

## TRIAL STATISTICAL ANALYSIS PLAN

**c29769550-01**

<b>BI Trial No.:</b>	1405-0009
<b>Title:</b>	Relative bioavailability of a single oral dose of BI 1323495 when administered alone or in combination with multiple oral doses of itraconazole in healthy male subjects (an open-label, one-way crossover study)  (including Protocol Amendment No.1 [c26556329-02])
<b>Investigational Product:</b>	BI 1323495
<b>Responsible trial statisticians:</b>	          Phone Fax:
<b>Date of statistical analysis plan:</b>	21OCT2019 SIGNED
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## 2. LIST OF ABBREVIATIONS

See Medicine Glossary:

website: [glossary](#)

Term	Definition / description
AESI	Adverse event of special interest
ALT	Alanine transaminase
ANOVA	Analysis of variance
AST	Aspartate transaminase
AUC <sub>0-tz</sub>	Area under the concentration-time curve of the analyte in plasma over the time interval from 0 to the last quantifiable data point
AUC <sub>0-∞</sub>	Area under the concentration-time curve of the analyte in plasma over the time interval from 0 extrapolated to infinity
BMI	Body mass index
BWU	Bioavailability/Bioequivalence, Within-Subject Design, uncontrolled w.r.t. time
CI	Confidence interval
C <sub>max</sub>	Maximum measured concentration of the analyte in plasma
CV	Arithmetic coefficient of variation
DBLM	Database lock meeting
DILI	Drug induced liver injury
gCV	Geometric coefficient of variation
gMean	Geometric mean
ITZ	Itraconazole
LLT	Lower level term
Max	Maximum
MedDRA	Medical Dictionary for Regulatory Activities
Min	Minimum
N	Number non-missing observations
P10	10th percentile
P90	90th percentile
PKS	PK parameter analysis set
Q1	1st quartile

Term	Definition / description
Q3	3rd quartile
R / Ref	Reference treatment
RAGe	Report Appendix Generator system
SD	Standard deviation
SOC	System organ class
T	Test treatment
TS	Treated set
ULN	Upper limit of normal
WHO-DD	World Health Organization Drug Dictionary
XPKISTAT	Library of SAS <sup>®</sup> Macros for PK analysis

### 3. INTRODUCTION

As per ICH E9 (1), the purpose of this document is to provide a more technical and detailed elaboration of the principal features of the analysis described in the protocol, and to include detailed procedures for executing the statistical analysis of the primary and secondary variables and other data.

This Trial statistical analysis plan (TSAP) assumes familiarity with the Clinical Trial Protocol (CTP), including protocol amendment. In particular, the TSAP is based on the planned analysis specification as written in CTP Section 7 “Statistical Methods and Determination of Sample Size”. Therefore, TSAP readers may consult the CTP for more background information on the study, e.g., on study objectives, study design and population, treatments, definition of measurements and variables, planning of sample size, randomisation.

Study data (including data entered in the RAVE EDC system and external data provided by suppliers) will be stored in a Clinical Data Repository (CDR).

Pharmacokinetic (PK) parameters will be calculated using Phoenix WinNonlin<sup>TM</sup> software (version 6.3 or higher, Certara USA Inc., Princeton, NJ, USA).

The statistical analyses will be performed within the validated working environment CARE, including SAS<sup>TM</sup> (current Version 9.4, by SAS Institute Inc., Cary, NC, USA), and a number of SAS<sup>TM</sup>-based tools (e.g., macros for the analyses of AE data or laboratory data; Report Appendix Generator system (RAGe) for compilation/formatting of the CTR appendices).

#### 4. CHANGES IN THE PLANNED ANALYSIS OF THE STUDY

All analyses as planned in the CTP will be performed and are described in more detail in this TSAP. The following change compared to the protocol will be made:

The PKS defined in the CTP will be called “Pharmacokinetic **parameter** analysis set” instead of “Pharmacokinetic analysis set”.



