

# Baseline biopsy characteristics in an IgA nephropathy clinical trial of ravulizumab: A prespecified analysis of the SANCTUARY trial

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## INTRODUCTION

- Immunoglobulin A (IgA) nephropathy is the most frequent primary glomerulonephritis and complement activation plays a key role in pathogenesis<sup>1,2</sup>
- Diagnosis of IgA nephropathy is based on kidney biopsy; the MEST-C Oxford classification provides information on disease severity and prognosis<sup>3</sup>
  - MEST-C: mesangial hypercellularity (M), endocapillary hypercellularity (E), segmental glomerulosclerosis (S), tubular atrophy/interstitial fibrosis (T), and crescents (C)<sup>3</sup>
- Glomerular complement component C3 staining also has prognostic value<sup>4</sup>
- Clinical trials for IgA nephropathy are often limited by the lack of a central pathology laboratory to minimize interpersonal variation in histological scoring
- Inclusion of a central pathologist to confirm the diagnosis and MEST-C are strengths of the SANCTUARY trial (NCT04564339)<sup>5</sup>
- Biopsies obtained >1 year before trial enrolment may be adequate for diagnostic purposes but information from MEST-C scoring is no longer current
  - A recent biopsy within 1 year of trial enrolment is important for histologic interpretation
- This prespecified analysis reports the baseline biopsy characteristics of patients with IgA nephropathy enrolled in the phase 2 SANCTUARY trial of ravulizumab

A phase 3 trial of ravulizumab in adults with IgA nephropathy (I CAN study, NCT06291376; EU CT 2023-507851-31-00) is ongoing

## OBJECTIVES

- To assess baseline MEST-C scores and complement C3 staining in patients enrolled in the SANCTUARY trial of ravulizumab in IgA nephropathy
- To assess agreement between local and central pathology biopsy readings
- To assess treatment effect at Week 26 by baseline local pathology MEST-C scores
- To assess baseline proteinuria by MEST-C scores and complement C3 staining

## CONCLUSIONS

- This prespecified analysis of the SANCTUARY trial reported MEST-C scoring and glomerular complement staining from biopsies prior to treatment, based on local pathology (38 sites) and central pathology, for which there was strong interobserver agreement on E, S, and C scores and comparatively lower agreement on M and T
- More than a third of patients had biopsy evidence of active inflammation, as indicated by E lesions. On central pathology, the vast majority had at least some degree of tubular atrophy/interstitial fibrosis. Most patients had positive complement staining
- Ravulizumab had a generally similar proteinuria reduction treatment effect across the MEST-C categories. However, sample sizes were small



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## METHODS

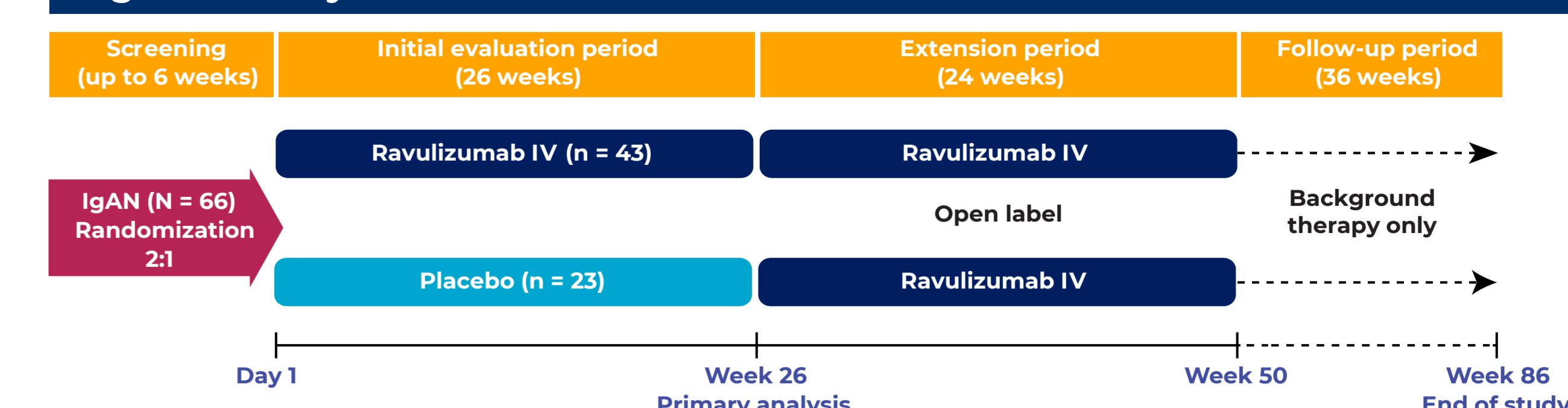
- The SANCTUARY trial design is depicted in **Figure 1**; patients were enrolled from 38 sites across 11 countries
- IgA nephropathy diagnosis was based on a kidney biopsy using a local pathology report obtained any time prior to Day 1
- A central pathology laboratory reviewed biopsies performed within 1 year prior to randomization
  - Light microscopy slides were sent to the central pathology lab; digital images were sent in cases where slide shipment was not possible
- Kidney biopsies were evaluated and scored based on MEST-C (Oxford) classification; glomerular complement C3 staining was also conducted
- Kappa statistics were used to evaluate the percent agreement between local and central pathology for MEST-C scores

