

Statistical Analysis Plan Addendum

Clinical Trial Protocol Identification No.	EMR 200017-014
Title:	A Phase II, randomized, double-blind, placebo-controlled, parallel-group, multicenter trial to evaluate the efficacy and safety of abituzumab in subjects with systemic sclerosis-associated interstitial lung disease (SSc-ILD)
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1 Signature Page

Statistical Analysis Plan Addendum: EMR 200017-014

A Phase II, randomized, double-blind, placebo-controlled, parallel-group, multicenter trial to evaluate the efficacy and safety of abituzumab in subjects with systemic sclerosis-associated interstitial lung disease (SSc-ILD)

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3 List of Abbreviations and Definition of Terms

Refer to the Study SAP.

4 Modification History

Unique Identifier for SAP Version	Date of SAP Version	Author	Changes from the Previous Version
1.0	05 April 2018	PPD	N/A – First version

5 Purpose of the Statistical Analysis Plan Addendum

The purpose of this statistical analysis plan (SAP) addendum is to document technical and detailed specifications for the Final Analysis for protocol EMR200017-014.

Due to the early termination of the study and consequently the limited number of subjects and data available, the full pre-planned analyses in the Study SAP (dated 15 November 2017) will not be carried out, since they may lead to unreliable conclusions. Hence this addendum references the applicable parts of the Study SAP and the Independent Data Monitoring Committee (IDMC) SAP (dated 29 June 2016) to produce the tables, listings, and figures (TLFs) to be included in the abbreviated clinical study report (CSR).

6 Summary of Clinical Study Features

Refer to the Study SAP.

7 Sample Size/Randomization

Refer to the Study SAP.

8 Overview of Planned Analyses

8.1 Final Analysis

Cut-off date:

The data cut-off date for the final analysis will be when all randomized subjects have completed the Week 12 Safety Follow-up visit or been lost to follow-up. No more data are expected to be entered into the database by the sites after this point.

A blinded data review meeting will be held prior to database lock in order to identify and confirm protocol deviations (PDs).

Dissemination of results:

The blinded Biostatistics study team will become unblinded after database lock. Results will be generated on the aggregate group level and delivered to the Sponsor as one package (no key TLFs will be produced) following the normal process.

TLF Shells:

Refer to [Appendix 1](#) for the TLF shells.



9 Changes to the Planned Analyses in the Clinical Study Protocol

See subsequent sections regarding the analyses that will be performed for the abbreviated CSR.

10 Protocol Deviations and Analysis Sets

Refer to the Study SAP.

Subject disposition will use the Screening Analysis Set, all baseline and safety summaries will use the Safety Analysis Set and all efficacy summaries will use the mITT Analysis Set. The other analysis sets will not be required for the following reasons:

- Intent-to-treat – All randomized subjects were treated so this is equivalent to the mITT analysis set.
- Per-Protocol – The purpose of this was for sensitivity analysis on the primary and key secondary endpoints, which will no longer be performed.
- PK – No PK analyses will be performed for the abbreviated CSR.
- Pharmacogenetics – No CC1 analyses will be performed for the abbreviated CSR.

Since no supportive analyses on the primary endpoint will be performed, no subgroups will be identified.

11 General Specifications for Statistical Analyses

Refer to the Study SAP. Only summary statistics will be computed.

Early Termination Visit:

Since subjects will be performing their early termination (ET) visits at different stages within the protocol schedule of assessments, for the efficacy by-visit summaries, the ET visits will reference the protocol visits in between which they fell chronologically.

For example, if a subject performed their Week 12 visit, then performed their ET visit at a later date (but before their scheduled Week 24 visit), their ET visit will be shown as “ET (W12 – W24)” to display the weeks that it fell between.

Missing data:

No imputation of missing efficacy data or duration of prior mycophenolate use at baseline will be performed.

12 Study Subjects

Subject disposition status will be summarized as per the Study SAP. Important PDs will be listed only and clinically important PDs will not be identified.

13 Demographics and Other Baseline Characteristics

Demographics, medical history, and disease history will be summarized as per the IDMC SAP.

14 Previous or Concomitant Medications/Procedures

Previous and concomitant medications will be summarized as per the IDMC SAP.

15 Treatment Compliance and Exposure

The following will be summarized using descriptive statistics:

- Treatment duration (weeks)
- Number of infusions of trial treatment received
- Treatment adherence (%)
- Infusion duration (hours) and volume (mL)
- Number of subjects with at least one dose adjustment (overall and by reason)
- Number of subjects with at least one infusion not completed as planned (overall and by reason)

Refer to the Study SAP for definitions.

16 Endpoint Evaluation

Summary statistics of absolute values and change from baseline will be provided for the following efficacy measurements. No statistical tests will be performed or confidence intervals presented.

Pulmonary function tests:

- FVC
- FVC % Predicted
- DLCO – eCRF result
- DLCO % Predicted – eCRF result
- KCO – eCRF result

HRCT results:

- QLF
- Total ILD

QoL questionnaires:

- Mahler TDI
- SGRQ
- Physician’s Global Assessment
- Patient’s Global Assessment
- SHAQ – VAS scores only
- EQ-5D – VAS score only

- Leicester Cough Index

Skin fibrosis:

- mRSS
- Digital Ulcer Count

Clinically meaningful disease progression and overall survival:

- Disease Progression
- Listing of Deaths

Refer to the Study SAP for details of these measurements and any derivations required.

17 Safety Evaluation

Refer to the IDMC SAP for the safety summaries that will be provided.



18 Appendix 1 – TLF shells

18.1 Tables and Figures

Programming note:

The shells use Treatment A, B & C.

The order of the treatment groups in the tables will be as follows:

Placebo	Abituzumab 500 mg	Abituzumab 1500 mg	Total
---------	-------------------	--------------------	-------

The same order will be used for listings.

This table of contents details which TLFs have been produced for the IDMCs and which are new for the final analysis:



EMR200017-014_Sy
noptic_CSR_TOC_for

Red text in the shells is where the IDMC TLF has been adjusted for the final analysis.

The headers will be:

Protocol: EMR 200017-014 (Abituzumab) - Cutoff date: DDMMYYYY (<Purpose>)

The footers will be:

Source: ADxx DDMMYYYY hh:mm <creation date of ADAM dataset>; Listing 16.2.xx; SDTM package: DDMMYYYY
Program_path\pgm_name.sas, DDMMYYYY hh:mm

Page 1 of x

15.1: Demographic and Baseline Data

15.1.1: Subject Disposition

Table 15.1.1.1: Subject Disposition Status

	Treatment A	Treatment B	Treatment C	Total
Number of screened subjects (a), n				xx
Number of screen failures, n				xx
Subject did not meet all eligibility criteria				xx
Withdrew informed consent				xx
Progressive disease				xx
Adverse event				xx
Lost to follow-up				xx
Death				xx
Other				xx
Missing				xx
Randomized subjects, n (%)	xx (100.0)	xx (100.0)	xx (100.0)	xx (100.0)
Baseline FVC <70% predicted	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Baseline FVC ≥70% predicted	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Received no treatment, n (%)*	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Treatment Ongoing, n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Treatment Completed, n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

Shell continues on the next page



15.1: Demographic and Baseline Data

15.1.1: Subject Disposition

Table 15.1.1.1: Subject Disposition Status

	Treatment A	Treatment B	Treatment C	Total
Subjects who discontinued trial treatment, n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Adverse Event	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Lost to Follow-up	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Protocol non-compliance	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Death	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Progressive disease	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Withdrew Consent	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Other	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Subjects who discontinued trial treatment and withdrew from the treatment period, n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Subjects who withdrew from the trial, n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Adverse Event	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Lost to Follow-up	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Protocol non-compliance	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Death	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Progressive disease	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Withdrew Consent	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Other	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

Note: Percentages for 'Subjects who discontinued trial treatment' are based on the number of subjects who received at least one infusion of trial treatment. All other percentages are based on the number of subjects randomized.

No subjects were randomized under the new stratification level of < 6 months prior mycophenolate duration, so this stratification factor has not been presented.

* Number of subjects who received no treatment based on current exposure data.