## 5. ENDPOINTS

### 5.1 PRIMARY ENDPOINTS

**Section 2.1.2 of the CTP:** *The following pharmacokinetic parameters will be determined for BI 1323495:*

- $AUC_{0-t_z}$  (area under the concentration-time curve of the analyte in plasma over the time interval from 0 to the last quantifiable data point)
- $C_{max}$  (maximum measured concentration of the analyte in plasma)

The pharmacokinetic parameters listed in Sections 2.1 and 2.2 of the protocol for BI 1323495 will be calculated according to the BI Standard Operating Procedure (SOP) ‘Standards and processes for analyses performed within Clinical Pharmacokinetics/Pharmacodynamics’ [001-MCS-36-472] ([7](#)).

### 5.2 SECONDARY ENDPOINTS

#### 5.2.1 Key secondary endpoints

This section is not applicable as no key secondary endpoints have been defined in the CTP.

#### 5.2.2 Secondary endpoints

**Section 2.1.3 of the CTP:** *The following pharmacokinetic parameter will be determined for BI 1323495:*

- $AUC_{0-\infty}$  (area under the concentration-time curve of the analyte in plasma over the time interval from 0 extrapolated to infinity)

### **5.3.2 Safety endpoints**

**Section 2.2.2.2 of the CTP:** *Safety and tolerability of the trial treatments will be assessed based on:*

- *Adverse events (including clinically relevant findings from the physical examination)*
- *Safety laboratory tests*
- *12-lead ECG*
- *Vital signs (blood pressure, pulse rate)*

## 6. GENERAL ANALYSIS DEFINITIONS

### 6.1 TREATMENTS

For basic study information on investigational products, assignment of treatment sequences and selection of doses, please see CTP, Sections 3 and 4.

The study will be performed as an open-label trial with two treatments (T and R) and one fixed sequence (R then T). There will be a washout period of at least 7 days between the administrations of BI 1323495 under Reference treatment and BI 1323495 under Test treatment.

In total, it was planned to assign 14 healthy male subjects.

For details of dosage and formulation see Table 6.1: 1 below:

Table 6.1: 1 Treatments and labels used in the analysis

Treatment			Short label
R	BI 1323495, 1*	tablet, qd	BI
T	BI 1323495, 1*	tablet +Itraconazole 20ml, 10mg/ml, qd	BI + ITZ

**Section 1.2.3 of the CTP:** *The Residual Effect Period (REP, i.e., the period after the last dose with measurable drug levels and/or pharmacodynamic effects still likely to be present) of BI 1323495 is approximately 7 days. [...]*

*For the use of itraconazole in Treatment T, the REP is defined as 9 days after last administration of itraconazole on Day 7 in Period 2.*

The following separate study phases will be defined for the analyses of AEs:

- **Screening** (ranging from 0:00 h on day of informed consent until first administration time of BI 1323495 alone in treatment period 1)
- **On treatment**
  - **BI alone** (labelled “**BI**”, including REP; i.e. ranging from the time of first administration of BI 1323495 alone in treatment period 1 until 0:00 h on the 7th day thereafter)
  - **Itraconazole alone** (labelled “**ITZ**”) (ranging from the time of first administration of Itraconazole until time of BI 1323495 administration in treatment period 2)
  - **BI + ITZ** (ranging from time of BI 1323495 administration in treatment period 2 until 0:00 h on the 9th day after the last administration of Itraconazole OR until 0:00h on the 7th day after the BI 1323495 administration in treatment period 2, whatever occurs later)
- **Follow-up (FU)** (labelled “**F/U BI**”, “**F/U BI+ITZ**”, “**F/U ITZ**”)
  - **F/U BI** (ranging from the end of REP of BI 1323495 in treatment period 1 until the first administration of itraconazole in period 2 or until trial termination, whatever occurs first)
  - **F/U BI+ITZ** (ranging from the end of REP of BI 1323495 or itraconazole (whatever occurs later) in treatment period 2 until trial termination)
  - **F/U ITZ** (ranging from the end of REP of itraconazole in treatment period 2 until trial termination if subject discontinues treatment prior to BI 1323495 drug administration – if applicable)

Two types of AE displays will be provided in the report:

- A) Section 15.3 and Appendix 16.1.13.1.8 (for ClinicalTrials.gov and EudraCT only) of the CTR displays:

In these displays, the on treatment phase will be analysed (labelled with the name of the study treatment (short label)). Screening and follow-up periods will not be included in this analysis.

