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- 2 EMA/CHMP/600958/2010
- 3 Committee of Medicines for Human Use (CHMP)
- 4 Appendix IV of the Guideline on the Investigation on
- 5 Bioequivalence (CPMP/EWP/QWP/1401/98 Rev.1):
- 6 Presentation of Biopharmaceutical and Bioanalytical Data
- ₇ in Module 2.7.1
- 8 Draft

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Comments should be provided using this <u>template</u>. The completed comments form should be sent to PKWPsecretariat@ema.europa.eu

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Keywords	Generic applications, bioequivalence data, BCS biowaiver documentation,
	Standardised presentation, CHMP, EMA, Guideline

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13	Ap	pendix IV of the Guideline on the Investigation on	
14	Bic	pequivalence (CPMP/EWP/QWP/1401/98 Rev.1):	
15	Pre	esentation of Biopharmaceutical and Bioanalytical Date	ta
16	in l	Module 2.7.1	
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1. Introduction

- 22 The objective of CTD Module 2.7.1 is to summarize all relevant information in the MAA dossier with
- 23 regard to biopharmaceutic studies and associated analytical methods
- 24 This Appendix contains a set of template forms to assist applicants in the preparation of Module 2.7.1
- 25 providing guidance with regard to data to be presented. Furthermore, it is anticipated that a
- 26 standardized presentation will facilitate the evaluation process. Applicants are therefore encouraged to
- use these template tables when preparing Module 2.7.1.
- 28 This Appendix is intended for generic applications according to Directive 2001/83/EC, Article 10(1), but
- 29 might be used in other applications including variations, fixed combinations, extensions and hybrid
- 30 applications.

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2. Instructions for completion and submission of the tables

- The tables should be completed only for the pivotal studies, as identified in the application dossier in
- 33 accordance with section 4.1 of the Bioequivalence guideline (CPMP/EWP/QWP/1401/98 Rev. 1). If
- 34 there is more than one pivotal bioequivalence study, then individual tables should be prepared for each
- 35 study. In addition, the following instructions for the tables should be observed:
 - Details of non-EU reference products are not needed.
 - Tables in Section 3 should be completed separately for each analyte per study. If there is more than one test product then the table structure could be adjusted.
 - Tables in Section 4 should only be completed for the method used in confirmatory (pivotal) bioequivalence studies. If more than one analyte was measured then Table 4.1 and potentially Table 4.3 should be completed for each analyte.
- 42 In general, applicants are encouraged using cross-references and footnotes for adding additional
- 43 information. Fields that do not apply should be completed as "Not applicable" together with an
- 44 explanatory footnote.
- 45 In addition, each section of the template should be cross-referenced to the location of supporting
- documentation or raw data in the application dossier.
- 47 It is recommended that the content of the tables is searchable and that the tables are not scanned.
- 48 Applicants are encouraged to provide Module 2.7.1 also in Word (.doc) or RTF format.

3. A note about BCS-based biowaiver documentation

- 50 Relevant data for justification of BCS-based biowaiver requests should be included in Module 5.3.1.2
- 51 "Comparative BA and Bioequivalence (BE) Study Reports". A summary of the data should be provided
- 52 in Module 2.7.1 with a justification for this biowaiver and a list of relevant references.

1. BIOWAIVER REQUEST

Table 1.1 Qualitative and quantitative composition of the Test product

Ingredient	Function	Strength (label claim)					
				XX mg (Production Batch Size)		XX mg (Production Batch Size)	
CORE		Quantity per unit	Quantity per %* Qu		%*	Quantity per unit	%*
TOTAL			100%		100%		100%
COATING							
TOTAL			100%		100%		100%

^{*}each ingredient expressed as a percentage of the total core or coating weight

<u>Instructions</u>

Include the composition of all strengths. Add additional columns if it is necessary.

Table 1.2 In vitro dissolution data for biowaiver request

Dissolution testing Site		Study Report Location (vol, page,link)
Dissolution Conditions	Apparatus	
	RPM	
	Volume	

	Collection Times (minutes or hours)				f2			
Dissolution Medium			5	10	15	20		
Strength 1	pH=							
# of units	pH=							
# Batch no	pH=							
Strength 2	pH =							
# of units	pH=							
# Batch no	pH=							
Strength 3	pH=							
# of units	pH=							
# Batch no	pH=							

<u>Instructions</u>

Fill this table only if biowaiver is requested for additional strengths besides the strength tested in the bioequivalence study. Only the mean dissolution values should be reported. f2 values should be computed relative to the strength tested in the bioequivalence study. Expand the table with additional rows if more than 3 strengths are requested. Fill a similar table for the reference product if sink condition could not be achieved.