15.1: Demographic and Baseline Data

15.1.4: Demographic Characteristics

Table 15.1.4.1: Demographic Characteristics - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)	Total N=xxx (100%)
Sex, n (%)				
Male	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Female	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Missing	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Race, n (%)				
White	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Black or African American	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Asian	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
American Indian or Alaska Native	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Native Hawaiian or Other Pacific Islander	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Other	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Missing (including not collected at site)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Ethnicity, n (%)				
Hispanic or Latino	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Not Hispanic or Latino	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Age (Years)				
n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Missing (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Mean ±SD	xx ±xx.xx	xx ±xx.xx	xx ±xx.xx	xx ±xx.xx
Median	xx	xx	xx	xx
Q1; Q3	xx; xx	xx; xx	xx; xx	xx; xx
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx

Shell continues on the next page

15.1: Demographic and Baseline Data

15.1.4: Demographic Characteristics

Table 15.1.4.1: Demographic Characteristics - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)	Total N=xxx (100%)
Age categories, n (%)				
< 35 years	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
>= 35 - < 50 years	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
>= 50 - < 76 years	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Missing	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

Note: Age is relative to the date of informed consent.

15.1: Demographic and Baseline Data

15.1.5: Medical History

Table 15.1.5.1: Medical History - SAF Analysis Set

Primary System Organ Class Preferred Term	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)	Total N=xxx (100%) n (%)
Subjects with at least one Medical History	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
SOC1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
PT1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
PT2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
SOC2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
PT1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
PT2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

MedDRA version xx.x.

Programming Note:

- Sort order by primary SOC and PT in alphabetical order. If SOC or PT is missing/not coded yet, then 'UNCODED' (or 'UNCODED + verbatim Investigator') will be indicated at the ADaM level. Uncoded SOC/PT should come first.

MedDRA version 19.1 or highest at time of data transfer.



15.1: Demographic and Baseline Data
15.1.6: Other Baseline Characteristics
Table 15.1.6.1: Disease History - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)	Total N=xxx (100%)
Time since date of diagnosis (years)				
n (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Missing (%)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Mean ±SD	xx ±xx.xx	xx ±xx.xx	xx ±xx.xx	xx ±xx.xx
Median	xx	xx	xx	xx
Q1; Q3	xx; xx	xx; xx	xx; xx	xx; xx
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx
Subject status/ disease history, n (%)				
Diffuse cutaneous SSc	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Limited cutaneous SSc	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
SSc sine scleroderma	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Systemic sclerosis symptom history (a), n (%)				
Skin thickening	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Digital ulcers	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Synovitis and/or joint contracture	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Tendon friction rubs	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
C .K. elevation and/or weakness	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Atrophy: Upper extremity	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Atrophy: Lower extremity	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Atrophy: Right extremity	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Atrophy: Left extremity	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Esophagus: Dysphagia	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Esophagus: Reflux	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Stomach: Early satiety	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Stomach: Vomiting	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Intestinal: Diarrhea	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Intestinal: Bloating	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

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15.1: Demographic and Baseline Data
15.1.6: Other Baseline Characteristics
Table 15.1.6.1: Disease History - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)	Total N=xxx (100%)
Intestinal: Constipation	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Incontinence	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Hypertension	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Renal crisis	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Dyspnea (significant)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Palpitations (significant)	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
ACR/EULAR Classification Criteria for Systemic Sclerosis (b), n (%)				
Skin thickening of the fingers of both hands extending proximal to the metacarpophalangeal joints	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Skin thickening of the fingers: Puffy fingers	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Skin thickening of the fingers: Sclerodactyly of the fingers	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Fingertip lesions: Digital tip ulcers	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Fingertip lesions: Fingertip pitting scars	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Telangiectasia	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Abnormal nailfold capillaries	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Lung involvement: Pulmonary arterial hypertension	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Lung involvement: Interstitial lung disease	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
Raynaud's Phenomenon	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
CC	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

III

Note: Partial date of diagnosis is imputed as follows: In case the day of diagnosis is missing, then the day will be imputed as the 1st. In case the month of diagnosis is missing, then the month will be imputed as January.

- (a) Counts are the number of subjects who reported 'Yes' to the questions.
- (b) Counts are the number of subjects who reported 'Present' to the questions.

15.1: Demographic and Baseline Data

15.1.7: Previous and Concomitant Medications, Procedures, Follow-up Treatments

Table 15.1.7.1: Previous Medication - SAF Analysis Set

ATC-2 nd Level Preferred Term	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)	Total N=xxx (100%) n (%)
Subjects with at least one Previous Medication	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
ATC1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
MED1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
MED2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
ATC2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
MED1	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)
MED2	xx (xx.x)	xx (xx.x)	xx (xx.x)	xx (xx.x)

WHO Drug Dictionary version Enhanced September 2016.

Note: If a subject has more than one medication within a ATC class/preferred term, the subject was counted once in that ATC class/preferred term.

In case multiple ATC-s are assigned to a medication, all ATC-s will be reported.

Programming Note:

- Sort order by ATC-2nd Level and Preferred Term in alphabetical order.

WHO Drug Dictionary version Enhanced September 2016 or highest at time of data transfer.

Repeat for table:

Table 15.1.7.2: Concomitant Medication - SAF Analysis Set

Change "Previous Medication" to "Concomitant Medication"



15.2: Efficacy Data

15.2.1: Annual Rate of Absolute FVC Change in Volume (mL)

Table 15.2.1.1: FVC Volume (mL) Observed and Change from Baseline by Visit - mITT Analysis Set

Visit / Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 12						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
Week 24						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx

Note: Baseline is defined as the last non-missing assessment value prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.
Percentages are based on the number of subjects present in the trial at that visit.

Programming note: Only display visits where FVC assessment is scheduled to be done and where there is data available.

Repeat table 15.2.1.1 for:

Table 15.2.1.2: FVC % Predicted Observed and Change from Baseline by Visit - mITT Analysis Set

15.2: Efficacy Data

15.2.2: Key Secondary Endpoints

Table 15.2.2.1: Change in Dyspnea from Baseline as Measured by the Mahler TDI - mITT Analysis Set

Visit / Statistic	Treatment A N=xxx	Treatment B N=xxx	Treatment C N=xxx
Baseline			
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx
Week 12			
Change from Baseline			
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx
.....			

Repeat table 15.2.1.1 for:

Table 15.2.2.2: mRSS Total Score Observed and Change from Baseline by Visit - mITT (dcSSc) Analysis Set

Programming note: Only display visits where mRSS assessment is scheduled to be done and where there is data available.

Add the following footnote:

Note: dcSSc = diffuse cutaneous systemic sclerosis. Only subjects with dcSSc at baseline are included in the mRSS summary.



15.2: Efficacy Data

15.2.2: Key Secondary Endpoints

Table 15.2.2.3: Absolute Change from Baseline in SGRQ Total Score - mITT Analysis Set

Visit/ Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 12						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
...						

15.2: Efficacy Data

15.2.2: Secondary Endpoints

Table 15.2.2.4: Clinically Meaningful Disease Progression - mITT Analysis Set

	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Number of Subjects with Clinically Meaningful Disease Progression	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Trial Treatment Discontinued	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SSc-ILD Progression	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Relative decrease from baseline in FVC % predicted >10% *	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Relative decrease from baseline in FVC % predicted of >5% to <10% and relative decrease in DLCO % predicted >15% *	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mycophenolate dose increased	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
New immunosuppressant or biologic drug initiated	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Trial Treatment Discontinued	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SSc Progression other than ILD	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Scleroderma renal crisis	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Left ventricular failure (defined as ejection fraction <45%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Pulmonary arterial hypertension requiring treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Other	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Trial Treatment Discontinued	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

* Confirmed on 2 occasions within 4 weeks.

Note: A subject may have both SSc-ILD progression and SSc progression other than ILD at the same time, and may progress more than once.

Repeat table 15.2.1.1 for:

Table 15.2.2.5: DLCO (mmol of CO/min/kPa) Observed and Change from Baseline by Visit - mITT Analysis Set

Programming note: Only display visits where DLCO assessment is scheduled to be done and where there is data available.



Repeat table 15.2.1.1 for:

Table 15.2.2.6: Absolute Change from Baseline in QLF - mITT Analysis Set

Table 15.2.2.7: DLCO % Predicted Observed and Change from Baseline by Visit - mITT Analysis Set

Programming note: Only display visits where DLCO assessment is scheduled to be done and where there is data available.

