UC San Diego

UC San Diego Previously Published Works

Title

Longitudinal patterns and predictors of response to standard-of-care therapy in lupus nephritis: data from the Accelerating Medicines Partnership Lupus Network.

Permalink

https://escholarship.org/uc/item/98r572bg

Journal

Arthritis Research & Therapy, 26(1)

Authors

Izmirly, Peter Kim, Mimi Carlucci, Philip et al.

Publication Date

2024-02-20

DOI

10.1186/s13075-024-03275-z

Copyright Information

This work is made available under the terms of a Creative Commons Attribution License, available at https://creativecommons.org/licenses/by/4.0/

Peer reviewed

RESEARCH Open Access

Longitudinal patterns and predictors of response to standard-of-care therapy in lupus nephritis: data from the Accelerating Medicines Partnership Lupus Network

Peter M. Izmirly^{1*†}, Mimi Y. Kim^{2†}, Philip M. Carlucci¹, Katherine Preisinger¹, Brooke Z. Cohen¹, Kristina Deonaraine¹, Devyn Zaminski¹, Maria Dall'Era³, Kenneth Kalunian⁴, Andrea Fava⁵, H. Michael Belmont¹, Ming Wu¹, Chaim Putterman⁶, Jennifer Anolik⁷, Jennifer L. Barnas⁷, Betty Diamond⁸, Anne Davidson⁸, David Wofsy³, Diane Kamen⁹, Judith A. James¹⁰, Joel M. Guthridge¹⁰, William Apruzzese¹¹, Deepak A. Rao¹², Michael H. Weisman¹³, The Accelerating Medicines Partnership in RA/SLE Network, Michelle Petri⁵, Jill Buyon^{1†} and Richard Furie^{8†}

Abstract

Background Leveraging the Accelerating Medicines Partnership (AMP) Lupus Nephritis (LN) dataset, we evaluated longitudinal patterns, rates, and predictors of response to standard-of-care therapy in patients with lupus nephritis.

Methods Patients from US academic medical centers with class III, IV, and/or V LN and a baseline urine protein/creatinine (UPCR) ratio \geq 1.0 (n=180) were eligible for this analysis. Complete response (CR) required the following: (1) UPCR < 0.5; (2) normal serum creatinine (\leq 1.3 mg/dL) or, if abnormal, \leq 125% of baseline; and (3) prednisone \leq 10 mg/day. Partial response (PR) required the following: (1) > 50% reduction in UPCR; (2) normal serum creatinine or, if abnormal, \leq 125% of baseline; and (3) prednisone dose \leq 15 mg/day.

Results Response rates to the standard of care at week 52 were CR = 22.2%; PR = 21.7%; non-responder (NR) = 41.7%, and not determined (ND) = 14.4%. Only 8/180 (4.4%) patients had a week 12 CR sustained through week 52. Eighteen (10%) patients attained a week 12 PR or CR and sustained their responses through week 52 and 47 (26.1%) patients achieved sustained PR or CR at weeks 26 and 52. Week 52 CR or PR attainment was associated with baseline UPCR > 3 (OR_{adj} = 3.71 [95%CI = 1.34–10.24]; p = 0.012), > 25% decrease in UPCR from baseline to week 12 (OR_{adj} = 2.61 [95%CI = 1.07–6.41]; p = 0.036), lower chronicity index (OR_{adj} = 1.33 per unit decrease [95%CI = 1.10–1.62]; p = 0.003), and positive anti-dsDNA antibody (OR_{adj} = 2.61 [95%CI = 0.93–7.33]; p = 0.069).

Conclusions CR and PR rates at week 52 were consistent with the standard-of-care response rates observed in prospective registrational LN trials. Low sustained response rates underscore the need for more efficacious therapies

 $^\dagger \text{Peter M. Izmirly, Mimi Y. Kim, Jill Buyon and Richard Furie contributed equally to this work.}$

*Correspondence:
Peter M. Izmirly
Peter.Izmirly@nyumc.org
Full list of author information is available at the end of the article



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

and highlight how critically important it is to understand the molecular pathways associated with response and non-response.

Keywords Lupus nephritis, Systemic lupus erythematosus (SLE), Outcome, Renal biopsy

Background

The Accelerating Medicines Partnership (AMP) RA/SLE Network was established with the goal of applying new technologies, such as single-cell RNA sequencing of diseased kidney tissue, to improve diagnostic and therapeutic tools that would ultimately enhance lupus nephritis (LN) outcomes [1]. The AMP LN cohort, initiated in the United States (US) through the multi-center enrollment of patients with LN undergoing standard-of-care kidney biopsies, reflects real-world management and outcomes of a diverse population. In a prior publication, Deonaraine et al. provided reassurances regarding the safety of obtaining kidney tissue for AMP research during clinically indicated biopsies [2]. In another analysis of the AMP dataset, Carlucci et al. noted a high frequency of proliferative as well as membranous nephritis in enrolled AMP patients with baseline levels of proteinuria lower (urine protein/creatinine ratios between 0.5 and 1) than the typical threshold required for inclusion in registrational LN clinical trials [3].

In this interrogation of the AMP dataset, we determined the percentages of patients who attained prespecified definitions of partial or complete responses at specific visits over 1 year of treatment follow-up and examined the longitudinal patterns of response. In addition, clinical and laboratory characteristics associated with clinical responses were identified. In contrast to global LN clinical trials, the AMP LN cohort affords an opportunity to generate outcome data representative of a US multicenter, multi-racial, multi-ethnic real-world experience.

Methods

Patient population

Patients with LN undergoing kidney biopsies as part of the standard of care were eligible to enroll in the prospective AMP LN study. The decision to biopsy was at the discretion of the treating rheumatologist or nephrologist to confirm suspected lupus nephritis de novo, an activity not responding to treatment, or relapse of disease. Inclusion in AMP required the following: (1) age \geq 18; (2) fulfillment of the revised American College of Rheumatology [4, 5] or the Systemic Lupus Erythematosus International Cooperating Clinics [6] classification criteria for SLE; (3) a urine protein/creatinine ratio (UPCR) > 0.5 at the time of biopsy. For the analyses reported herein,

the classification of responder status was restricted to patients with baseline random or 24-h UPCR \geq 1.0 since for patients with ratios between 0.5 and 0.999, proteinuric response has not been defined. Only patients with renal biopsies that demonstrated the International Society of Nephrology/Renal Pathology Society (ISN/RPS) classes III, IV, V, or combined III or IV with V read by the pathologist at each participating site were considered in this analysis [7, 8]. Exclusion criteria included the following: (1) a history of kidney transplant, (2) rituximab treatment within 6 months of biopsy, (3) pregnancy at the time of biopsy. The study protocol was approved by the institutional review boards and ethics committees of participating sites in adherence with the Declaration of Helsinki.

