed relationships changed. We also compared those taking NSAIDS ($n = 68$) with those not taking NSAIDs $(n = 153)$. This analysis revealed only two statistically significant differences between NSAID users and non-users: NSAID users were less likely to have a personal history of hypertension $(39\% \text{ vs } 24\%, p = 0.033)$ and less likely to have a family history of hyperlipidemia (33% vs 50% , $p = 0.019$). As arthritis is a common indication for NSAID use, the relationship between arthritis and hs-CRP was further explored. On univariable analysis, the presence of arthritis (identified as present on the SLEDAI) was not independently associated with hs-CRP (slope 0.606; SE 0.394; $p =$ 0.126), and adding arthritis to the hs-CRP model did not change the model results.

Discussion

The majority of APPLE subjects had normal or borderline values of LDL, HDL, TG, and Lpa according to definitions developed for the general pediatric and adolescent population in the United States. Optimal levels of hs-CRP and homocysteine are not well defined for pediatric populations, and hs-CRP guidelines are extrapolated from recommendations developed for the general adult population. Screening benchmarks developed for the general population may not be appropriate for children and adolescents with SLE, a group whose high atherogenic potential may warrant a more aggressive prevention strategy, similar to that

used for people with diabetes. The exclusion of patients with a family history of familial hypercholesterolemia or total cholesterol >350 mg/dl, nephrotic syndrome, and significant renal insufficiency likely resulted in a sample with lower rates of dyslipidemia than in other pediatric and adult SLE cohorts. In addition, all subjects were required to take hydroxychloroquine, which favorably impacts lipid profiles in adults and children with SLE.33,34

Chronic steroid use has conventionally been viewed as pro-atherogenic; however, in epidemiologic studies of adults with SLE, individuals treated with lower steroids (and other immunosuppressives) actually had higher carotid plaque burden.²³ In the baseline APPLE cohort, higher prednisone was associated with higher LDL and triglyceride levels. Given the cross-sectional nature of the data, we were not able to assess cumulative steroid exposure, which is likely important in cardiovascular risk and will be a focus of longitudinal data analysis once the APPLE trial is complete.

While this analysis corroborated several known associations – including the relationships between Lpa and Hispanic or non-White status; between BMI, triglycerides, and hs-CRP; between triglycerides and homocysteine; and between HDL and hs-CRP – other observations were novel. Higher SLE disease activity, as measured by the SLEDAI, was associated with higher LDL levels in this cohort. We did not find a significant relationship between higher disease activity and lower HDL levels, as frequently reported in the literature.34–36 This difference may be explained by the exclusion of individuals with nephrotic syndrome and significant baseline hypercholesterolemia or the effects of treatment with glucocorticoids (although an independent relationship between HDL and prednisone use was not found in multivariable analysis). In addition, APPLE subjects had low SLE disease activity at baseline, which may have limited the ability to identify effects of active disease on lipid profiles.

The association between NSAID use and higher levels of hs-CRP was surprising and intriguing, particularly given the association between NSAID use and cardiovascular disease. However, this relationship proved to exist primarily at two of 21 sites, and thus is of unclear significance and may represent a spurious finding in the setting of multiple comparisons. The baseline APPLE data did not include information concerning the relative cyclo-oxygenase-selectivity, dose, or frequency of NSAID use. No previous studies have addressed NSAID use and its relationship with hs-CRP in the setting of SLE.

In this cohort, hs-CRP and SLE disease activity did not show a significant association with each other, but longer disease duration did predict higher hs-CRP. The existing literature demonstrates mixed results regarding the relationship between SLE disease activity and hs-CRP, bringing into question the utility of this laboratory measure as a marker of disease activity.37–42 Two large studies have evaluated hs-CRP in adults with SLE. Bertoli *et al*. studied 588 subjects with SLE in the LUMINA cohort and found that hs-CRP was associated with disease activity (as measured by Systemic Lupus Activity Measure-Revised (SLAM-R).37 Lee *et al*. examined 610 patients with SLE and found in multivariable analysis that higher SELENA-SLEDAI scores predicted higher hs-CRP, but duration of SLE was not an independent predictor of hs-CRP.38 Other studies have not demonstrated any relationship

between hs-CRP and SLE disease activity.40–42 Antibodies to C-reactive protein (not measured in the APPLE cohort) can lower hs-CRP levels in SLE and may represent an important confounder.⁴³

Even in the absence of frank renal insufficiency and nephrotic syndrome, evidence of renal injury (proteinuria, creatinine clearance, and history of hypertension) was significantly associated with markers of cardiovascular risk, especially HDL and homocysteine levels. Lower creatinine clearance predicted higher levels of homocysteine and HDL. These seemingly disparate results may in part be due to the known relationship between hyperglomerulofiltration and markers of the metabolic syndrome (including low HDL) in the setting of chronic kidney disease.⁴⁴ Proteinuria was associated with higher levels of homocysteine, consistent with findings in the general population, chronic kidney disease and diabetes.^{45,46} In the general population, even small decrements of renal function are associated with significant increases in cardiovascular morbidity. 47 Taken together, these results suggest that even modest renal injury is related to homocysteine and HDL concentrations, and highlight the importance of renal status in cardiovascular health for children and adolescents with SLE.