Table 15.2.3.1: KCO Observed and Change from Baseline by Visit - mITT Analysis Set

Table 15.2.3.2: Extent of total ILD Observed and Change from Baseline by Visit - mITT Analysis Set

15.2: Efficacy Data
15.2.4: QoL Endpoints

Table 15.2.4.1: Leicester Cough Index Total Score - Observed and Change from Baseline - mITT Analysis Set

Visit/ Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 24						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
...						

Note: Baseline is defined as the last non-missing measurement prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.

Percentages are based on the number of subjects present in the trial at that visit.

Note: The domain score is the sum of all scores in that domain divided by the number of questions in that domain (ranging from 1 to 7). The total score is the sum of all domain scores (ranging from 3 to 21).

Repeat table 15.2.4.1 for:

Table 15.2.4.2: Physician's Global Assessment - Observed and Change from Baseline - mITT Analysis Set

Update footnotes:

Note: Baseline is defined as the last non-missing measurement prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.

Percentages are based on the number of subjects present in the trial at that visit.

Note: The Physician's Global Assessment consists of one question that is answered on a VAS from of 0 to 100,

where 0 is the most positive response and 100 is the most negative.

Repeat table 15.2.4.1 for:

Table 15.2.4.3: Patient's Global Assessment - Observed and Change from Baseline - mITT Analysis Set

Update footnote to say 'Patient's Global Assessment' instead of 'Physician's Global Assessment'

15.2: Efficacy Data
15.2.4: QoL Endpoints

Table 15.2.4.4: SHAQ - VAS Scores - Observed and Change from Baseline - mITT Analysis Set

VAS category/ Visit/ Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Pain:						
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 24						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
...						

Note: Baseline is defined as the last non-missing measurement prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.

Percentages are based on the number of subjects present in the trial at that visit.

Note: The VAS are scored from 0 to 100, where 0 is the most positive response and 100 is the most negative.

15.2: Efficacy Data
15.2.4: QoL Endpoints

Table 15.2.4.5: EQ-5D VAS Score - Observed and Change from Baseline - mITT Analysis Set

Visit/ Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 24						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
...						

Note: Baseline is defined as the last non-missing measurement prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.
Percentages are based on the number of subjects present in the trial at that visit.

Note: The EQ-5D VAS is numbered from 0 to 100 where the endpoints are labelled 'Best health you can imagine' (100) and 'Worst health you can imagine' (0).

15.2: Efficacy Data

15.2.5: Skin Fibrosis Endpoints

Table 15.2.5.1: Digital Ulcer Count - Observed and Change from Baseline - mITT Analysis Set

Ulcer Type/ Visit/ Statistic	Treatment A N=xxx		Treatment B N=xxx		Treatment C N=xxx	
	Observed	Change from Baseline	Observed	Change from Baseline	Observed	Change from Baseline
Active:						
Baseline						
n (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Missing (%)	xxx (xx.x)		xxx (xx.x)		xxx (xx.x)	
Mean ±SD	xx.x ±xx.xx		xx.x ±xx.xx		xx.x ±xx.xx	
Median	xx.x		xx.x		xx.x	
Q1; Q3	xx.x; xx.x		xx.x; xx.x		xx.x; xx.x	
Min; Max	xx; xx		xx; xx		xx; xx	
Week 12						
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missing (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx	xx; xx
...						

Note: Baseline is defined as the last non-missing measurement prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.
Percentages are based on the number of subjects present in the trial at that visit.

Note: Only subjects with digital ulcers at baseline continue with the digital ulcer assessment throughout the trial. The total count is the sum of all active, undetermined and healed ulcers.

15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.1: Duration of Treatment (weeks) - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)
Duration of treatment (weeks), n (%)			
>=4 and <=28 weeks	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>28 and <=52 weeks	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>52 and <=76 weeks	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>76 and <=100 weeks	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>100 weeks	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx

Note: Duration of treatment (weeks) is calculated as:

(Last infusion date prior to data cut-off - first infusion date + 28) / 7 days.

Programming note:

- **Summary statistics are based on subjects who received at least one infusion of trial treatment → Min>0 (this should be equivalent to the SAF analysis set)**



15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.2: Number of Infusions - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)
Total number of infusions received, n (%)			
1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>1 and <=8	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>8 and <=14	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>14 and <=20	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>20 and <=26	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>26	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx
Total number of infusions across all subjects, n	xxx	xxx	xxx

Programming note:

- **Summary statistics are based on subjects who received at least one infusion of trial treatment → Min>0 (this should be equivalent to the SAF analysis set)**

15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.2b: Treatment **Adherence** - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)
Adherence with Treatment, n (%)			
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
<80%	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
80% - 100%	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>100% - 120%	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
>120%	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx

Note: **Adherence** by subject is calculated as: (sum of actual volume received/ sum of planned volume, up to and including their last dose recorded on their treatment termination visit)*100.
For the continuous summary, values >100% are set to 100%.

Programming note:

- **Summary statistics are based on subjects who received at least one infusion of trial treatment (this should be equivalent to the SAF analysis set)**
- **The Treatment Compliance definition for the IDMC is equivalent to the Treatment Adherence definition in the Study SAP, for the Primary & Final analyses.**



15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.3: Infusion Duration and Volume - SAF Analysis Set

	Treatment A N=xxx (100%)	Treatment B N=xxx (100%)	Treatment C N=xxx (100%)
Mean infusion duration (hours)			
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx
Mean actual volume infused (mL)			
n (%)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Mean ±SD	xx.x ±xx.xx	xx.x ±xx.xx	xx.x ±xx.xx
Median	xx.x	xx.x	xx.x
Q1; Q3	xx.x; xx.x	xx.x; xx.x	xx.x; xx.x
Min; Max	xx; xx	xx; xx	xx; xx

Note: Summary statistics are based on the mean infusion duration and mean volume received per subject.

Programming note:

- **Summary statistics are based on subjects who received at least one infusion of trial treatment → Min>0 (this should be equivalent to the SAF analysis set)**



15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.4: Dose Adjustments - SAF Analysis Set

	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Number of subjects without any dose adjustment (a)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Number of subjects with at least one dose adjustment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Dose adjusted:			
Adverse Event	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Other	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
No dose:			
Adverse Event	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Missed Dose	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Other	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Number of subjects with at least one missed visit (b)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

Note: A subject may have more than one reason if they have more than one dose adjustment, so will be counted under each reason.

(a) Subjects without any dose adjustment do not include subjects with missed visits.

(b) Subjects with missed visits may not be counted under 'Dose adjusted' or 'No dose' due to not completing the 'STUDY TREATMENT ADMINISTRATION DETAILS' eCRF page at the missed visit.

Programming note: Summary statistics are based on subjects who received at least one infusion of trial treatment (this should be equivalent to the SAF analysis set)



15.3: Safety Data

15.3.0: Treatment Compliance and Exposure

Table 15.3.0.5: Infusion Outcome - SAF Analysis Set

	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Number of subjects with all infusions completed as planned	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Number of subjects with at least one infusion not completed as planned	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Adverse Event (other than infusion-related reaction or overdose)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Infusion-related reaction	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Overdose	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Other	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

Note: A subject may have more than one reason if they have more than one infusion not completed as planned, so will be counted under each reason.

Programming note:

- **Summary statistics are based on subjects who received at least one infusion of trial treatment (this should be equivalent to the SAF analysis set)**



15.3: Safety Data

15.3.1: Display of Adverse Events

Table 15.3.1.1: Overview of Treatment Emergent Adverse Events (TEAEs) - SAF Analysis Set

Number of Subjects with:	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Any TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any serious TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related serious TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any grade >=3 TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any grade >=4 TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related grade >=3 TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related grade >=4 TEAE	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any TEAE leading to death	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related TEAE leading to death	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any serious treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

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15.3: Safety Data

15.3.1: Display of Adverse Events

Table 15.3.1.1: Overview of Treatment Emergent Adverse Events (TEAEs) - SAF Analysis Set

Number of Subjects with:	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Any trial treatment related serious treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any grade >=3 treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any grade >=4 treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related grade >=3 treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related grade >=4 treatment emergent AESI	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any treatment emergent AESI leading to death	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related treatment emergent AESI leading to death	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

NCI-CTCAE version 4.03.

Note: TEAEs are defined as events that start within the day of first dose of trial treatment until 28 days after last dose of treatment.

Related TEAEs are events with relationship missing, unknown or related.

AESI = Adverse Event of Special Interest.