2. BIOEQUIVALENCE TRIAL INFORMATION

Table 2.1 Test and reference product information

Product Characteristics	Test product	Reference Product
Name		
Strength		
Dosage form		
Manufacturer ¹		
Batch #		
Batch size (Biobatch)		
Assay %		
Commercial Batch Size		
Expiry date (Retest date)		
Location of product certificate	Vol/page, link	Vol/page, link
Member State where the reference product is purchased from:		

Table 2.2 Study site

Study ID (vol/page,link)	Study title			Has this site been inspected by an EU Authority?
Clinical Study Site	Name	Address	Study period:	Authority: Year:
Bioanalytical Facility	Name	Address	Study period:	Authority: Year:
Data analysis	Name	Address		Authority: Year:
Sponsor of the study	Name	Address		

Table 2.3 Study design

Study ID:	
Report Location (vol/page, link):	
Design:	
Dose:	
Single/Multiple dose:	
Number of periods:	
Two-stage design:	(yes/no)
Fasting, Fed:	
# Volunteers (dosed):	# Volunteers (evaluated):

3. RESULTS

Table 3.1 Pharmacokinetic data for <analyte>

Pharmacokinetic	**Arithmetic Means (±SD)			
parameter	Test Reference product			
AUC _(0-t) 1				
AUC _(0-∞) ²				
Cmax				
Tmax				
Others				

 $^{^1}$ AUC_(0-72h) can be reported instead of AUC_(0-t), in studies with a sampling period of 72 h, and where the concentration at 72 h is quantifiable 2 AUC_(0- ∞) does not need to be reported when AUC_(0-72h) is reported instead of AUC_(0-t).

Table 3.2 Additional pharmacokinetic data for <analyte>

Records ¹ where $AUC_{(0-t)}/AUC_{(0-\infty)} < 0.8^2$	List patient IDs and period numbers
Records where Cmax is the first point	List patient IDs and period numbers
Records with significant carry-over	List patient IDs, period numbers, treatment,
(pre-dose sample > 5% Cmax)	pre-dose concentration

¹ A subject record is a list of concentrations after one administration

Table 3.3 Bioequivalence evaluation of <analyte>

Pharmacokinetic parameter	Ratio Test/Ref	Confidence Intervals	CV%¹
AUC _(0-t)			
Cmax			
Others			

¹Estimated from the Residual Mean Squares. For replicate design studies report the within-subject CV% using only the reference product data.

 $^{^{2}}$ Only if the last sampling point of AUC_(0-t) is less than 72h

4. BIOANALYTICS

Table 4.1 Bioanalytical method validation

Analytical Validation Report	<study number<="" report="" th=""></study>
	Locations (vol/page, link)>
This analytical method was used in the following studies:	<study ids=""></study>
Short description of the method	<e.g. binding="" gc="" hplc="" ligand="" ms="" ms,=""></e.g.>
Analyte	<name, (vol="" certificate="" link)="" location="" of="" page,="" product=""></name,>
Internal standard (IS)	<name, (vol="" certificate="" link)="" location="" of="" page,="" product=""></name,>
Calibration range	
Linearity	<r></r>
Lower limit of quantification (LOQ)	<accuracy, precision=""></accuracy,>
Standard curve concentrations (units/mL)	
QC concentrations (units/mL, vol/page link)	
Between-run accuracy	<range by="" or="" qc=""></range>
Between-run precision	<range by="" or="" qc=""></range>
Within-run accuracy	<range by="" or="" qc=""></range>
Within-run precision	<range by="" or="" qc=""></range>
Matrix Factor % (Analyte and IS)	Trange or by Qe>
CV% of IS normalized matrix factor	
Short term stability of the stock solution and working	<confirmed (time).<="" td="" to="" up=""></confirmed>
solutions	Observed change %
	at Temperature °C>
Long term stability of the stock solution and working	<confirmed (time).<="" td="" to="" up=""></confirmed>
solutions	Observed change %
	at Temperature °C>
Short-term stability in plasma at room	<confirmed (time).="" observed<="" td="" to="" up=""></confirmed>
temperature (QC)	change %>
Post-preparative stability (dry extract stability)	<confirmed (time).="" observed<="" td="" to="" up=""></confirmed>
, , , , , , , , , , , , , , , , , , , ,	change %>
Long term stability in plasma (vol/page,link)	<confirmed (time).<="" td="" to="" up=""></confirmed>
	Observed change %
	at Temperature °C>
Autosampler storage stability	<confirmed (time).="" observed<="" td="" to="" up=""></confirmed>
· · · · · · · · · · · · · · · · · · ·	change %>
Freeze and thaw stability (-Temperature °C)	<pre><# cycles, Observed change %></pre>
Dilution integrity	<concentration diluted="" td="" x-fold,<=""></concentration>
	Accuracy % Precision %>
Partial validations 1	<describe of<="" reason="" shortly="" td="" the=""></describe>
Location (vol/page, link)	revalidation>
Partial validations 2	<describe of<="" reason="" shortly="" td="" the=""></describe>
Location (vol/page, link)	revalidation>
Cross validation	<describe of<="" reason="" shortly="" td="" the=""></describe>
	crossvalidation>

¹Might not be applicable for the given analytical method

Table 4.2 Storage period of study samples

Study ID	Longest storage period

Table 4.3 Study sample analysis

Study ID		
Total numbers of collected samples	# collected samples	
Total number of samples with valid results	# samples	
Total number of reassayed samples ^{1,2}	# reassayed samples	
Total number of analytical runs ¹	# runs	
Total number of valid analytical runs ¹	# valid runs	
Incurred sample reanalysis		
Number of samples	# samples	
Percentage of samples where the difference between the two values was less than 20% of the mean"	% samples	

<u>Instructions</u>
Fill Table 4.3 for each pivotal trial.

¹ Without incurred samples ² Due to other reasons than not valid run