

IN VITRO CHARACTERIZATION OF THE GENERIC PEPTIDE THERAPEUTIC: THE GLP-1R AGONIST CASE STUDY

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INTRODUCTION

Newer generation of anti-diabetics such as glucagon-like peptide-1 receptor agonists (GLP-1RA) have shown to be an efficient adjunctive therapy in patients with diabetes type 2 who failed to respond to oral antihyperglycemic agents [1, 2]. By the 2045 it has been estimated that 700 million people will have diabetes, chronic disease which has risen to major global health issue [3]. Generic version of GLP-1RA would decrease drug price vastly and would make newer diabetes medications widely accessible.

In order to meet regulatory requirements [4] a comprehensive analytical characterization needs to be performed and an equivalency of the generic Exenatide injection (Gx) to the innovator drug must be demonstrated. Exenatide is a 39-amino acid peptide which stimulates GLP-1R to cause insulin release [1]. In this work, the extensive comparability study, which was aimed to ensure quality, safety and efficacy of the generic peptide product similar to the Reference Listed Drug (RLD) Byetta®, is described. *In vitro* characterization of the generic drug included various analyses of physicochemical properties such as identity, structure, self-association and aggregation studies, purity and peptide content determination, whereby different orthogonal analytical techniques were employed. Most of the analytical methods were developed and optimized in-house. The comparison study was performed on different batches of the Exenatide injection and RLD samples.

WORLD
2045 700 million
2030 578 million
2019 463 million
51 % increase



Physicochemical properties: appearance, pH, osmolality
Particulate matter: visible and subvisible particles
Molecular mass: HR-MS (intact mass)
Amino acid composition: LC-UV (AAA)
Charged variants: cIEF
Oligomers and aggregates: SDS-PAGE, Native-PAGE, SEC-UV, SEC-MALS, AUC, MFI, AF4
Impurity profile: SCX-HPLC, RP-HPLC, HPLC-HRMS
Optical purity: GC-MS
Primary sequence: peptide mapping (UPLC-ESI-MS/MS) and peptide sequencing (HRMS/MS)
Secondary structure: CD (Far-UV), IF, NMR, Raman and FTIR
Tertiary structure: CD (Near-UV)
Biological activity: Cell based potency

1 Exenatide primary structure

HGEGFTTSLSKQMEEEAVRLFIEWLKNKGGPSSGAPPPS-[NH₂]

- Impact on safety and efficacy
- Sameness of Gx and RLD product by peptide mapping (UPLC-HRMS), peptide sequencing (HRMS/MS) and intact mass (MS)

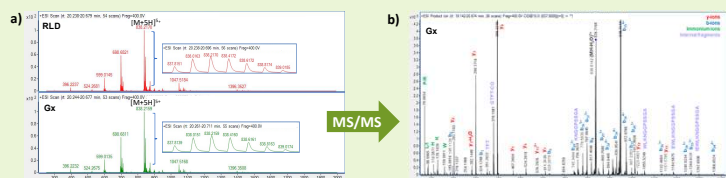


Figure 1. Intact mass and peptide sequencing by ESI-MS/MS: a) Full scan MS spectra of the RLD Byetta® and the Exenatide injection (Gx) with assigned [M+5H]⁵⁺ ions and zoomed-in isotopic distribution; b) Assigned MS/MS spectra of the [M+5H]⁵⁺ (m/z 838) of Gx sample

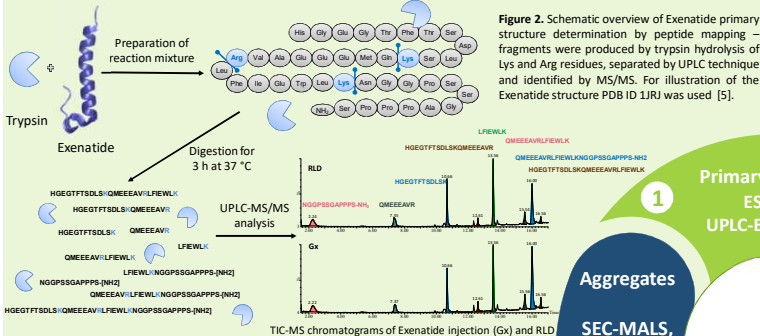


Figure 2. Schematic overview of Exenatide primary structure determination by peptide mapping – fragments were produced by trypsin hydrolysis of lys and Arg residues, separated by UPLC technique and identified by MS/MS. For illustration of the Exenatide structure PDB ID 1JRI was used [5].