15.3: Safety Data

15.3.1: Display of Adverse Events

Table 15.3.1.2: Overview of TEAEs Leading to Discontinuation / Dose Reduction of Treatment - SAF Analysis Set

Number of Subjects with:	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Any TEAE leading to temporary discontinuation of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related TEAE leading to temporary discontinuation of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any TEAE leading to permanent discontinuation of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related TEAE leading to permanent discontinuation of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any TEAE leading to dose reduction of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Any trial treatment related TEAE leading to dose reduction of trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

Note: TEAEs are defined as events that start within the day of first dose of trial treatment until 28 days after last dose of treatment.
Related TEAEs are events with relationship missing, unknown or related.



15.3: Safety Data

15.3.1: Display of Adverse Events

Table 15.3.1.3: TEAE by Primary System Organ Class (SOC) and Preferred Term (PT) - SAF Analysis Set

Primary System Organ Class Preferred Term	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Subjects with at least one event	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SOC1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT11	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT12	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT13	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SOC2	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT21	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT22	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

MedDRA version 19.1.

Note: TEAEs are defined as events that start within the day of first dose of trial treatment until 28 days after last dose of treatment.

Programming Note: sorted by primary SOC and PT in alphabetical order.
MedDRA version 19.1 or highest at time of data transfer.



Repeat table 15.3.1.3 for:

Table 15.3.1.4: Trial Treatment Related TEAEs by SOC and PT - SAF Analysis Set

Add footnote: Related TEAEs are events with relationship missing, unknown or related.

Table 15.3.1.5: Serious TEAEs by SOC and PT - SAF Analysis Set

Table 15.3.1.6: Trial Treatment Related Serious TEAEs by SOC and PT - SAF Analysis Set

Add footnote: Related TEAEs are events with relationship missing, unknown or related.

Table 15.3.1.7: Non-serious TEAEs by SOC and PT at a Frequency Threshold of 5% - SAF Analysis Set

Add footnote: Frequency threshold of 5% in any treatment group.

15.3: Safety Data

15.3.1: Display of Adverse Events

Table 15.3.1.8: TEAEs by Worst Grade, SOC and PT - SAF Analysis Set

Primary System Organ Class Preferred Term	Treatment A N=xxx (100%)				Treatment B N=xxx (100%)				Treatment C N=xxx (100%)			
	Any Grade n (%)	Grade >=3 n (%)	Grade >=4 n (%)	Grade 5 n (%)	Any Grade n (%)	Grade >=3 n (%)	Grade >=4 n (%)	Grade 5 n (%)	Any Grade n (%)	Grade >=3 n (%)	Grade >=4 n (%)	Grade 5 n (%)
Subjects with at least one event	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SOC1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT11	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT12	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
SOC2	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
PT21	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
...												

MedDRA version 19.1, NCI-CTCAE version 4.03.

Note: Any grade includes 'Missing' grade. Worst grade per subject/SOC/PT is reported.

TEAEs are defined as events that start within the day of first dose of trial treatment until 28 days after last dose of treatment.

Programming Note: MedDRA version 19.1 or highest at time of data transfer.



Repeat table 15.3.1.8 for:

Table 15.3.1.9: Trial Treatment Related TEAEs by Worst Grade, SOC and PT - SAF Analysis Set

Add footnote: Related TEAEs are events with relationship missing, unknown or related.

Repeat table 15.3.1.3 for:

Table 15.3.1.10: TEAEs Leading to Death by SOC and PT - SAF Analysis Set

Table 15.3.1.11: TEAEs Leading to Trial Treatment Discontinuation by SOC and PT - SAF Analysis Set

Table 15.3.1.12: TEAEs Leading to Dose Reduction of Trial Treatment by SOC and PT - SAF Analysis Set

Table 15.3.1.13: TEAEs Leading to Withdrawal from Trial by SOC and PT - SAF Analysis Set

Table 15.3.1.14: Treatment Emergent AESIs by SOC and PT - SAF Analysis Set

Add footnote: AESI = Adverse Event of Special Interest.

15.3: Safety Data

15.3.2: Listings of Deaths, Other Serious and Significant Adverse Events

Table 15.3.2.1: Deaths by Primary Reason - SAF Analysis Set

	Treatment A N=xxx (100%) n (%)	Treatment B N=xxx (100%) n (%)	Treatment C N=xxx (100%) n (%)
Subjects who died	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Primary reason for death			
Progressive disease and/or disease related condition	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Event unrelated to trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Event related to trial treatment	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)
Unknown	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)

Note: Reason as reported on death CRF page.

15.3: Safety Data

15.3.2: Listings of Deaths, Other Serious and Significant Adverse Events

Table 15.3.2.3: Listing of Serious TEAEs - SAF Analysis Set

AE No	Preferred Term/ Investigator Term	Start/ End Date of AE	Dur. of AE (days)	Onset Rel.to Trial Treat.	Rel. to Trial Treat.	Grade	Action on Trial Treat. (a)	Outcome (b)	Rel. Day from First Admin.
-------	--------------------------------------	--------------------------	----------------------	---------------------------------	----------------------------	-------	----------------------------------	-------------	----------------------------------

Treatment Arm: Treatment A, Subject ID: XXXXXXXXXX, Age: xx, Sex: X, Race: XXXXXXXX

Treatment Administration: First Date: DDMONYYYY, Last Date: DDMONYYYY, Number of infusions: xx

X	XXXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXXX	DDMONYYYY/ DDMONYYYY	xx	Before	Unrelated	1	No change	Res.	xx
---	---	-------------------------	----	--------	-----------	---	-----------	------	----

Treatment Arm: Treatment A, Subject ID: XXXXXXXXXX, Age: xx, Sex: X, Race: XXXXXXXX

Treatment Administration: First Date: DDMONYYYY, Last Date: DDMONYYYY, Number of infusions: xx

X	XXXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXXX	DDMONYYYY/ DDMONYYYY	xx	During	Related	2	Reduced	Ong.	xx
---	---	-------------------------	----	--------	---------	---	---------	------	----

Treatment Arm: Treatment B, Subject ID: XXXXXXXXXX, Age: xx, Sex: X, Race: XXXXXXXX

Treatment Administration: First Date: DDMONYYYY, Last Date: DDMONYYYY, Number of infusions: xx

X	XXXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXXX	DDMONYYYY/ DDMONYYYY	xx	After	Unrelated	3	Interrupted	Res. w. sequ.	xx
---	---	-------------------------	----	-------	-----------	---	-------------	---------------	----

MedDRA version 19.1, NCI-CTCAE version 4.03.

Note: (a) Action: No change = Dose not changed, Reduced = Dose reduced, Interrupted = Drug interrupted, Withdrawn = Drug withdrawn, N/A = Not applicable.

(b) Outcome: Change = Change in severity, Res. = Resolved, Res. w. sequ. = Resolved with sequelae, Ong. = Ongoing, Fatal = Fatal, Unknown = Unknown.

Programming note: MedDRA version 19.1 or highest at time of data transfer.



15.3: Safety Data

15.3.5: Clinical Laboratory Evaluations

Table 15.3.5.1: Hematology - Shift from Baseline to Worst On-treatment Value - SAF Analysis Set

Laboratory Test	Treatment	Baseline Classification	Worst On-treatment Value				Total n (%)
			Low n (%)	Normal n (%)	High n (%)	Missing n (%)	
PARAM1 (Unit1)	Treatment A N=XXX	Low	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Normal	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		High	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Missing	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Total	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
	Treatment B N=XXX	Low	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Normal	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		High	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Missing	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Total	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
.....							

Note: Baseline is the last measurement prior to the first dose of any trial treatment.
Normal category includes low values for high parameters and high values for low parameters.

Repeat table 15.3.5.1 for:

Table 15.3.5.2: Clinical Chemistry - Shift from Baseline to Worst On-treatment Value - SAF Analysis Set

Programming Note: All parameters and treatments are reported.



- For Hematology: Hematocrit, Hemoglobin, Mean cellular hemoglobin, Mean cellular hemoglobin concentration, Mean cellular volume, Platelet count, Red blood cell count, Red cell distribution width, White blood cell count and differential.

- For Clinical Chemistry: Gamma glutamyl transferase, ALT, Albumin, AP, AST, Bilirubin – direct (only if total bilirubin is outside the normal range), Bilirubin – total, Calcium, Chloride, Bicarbonate, Serum creatinine, Glucose, Potassium, Protein – total, Sodium, Urea, Uric acid, C-reactive protein.