– Kappa values ranged from 0 to 1 (no agreement:  $\leq 0$ , slight agreement: 0.01 – 0.20, fair agreement: 0.21 – 0.40, moderate agreement: 0.41 – 0.60, strong agreement: 0.61 – 0.80, almost perfect agreement: 0.81 – 1.00)

### Assessments

- Baseline biopsy characteristics for MEST-C and glomerular C3 staining by local and central pathology assessment
- Percent agreement between local and central pathology MEST-C scores
- Treatment effect on change in proteinuria at Week 26 by baseline local pathology MEST-C scores
- Post hoc: baseline proteinuria by biopsy characteristics based on central pathology scores

Figure 1. Study schematic\*



\*Key eligibility criteria for the phase 2 SANCTUARY trial: aged 18 – 75 years, biopsy-proven IgA nephropathy diagnosis, eGFR  $\geq 30$  mL/min/1.73 m<sup>2</sup>, mean proteinuria  $\geq 1$  g/day at screening (based on 2 complete 24-hour urine collections during the screening period), and on stable maximally tolerated renin-angiotensin system inhibitor.

## RESULTS AND INTERPRETATION

- Local pathology MEST-C score and C3 staining on biopsies prior to Day 1 are shown in **Table 1**

Table 1. Baseline biopsy characteristics – local pathology\*

Category	Ravulizumab n (%)	Placebo n (%)	Total n (%)
<b>MEST-C score</b>			
<b>M-score</b>	n = 37	n = 20	n = 57
M0	14 (37.8)	7 (35.0)	21 (36.8)
M1	23 (62.2)	13 (65.0)	36 (63.2)
<b>E-score</b>	n = 36	n = 20	n = 56
E0	19 (52.8)	14 (70.0)	33 (58.9)
E1	17 (47.2)	6 (30.0)	23 (41.1)
<b>S-score</b>	n = 37	n = 20	n = 57
S0	5 (13.5)	2 (10.0)	7 (12.3)
S1	32 (86.5)	18 (90.0)	50 (87.7)
<b>T-score</b>	n = 36	n = 20	n = 56
T0	22 (61.1)	11 (55.0)	33 (58.9)
T1	14 (38.9)	8 (40.0)	22 (39.3)
T2	0	1 (5.0)	1 (1.8)
<b>C-score</b>	n = 27	n = 20	n = 47
C0	15 (55.6)	14 (70.0)	29 (61.7)
C1	11 (40.7)	6 (30.0)	17 (36.2)
C2	1 (3.7)	0	1 (2.1)
<b>Glomerular C3 staining</b>			
0/1+	18 (48.6)	6 (28.6)	24 (41.4)
2+/3+	19 (51.4)	15 (71.4)	34 (58.6)

\*Full analysis set.