Baseline demographics from a predetermined set of categories, including self-reported race (Asian, Black, White, Other)/ethnicity (Hispanic, non-Hispanic) as required for NIH-funded studies, and clinical characteristics were recorded at the time of biopsy. Laboratory tests and medications were documented at each visit (baseline, week 12, week 26, and week 52) and were performed at the participating sites. Given medication changes occurred after the baseline visit in response to receipt of the kidney biopsy results, we chose the week 12 treatment to represent the induction regimen. For steroids, the higher dose at either baseline or week 12 was considered the induction dose for similar reasons. Pulse steroids were also captured separately.

Outcomes

Complete response (CR) required the following: (1) UPCR < 0.5; (2) normal creatinine (\leq 1.3 mg/dL) or, if abnormal, \leq 125% of baseline; and (3) prednisone \leq 10 mg/day at the time of the study visit. Partial response required the following: (1) > 50% reduction in UPCR; (2) normal creatinine (\leq 1.3 mg/dL) or, if abnormal, \leq 125% of baseline; and (3) prednisone dose \leq 15 mg/day at the time of the study visit. Patients who did not achieve a CR or PR at the specific timepoints were considered non-responders (NR) or not determined (ND) if data were missing. These response definitions were based on the ACCESS Trial [9]. In agreement with the ACCESS trial, we specifically decided not to include the microscopic review of the urine sediment given the absence of uniformity across sites in assessing urinary sediment

and the challenge of attribution especially in a population of young women. The prednisone threshold for CR at ≤ 10 mg prednisone was also based on the ACCESS trial. However, the ≤ 15 mg prednisone maximum for defining PR was agreed upon unanimously by the site investigators.

Although proteinuria was measured by either a UPCR on spot urine or a timed urine collection, consistency of the method across the study for an individual was required. While determination from a timed urine collection was preferred, if this method was not performed at all time points for an individual participant, calculations from spot urine were utilized.

Statistical analysis

Descriptive statistics are presented as mean and standard deviation or median and interquartile range for continuous variables and frequencies for categorical variables. Pairwise agreement between response status at different time points was estimated by computing the kappa statistic. Logistic regression was performed to identify variables that independently discriminated persistent responders and never responders and estimate adjusted odds ratios (OR_{adi}). Given the small number of patients who had a CR or PR at all three follow-up visits, persistent responders were defined as those patients who achieved CR or PR at both 26 and 52 weeks; never responders were patients who did not achieve either CR or PR at any visit. In addition, logistic and multinomial logistic regression models were fit to the data to identify independent predictors of response status at 52 weeks only. Variable selection during model development was based on both statistical and clinical considerations, but the final model included only those variables that remained significant at the p < 0.10 level (a more liberal threshold for retaining variables in the final model was applied given the limited number of events). In addition to those variables listed in Table 1, potential predictors included baseline creatinine (>1.3 vs \leq 1.3), protein decreasing by 25% at 12 weeks, membranous vs proliferative and class III + V/IV + V biopsies, and induction prednisone dose (≥ 30 mg, < 30 and > 10 mg, and ≤ 10 mg). Missing data in the logistic regression analysis was handled using list-wise deletion. Sensitivity analysis was also performed based on non-responder imputation and multiple imputation (MI) with 40 imputed data sets. The MI model included the outcome variable, predictors from all logistic regression models, and several additional auxiliary variables (prednisone use, activity index, creatine level). All analyses were performed in SAS, version 9.4.

Table 1 Demographics and baseline characteristics of patients with baseline UPCR > 1