Worst value for non-CTCAE parameters need to be defined with the team for each parameter (e.g. Blood urea nitrogen high). In case worse value could be either low or high for some tests, then report both PARAM low and PARAM high (e.g. Hematocrit low and Hematocrit high).

In case, there is not enough space to display the table on one page, merge the columns for parameter and treatment arm.

15.3: Safety Data

15.3.5: Clinical Laboratory Evaluations

Table 15.3.5.3: Hematology - Shift in Toxicity Grading from Baseline to Highest Grade - SAF Analysis Set

Laboratory Test	Treatment	Baseline NCI-CTCAE Grade	Worst NCI-CTCAE Grade							Total n (%)
			Grade 0 n (%)	Grade 1 n (%)	Grade 2 n (%)	Grade 3 n (%)	Grade 4 n (%)	Missing n (%)		
PARAM1 (Unit1)	Treatment A N=xxx	Grade 0	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 2	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 3	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 4	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Missing	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Total	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)

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15.3: Safety Data

15.3.5: Clinical Laboratory Evaluations

Table 15.3.5.3: Hematology - Shift in Toxicity Grading from Baseline to Highest Grade - SAF Analysis Set

Laboratory Test	Treatment	Baseline NCI-CTCAE Grade	Worst NCI-CTCAE Grade							Total n (%)
			Grade 0 n (%)	Grade 1 n (%)	Grade 2 n (%)	Grade 3 n (%)	Grade 4 n (%)	Missing n (%)		
	Treatment B N=xxx	Grade 0	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 2	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 3	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 4	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Missing	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Total	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)

Shell continues on the next page



15.3: Safety Data

15.3.5: Clinical Laboratory Evaluations

Table 15.3.5.3: Hematology - Shift in Toxicity Grading from Baseline to Highest Grade - SAF Analysis Set

Laboratory Test	Treatment	Baseline NCI-CTCAE Grade	Worst NCI-CTCAE Grade							Total n (%)
			Grade 0 n (%)	Grade 1 n (%)	Grade 2 n (%)	Grade 3 n (%)	Grade 4 n (%)	Missing n (%)		
	Treatment C N=xxx	Grade 0	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 1	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 2	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 3	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Grade 4	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Missing	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)
		Total	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (xx.x)	xxx (100.0)

NCI-CTCAE version 4.03.

Note: Baseline is the last measurement prior to the first dose of any trial treatment.

Repeat table 15.3.5.3 for:

Table 15.3.5.4: Clinical Chemistry - Shift in Toxicity Grading from Baseline to Highest Grade - SAF Analysis Set

Programming Note: Only gradable parameters are reported, examples are provided below – present all gradable parameters that are available.

- For Hematology: Leukocytes high (10E9/L), Leukocytes low (10E9/L), Hemoglobin high (g/L), Hemoglobin low (G/L), Platelets (10E9/L), Lymphocytes high (10E9/L), Lymphocytes low (10E9/L), Neutrophils (10E9/L)...

- For Biochemistry: Bilirubin (µmol/L), AST (U/L), ALT (U/L), Alkaline Phosphatase (U/L), Creatinine (µmol/L), Albumin (G/L), Calcium low (mmol/L), Calcium high (mmol/L), Sodium low (mmol/L), Sodium high (mmol/L), Potassium low (mmol/L), Potassium high (mmol/L), Glucose high (mmol/L), Glucose low (mmol/L), Uric acid high (mmol/L)...

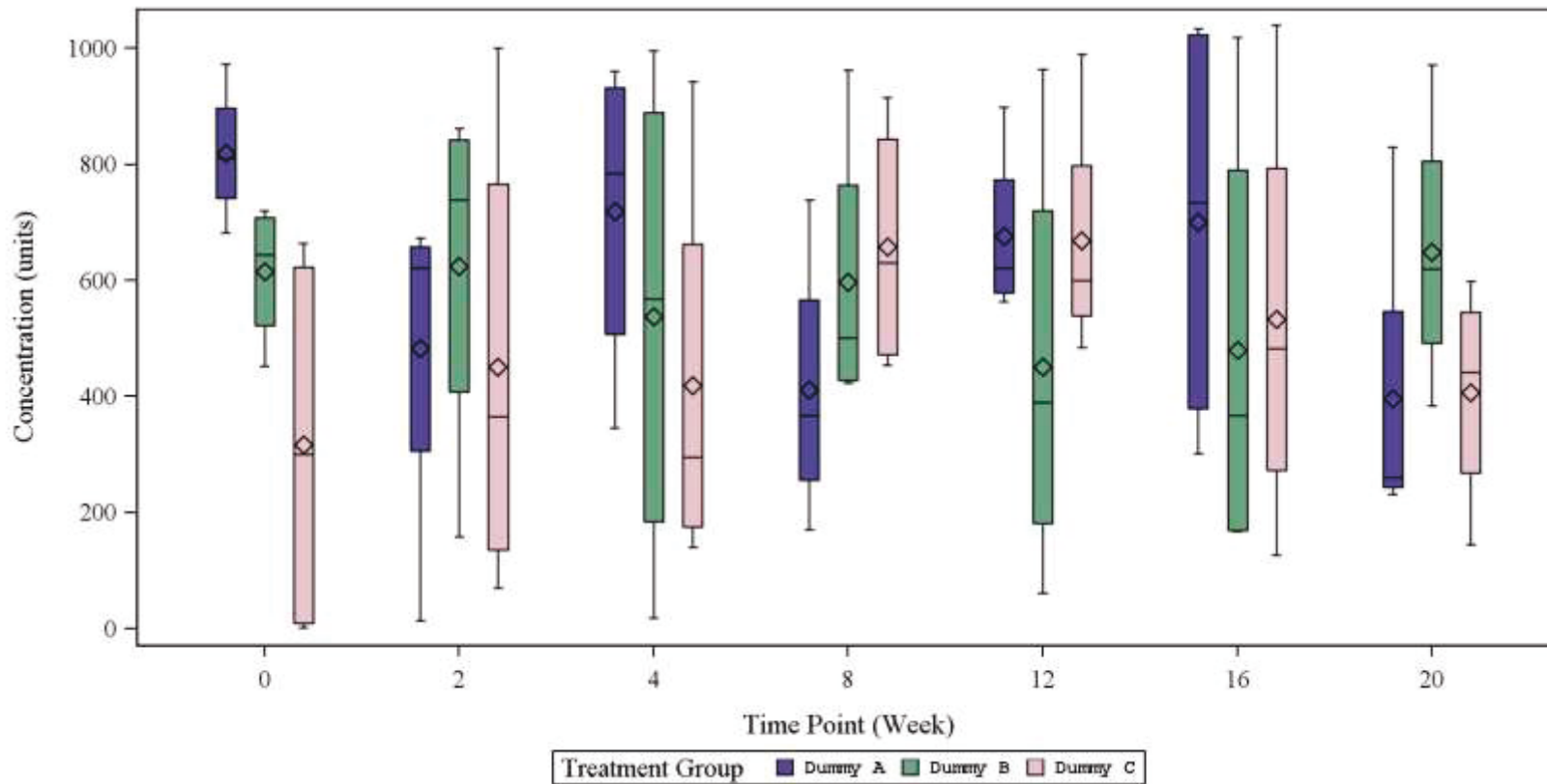
In case there is not enough space to display the table on one page, merge the columns for parameter and treatment.



15.3: Safety Data

15.3.5: Clinical Laboratory Evaluations

Figure 15.3.5.5: Boxplot of Change from Baseline for ALT Values by Visit - SAF Analysis Set



Programming Note: Split figure across multiple pages as the number of weeks increases.

Programming note: The y-axis label should be "<PARAM> Change from Baseline (<units>). The x-axis label should be "Visit". For the partially unblinded outputs, the Treatment Group legend will show "Treatment A" etc.

Use only the following colors (applies to all figures): Cyan, Magenta, Yellow & Black. Also use different symbols for each treatment group.