The following total will be provided in addition (Section 15.3 only):

- a total over all on treatment phases involving BI ("**BI Total on treatment**")
- a total over all on treatment phases included in this analysis ("**Total on treatment**")

**B)** Section 15.4 and Appendix 16.1.13.1.8 (except for ClinicalTrials.gov and EudraCT) of the CTR displays:

- Screening
- On treatment (labelled with the name of the study treatment (short label))
- Follow-up BI (labelled "**FU BI**")
- Follow-up BI+ITZ (labelled "**FU BI+ITZ**")
- Follow-up ITZ (labelled "**F/U ITZ**") – if applicable

In Section 16.1.13.1.8 AE tables, the following totals will be provided in addition:

- a total over all on treatment phases involving BI ("**BI Total**")
- a total over all study phases ("**Total**")

For detailed information on the handling of the treatments refer to Technical TSAP ADS plan and Analysis Data Reviewers guide.

## 6.2 IMPORTANT PROTOCOL DEVIATIONS

Data discrepancies and deviations from the CTP will be identified for all treated subjects.

**Section 7.3 of the CTP:** *Adherence to the protocol will be assessed by the trial team.*

*Important protocol deviation (IPD) categories will be suggested in the IQRMP plan, IPDs will be identified no later than in the Report Planning Meeting, and the IPD categories will be updated as needed.*

Consistency check listings (for identification of deviations of time windows) and a list of protocol deviations (e.g. deviations in drug administration, in blood sampling times, etc.) will be provided to be discussed at the report planning meeting (RPM). At this meeting, all manual deviations identified at the sites by the CRAs and deviations too complex to program will be reviewed by the trial team to decide which are considered important. For definition of important protocol deviations (iPD), and for the process of identification of these, refer to the Boehringer Ingelheim (BI) SOP "Identify and Manage Important Protocol Deviations (iPD)" [\(2\)](#).

If any iPDs are identified, they are to be summarised into categories and will be captured in an accompanying Excel spreadsheet [\(3\)](#). Categories which are considered to be iPDs in this trial are defined in the integrated quality and risk management plan (IQRMP). If the data show other iPDs, the definition in the IQRMP will be supplemented accordingly by the time of the Report Planning Meeting.

The iPDs will be summarised and listed. [Table 6.2: 1](#) below specifies which kind of iPDs could potentially lead to exclusion from which analysis set. The decision on exclusion of

patients from analysis sets will be made at the latest at the Report Planning Meeting, after discussion of exceptional cases and implications for analyses.

Table 6.2: 1 Handling of iPDs

iPD code	iPD Category & Brief Description	Excluded from which analysis set
iPD B1	Informed consent not available/not done	All
iPD B2	Informed consent too late	None
iPD A1	Inclusion Criteria Not Met	PKS
iPD A2	Exclusion Criteria Met	PKS
iPD C1	Incorrect trial medication taken	PKS
iPD C2	Randomisation not followed	PKS
iPD C3	Non-compliance	PKS
iPD C4	Medication code broken inappropriately	PKS
iPD C5	Incorrect intake of trial medication	PKS
iPD C6	Improper washout between treatments	PKS
iPD D1	Prohibited medication use	PKS
iPD D2	Mandatory medication not taken	PKS
iPD D3	Improper washout of concomitant medication	PKS
iPD E1	Certain violations of procedures used to measure primary or secondary data	PKS
iPD F1	Certain violations of time schedule used to measure primary or secondary data	PKS
iPD G1	Incorrect intake of meal	PKS
iPD G2	PDs affecting safety and rights of subjects	None