Age, mean (SD) Ethnicity: Hispanic Race Asian Black Asian Black Other/unknown First biopsy UPCR, mean [IQR] Nephrotic proteinuria Serum creatinine mg/dL, mean [range] (n = 179) High serum creatinine (n = 179) Low C3 (n = 178) Low C4 (n = 178) Serum albumin g/dL, mean [range] (N = 171) Positive anti-dsDNA (n = 176) Biopsy class [III] [IV] [IV] [IV] S1 (28.3%) [III][V] [IV] Activity Index, mean [range] (n = 143) S9 (35.2 (11.4) 29 (16.1%) 59 (32.7%) 84 (22.2%) 62 (34.0%) 92 (12.2%) 62 (34.0%) 92 (45.6%) 82 (45.6%) 82 (45.6%) 82 (45.6%) 82 (45.6%) 82 (45.6%) 82 (45.6%) 82 (45.6%) 83 (1.0-4.74] 94 (25.7%) 116 (65.2%) 117 (10-4.7] 95 (10-4.7] 96 (20.0%) 118 (20.0%) 119 (10-18) 84 (20.0%) 119 (10-18) 85 (11.4) 85 (11.4) 86 (20.0%) 86 (20.0%) 87 (10-18)	with baseline open 2 i	
Age, mean (SD) 35.2 (11.4) Ethnicity: Hispanic 59 (32.7%) Race Asian 29 (16.1%) Black 76 (42.2%) White 53 (29.4%) Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60-4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n=179) 1.25 [0.4-7.4] High serum creatinine (n = 179) ^a 46 (25.7%) Low C3 (n = 178) ^a 116 (65.2%) Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [IV] 35 (19.4%) [V] 51 (28.3%) [IV] 3 (1.7%) [IV][V] 3 (20.0%) [V] 51 (28.3%) [IV][V] 3 (20.0%) [V] 51 (28.3%) [IV][V] 3 (40.7%) [V] 51 (Demographics (n = 180)	
Ethnicity: Hispanic 59 (32.7%) Race Asian 29 (16.1%) Black 76 (42.2%) White 53 (29.4%) Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60-4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n=179) 1.25 [0.4-7.4] High serum creatinine (n = 179) ^a 46 (25.7%) Low C3 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [IV] 3 (1.7%) [IV] 3 (20.0%) [IV]	Sex: female	156 (86.7%)
Asian 29 (16.1%) Black 76 (42.2%) White 53 (29.4%) Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60–4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n=179) 1.25 [0.4–7.4] High serum creatinine (n=179) 46 (25.7%) Low C3 (n=178) 116 (65.2%) Low C4 (n=178) 116 (65.2%) Low C4 (n=178) 116 (65.2%) Serum albumin g/dL, mean [range] (N=171) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [IIII] 30 (16.7%) [IV] 51 (28.3%) [IIII][V] 36 (20.0%) [IV][V] 36 (20.0%) [IV][V] Activity Index, mean [range] (n=143) 5.4 [0–18] Chronicity Index, mean [range] (n=143) 3.3 [0–10] Extra renal activity on hybrid SELENA-SLEDAI* Medications* Hydroxychloroquine Daily average dose [range] 718 (24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Age, mean (SD)	35.2 (11.4)
Asian 29 (16.1%) Black 76 (42.2%) White 53 (29.4%) Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60–4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n = 179) 1.25 [0.4–7.4] High serum creatinine (n = 179)a 46 (25.7%) Low C3 (n = 178)a 116 (65.2%) Low C4 (n = 178)a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0–4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 36 (20.0%) [IV][V] 36 (20.0%) [IV][V] Activity Index, mean [range] (n = 143) 5.4 [0–18] Chronicity Index, mean [range] (n = 143) 3.3 [0–10] Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications Hydroxychloroquine 137 (76.1%) Daily average dose [range] 135 (11.7%) Puse steroids 116 (6.44%) Daily average dose [range] Mycophenolate mofetil 15 aliy average dose [range] Mycophenolate mofetil 16 (6.44%) Daily average dose [range] Mycophenolate mofetil 17 fing [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus Belimumab 4 (2.2%)	Ethnicity: Hispanic	59 (32.7%)
Black White Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n = 179) 1.25 [0.4–7.4] High serum creatinine (n = 179) 1.26 (57.9%) Low C3 (n = 178) ^a 116 (65.2%) Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) Positive anti-dsDNA (n = 176) Biopsy class [III] 30 (16.7%) [IV] (V] 51 (28.3%) [III][V] (V] 51 (28.3%) [III][V] (V] (V] (V] (V] (V] (V] (V] (V] (V] (Race	
White 53 (29.4%) Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60−4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n=179) 1.25 [0.4−7.4] High serum creatinine (n=179)³ 46 (25.7%) Low C3 (n=178)³ 102 (57.3%) Serum albumin g/dL, mean [range] (N=171) 3.1 [1.0−4.7] Positive anti-dsDNA (n=176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [IV] 51 (28.3%) [IV] 3 (1.7%) [IV][V] 3 (20.0%) [V] 25 (13.9%) Activity Index, mean [range] (n=143) 5.4 [0-18] Chronicity Index, mean [range] (n=143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications* Hydroxychloroquine 137 (76.1%) Daily average dose [range] 24.4 mg [2.5-120] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120]	Asian	29 (16.1%)
Other/unknown 22 (12.2%) First biopsy 62 (34.0%) UPCR, mean [IQR] 3.5 [1.60–4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n=179) 1.25 [0.4–7.4] High serum creatinine (n=179)a 46 (25.7%) Low C3 (n=178)a 116 (65.2%) Low C4 (n=178)a 102 (57.3%) Serum albumin g/dL, mean [range] (N=171) 3.1 [1.0–4.7] Positive anti-dsDNA (n=176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 36 (20.0%) [IV][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n=143) 5.4 [0–18] Chronicity Index, mean [range] (n=143) 3.3 [0–10] Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications* Hydroxychloroquine 137 (76.1%) Daily average dose [range] 24.4 mg [2.5–120] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.5 mg [2.5–120] Pulse steroids 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Black	76 (42.2%)
First biopsy UPCR, mean [IQR] Nephrotic proteinuria Serum creatinine mg/dL, mean [range] (n = 179) High serum creatinine (n = 179) ^a Low C3 (n = 178) ^a Low C4 (n = 178) ^a Low C4 (n = 178) ^a Serum albumin g/dL, mean [range] (N = 171) Positive anti-dsDNA (n = 176) Biopsy class [III] [IV] [IV] [V] [S1 (28.3%) [III][V] [IV] Activity Index, mean [range] (n = 143) Chronicity Index, mean [range] (n = 143) Extra renal activity on hybrid SELENA-SLEDAl ^b Medications ^c Hydroxychloroquine Daily average dose [range] Prednisone/methylprednisolone Daily average dose [range] Pulse steroids Mycophenolate mofetil Daily average dose [range] Mycophenolic acid Daily average dose [range] Cyclophosphamide Azathioprine Facrolimus Belimumab 4 (2.2%)	White	53 (29.4%)
UPCR, mean [IQR] 3.5 [1.60–4.38] Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n = 179) 1.25 [0.4–7.4] High serum creatinine (n = 179) ^a 46 (25.7%) Low C3 (n = 178) ^a 116 (65.2%) Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0–4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 36 (20.0%) [IV][V] 37 (76.1%) [IV][V] 37 (48.3%) Medications ^c 137 (76.1%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] <td< td=""><td>Other/unknown</td><td>22 (12.2%)</td></td<>	Other/unknown	22 (12.2%)
Nephrotic proteinuria 82 (45.6%) Serum creatinine mg/dL, mean [range] (n = 179) 1.25 [0.4-7.4] High serum creatinine (n = 179) ^a 46 (25.7%) Low C3 (n = 178) ^a 116 (65.2%) Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 36 (20.0%) [IV][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 5.4 [0-18] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 24.4 mg [2.5-120] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000]	First biopsy	62 (34.0%)
Serum creatinine mg/dL, mean [range] (n = 179) High serum creatinine (n = 179)a 46 (25.7%) Low C3 (n = 178)a 116 (65.2%) Low C4 (n = 178)a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 3 (1.7%) [III][V] 3 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7-800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000] Mycophenolic acid 8 (4.4%) Daily average dose	UPCR, mean [IQR]	3.5 [1.60-4.38]
High serum creatinine (n = 179) ^a 46 (25.7%) Low C3 (n = 178) ^a 116 (65.2%) Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAl ^b 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) 356.1 mg [85.7-800] Prednisone/methylprednisolone 135 (75.0%) 24.4 mg [2.5-120] Pulse steroids 116 (6.44%) 2435.3 mg [500-3000] Mycophenolic acid 2435.3 mg [500-3000] Mycophenolic acid 2445.3 mg [500-2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab	Nephrotic proteinuria	82 (45.6%)
Low C3 (n = 178) ^a Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) Positive anti-dsDNA (n = 176) Biopsy class [III] 30 (16.7%) [IV] 51 (28.3%) [III][IV] 36 (20.0%) [IV][V] Activity Index, mean [range] (n = 143) Chronicity Index, mean [range] (n = 143) Extra renal activity on hybrid SELENA-SLEDAl ^b Medications ^c Hydroxychloroquine Daily average dose [range] Prednisone/methylprednisolone Daily average dose [range] Pulse steroids Mycophenolate mofetil Daily average dose [range] Mycophenolic acid Daily average dose [range] Mycophenolic acid Daily average dose [range] Cyclophosphamide Azathioprine Tacrolimus Belimumab 116 (65.2%) 102 (57.3%) 3.1 [1.0-4.7] 3.1 [1.0-4.7] 3.1 [1.0-4.7] 4.4 (1.1.3%) 1.5 (28.3%) 1.7 (6.1%) 3.3 [0-10] 1.7 (76.1%) 3.3 [0-10] 1.7 (76.1%) 1.7 (7	Serum creatinine mg/dL, mean [range] $(n = 179)$	1.25 [0.4–7.4]
Low C4 (n = 178) ^a 102 (57.3%) Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0−4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][IV] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0−18] Chronicity Index, mean [range] (n = 143) 87 (48.3%) Medications ^c 448.3%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	High serum creatinine $(n = 179)^a$	46 (25.7%)
Serum albumin g/dL, mean [range] (N = 171) 3.1 [1.0-4.7] Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][IV] 36 (20.0%) [IV][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 87 (48.3%) Medications ^c 49/4 (48.3%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7-800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360-2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab	Low C3 $(n = 178)^a$	116 (65.2%)
Positive anti-dsDNA (n = 176) 124 (70.5%) Biopsy class [III] 30 (16.7%) [IV] 35 (19.4%) [V] 51 (28.3%) [III][IV] 3 (1.7%) [III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 87 (48.3%) Medications ^c 47 (48.3%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7-800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360-2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Low C4 $(n = 178)^a$	102 (57.3%)
Biopsy class	Serum albumin g/dL, mean [range] ($N = 171$)	3.1 [1.0-4.7]
[III] [IV] [IV] [ST 1 (28.3%) [III] [IV] [IV] [III] [V] [III] [V] [III] [V] [IV] [V] [IV] [V] [V] [V] [V] [V] [V] [V] [V] [V] Activity Index, mean [range] (n = 143) [Chronicity Index, mean [range] (n = 143) [Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications Hydroxychloroquine Daily average dose [range] Prednisone/methylprednisolone Daily average dose [range] Pulse steroids Mycophenolate mofetil Daily average dose [range] Mycophenolic acid Daily average dose [range] Cyclophosphamide Azathioprine Facrolimus Belimumab As (16.7%) As (19.4%) As (10.5%) As (19.4%) As	Positive anti-dsDNA ($n = 176$)	124 (70.5%)
[IV] 35 (19.4%) [V] 51 (28.3%) [III][IV] 3 (1.7%) [III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 16 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Biopsy class	
[V] 51 (28.3%) [III][IV] 3 (1.7%) [III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAlb 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	[III]	30 (16.7%)
[III][IV] 3 (1.7%) [III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c 448.3%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	[IV]	35 (19.4%)
[III][V] 36 (20.0%) [IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c 48.3%) Hydroxychloroquine Daily average dose [range] 137 (76.1%) Daily average dose [range] 356.1 mg [85.7-800] Prednisone/methylprednisolone Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil Daily average dose [range] 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000] Mycophenolic acid Baily average dose [range] 36 (4.4%) Daily average dose [range] 1215 mg [360-2880] Cyclophosphamide Azathioprine Gazenius 24 (13.3%) Tacrolimus Belimumab 4 (2.2%)	[V]	51 (28.3%)
[IV][V] 25 (13.9%) Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c 137 (76.1%) Hydroxychloroquine 135 (76.1%) Daily average dose [range] 356.1 mg [85.7-800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5-120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500-3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360-2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	[III][IV]	3 (1.7%)
Activity Index, mean [range] (n = 143) 5.4 [0-18] Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAI ^b 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 16.6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 224 (13.3%) Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	[III][V]	36 (20.0%)
Chronicity Index, mean [range] (n = 143) 3.3 [0-10] Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c 137 (76.1%) Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	[IV][V]	25 (13.9%)
Extra renal activity on hybrid SELENA-SLEDAIb 87 (48.3%) Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Activity Index, mean [range] (n = 143)	5.4 [0-18]
Medications ^c Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Chronicity Index, mean [range] (n = 143)	3.3 [0-10]
Hydroxychloroquine 137 (76.1%) Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Extra renal activity on hybrid SELENA-SLEDAI b	87 (48.3%)
Daily average dose [range] 356.1 mg [85.7–800] Prednisone/methylprednisolone 135 (75.0%) Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Medications ^c	
Daily average dose [range] 24.4 mg [2.5–120] Pulse steroids 21 (11.7%) Mycophenolate mofetil 116 (6.44%) Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	, , ,	
Daily average dose [range] 2435.3 mg [500–3000] Mycophenolic acid 8 (4.4%) Daily average dose [range] 1215 mg [360–2880] Cyclophosphamide 24 (13.3%) Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Daily average dose [range]	24.4 mg [2.5-120]
Azathioprine 6 (3.3%) Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Daily average dose [range] Mycophenolic acid	2435.3 mg [500–3000] 8 (4.4%)
Tacrolimus 19 (10.6%) Belimumab 4 (2.2%)	Cyclophosphamide	24 (13.3%)
Belimumab 4 (2.2%)	Azathioprine	6 (3.3%)
(1.17)	Tacrolimus	19 (10.6%)
Leflunomide 1 (0.6%)	Belimumab	4 (2.2%)
	Leflunomide	1 (0.6%)