Repeat figure 15.3.5.5 for:

Figure 15.3.5.6: Boxplot of Change from Baseline for AST Values by Visit - SAF Analysis Set

Figure 15.3.5.7: Boxplot of Change from Baseline for Total Bilirubin Values by Visit - SAF Analysis Set

Figure 15.3.5.8: Boxplot of Change from Baseline for Total White Blood Count Values by Visit - SAF Analysis Set

Figure 15.3.5.9: Boxplot of Change from Baseline for Neutrophils Values by Visit - SAF Analysis Set

18.2 Listings

16.2.1: Subject Disposition

Listing 16.2.1.1: Subject Disposition and Withdrawals - Screening Analysis Set

Treatment Group	Subject Identifier	Screening Date	Randomized?/ Date Randomized/ Stratification Factor (a)	Withdrew Prior to Being Treated?	In SAF Analysis Set?	Withdrew from trial treatment/ Reason for Withdrawal	Date of Last Dose	Discont- inued Trial?/ Primary Reason	Date Discont- inued
Treatment A	XXXXXXX	DDMMYYYY	Yes/DDMMYYYY/ FVC <70%	XXX	XXX	Yes/XXXXXXXX XXXX	DDMMYYYY	Yes/ xxxxxxx	DDMMYYYY
	XXXXXXX	DDMMYYYY	Yes/DDMMYYYY/ FVC ≥70%	XXX	XXX	No		No	
	XXXXXXX	DDMMYYYY	Yes/DDMMYYYY/ XXXXXX	XXX	XXX	No		No	
								
Treatment B	XXXXXXX	DDMMYYYY	Yes/DDMMYYYY/ XXXXXX	XXX	XXX	No		No	
	...								
<i>Etc.</i>									

(a) Stratification Factor from eCRF: FVC <70% = Baseline FVC <70% predicted, FVC ≥70% = Baseline FVC ≥70% predicted.
Note: Baseline is defined as the last non-missing assessment value prior to randomization.

Programming Note: If a subject is screened but not yet randomized, treatment group will be Not Randomized. If they are screen failed then treatment group is Screen Failures.
If there are subjects in the Not Randomized group then add the following footnote (note indentation to line up with baseline footnote):

'Not Randomized' treatment group contains subjects who are still in screening at time of data cutoff.
Ensure any "Other" reasons for treatment and trial termination are specified, "Other: xxxxxx".



16.2.1: Subject Disposition
Listing 16.2.2.1: Important Protocol Deviations - ITT Analysis Set

Treatment Group	Subject Identifier	Type	Deviation details
Treatment A	XXXXXXX	Eligibility and Entry Criteria	XXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXXXXXX
		Efficacy Criteria	XXXXXXXXXXXXXXXXXXXXX
	XXXXXXX	XXXXXXXXXXXXX	XXXXXXXXXXXXXXXXXXXXX
	XXXXXXX	XXXXXXXXXXXXX	XXXXXXXXXXXXXXXXXXXXX
		XXXXXXXXXXXXXXXXXXXXX
Treatment B	XXXXXXX	XXXXXXXXXXXXX XXXXXXXXXXXXX	XXXXXXXXXXXXXXXXXXXXX XXXXXXXXXXXXXXXXXXXXX
<i>Etc.</i>	...		

Programming note: Present each deviation on a new line.



16.2.4: Demographic Data
Listing 16.2.4.1: Subject Demographics - SAF Analysis Set

Treatment Group	Subject Identifier	Sex	Birth Date	Age (years)	Race	Ethnicity
Treatment A	XXXXXXX	Male	DDMMYYYY	XX	XXXXXXX	XXXXX
	XXXXXXX	Female	YYYY	XX	XXXXXXX	XXXXX
	XXXXXXX	XXXX	DDMMYYYY	XX	XXXXXXX	XXXXX
					
Treatment B	XXXXXXX	XXXX	DDMMYYYY	XX	XXXXXXX	XXXXX
	...					
<i>Etc.</i>						

Note: Age is relative to the date of informed consent.



16.2.4: Demographic Data

Listing 16.2.4.2: Subject Medical History - SAF Analysis Set

Treatment Group	Subject Identifier	System Organ Class	Preferred Term	Verbatim Text	Start Date	End Date	Related to Study Condition?
Treatment A	XXXXXXX	XXXXXXX	XXXXXXX	XXXXXXX	DDMMYYYY	Ongoing	Yes
		XXXXXXX	XXXXXXX	XXXXXXX	DDMMYYYY	DDMMYYYY	No
		XXXXXXX	XXXXXXX	XXXXXXX	DDMMYYYY	XXXXXXXX	Unknown
.....							
Treatment B	XXXXXXX	XXXXXXX	XXXXXXX	XXXXXXX	DDMMYYYY	XXXXXXXX	
...							
<i>Etc.</i>							

MedDRA version xx.x.

Programming note: MedDRA version 19.1 or highest at time of data transfer.

16.2.4: Demographic Data

Listing 16.2.4.3: Subject Disease History - SAF Analysis Set

Treatment Group	Subject Identifier	Date of Diagnosis	Time Since Date of Diagnosis (years)	Status	Systemic Sclerosis Symptom History (a)	ACR/EULAR Classification Criteria for Systemic Sclerosis (b)
Treatment A	XXXXXXX	DDMMYYYY	XX.XX	Diffuse cutaneous SSc	Skin Thickening/ Tendon friction rubs/ XXXXXXX	Fingertip lesions: Fingertip pitting scars/ Abnormal nailfold capillaries/ XXXXXX
	XXXXXXX	DDMMYYYY	XX.XX	Limited cutaneous SSc	XXXXXXX/ XXXXXXXX / XXXXXXXXXX/ XXXXXXX	XXXXXXX/ XXXXXXXX / XXXXXXXXXX/ XXXXXXX
	XXXXXXX	DDMMYYYY	XX.XX	XXXXXXX	XXXXXXX/ XXXXXXXX / XXXXXXXXXX/ XXXXXXX	XXXXXXX/ XXXXXXXX / XXXXXXXXXX/ XXXXXXX
					
Treatment B	XXXXXXX	DDMMYYYY	XX.XX	XXXXXXXXX	XXXXXXX/ XXXXXXXX / XXXXXXXXXX/ XXXXXXX	
	...					

Etc.

(a) Where 'Yes' was selected on the eCRF.

(b) Where 'Present' was selected on the eCRF.

Note: Time since date of diagnosis is relative to date of informed consent.

Programming note: Concatenate all Systemic Sclerosis Symptom History sub-categories where 'Yes' is ticked on the eCRF, concatenate all ACR/EULAR Classification Criteria for Systemic Sclerosis sub-categories where 'Present' is ticked on the eCRF.



16.2.4: Demographic Data

Listing 16.2.4.4: Previous Medications - SAF Analysis Set

Treatment Group	Subject Identifier	ATC Level 2/ Preferred Term/ Reported Name	Start Date/ End Date (Trial Day)	Dose (Unit)	Frequency	Route	Reason (Indication)	Reason(s)
Treatment A	XXXXXXX	XXXXXXXXXXXXX/ XXXXXXXXX/ XXXXXXXXX	DDMMYY (xx) / DDMMYY (xx)	XXX xx	XX	XX	XXXXXXXXX	Adverse event/ Disease related condition
		XXXXXXXXXXXXX/ XXXXXXXXX/ XXXXXXXXX	DDMMYY (xx) / DDMMYY (xx)	XXX xx	Other: XXX	XX	XXXXXXXXX	XXXXXXXXX
							
Treatment B	XXXXXXX	XXXXXXXXXXXXX/ XXXXXXXXX/ XXXXXXXXX	DDMMYY (xx) / DDMMYY (xx)	XXX	XX	Other: XX	XXXXXXXXXXXXX	XXXXXX
	...							
<i>Etc.</i>								

WHO Drug Dictionary version Enhanced September 2016.

Note: In case multiple ATCs are assigned to a medication, all ATCs will be reported.

Programming note: WHO Drug Dictionary version Enhanced September 2016 or highest at time of data transfer.

Repeat listing 16.2.4.4 for:

Listing 16.2.4.5: Concomitant Medications - SAF Analysis Set

Change "Previous Medication" to "Concomitant Medication"

Listing 16.2.4.6: Background Therapy - SAF Analysis Set

Change "Previous Medication" to "Background Therapy".