- 30 of the 66 patients (45%) had a biopsy within 1 year prior to randomization
  - Within 3 months prior: 9 patients (30%); 3 to 6 months prior: 9 patients (30%); between 6 and 12 months prior: 12 patients (40%)
- Table 2** reports MEST-C score and C3 staining among patients with a biopsy within 1 year prior to randomization and with central pathology data available

Table 2. Baseline biopsy characteristics – central pathology – among patients with biopsy  $\leq 1$  year prior to randomization

Category	Ravulizumab n (%)	Placebo n (%)	Total n (%)
<b>MEST-C score</b>			
<b>M-score</b>	n = 11	n = 8	n = 19
M0	6 (54.5)	4 (50.0)	10 (52.6)
M1	5 (45.5)	4 (50.0)	9 (47.4)
<b>E-score</b>	n = 11	n = 8	n = 19
E0	5 (45.5)	7 (87.5)	12 (63.2)
E1	6 (54.5)	1 (12.5)	7 (36.8)
<b>S-score</b>	n = 11	n = 8	n = 19
S0	0	0	0
S1	11 (100)	8 (100)	19 (100)
<b>T-score</b>	n = 11	n = 8	n = 19
T0	2 (18.2)	0	2 (10.5)
T1	8 (72.7)	8 (100)	16 (84.2)
T2	1 (9.1)	0	1 (5.3)
<b>C-score</b>	n = 11	n = 8	n = 19
C0	6 (54.5)	6 (75.0)	12 (63.2)
C1	5 (45.5)	2 (25.0)	7 (36.8)
C2	0	0	0
<b>Glomerular C3 staining</b>			
0/1+	7 (63.6)	2 (22.2)	9 (45.0)
2+/3+	4 (36.4)	7 (77.8)	11 (55.0)

- Both local and central pathology MEST-C scores were available for 24 patients (except for C-scores), regardless of biopsy age (**Table 3**)
  - IgA nephropathy was confirmed as the primary diagnosis by central pathology in all cases
- Strong agreement between local and central pathology scores was noted for E, S, and C (**Table 3**)
  - Agreement for M and T was comparatively lower

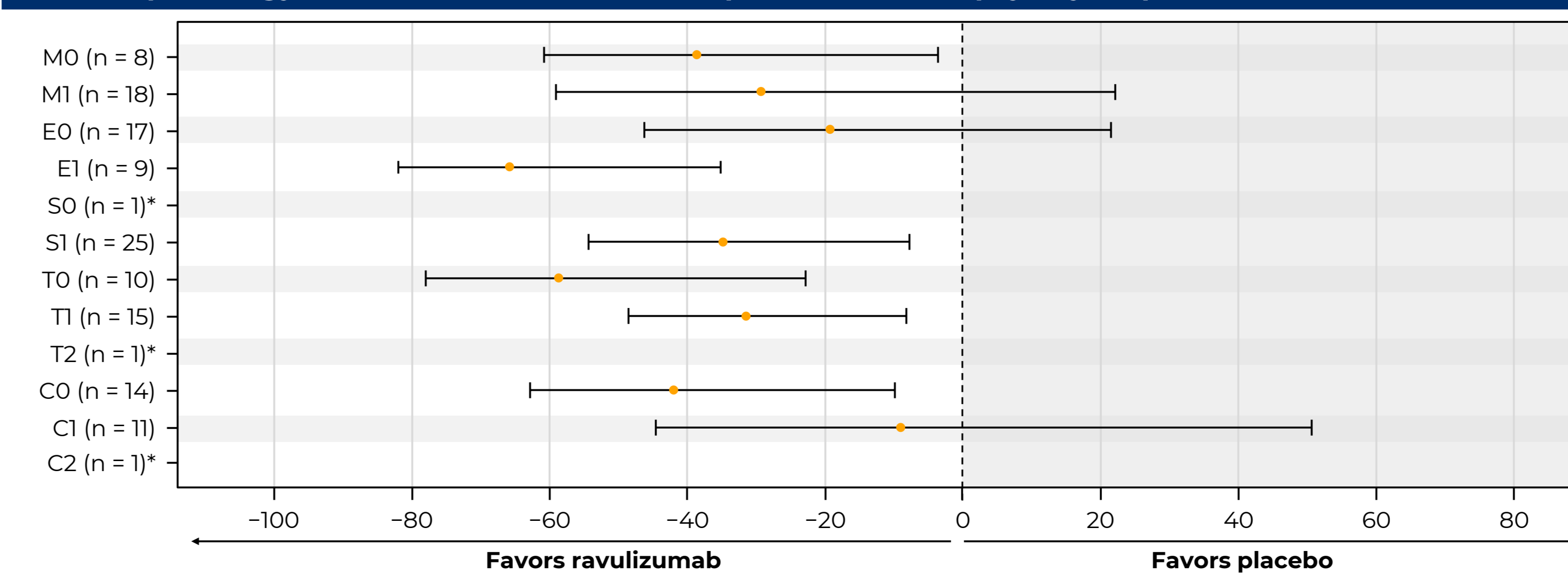
Table 3. Agreement between local and central pathology for MEST-C scores