Unless otherwise indicated, variables had data available for all 180 patients *UPCR* urine protein/creatinine ratio, *anti-dsDNA* anti-double-stranded DNA autoantibodies, *SELENA-SLEDAI* Safety of Estrogens in Lupus Erythematosus: National Assessment- Systemic Lupus Erythematosus Disease Activity Index

^a Classified by local laboratory cutoff

^b Includes all hybrid SELENA-SLEDAI domains that include clinical activity excluding serologic and renal urine activity

^c Captured at week 12 visits, for steroids the higher dose at two visits (baseline and week 12) was considered induction dosing given patients who had their doses increased after the biopsy would not be captured at the baseline visit

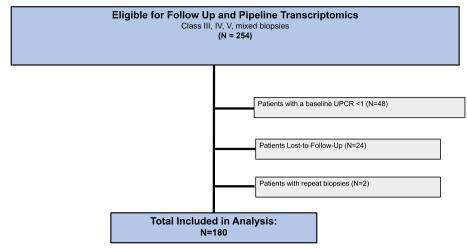


Fig. 1 Flow diagram of study enrollment

Results

Baseline characteristics

One hundred eighty patients met the inclusion criteria (Fig. 1). Of these, 86.7% were women, 29.4% were White, and 32.7% were Hispanic (Table 1). The mean age was 35.2 (SD 11.4) years. Using administered medications at week 12 to capture induction therapy, the majority (64%) were treated with mycophenolate mofetil, and 13% with cyclophosphamide. Seventy-five percent of the cohort received steroids with an average dose of prednisone equivalent of 24.4 mg, and 76% were taking hydroxychloroquine. Biopsy classes were as follows: III = 16.7%, IV = 19.4%, V = 28.3%, and III + V/IV + V = 33.9%. Sixtysix percent of patients had a previous biopsy. Average baseline creatinine was 1.25. A positive anti-dsDNA antibody (measured locally) was present in 70.5%, 65.2% had a low C3 level, and 57.3% had a low C4 level. The average baseline UPCR was 3.5. Overall, 48.3% of the 180 patients had extra renal activity on the hybrid SELENA- SLEDAI at baseline.

Longitudinal patterns of response

The response rates and graphical heat map displays of responses at each visit are shown in Fig. 2A. Response rates at week 52 were as follows: CR=22.2%; PR=21.7%; NR=41.7%; and ND=14.4%. Only 8/180 (4.4%) of patients had a confirmed week 12 CR response sustained through week 52. Eighteen (10%) patients attained a PR or CR at week 12 and sustained their responses through week 52, and 47 (26.1%) patients achieved a PR or CR at week 26, which was sustained at week 52. Overall, 40/180 (22.2%) were confirmed NR at all time points, which increased to 67 (37.2%) when non-responder imputation (NRI) was applied for missing data (Supplemental

Fig. 1A). Figure 2B is a display restricted to patients ($n\!=\!118$) for whom responder status was available at all time points. Although not used in further analysis of renal responder status, applying less stringent definitions of proteinuric responses, independent of creatinine or prednisone dose at 52 weeks, 69/180 (38.3%) had a UPCR \leq 0.8 and 62/180 (34.4%) had a UPCR \leq 0.7 compared to 48/180 (26.7%) attaining a UPCR of \leq 0.5. The most common reason for regressing at 52 weeks from an initial CR/PR was the return of proteinuria above the response definition.