16.2.5: Treatment Compliance and Exposure

Listing 16.2.5.1: Trial Treatment Administration Details - SAF Analysis Set

Treatment Group	Subject Identifier	Visit	Preparation Date (Time) / Kit Number	Admin. Start Date (Time) / End Date (Time)	Duration of infusion (hours)	Change in Dose?	Planned Volume (mL)	Actual Volume (mL)	Infusion Completed as Planned?
A	XXXXXXX	XXXXXXX	DDMMYY (HH:MM) / XXXX	DDMMYY (HH:MM) / DDMMYY (HH:MM)	XX.XX	No	XXX	XXX	No: Adverse event
		XXXXXXX	DDMMYY (HH:MM) / XXXX	DDMMYY (HH:MM) / DDMMYY (HH:MM)	XX.XX	Dose adjusted: Adverse event	XXX	XXX	Yes
		XXXXXXX	DDMMYY (HH:MM) / XXXX	DDMMYY (HH:MM) / DDMMYY (HH:MM)	XX.XX	XXXXXXX	XXX	XXX	XXX
		XXXXXXX*				No dose: Missed visit	XXX		No: Other, No dose: missed visit

Etc.

* = Visit derived due to the subject not completing the "STUDY TREATMENT ADMINISTRATION DETAILS" eCRF page at the missed visit.

Note: Admin. = Administration.

Duration of infusion is calculated as: End time of administration - start time of administration.



16.2.5: Treatment Compliance and Exposure

Listing 16.2.5.2: Overall Exposure to Trial Treatment - SAF Analysis Set

Treatment Group	Subject Identifier	Treatment Duration (weeks)	Number of Infusions	Number of Dose Adjustments	Number of Infusions Not Completed as Planned	Overall Treatment Adherence (%)
Treatment A	XXXXXXX	XXX	XX	X	X	XX

Etc.

Note: Treatment duration is calculated as: (Last infusion date prior to data cut-off - first infusion date + 28) / 7 days.

Subject overall treatment adherence is calculated as: (sum of actual volume received/ sum of planned Volume, up to and including their last dose recorded on their treatment termination visit)*100.

16.2.6: Efficacy Data

Listing 16.2.6.1: FVC Volume (mL) and FVC % Predicted - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	FVC (mL)		FVC % Predicted	
				Observed	Change from Baseline	Observed	Change from Baseline
Treatment A	XXXXXXXX	XXXXXXXX	DDMMYYYY (xx)	XX*		XX*	
		XXXXXXXX	DDMMYYYY (xx)	XX	XX	XX	XX
		XXXXXXXX	DDMMYYYY (xx)	XX	XX	XX	XX

Etc.

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.

Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data

Listing 16.2.6.2: mRSS Total Score - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	mRSS Total Score	
				Observed	Change from Baseline
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	XX*	
		XXXXXXX	DDMMYYYY (xx)	XX	XX
		XXXXXXX	DDMMYYYY (xx)	XX	XX

Etc.

* Baseline value.

^ = diffuse cutaneous systemic sclerosis at baseline.

Note: Baseline is defined as the last non-missing assessment value prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.
Trial day is defined relative to the date of randomization.

16.2.6: Efficacy Data

Listing 16.2.6.3: Clinically Meaningful Disease Progression - mITT Analysis Set

Treatment Group	Subject Identifier	Date of Progression (Trial Day)	Progression Criteria Fulfilled	SSc-ILD progression		Was the Trial Treatment Discontinued?
				Was Mycophenolate Dose Increased?	Was a New Immunosuppressant or Biologic Drug Initiated?	
Treatment A	XXXXXXX	DDMMYYYY (xx)	SSc-ILD: Relative decrease from baseline in FVC % predicted >10% SSc other than ILD: Scleroderma renal crisis	No	No	No
		DDMMYYYY (xx)				XX

Etc.

Note: Trial day is defined relative to the date of randomization.

Programming note: Only display subjects with clinically meaningful disease progression. Concatenate all details selected for SSc other than ILD (including the free text details for 'Other'). If both SSc-ILD and SSc other than ILD were selected on the same date, then display on a new line.



16.2.6: Efficacy Data
Listing 16.2.6.4: DLCO - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	DLCO as recorded on eCRF		DLCO Result in SI Unit (mmol of CO/min/kPa)		DLCO % Predicted	
				Observed (unit)		Observed	Change from Baseline	Observed	Change from Baseline
Treatment A	XXXXXXXX	XXXXXX	DDMMYYYY (xx)	XX (xxx)		XX*		XX*	
		XXXXXX	DDMMYYYY (xx)	XX (xxx)		XX	XX	XX	XX
		XXXXXX	DDMMYYYY (xx)	XX (xxx)		XX	XX	XX	XX

Etc.

* Baseline value.
Note: Baseline is defined as the last non-missing assessment value prior to randomization. If such a value is missing then the last value prior to the first trial treatment administration will be used.
Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data
 Listing 16.2.6.5: KCO - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Test overread grading	VA	DLCO/ VA	KCO	
							Observed	Change from baseline
Treatment A	XXXXXXXX	XXXXXXXX	DDMMYYYY (xx)	Acceptable	xx	xx+	xx*	
		XXXXXXXX	DDMMYYYY (xx)	Unacceptable	xx	xx	xx	xx
		XXXXXXXX	DDMMYYYY (xx)	xxxxx				

Etc.

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to randomization.
 Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data
Listing 16.2.6.6: Digital Ulcers - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Completed Status	Date of Assessment (Trial Day)	Number of						Total Digital Ulcer count
					Active Ulcers		Undetermined Ulcers		Healed Ulcers		
					Absolute Value	CFB	Absolute Value	CFB	Absolute Value	CFB	
Treatment A	XXXXXXXX	XXXXXX	Done	DDMMYYYY (xx)	XXX*		XXX*		XXX*		XXX
		XXXXXX	Not Done								
		XXXXXX	XXXXXXXX	DDMMYYYY (xx)	XXX	XXX	XXX	XXX	XXX	XXX	XXX

Etc.

* Baseline value.
Note: Baseline is defined as the last non-missing assessment value prior to randomization.
CFB = Change from Baseline.
Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data

Listing 16.2.6.7: Mahler BDI/TDI - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Question	Score		
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	Done	Functional Impairment	x		
					Magnitude of Task	x		
					Magnitude of Effort	x		
							Total	x
			XXXXXXX	DDMMYYYY (xx)	Done	Changes in Functional Impairment	x	
		Changes in Magnitude of Task				x		
		Changes in Magnitude of Effort				x		
						Total	x	
			XXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill			

Etc.

Note: Baseline is defined as the last non-missing assessment value prior to randomization. The BDI is completed at baseline and the TDI is completed at subsequent visits.
Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data

Listing 16.2.6.8: SGRQ - Question Scores - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Question (component)	Answer(s)	Weight	
Treatment A	XXXXXXXX	XXXXXXXX	DDMMYYYY (xx)	Done	xxxxxx (S)	xxxx	x	
					xxxxxx (I)	xxxx	x	
					xxxxxx	xxxx	x	
						...	xxxxxx	x
		XXXXXXXX	DDMMYYYY (xx)	Done	xxxxxx	xxxx	x	
					xxxxxx	xxxxxx	x	
					xxxxxx	xxxx	x	
		XXXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill	xxxxxx	xxxxxx	x	

Etc.

Note: Component abbreviations: S = Symptoms, I = Impacts, A = Activity.
Trial day is defined relative to the date of randomization.



16.2.6: Efficacy Data

Listing 16.2.6.9: SGRQ - Component and Total Scores - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Component	Score	Change from baseline
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	Done	Symptoms	x	
					Activity	x	
					Impacts	x	
					Total	x*	
		XXXXXXX	DDMMYYYY (xx)	Done	Symptoms	x	
					Activity	x	
XXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill	Impacts	x			
			Total	x	x		

Etc.

* Baseline value (total scores only).

Note: Baseline is defined as the last non-missing measurement prior to randomization.

Trial day is defined relative to the date of randomization.

The component score is 100 x summed weights from all positive items in that component divided by the sum of weights for all items in that component. The total score is 100 x summed weights from all positive items in the questionnaire divided by the sum of weights for all items in the questionnaire.



16.2.6: Efficacy Data

Listing 16.2.6.10: Leicester Cough Index - Question Scores - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Question (domain)	Score
Treatment A	XXXXXXXX	XXXXXXXX	DDMMYYYY (xx)	Done	xxxxxx (P)	x
					xxxxxx (Psy)	x
					xxxxxx (S)	x
		XXXXXXXX	DDMMYYYY (xx)	Done	xxxxxx	x
					xxxxxx	x
						x
XXXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill		x		

Etc.

Note: Trial day is defined relative to the date of randomization.
 Domain abbreviations: P = Physical, Psy = Psychological, S = Social.
 1 is the most negative score, 7 is the most positive score.