Category	Local pathology n (%)	Central pathology n (%)	Percent agreement (%)	Kappa statistic* (95% CI)
<b>M-score</b>				
M0	n = 24	n = 24	75.0	0.50 (0.20, 0.80)
M1	18 (75.0)	12 (50.0)		
<b>E-score</b>				
E0	n = 24	n = 24	87.5	0.75 (0.48, 1)
E1	13 (54.2)	14 (58.3)		
E1	11 (45.8)	10 (41.7)		
<b>S-score</b>				
S0	n = 24	n = 24	91.7	0.63 (0.16, 1)
S1	4 (16.7)	2 (8.3)		
S1	20 (83.3)	22 (91.7)		
<b>T-score</b>				
T0	n = 24	n = 24	66.7	NA
T1	10 (41.7)	4 (16.7)		
T1	14 (58.3)	18 (75.0)		
T2	0	2 (8.3)		
<b>C-score</b>				
C0	n = 22	n = 22	90.9	0.81 (0.57, 1)
C0	12 (54.5)	14 (63.6)		
C1	10 (45.5)	8 (36.4)		

\*An unweighted kappa statistic and corresponding 95% CI is represented.

- In general, a similar treatment effect on proteinuria was observed across MEST-C categories (**Figure 2**)

Figure 2. Treatment effect (90% CI) on change in proteinuria (24-hour UP) from baseline to Week 26 based on local pathology MEST-C score at baseline, in patients with biopsy  $\leq 1$  year prior to randomization



\*Insufficient sample size.

- Baseline mean proteinuria was similar across MEST-C and glomerular C3 staining categories (**Table 4**)

Table 4. Baseline proteinuria by biopsy characteristics (central pathology scores)\*

Category	Baseline 24-hour UPCR (g/g)	
	Mean (SD)	Median (Q1, Q3)
<b>MEST-C score</b>		
M0 (n = 10)	1.29 (0.53)	1.16 (0.93, 1.64)
M1 (n = 9)	1.21 (0.39)	1.26 (0.97, 1.38)
E0 (n = 12)	1.24 (0.54)	1.09 (0.84, 1.64)
E1 (n = 7)	1.28 (0.31)	1.26 (1.09, 1.64)
S1 (n = 19)	1.25 (0.46)	1.23 (0.93, 1.64)
T1 (n = 16)	1.27 (0.50)	1.25 (0.84, 1.64)
C0 (n = 12)	1.28 (0.53)	1.30 (0.86, 1.64)
C1 (n = 7)	1.20 (0.34)	1.15 (0.93, 1.64)
<b>Glomerular C3 staining</b>		
0/1+ (n = 9)	1.22 (0.52)	1.23 (0.76, 1.38)
2+/3+ (n = 11)	1.28 (0.40)	1.26 (0.97, 1.64)

\*Data are included for categories with  $\geq 5$  patients, who had biopsy within 1 year before randomization.

### Abbreviations

C3, complement component 3; CI, confidence interval; eGFR, estimated glomerular filtration rate; IgAN, immunoglobulin A nephropathy; IV, intravenous; MEST-C, mesangial hypercellularity, endocapillary hypercellularity, segmental glomerulosclerosis, tubular atrophy/interstitial fibrosis, and crescents; NA, not available; Q1, first quartile; Q3, third quartile; SD, standard deviation; UP, urine protein; UPCR, urine protein:creatinine ratio.

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### Disclosures

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