Based on the observed data, there was a fair agreement between response status at weeks 12 and 26 (kappa=0.41 [95% CI 0.27–0.56]) and between weeks 26 and 52 (kappa=0.36 [95% CI=0.21–0.51]). As expected, agreement in response status between weeks 12 and 52 was weaker (kappa=0.16 [95% CI=0.015–0.30]) (Table 2). When NRI was used to handle missing data, agreement in response status across visits was similar or slightly lower (Supplementary Table 1).

Patient characteristics associated with persistent responses at weeks 26 and 52

As shown in Table 3, logistic regression analysis indicated that the following patient characteristics independently favored CR or PR responses at both weeks 26 and 52 (persistent responders) compared to NR at all time points: a > 25% decrease in UPCR between baseline and week 12 (OR_{adj}=7.37 [95% CI=2.31–23.49]; p<0.001), positive anti-dsDNA antibody (OR_{adj}=4.70 [95% CI=1.19–18.51]; p=0.027), first biopsy (OR_{adj}=3.12 [95% CI=0.89–10.89]; p=0.075) and no use of cyclophosphamide for induction (OR_{adj}=5.08 [95% CI=0.80–32.26];

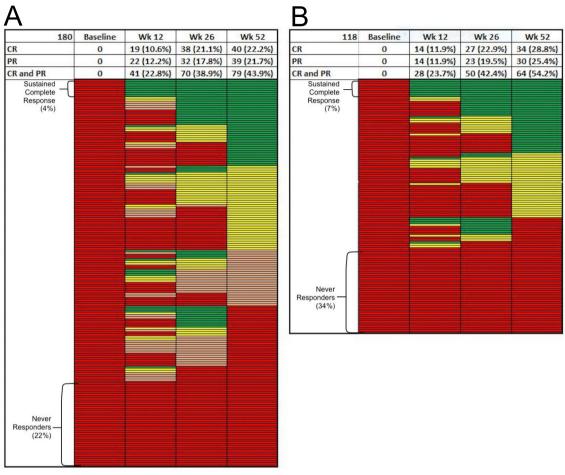


Fig. 2 Response rates and graphical heat map displays of responses at each visit

 Table 2
 Agreement in response status (complete or partial) across visits

	% Response at both visits	% Non-response at both visits	% Discordant response status	Kappa (95% CI) for agreement in response status across visits
Week 12 and week 26	20.8%	51.5%	27.7%	0.41 (0.27, 0.56)
Week 12 and week 52	16.5%	40.9%	42.5%	0.16 (0.015, 0.30)
Week 26 and week 52	33.8%	33.8%	32.4%	0.36 (0.21, 0.51)

Assuming not determined = missing

p=0.084). Estimated odds ratios observed in sensitivity analyses in which missing data were addressed with non-response imputation and multiple imputation showed similar results (Supplementary Table 2).

Patient characteristics associated with response at week 52 Patient characteristics favoring CR or PR responses

compared to NR at week 52 from logistic regression analysis were as follows: UPCR>3 at baseline $(OR_{adj}=3.71 \quad [95\%CI=1.3-10.24]; \quad p=0.012),>25\%$ decrease in UPCR from baseline to week 12 $(OR_{adj}=2.61)$

[95%CI=1.07–6.41]; p=0.036), lower chronicity index (OR_{adj}=1.33 per unit decrease [95%CI=1.10–1.62]; p=0.003), and a positive anti-dsDNA antibody (OR_{adj}=2.61 [95%CI=0.93–7.33]; p=0.069) (Table 4). Sensitivity analyses using methods to address missing data again showed similar trends, but the estimated odds ratio of UPCR > 3 was lower with multiple imputation (Supplementary Table 3). Limiting these analyses to Class V only, estimated odds ratios of predictor variables were larger but less statistically significant because of the smaller sample size (Supplementary Table 4).

Table 3 Predictors of response (complete or partial) at both weeks 26 and 52 versus no response at all visits from logistic regression

Predictor variable	Odds ratio estimate (95% confidence interval)	<i>P</i> value
First biopsy	3.12 (0.89–10.89)	0.075
Anti-dsDNA antibody positive	4.70 (1.19-18.51)	0.027
No Cyclophosphamide induction	5.08 (0.80-32.26)	0.084
UPCR > 25% decrease from baseline to week 12	7.37 (2.31–23.49)	< 0.001

Based on available data for responder adjudication and all covariates *UPCR* urine protein/creatine ratio, *anti-dsDNA* antibody anti-double-stranded DNA autoantibody

In exploratory analyses, multinomial logistic regression with week 52 response status considered as three separate categories—CR, PR, and NR (in contrast to combining CR and PR)—suggested baseline positive anti-dsDNA antibody, > 25% decrease in UPCR from baseline to week 12, and chronicity index discriminated CR versus NR, while UPCR > 3 at baseline discriminated PR versus NR (Supplementary Table 5).

Discussion

The AMP LN cohort provided outcome data representative of a large US multicenter, multi-racial, multi-ethnic real-world experience. In 180 patients, the response rates at week 52 were similar to those observed in pivotal FDA trials with complete response in only a fifth of the cohort and nearly half non-responders. Very few patients had a week 12 CR response sustained through the entire year of the study, and only 26% attained a PR or CR at both week 26 and week 52. Agreement in response status between 12 and 52 weeks was low. A>25% decrease in UPCR from baseline to week 12 and/or a baseline positive antidsDNA antibody predicted both persistent CR or PR responses at weeks 26 and 52 and a CR or PR at 52 weeks only. First biopsy and/or no use of cyclophosphamide induction was only associated with sustained responses at weeks 26 and 52, whereas a baseline UPCR>3 and lower chronicity index were only associated with CR or PR responses at 52 weeks.

In BLISS-LN [10], a phase III 2-year study of belimumab in patients with proliferative and/or membranous nephritis, the probabilities of achievement of the primary endpoint (Primary Efficacy Renal Response) as well as secondary endpoint (Complete Renal Response) were determined. While entry criteria, endpoints, and treatment interventions differed from the AMP study, achievement in BLISS-LN of sustained CRR, which most closely approximates the AMP endpoint, was approximately 13% at 1 year in the placebo group.

The CR rate (22.2%) in AMP was very similar to those reported in LN clinical trials despite the differences in definitions across studies. In recently published clinical trials of belimumab, voclosporin, and obinutuzumab, CR rates of 20% (week 104), 23% (week 52), and 23% (week 52) in the placebo/standard of care arms, respectively, were observed [10–12]. PR rates of 17% (week 104), 50% (week 52), and 13% (week 52) in the placebo/standard of care arms were observed in the belimumab, voclosporin, and obinutuzumab studies, respectively, compared with 21.7% in AMP [10–12].