## 6.3 SUBJECT SETS ANALYSED

### Section 7.3 of the CTP:

- *Treated set (TS): The treated set includes all subjects who were entered and treated with at least one dose of study drugs. The treated set will be used for safety analyses.*
- *Pharmacokinetic parameter analysis set (PKS): This set includes all subjects in the treated set (TS) who provide at least one PK endpoint that was defined as primary or secondary and was not excluded due to a protocol deviation relevant to the evaluation of PK or due to PK non-evaluability (as specified in the CTP subsection 'Pharmacokinetics'). Thus, a subject will be included in the PKS, even if he contributes only one PK parameter value for one period to the statistical assessment. Descriptive and model based analyses of PK parameters will be based on the PKS.*

[...]

*Plasma concentration data and parameters of a subject will be included in the statistical pharmacokinetic (PK) analyses if they are not flagged for exclusion due to a protocol deviation relevant to the evaluation of PK (to be decided no later than in the Report Planning Meeting) or due to PK non-evaluability (as revealed during data analysis, based on the criteria specified below). Exclusion of a subject's data will be documented in the CTR.*

*Relevant protocol deviations may be*

- *Incorrect trial medication taken, i.e. the subject received at least one dose of trial medication the subject was not assigned to*
- *Incorrect dose of trial medication taken*
- *Use of restricted medications*

*Plasma concentrations and/or parameters of a subject will be considered as non-evaluable, if for example*

- *The subject experienced emesis that occurred at or before two times median  $t_{max}$  of the respective treatment (Median  $t_{max}$  is to be determined excluding the subjects experiencing emesis),*
- *A pre-dose concentration of BI 1323495 is  $>5\%$   $C_{max}$  value of that subject in the respective treatment period*
- *Missing samples/concentration data at important phases of PK disposition curve*

The descriptive analysis of PK concentrations will be based on the ADS ADPC as described at the beginning of [Section 7](#).

Table 6.3: 1 Subject sets analysed

Class of endpoint	Subject set	
	TS	PKS
Analyses of PK endpoints		X
Safety parameters	X	
Demographic/baseline parameters	X	
Important protocol deviations	X	
Disposition	X	

## 6.5 POOLING OF CENTRES

This section is not applicable, because the study was performed in only one centre.

## 6.6 HANDLING OF MISSING DATA AND OUTLIERS

Handling of missing data and outliers will be performed as described in the CTP, Section 7.5.

The only exceptions where imputation might be necessary for safety evaluation are AE dates. Missing or incomplete AE dates are imputed according to BI standards (see 001-MCG-156\_RD-01 (4)).

Missing data and outliers of PK data are handled according to BI standards (see 001-MCS-36-472\_RD-01) (5).

## 6.7 BASELINE, TIME WINDOWS AND CALCULATED VISITS

The baseline value is defined as the last measurement before administration of BI 1323495 in each period.

**Section 6.1 of the CTP:** *Exact times of measurements outside the permitted time windows will be documented. The acceptable time windows for screening, the end of trial examination, and measurements and assessments scheduled to occur 'before' trial medication administrations are provided in the CTP Flow Chart.*

*If not stated otherwise in the Flow Chart, the acceptable deviation from the scheduled time for the assessment of safety (e.g. vital signs, ECG, laboratory tests) will be  $\pm 30$  min.*



Adherence to time windows will be checked via the consistency check listings at the RPM.

Unscheduled measurements of laboratory data and vital signs data will be assumed to be repeat measurements of the most recent scheduled measurement (e.g. for follow-up or confirmation of a particular value). Therefore, unscheduled measurements will be assigned to the planned time point of the previous scheduled measurement.

## 7. PLANNED ANALYSIS

Safety analysis (refer to [Section 7.8](#)) will be performed by \_\_\_\_\_ and will be presented in Sections 15.1 to 15.4 of the CTR and in Appendix 16.2 and 16.1.13.1.