16.2.6: Efficacy Data

Listing 16.2.6.11: Leicester Cough Index - Domain and Total Scores - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Domain	Score	Change from baseline
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	Done	Physical	x	
					Psychological	x	
					Social	x	
					Total	x*	
		XXXXXXX	DDMMYYYY (xx)	Done	Physical	x	
					Psychological	x	
					Social	x	
		XXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill	Total	x	x

Etc.

* Baseline value (total score only).

Note: Baseline is defined as the last non-missing assessment value prior to randomization.

Trial day is defined relative to the date of randomization.

Domain scores are the sum of all scores in the domain divided by the number of questions within that domain.



16.2.6: Efficacy Data

Listing 16.2.6.12: Physician's Global Assessment - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	Done/ Not done - Reason	Score	Change from baseline
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	Done	x*	
		XXXXXXX	DDMMYYYY (xx)	Done	x	x
		XXXXXXX	DDMMYYYY (xx)	Not done - Subject felt too ill		

Etc.

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to randomization. Trial day is defined relative to the date of randomization.

The score is based on the question: "On a scale of 0-100, where would you rate the overall effect systemic sclerosis has on your patient at this time", where 0 = has no effect at all and 100 = worst possible effect.

Repeat listing 16.2.6.12 for:

Listing 16.2.6.13: Patient's Global Assessment - mITT Analysis Set

Update footnote: The score is based on the question: "On a scale of 0-100, where would you rate the overall effect systemic sclerosis has on you at this time", where 0 = has no effect at all and 100 = worst possible effect.

16.2.6: Efficacy Data

Listing 16.2.6.14: SHAQ - VAS Scores - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Overall disease activity		Raynaud's phenomenon		Finger ulcers		Breathing problems		Intestinal problems	
			Observed	CFB	Observed	CFB	Observed	CFB	Observed	CFB	Observed	CFB
Treatment A	XXXXXXXX	XXXXXX	xx*		xx*		xx*		xx*		xx*	
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		...										
	XXXXXXXX	XXXXXX	xx^		xx^		xx^		xx^		xx^	
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		XXXXXX	xx	xx	xx	xx	xx	xx	xx	xx	xx	xx
		...										
	XXXXXXXX	XXXXXX	xx		xx		xx		xx		xx	

Etc.

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to randomization.

The VAS scores range from 0 (most positive response) to 100 (most negative response).

CFB = Change from baseline.

16.2.6: Efficacy Data

Listing 16.2.6.15: EQ-5D VAS Score - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	VAS Score	
				Observed	CFB
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	X*	
				X	X
				X	X
	...				
...	XXXXXXX		DDMMYYYY (xx)	X*	
				X	X
				X	X
	...				

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to randomization.

Trial day is defined relative to the date of randomization.

CFB = Change from Baseline.



16.2.6: Efficacy Data

Listing 16.2.6.16: HRCT Results - mITT Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Assessment (Trial Day)	QLF (%) (a)		Extent of ILD (%)	
				Observed	Change from Baseline	Observed	Change from Baseline
Treatment A	XXXXXXX	XXXXXXX	DDMMYYYY (xx)	X*		X*	
			DDMMYYYY (xx)	X	X	X	X
			DDMMYYYY (xx)	X	X	X	X
	...						
...	XXXXXXX		DDMMYYYY (xx)	X*		X*	
			DDMMYYYY (xx)	X	X	X	X
			DDMMYYYY (xx)	X	X	X	X
	...						

* Baseline value.

(a) QLF in region of highest severity.

Note: Baseline is defined as the last non-missing assessment value prior to randomization.

Trial day is defined relative to the date of randomization.

16.2.7: Display of Adverse Events
Listing 16.2.7.1: All Adverse Events - SAF Analysis Set

Treatment Group	Subject Identifier	System Organ Class/ Preferred Term/ Verbatim Text	Start Date/ End Date (Trial Day)	Relationship /Timing to Trial Trt	Toxic- ity Grade	Causal- ity Factors	Action Taken With Trial Trt/Other Action	Outcome
Treatment A	XXXXXXX	XXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXX+*	DDMMYYYY (xx) / DDMMYYYY (xx)	XXXXXXXXXXXXX/ XXXXXXX	X	XX	XXXXXXXXXXXXX/ XXXXXXXXXXXXX	XXXXXXXXXXXXX

Etc.

+ Serious Adverse Event.
Adverse Event of Special Interest.
* Treatment Emergent Adverse Event.
Note: Trial day is defined relative to the date of randomization.



16.2.7: Display of Adverse Events
Listing 16.2.7.2: Serious Adverse Events - SAF Analysis Set

Treatment Group	Subject Identifier	System Organ Class/ Preferred Term/ Verbatim Text	Start Date/ End Date (Trial Day)	Seriousness Criteria	Relationship/Action Taken with Trial Trt
Treatment A	XXXXXXX	XXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXXXXXX #	DDMMYYYY (xx) / DDMMYYYY (xx)	Requires/prolongs hospitalization	XXXXXXXXXXXXXXXXXXXXX/ XXXXXXXXXXXXX

. . .

Etc.

Adverse Event of Special Interest.
\$ Non-Treatment Emergent Adverse Event.
Note: Trial day is defined relative to the date of randomization.



16.2.7: Display of Adverse Events
Listing 16.2.7.3: Listing of Deaths - SAF Analysis Set

Treatment Group	Subject Identifier	Informed Consent Date	Randomization Date	First/Last Trial Trt. Date	Death Date (days since last dose)	Primary Reason for Death
Treatment A	XXXXXXX	DDMMYYYY	DDMMYYYY	DDMMYYYY/ DDMMYYYY	DDMMYYYY (XXX)	XXXXXXXXXXXXXXXXXXXXXXXXXX
	. . .					

Etc.



16.2.8: Clinical Laboratory Evaluations
Listing 16.2.8.1: Hematology Parameters - SAF Analysis Set

Treatment Group	Subject Identifier	Visit	Date of Sample (Trial Day)	Comment	Parameter (Unit)	Normal Range	Result	NCI-CTC grade (a)
Treatment A	XXXXXXX	XXXXXXXX	DDMMYYYY (xx)	XXXXXXXXXX	XXXXXXXXXX	XX - XX	XXX*	x/x
					XXXXXXXXXX	XX - XX	XXX L	x
					XXXXXXXXXX	XX - XX	XXX H	
	. . .							

Etc.

(a) NCI-CTCAE version 4.03 for gradeable parameters. For parameters with both low and high grades, the grades are presented as Low/High.

L = value below normal range, H = value above normal range.

* Baseline value.

Note: Baseline is defined as the last non-missing assessment value prior to first dose of trial treatment.

Trial day is defined relative to the date of randomization.

Programming Note: Repeat listing 16.2.8.1 for:

Listing 16.2.8.2: Clinical Chemistry Parameters - SAF Analysis Set



16.2.9: Other Safety Evaluations
Listing 16.2.9.1: Vital Signs - SAF Analysis Set

Treatment Group	Subject Identifier	Visit	Assessment Date (Trial Day)	Blood Pressure (mmHg)		Pulse Rate (beats/min)	Temperature (°C) (Location)	Height (cm)	Weight (kg)
				Systolic	Diastolic				
Treatment A	XXXXXXX	XXXXXXXX	DDMMYYYY (***)	XXX	XXX	XXX	XX (XXXXXX)	XX	XX
		XXXXXXXX	DDMMYYYY (***)	XXX	XXX	XXX	XX (XXXXXX)		
	. . .								

Etc.

~~Note: Trial day is defined relative to the date of randomization.~~



16.2.9: Other Safety Evaluations
Listing 16.2.9.2: Electrocardiogram Data - SAF Analysis Set

Treatment Group	Subject Identifier	Visit	Assessment Date	(Trial Day)	Result	Clinically Significant? (a)
Treatment A	XXXXXXXX	XXXXXXXX	DDMMYYYY	(**)	Normal	
		XXXXXXXX	DDMMYYYY	(**)	Abnormal	No
		XXXXXXXX	DDMMYYYY	(**)	XXXXXXXXXX	xxxxxx
. . .						

Etc.

(a) If result is 'Abnormal'.

~~* Baseline value.~~

~~Note: Baseline is defined as the last non-missing assessment value prior to first dose of trial treatment.~~

~~— Trial day is defined relative to the date of randomization.~~