There are several limitations that could have influenced the results of this study. Doses of medications were recorded only at the respective visits, and thus there was likely an underestimation of the highest dose of administered steroids. Furthermore, potential changes in immunosuppression between visits such as intravenous regimens may not have been captured. As a result, changes in medications between visits, particularly after 26 weeks when a patient could have been considered an induction responder, were not analyzed in predictors of responses. The upper limit of normal for creatinine in some laboratories may be lower than 1.3 mg/dL, and it is acknowledged that given the high frequency of young adult females, the level chosen may be abnormal in this population. Applying a lower normal value would have resulted in even lower response outcomes. The small number of patients achieving a sustained CR or sustained PR precluded analyses of predictors of persistent response. Although of interest, there were too few patients to analyze those that initially responded but lost

Table 4 Predictors of week 52 response (complete or partial) versus no response from logistic regression analysis

Predictor variable	Odds ratio estimate (95% confidence interval)	P value
Anti-dsDNA antibody positive	2.61 (0.93–7.33)	0.069
UPCR > 25% decrease from baseline to week 12	2.61 (1.07–6.41)	0.036
Chronicity Index per unit decrease	1.33 (1.10–1.62)	0.003
UPCR > 3 at baseline	3.71 (1.34–10.24)	0.012

Based on available data for responder adjudication and all covariates

response at 52 weeks. Missing data is also a limitation although this was addressed using methods as previously described [13]. The negative association of cyclophosphamide with renal response may have been due to confounding by indication, especially in a cohort where the majority of patients had a prior history of LN. Complete response with proteinuria < 0.5 was a predefined outcome at the start of this study which began in 2014 to be consistent with current clinical trials at that time and will be used for future AMP biomarker studies [14]. Since then, there has been emerging evidence that proteinuria < 0.8 at 12 months is predictive of favorable long-term renal outcomes [15-17]. In this study, even liberalizing the definition of response to < 0.8 independent of creatinine or prednisone dose still resulted in a poor response rate at 38%.

The strengths of this study are that data were generated from academic institutions with familiarity in the treatment of lupus nephritis. This study represents real-world standard of care and includes sicker patients who otherwise would be excluded from clinical trials. In addition, the AMP cohort comprised a diverse racial and ethnic group of patients. This study also evaluated sustained response [18] as well as predictors of response, items which have not been commonly evaluated in LN trials.

Conclusions

In summary, clinical data from the AMP Lupus Network revealed rates of 52-week CR, PR, and CR and PR that were consistent with standard of care/placebo response rates from recently conducted LN trials. Low sustained CR rates not only underscore the need for more efficacious therapies but highlight how critically important it is to understand the molecular pathways that are associated with response and non-response.

Abbreviations

AMP Accelerating Medicines Partnership

CR Complete response
dsDNA Double-stranded DNA
LN Lupus nephritis
NR Non-responder
ND Not determined
PR Partial response

SELENA-SLEDAI Safety of Estrogens in Lupus Erythematosus: National

Assessment-Systemic Lupus Erythematosus Disease Activ-

ity Index

UPCR Urine protein to creatinine ratio

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s13075-024-03275-z.

Additional file 1: Supplemental Table 1. Agreement in response status (complete or partial) across visits using non-responder imputation for

missing data. Supplemental Table 2. Predictors of response (complete or partial) at both weeks 26 and 52 versus no response at all visits from logistic regression using non-responder imputation for missing response data and multiple imputation for missing covariate data. Supplemental **Table 3.** Predictors of response (complete or partial) at week 52 versus no response from logistic regression analysis using non-responder imputation for missing response data and multiple imputation for missing covariate data. Supplemental Table 4. Predictors of response (complete or partial) at week 52 versus no response from logistic regression analysis for Class V cases only. Supplemental Table 5. Predictors of week 52 response using multinomial regression with available data. Supplemental Figure 1. Temporal patterns in the response status of patients with systemic lupus erythematosus receiving standard of care therapy employing nonresponder imputation for missing data for 180 patients included. Green indicates complete response, yellow indicates partial response and red indicates no response.

Acknowledgements

The authors would like to thank Benjamin Wainwright for assistance in preparing this manuscript. Some data were previously presented in preliminary form as an abstract at the 2021 annual meeting of the American College of Rheumatology. The funding source had no role in the drafting of this manuscript or the analyses presented therein.

The Accelerating Medicines Partnership in RA/SLE Network

AMP RA Network

Rochester

Jennifer Anolik Darren Tabechian Ralf Thiele Jennifer Hossler Brendan Boyce Nida Meednu Javier Rangel-Moreno Christopher Ritchlin

Hospital for Special Surgery (HSS)

Vivian Bykerk
Laura Donlin
Susan Goodman
Lionel Ivashkiv
Alessandra Pernis
Ed DiCarlo
Dana Orange
John Carrino
Oganna (Kenny) Nwawka
Endo Yoshimi
Rahul Satija
Lionel Ivashkiv
Robert Darnell

University of Pittsburgh

Michael McNamara

Mark Figgie

Larry W. Moreland Mandy J. McGeachy Jay Kolls Aaron Wise Andrew Cordle

Feinstein/Northwell

Peter Gregersen Diane Horowitz

UK Birmingham (under Feinstein/Northwell)

Andrew D. Filer Jason Turner Holly Adams

UK London (under Feinstein/Northwell)

Costantino Pitzalis Stephen Kelly Rebecca Hands

Brigham and Women's Hospital

Michael Brenner Derrick Todd Kevin Wei Deepak Rao Fumitaka Mizoguchi

University of Colorado (EMORA)

V. Michael Holers Kevin D. Deane Jennifer A. Seifert Nirmal K. Banda

University of California San Diego (EMORA)

Gary S. Firestein David Boyle

Cedars Sinai (EMORA)

Michael H. Weisman Ami Ben-Artzi Lindsy Forbess

University of Massachusetts (EMORA)

Ellen Gravallese Karen Salomon-Escoto

Northwestern University (REASON under EMORA Network)

Harris Perlman Arthur Mandelin Emily Bacalao

Washington University (REASON)

Deborah Parks John Atkinson

Columbia University (REASON)

Joan Bathon

Mayo Clinic (REASON)

Eric Matteson

University of Alabama (REASON)

Louis Bridges Laura B. Hughes

Michigan (REASON)

David Fox Robert Ike

AMP SLE Network

Johns Hopkins

Michelle Petri Chun-Hao Lee Derek Fine Manny Monroy-Trujillo

Rochester

Jennifer Anolik Ummara Shah

Cedars

Michael Weisman Mariko Ishimori

New York University (NYU) (METRO)

Jill P. Buyon Robert M. Clancy Peter Izmirly Michael Belmont Amit Saxena Ming Wu Nicole Bornkamp

Albert Einstein College of Medicine (METRO)

Chaim Putterman Evan Der Beatrice Goilav Nicole Jordan Daniel Schwartz James Pullman

University of California San Francisco (PEARL)