Inferential statistical analyses of PK endpoints (refer to [Section 7.4](#) and [Section 7.5.2](#)) will also be performed by \_\_\_\_\_ and will be presented in Section 15.5 of the CTR and in Appendix 16.1.13.3.

Descriptive data analysis of PK endpoints and concentrations will be performed by the department Translational Medicine and Clinical Pharmacology (TMCP) at BI and will be presented in Section 15.6 of the CTR.

The format of the listings and tables will follow the standards defined in the BI corporate guideline “Reporting of Clinical Trials and Project Summaries” [001-MCG-159] ([6](#)) with the exception of those generated for PK-calculations ([7](#)).

The individual values of all subjects will be listed, sorted by treatment sequence, subject number, visit and actual treatment (if appropriate).

The listings will be included in Appendix 16.2 of the CTR.

For end-of-text tables, the set of summary statistics for non-PK parameters is:

N	number non-missing observations
Mean	arithmetic mean
SD	standard deviation
Min	minimum
Median	median
Max	maximum

For analyte concentrations, the following descriptive statistics will additionally be calculated:

CV	arithmetic coefficient of variation
gMean	geometric mean
gCV	geometric coefficient of variation

For PK parameters, the following descriptive statistics will additionally be calculated:

CV	arithmetic coefficient of variation
gMean	geometric mean
gCV	geometric coefficient of variation
P10	10th percentile
Q1	1st quartile
Q3	3rd quartile
P90	90th percentile

The data format for descriptive statistics of concentrations will be identical to the data format of the respective concentrations. The descriptive statistics of PK parameters will be calculated using the individual values with the number of decimal places as provided by the evaluation program. Then the individual values as well as the descriptive statistics will be reported with three significant digits in the CTR.

Tabulations of frequencies for categorical data will include all possible categories and will display the number of observations in a category, as well as the percentage (%) for each treatment sequence/group. Percentages will be rounded to one decimal place and will be based on all subjects in the respective subject set whether they have non-missing values or not. The category 'missing' will be displayed only if there are actually missing values.

Units of variables should be given in the titles or column/row descriptors in square brackets (e.g. [mg]).

#### Exclusion of PK parameters

The ADS ADPP (PK parameters) contains column variables indicating inclusion/exclusion (APEXC) of a PK parameter and an analysis flag comment (APEXCO). All analyses based on the PKS will include parameters if they are not flagged for exclusion, that is APEXCO is equal to "Included".

#### Exclusion of PK concentrations

The ADS ADPC (PK concentrations per time-point or per time-interval) contains column variables ACEXC and ACEXCO indicating inclusion/exclusion (ACEXC) of a concentration and an analysis flag comment (ACEXCO). Exclusion of a concentration depends on the analysis flag comment ACEXCO. For example, if ACEXCO is set to 'ALL CALC', the value will be excluded for all types of analyses based on concentrations. If ACEXCO is set to 'DESC STATS' the value will be excluded from descriptive evaluations per planned time point/time interval. If ACEXCO contains the addition 'TIME VIOLATION' or 'TIME DEVIATION' the value can be used for further analyses based on actual times. If ACEXCO is set to 'HALF LIFE', the value will be excluded from half-life calculation (and, as a consequence, any calculation that relies on  $\lambda_z$ ) only; the value is included for all other analyses. Concentrations excluded from analysis will be listed in the CTR associated with an appropriate flag.

Further details are given in 001-MCS-36-472\_RD-01 "Noncompartmental Pharmacokinetic / Pharmacodynamic Analyses of Clinical Studies" ([5](#)) and 001-MCS-36-472\_RD-03 "Description of Analytical Transfer Files and PK/PD Data Files" ([11](#)).

## **7.1 DEMOGRAPHIC AND OTHER BASELINE CHARACTERISTICS**

Only descriptive statistics are planned for this section of the report, based on the TS. The data will be summarised in total.