There are several limitations to this study, including its cross-sectional nature. The sample size was powered to detect change in the primary outcome of the APPLE trial, progression rate of CIMT; thus it may not have had sufficient power to identify meaningful associations between clinical features and markers of cardiovascular risk. In addition, we performed many comparisons in these analyses, and our chances of finding some statistically significant relationships were high. Thus, the results of this exploratory analysis must be interpreted with caution and confirmed in future prospective studies. Despite these limitations, these results are derived from the largest, most diverse pediatric SLE cohort in which fasting LDL, HDL, Lpa, hs-CRP, TG and homocysteine have been measured. The potential relationships reported here are hypothesis-generating and need to be confirmed by future, longitudinal studies of the APPLE and other cohorts. As in any clinical trial, there is risk of selection bias, limiting the generalizability of the APPLE cohort to pediatric lupus patients as a whole. As examples of this selection bias, some obese patients were excluded due to inability to obtain high-quality CIMT images. Patients with nephrotic syndrome and significant renal insufficiency were also excluded. As in all studies, historical variables are subject to recall bias. Race and ethnicity are highly complex, and any categorization introduces bias and limitations. In this study, race and ethnicity were separately selfreported. Given the small sample size, we elected to limit analysis to the largest groups, non-Hispanic Whites, Hispanics, and non-Whites.

The presented models explain only 7–23% of the variability in these markers of cardiovascular risk, suggesting that other important measured or unmeasured risk factors were not identified in the current study. Few studies in adult or pediatric SLE have examined LDL, HDL, TG, Lpa, homocysteine, or hs-CRP individually as outcomes using multivariable linear regression models, and therefore it is difficult to compare the fit of our models with others in the literature. Silverman *et al*. described models with significantly higher model adjusted R^2 values for HDL (0.577) and TG (0.629); however, these analyses involved a more homogeneous sample of recently diagnosed and untreated pediatric SLE

patients, suggesting that treatment effects and variations in disease activity likely influence these markers of cardiovascular risk.³⁵

Taken together, the results of these analyses suggest that low SLE disease activity and preserved renal function may correlate with improvements in markers of cardiovascular risk, particularly LDL, HDL, and homocysteine. Because cardiovascular screening recommendations developed for the general population may not be adequate or optimal for children and adults with SLE, further research is needed to determine which markers of cardiovascular risk reliably predict cardiovascular events in children and adolescents with SLE.

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Figure 1.

Proportion of APPLE subjects with normal, abnormal or borderline levels of laboratory markers of cardiovascular risk. ^aAdapted from references Pearson *et al.*, ¹⁰ NCEP, ³⁰ and Obisesan.³¹ ^aThere is no accepted borderline range for lipoprotein A. HDL: high-density lipoprotein, hs-CRP: high-sensitivity C-reactive protein; LDL: low-density lipoprotein, Lpa: Lipoprotein A, TG: triglycerides, Tot. Chol: total cholesterol.

Table 1

Baseline characteristics of the APPLE subjects

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ACE: angiotensin converting enzyme, Apo-A1: apolipoprotein A1, Apo-B: apolipoprotein B, HDL: high-density lipoprotein, LDL: low-density lipoprotein, hs-CRP: high-sensitivity C-reactive protein, Hx: history of, SLE: systemic lupus erythematosus, SLEDAI: Systemic Lupus Erythematosus Disease Activity Index, SDI: Systemic Lupus International Collaborating Clinics/American College of Rheumatology Damage Index.

a Some subjects self-reported more than one race.

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Table 2

Univanable relationships between clinical variables and laboratory markers of cardiovascular risk in the baseline APPLE cohort Univanable relationships between clinical variables and laboratory markers of cardiovascular risk in the baseline APPLE cohort

BMI: body mass index, CVD: cardiovascular disease, HDL: high-density lipoprotein, hs-CRP: high sensitivity C-reactive protein, Hx: history of, LDL: low-density lipoprotein, Lpa: lipoprotein A, NI: not
included in univariab BMI: body mass index, CVD: cardiovascular disease, HDL: high-density lipoprotein, hs-CRP: high sensitivity C-reactive protein, Hx: history of, LDL: low-density lipoprotein, Lpa: lipoprotein A, NI: not included in univariable analysis due to predicted collinearity, SE: standard error, SDI: Systemic Lupus International Collaborating Clinics/American College of Rheumatology Damage Index, SLE: systemic lupus erythematosus, TG: triglycerides.

Table 3

Multivanable adjusted relationships between clinical variables and laboratory markers of cardiovascular risk in the baseline APPLE cohort

BMI: body mass index, HDL: high-density lipoprotein, hs-CRP: high-sensitivity C-reactive protein, Lpa: lipo-protein a, HTN: hypertension, Hx: history of, MVI: multivitamin, NSAID: non-steroidal anti-inflammatory drug, SE: standard error, SLEDAI: Systemic Lupus Erythematosus Disease Activity Index, TG: triglyceride.