David Wofsy Dawn Smilek Patti Tosta

Feinstein/Northwell (PEARL)

Betty Diamond

Michigan (PEARL)

Matthias Kretzler Celine C. Berthier

University of Cincinnati (PEARL)

F. Steve Woodle Dave Hildeman

Brigham and Women's Hospital (PEARL)

Michael Brenner Deepak Rao

Technology sites

STAMP (Stanford)

William Robinson Garry Nolan Veronica Gonzales

Brigham and Women's Hospital

Michael Brenner Deepak Rao Kevin Wei Jim Lederer Joshua Keegan Adam Chicoine Yanyan Liu Gerald Watts

Broad Institute

Nir Hacohen Arnon Arazi David Lieb Thomas Eisenhaure

Rockefeller (METRO)

Thomas Tuschl

AMP Operations Network

William Apruzzese (NIAMS)

Leadership Center (Stanford)

PJ Utz

Mina Rohani-Pichavant

DCMG

Rohit Gupta Holden Maecker

TRG (at Oklahoma Medical Research Foundation)

Judith A. James Joel M. Guthridge Wade DeJager Susan Macwana

SBG

Soumya Raychaudhuri Yvonne Lee Kamil Slowikowski Chamith Fonseka Fan Zhang Maria Guitierrez-Arcelus

NIH/NIAMS

Justine Buschman Jennifer Chi Su-Yau Mao Susana Serrate-Sztein Yan Wang

NIH/NIAID

Quan Chen John Peyman Ellen Goldmuntz

ImmPort

Patrick Dunn

Authors' contributions

P.M.I. conception, design, interpretation of data, drafter of manuscript and substantively revised it. M.Y.K. conception, design, interpretation of data, statistical analysis, substantively revised manuscript. prepared Tables 2, 3 and 4 and supplemental Tables. P.M.C. conceptualization; substantively revised manuscript. K.P. data acquisition, helped prepare Table 1 and prepared Fig. 1. B.Z.C. data acquisition, helped prepare Table 1. K.D. data acquisition, helped prepare Table 1. D.Z. data acquisition, prepared Table 1. M.D. conception, data acquisition. K.K. conception, data acquisition. A.F. data acquisition. H.M.B. data acquisition. M.H.W. data acquisition. C.P. data acquisition. J.A. data acquisition. J.B. data acquisition. B..D. data acquisition. A.D. data acquisition. D.W. data acquisition. D.K. data acquisition. J.A.J. data acquisition. J.M.G. data acquisition. W.A. data acquisition. DR data acquisition. M.W. data acquisition. M.P. data acquisition. J.P.B. conception, design, interpretation of data, substantively revised manuscript. R.F. conception, design, interpretation of data, substantively revised manuscript, prepared Fig. 2 and supplemental Figure. All authors have approved the submitted version of the manuscript and have agreed both to be personally accountable for the author's own contributions and to ensure that questions related to the accuracy or integrity of any part of the work, even ones in which the author was not personally involved, are appropriately investigated, resolved, and the resolution documented in the literature.

Funding

This work was supported by the Accelerating Medicines Partnership (AMP) in Rheumatoid Arthritis and Lupus Network. AMP is a public–private partnership (AbbVie Inc., Arthritis Foundation, Bristol-Myers Squibb Company, Foundation for the National Institutes of Health, GlaxoSmithKline, Janssen Research and Development, LLC, Lupus Foundation of America, Lupus Research Alliance, Merck Sharp & Dohme Corp., National Institute of Allergy and Infectious Diseases, National Institute of Arthritis and Musculoskeletal and Skin Diseases, Pfizer Inc., Rheumatology Research Foundation, Sanofi and Takeda Pharmaceuticals International, Inc. created to develop new ways of identifying and validating promising biological targets for diagnostics and drug development. Funding was provided through grants from the National Institutes of Health (UH2-AR067676, UH2-AR067677, UH2-AR067691, UH2-AR067681, UH2-AR067689, UH2-AR067690, UH2-AR067691, UH2-AR067694, and UM2-AR067678).

Availability of data and materials

The NIH is in the process of releasing the clinical datasets analyzed during the current study.

Declarations

Ethics approval and consent to participate

The study protocol was approved by the institutional review boards and ethics committees of participating sites in adherence with the Declaration of Helsinki.

Consent for publication

Not applicable.

Competing interests

The authors would like to disclose the following financial relationships: Izmirly, Peter Consultant: GSK, Momenta/Janssen; Kim, Mimi: None; Carlucci, Philip: none; Preisinger, Katherine: none; Zaminski, Devyn: none; Dall'Era, Maria: Consultant: GSK, Aurinia, AstraZeneca, Lilly, Biogen; Data Monitoring Committee: Janssen, Pfizer, Amgen; Kalunian, Kenneth: Advisor: AstraZeneca, RemeGen, BMS, Amgen, EquilliumBio, KezarBio, Aurinia, GSK, Roche/Genentech, Novarits, Horizon, Idorsia, Artivabio, Cabaletta, Merck, Alpine Immune Sciences, AbbVie, Vera Therapeutics, Aclaris Therapeutics, Gilead, Pfizer; Grants: Amgen, UCB, Acceleron, PreventionBio, Feinstein Institute, Alexion, NIH/MUSC, NIH/Immune Tolerance Network, Cugene, Kyowa Kirin, Daichi Sankyo; Fava, Andrea: none; Belmont, Michael: none; Wu, Ming: none; Putterman, Chaim: none; Anolik, Jennifer: Pending; Barnas, Jennifer: none; Diamond, Betty: Consultant: Nextcure, DBV, Alpine, iCell, Bayer, Magnolia, Kyverna, Nighthawk; Davidson, Anne: none; Wofsy, David: none; Kamen, Diane: Data safety and monitoring committee member: Alpine Immune Sciences; Data And Safety Monitoring Chair: Equillium; James, Judith: Advisor: GSK, Novartis; IP licensing and grant support: Progentec Biosciences; Guthridge, Joel: none; Apruzzese, William: Employment: Pfizer (began after this work concluded); Rao, Deepak: Consultant: GSK, AstraZeneca, Bristol-Myers Squibb, Pfizer; grant support unrelated to this work: Janssen, Bristol-Myers Squibb, Merck; Weisman, Michael: none; Petri, Michelle Consultant: Alexion, Amgen, AnaptysBio, Annexon Bio, Argenx, AstraZeneca, Axdev, Biogen, Boxer Capital, Cabaletto Bio, Caribou Biosciences Inc, CVS Health, Exo Therapeutics, Gilead Biosciences, GSK, Horizon Therapeutics, Idorsia Pharmaceuticals, Janssen, Kira Pharmaceuticals, Eli Lilly, MedShr, Momenta Pharmaceuticals, Nexstone Immunology, Nimbus Lakshmi, Proviant, Sanofi, Sinomab Biosciences, UCB; Invited speaker: Arthros-FocusMedEd, Aurinia; Data safety and monitoring board member: Emergent Biosolutions, IQVIA, Merck EMD Serono; grant support: Eli Lilly, Exagen, AstraZeneca, GSK, Thermofisher, Janssen, Aurinia; Buyon, Jill, Consultant: Bristol-Myers Squibb; GlaxoSmith Kline, Related Sciences, Ventus Therapeutics; Furie, Richard: Consultant: GSK, Genetech, Aurinia, Kezar.