## 7.2 CONCOMITANT DISEASES AND MEDICATION

Frequency tables are planned for this section of the report, based on the TS.

Concomitant diseases will be coded using the coding system of the Medical Dictionary for Drug Regulatory Activities (MedDRA). Medications will be coded using the World Health Organization Drug Dictionary (WHO-DD). The coding version number will be displayed as a footnote in the respective output.

The diagnoses and medications will be listed. Subjects without any concomitant diagnoses or concomitant therapies should be marked with a “No” in the respective column.

A medication will be considered concomitant, if it

- is ongoing at the time of the first administration of the respective treatment
- starts within the analysis phase of the respective treatment (see [Section 6.1](#) for a definition of treatments and analysis phases)

The relevance of the concomitant therapies to the evaluation of PK will be decided no later than at the RPM.

## 7.3 TREATMENT COMPLIANCE

**Section 4.3 of the CTP:** *Compliance will be assured by administration of all trial medication in the study centre under supervision of the investigating physician or a designee. The measured plasma concentrations of trial medication will provide additional confirmation of compliance.*

It is not intended to list the compliance separately. Any deviations from complete intake will be addressed in the RPM (cf. TSAP [Section 6.2](#)) and described in the CTR.

## 7.4 PRIMARY ENDPOINTS

Relative bioavailability of BI 1323495 in plasma (i.e. the effect of itraconazole on the PK of BI 1323495) is to be determined on the basis of the primary pharmacokinetic endpoints  $AUC_{0-tz}$  and  $C_{max}$  (see [Section 5.1](#)).

### 7.4.1 Primary analysis of the primary endpoints

**Section 7.3.1 of the CTP:** *The statistical model used for the analysis of the primary endpoints will be an analysis of variance (ANOVA) model on the logarithmic scale. That is, the PK endpoints will be log-transformed (natural logarithm) prior to fitting the ANOVA model. This model will include effects accounting for the following sources of variation: subject and treatment. The effect ‘subject’ will be considered as random, whereas the effect ‘treatment’ will be considered as fixed. The model is described by the following equation:*

$y_{km} = \mu + \tau_k + s_m + e_{km}$ , where

$y_{km}$  = logarithm of response measured on subject  $m$  receiving treatment  $k$ ,

$\mu$  = the overall mean,

$s_m$  = the effect associated with the  $m^{\text{th}}$  subject,  $m = 1, 2, \dots, n$

$\tau_k$  = the  $k^{\text{th}}$  treatment effect,  $k = 1, 2$ ,

$e_{km}$  = the random error associated with the  $m^{\text{th}}$  subject who received treatment  $k$ .

where  $s_m \sim N(0, \sigma_B^2)$  i.i.d.,  $e_{km} \sim N(0, \sigma_W^2)$  i.i.d. and  $s_m, e_{km}$  are independent random variables. The indices 'B' and 'W' correspond to 'between' and 'within' variability, respectively.

Point estimates for the ratios of the geometric means (test/reference) for the primary endpoints (see [Section 5.1](#)) and their two-sided 90% confidence intervals (CIs) will be provided.

For each endpoint, the difference between the expected means for  $\log(T)$ - $\log(R)$  will be estimated by the difference in the corresponding adjusted means (Least Squares Means). Additionally their two-sided 90% confidence intervals will be calculated based on the residual error from the ANOVA and quantiles from the  $t$ -distribution. These quantities will then be back-transformed to the original scale to provide the point estimate and 90% CIs for each endpoint.

The implementation for this analysis will be accomplished by using the XPKISTAT macro, based on PKS, and option BWU (Bioavailability/Bioequivalence, within-subject design, uncontrolled w.r.t. time).

## **7.5 SECONDARY ENDPOINTS**

### **7.5.1 Key secondary endpoints**

This section is not applicable as no key secondary endpoint has been specified in the protocol.

### **7.5.2 Secondary endpoints**

The secondary PK parameter  $AUC_{0-\infty}$  ( $AUC_{INF\_pred}$ ) will be assessed using the same methods as described for the primary endpoints.

## **7.7 EXTENT OF EXPOSURE**

Descriptive statistics are planned for this section of the report based on the TS. The date and time of drug administration will be listed for each subject.

## 7.8 SAFETY ANALYSIS

All safety analyses will be performed on the TS.

The safety data for treated subjects who failed to complete the study (dropouts or withdrawals) will be reported as far as their data are available. All withdrawals will be documented and the reason for withdrawal recorded.

### 7.8.1 Adverse Events

AEs will be coded with the most recent version of MedDRA.

The analyses of AEs will be descriptive in nature and will be based on BI standards as presented in the corporate guideline: “Analysis and Presentation of Adverse Event Data from Clinical Trials” [001-MCG-156] ([8](#)).

The standard AE analyses will be based on the number of subjects with AEs (and not on the number of AEs).

For analysis, multiple AE occurrence data on the case report form (CRF) will be collapsed into one AE provided that all of the following applies:

- All AE attributes are identical (lower level term (LLT), intensity, action taken, therapy required, seriousness, reason for seriousness, relationship, outcome, AE of special interest (AESI))
- The occurrences were time-overlapping or time-adjacent (time-adjacency of two occurrences is given if the second occurrence started within one hour after end of the first occurrence).

For further details on summarization of AE data, please refer to [001-MCG-156] ([8](#)).

The analysis of AEs will be based on the concept of treatment-emergent AEs. That means that all AEs will be assigned to the screening, treatment or follow-up phases as defined in [Section 6.1](#). Therefore, all AEs which occur through the treatment phase and throughout the REP will be considered as on treatment.

**Section 5.2.6.1.4 of the CTP:** *The following are considered as AESIs:*

- Hepatic injury  
*A hepatic injury is defined by the following alterations of hepatic laboratory parameters:*

- *An elevation of AST (aspartate transaminase) and/or ALT (alanine transaminase)  $\geq 3$ -fold ULN combined with an elevation of total bilirubin  $\geq 2$ -fold ULN measured in the same blood sample, or*
- *Aminotransferase (ALT, and/or AST) elevations  $\geq 10$  fold ULN*

*These lab findings constitute a hepatic injury alert and the subjects showing these lab abnormalities need to be followed up according to the 'DILI checklist' provided in the ISF. In case of clinical symptoms of hepatic injury (icterus, unexplained encephalopathy, unexplained coagulopathy, right upper quadrant abdominal pain, etc.) without lab results (ALT, AST, total bilirubin) available, the Investigator should make sure that these parameters are analysed, if necessary in an unscheduled blood test. Should the results meet the criteria of hepatic injury alert, the procedures described in the DILI checklist should be followed.*

**Section 7.3.4 of the CTP:** *Note that AEs occurring after the last per protocol contact but entered before final database lock will be reported to Pharmacovigilance only and will not be captured in the trial database.*

For more detail see the TSAP ADS plan.

According to ICH E3 (9), AEs classified as 'other significant' need to be reported and will include those non-serious and non-significant adverse events with

- (i) 'action taken = discontinuation' or 'action taken = reduced', or
- (ii) marked haematological and other lab abnormalities or lead to significant concomitant therapy as identified by the Clinical Monitor/Investigator at the Report Planning Meeting.

An overall summary of AEs (including AESIs) will be presented.

The frequency of subjects with AEs will be summarised by treatment, primary system organ class (SOC) and preferred term (PT). Separate tables will be provided for subjects with other significant AEs according to ICH E3 (9), for subjects with serious AEs, for subjects with drug-related AEs, for subjects with drug related serious adverse events and for subjects with AESIs.

The SOC and PTs will be sorted by frequency (within SOC). The MedDRA version number will be displayed as a footnote in the respective output.

In addition, frequencies of subjects with non-serious AEs that had an incidence of  $> 5\%$  for at least one treatment will be summarised by treatment, primary SOC and PT.