Author details

¹New York University Grossman School of Medicine, 550 First Avenue, MSB 593D, New York, NY 10016, USA. ²Albert Einstein College of Medicine, Bronx, New York, NY, USA. ³University of California San Francisco, San Francisco, CA, USA. ⁴University of California San Diego, San Diego, CA, USA. ⁵Johns Hopkins University, Baltimore, MD, USA. ⁶Azrieli Faculty of Medicine, Zefat, Israel. ⁷University of Rochester Medical Center, Rochester, NY, USA. ⁸Zucker School of Medicine at Hofstra/Northwell, Manhasset, NY, USA. ⁹Medical University of South Carolina, Charleston, SC, USA. ¹⁰Oklahoma Medical Research Foundation, Oklahoma City, OK, USA. ¹¹Pfizer Inc., New York, NY, USA. ¹²Brigham and Women's Hospital, Boston, MA, USA. ¹³Stanford University, Palo Alto, CA, USA.

Received: 27 October 2023 Accepted: 23 January 2024 Published online: 20 February 2024

References

 Hoover P, Der E, Berthier CC, Arazi A, Lederer JA, James JA, Buyon J, Petri M, Belmont HM, Izmirly P, et al. Accelerating Medicines Partnership: organizational structure and preliminary data from the phase 1 studies of lupus nephritis. Arthritis Care Res (Hoboken). 2020;72(2):233–42.

- Deonaraine KK, Carlucci PM, Fava A, Li J, Wofsy D, James JA, Putterman C, Diamond B, Davidson A, Fine DM, et al. Safety of procuring research tissue during a clinically indicated kidney biopsy from patients with lupus: data from the Accelerating Medicines Partnership RA/SLE Network. Lupus Sci Med. 2021;8(1):e000522.
- 3. Carlucci P, Li J, Fava A, Deonaraine K, Wofsy D, James J, Putterman C, Diamond B, Davidson A, Fine DM et al: High incidence of proliferative and membranous nephritis in SLE patients with low proteinuria in the Accelerating Medicines Partnership. Rheumatology (Oxford) 2022, in press.
- Hochberg MC. Updating the American College of Rheumatology revised criteria for the classification of systemic lupus erythematosus. Arthritis Rheum. 1997;40(9):1725.
- Tan EM, Cohen AS, Fries JF, Masi AT, McShane DJ, Rothfield NF, Schaller JG, Talal N, Winchester RJ. The 1982 revised criteria for the classification of systemic lupus erythematosus. Arthritis Rheum. 1982;25(11):1271–7.
- Petri M, Orbai A-M, Alarcón GS, Gordon C, Merrill JT, Fortin PR, Bruce IN, Isenberg D, Wallace DJ, Nived O, et al. Derivation and validation of the Systemic Lupus International Collaborating Clinics classification criteria for systemic lupus erythematosus. Arthritis Rheum. 2012;64(8):2677–86.
- Weening JJ, D'Agati VD, Schwartz MM, et al. The classification of glomerulonephritis in systemic lupus erythematosus revisited. Kidney international. 2004;65(2):521-30. https://doi.org/10.1111/j.1523-1755.2004. 00443 x.
- Bajema IM, Wilhelmus S, Alpers CE, et al. Revision of the International Society of Nephrology/Renal Pathology Society classification for lupus nephritis: clarification of definitions, and modified National Institutes of Health activity and chronicity indices. Kidney international. 2018;93(4):789-96. https://doi.org/10.1016/j.kint.2017.11.023.
- Access Trial Group. Treatment of lupus nephritis with abatacept: the Abatacept and Cyclophosphamide Combination Efficacy and Safety Study. Arthritis Rheumatol. 2014;66(11):3096–104.
- Furie R, Rovin BH, Houssiau F, Malvar A, Teng YKO, Contreras G, Amoura Z, Yu X, Mok C-C, Santiago MB, et al. Two-year, randomized, controlled trial of belimumab in lupus nephritis. N Engl J Med. 2020;383(12):1117–28.
- Rovin BH, Teng YKO, Ginzler EM, Arriens C, Caster DJ, Romero-Diaz J, Gibson K, Kaplan J, Lisk L, Navarra S, et al. Efficacy and safety of voclosporin versus placebo for lupus nephritis (AURORA 1): a double-blind, randomised, multicentre, placebo-controlled, phase 3 trial. Lancet. 2021;397(10289):2070–80.
- Furie RA, Aroca G, Cascino MD, Garg JP, Rovin BH, Alvarez A, Fragoso-Loyo H, Zuta-Santillan E, Schindler T, Brunetta P, et al. B-cell depletion with obinutuzumab for the treatment of proliferative lupus nephritis: a randomised, double-blind, placebo-controlled trial. Ann Rheum Dis. 2022;81(1):100–7.
- Kim M, Merrill JT, Wang C, Viswanathan S, Kalunian K, Hanrahan L, Izmirly P. SLE clinical trials: impact of missing data on estimating treatment effects. Lupus Sci Med. 2019;6(1):e000348.
- Fava A, Rao DA, Mohan C, Zhang T, Rosenberg A, Fenaroli P, Belmont HM, Izmirly P, Clancy R, Trujillo JM, et al. Urine proteomics and renal single-cell transcriptomics implicate interleukin-16 in lupus nephritis. Arthritis Rheumatol. 2022;74(5):829–39.
- Ugolini-Lopes MR, Seguro LPC, Castro MXF, Daffre D, Lopes AC, Borba EF, Bonfa E. Early proteinuria response: a valid real-life situation predictor of long-term lupus renal outcome in an ethnically diverse group with severe biopsy-proven nephritis? Lupus Sci Med. 2017;4(1):e000213.
- Tamirou F, Lauwerys BR, Dall'Era M, Mackay M, Rovin B, Cervera R, Houssiau FA, Investigators MNT. A proteinuria cut-off level of 0.7 g/day after 12 months of treatment best predicts long-term renal outcome in lupus nephritis: data from the MAINTAIN Nephritis Trial. Lupus Sci Med. 2015;2(1):e000123.
- Dall'Era M, Cisternas MG, Smilek DE, Straub L, Houssiau FA, Cervera R, Rovin BH, Mackay M. Predictors of long-term renal outcome in lupus nephritis trials: lessons learned from the Euro-Lupus Nephritis cohort. Arthritis Rheumatol. 2015;67(5):1305–13.
- 18. Kim M, Merrill J, Kalunian K, Hahn B, Roach A, Izmirly P. Lupus Foundation of America Collective Data Analysis Initiative G: Brief report: Longitudinal patterns of response to standard of care therapy for systemic lupus erythematosus: implications for clinical trial design. Arthritis Rheumatol. 2017;69(4):785–90.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.