For disclosure of adverse events on EudraCT, additional information not included in a standard AE analysis will be performed. The following three entries will be created:

- Adverse Events per arm for disclosure on EudraCT
- Non-serious Adverse Events for disclosure on EudraCT
- Serious Adverse Events for disclosure on EudraCT



### **7.8.2 Laboratory data**

The analyses of laboratory data will be descriptive in nature and will be based on BI standards [001-MCG-157] ([10](#)).

Laboratory data will be analysed qualitatively via comparison of laboratory data to their reference ranges. Values outside the reference range as well as values defined as possibly clinically significant will be flagged in the data listings.

Clinically relevant findings in laboratory data will be reported as baseline conditions (at screening) or as AEs (during the trial) if judged clinically relevant by the investigator, and will be analysed as such.

It is the investigator's responsibility to decide whether a lab value is clinically significantly abnormal or not (at the RPM at the latest).

Descriptive statistics of laboratory data will be calculated by planned time point based on the worst value of the subject at that planned time point (or assigned to that planned time point).

### **7.8.3 Vital signs**

Descriptive statistics over time including change from baseline will be performed for vital signs (blood pressure and pulse rate). In the listing the difference from baseline will also be displayed.

For vital signs, descriptive statistics will be calculated by planned time point based on the last value of the subject at that planned time point (or assigned to that planned time point).

Clinically relevant findings in vital signs will be reported as AEs.

### **7.8.4 ECG**

ECG recordings will be checked by the investigator for pathological results. Clinically relevant abnormal findings for ECG will be listed under 'Relevant Medical History / Baseline Conditions' (when they occurred during screening) or will be reported as AEs (when they occurred during treatment), and will be analysed as such.

No separate ECG listing will be provided.

### **7.8.5 Others**

#### Physical examination

Physical examination findings will be reported as relevant medical history/baseline condition (i.e., a condition already existent before intake of study drug) or as AE and will be summarised as such.

No separate listing or analysis of physical examination findings will be prepared.

## 8. REFERENCES

1.	<i>CPMP/ICH/363/96</i> : "Statistical Principles for Clinical Trials", ICH Guideline Topic E9, Note For Guidance on Statistical Principles for Clinical Trials, current version.
2.	<i>001-MCS-40-413</i> : "Identify and Manage Important Protocol Deviations (iPD) ", current version, IDEA for CON.
3.	<i>BI-KMED-COPS-TMP-0001</i> : "iPD log", current version; IDEA for GEN.
4.	<i>001-MCG-156_RD-01</i> : "Handling of Missing and Incomplete AE Dates", current version; IDEA for CON.
5.	<i>001-MCS-36-472_RD-01</i> : "Noncompartmental Pharmacokinetic/Pharmacodynamic Analyses of Clinical Studies", current version; IDEA for CON.
6.	<i>001-MCG-159</i> : "Reporting of Clinical Trials and Project Summaries", current version; IDEA for CON.
7.	<i>001-MCS-36-472</i> : "Standards and processes for analyses performed within Clinical Pharmacokinetics/Pharmacodynamics", current version; IDEA for CON.
8.	<i>001-MCG-156</i> : "Analysis and Presentation of Adverse Event Data from Clinical Trials", current version; IDEA for CON.
9.	<i>CPMP/ICH/137/95</i> : "Structure and Content of Clinical Study Reports", ICH Guideline Topic E3; Note For Guidance on Structure and Content of Clinical Study Reports, current version.
10.	<i>001-MCG-157</i> : "Handling, Display and Analysis of Laboratory Data", current version; IDEA for CON.
11.	<i>001-MCS-36-472_RD-03</i> : "Description of Analytical Transfer Files and PK/PD Data Files", current version; IDEA for CON.



## 10. HISTORY TABLE

Table 10: 1 History table

Version	Date (DD-MMM-YY)	Author	Sections changed	Brief description of change
Final	21-OCT-2019		None	This is the final TSAP without any modification