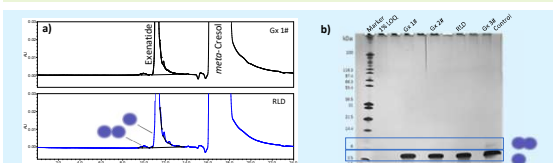


Figure 6. Irreversible aggregates determination by orthogonal techniques: a) SEC-UV chromatogram of the Exenatide injection (Gx) 1# and RLD Byetta® samples; b) SDS-PAGE comparison of different Gx batches and RLD

Aggregates

- Impact on safety and efficacy
- Enhance immune response
- Irreversible (SEC-UV, SDS-PAGE) and reversible (AF4, SEC-MALS)
- Different methods required for a wide range of aggregates
- Sensitive, orthogonal methods to assure accuracy and robustness

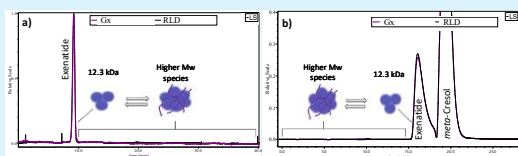


Figure 7. Self-association and reversible aggregates comparison of the Exenatide injection (Gx) and the RLD Byetta® by orthogonal techniques a) AF4 and b) SEC-MALS

Table 1. Exenatide process related (API) and degradation impurities analyzed by orthogonal techniques – RP-HPLC and SCX-HPLC. Impurities are marked in chromatograms in the Figure 3.

Impurity	Modification
D, E, G, H, N, Q, S	process (API)
A, C	deamidation
I	deamidation
B	oxidation
F, G, K, M	truncation
G	acetylation
J, L	dehydration
R, T	isomerization
P	spinacine formation

Impurities O, P, R, S and T co-eluted with main Exenatide peak in RP-HPLC (Figure 3 a) and were resolved from the main peak in SCX-HPLC (Figure 3 b) chromatograms.

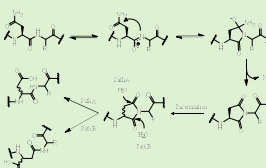


Figure 4. Mechanism of asparagine deamidation – example of common degradation impurity formation

Peptide - related impurities

- Impact on safety and efficacy
- Risk of immunogenicity
- Sensitive, high-resolution (HPLC-HRMS) and orthogonal analytical chromatographic techniques e.g. RP, IEX, HILIC, HIC...

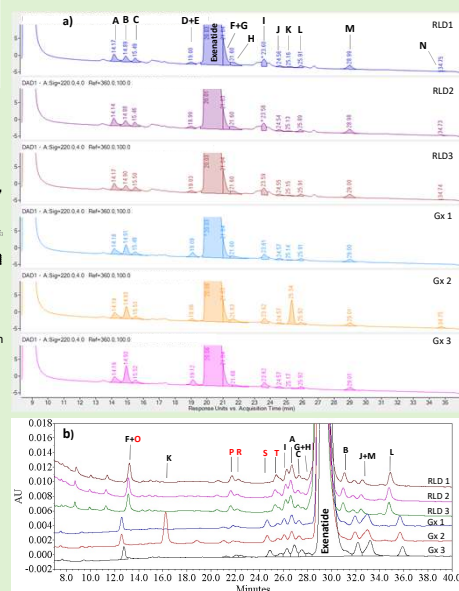
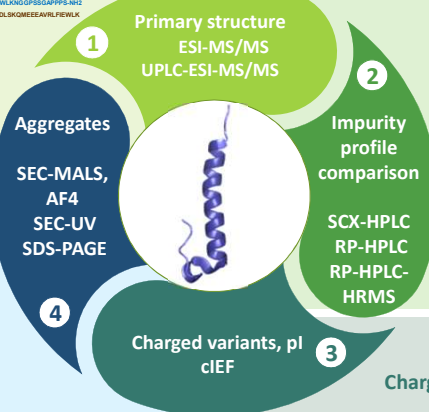


Figure 3. Exenatide impurity profile comparison of the Exenatide injection (Gx) and the RLD Byetta® samples by the means of a) RP-HPLC and b) SCX-HPLC methods; different selectivity achieved by orthogonal chromatographic techniques – red marked impurities co-eluted with the main Exenatide peak in RP-HPLC (Figure a) and were separated in SCX-HPLC (Figure b) chromatograms



- Impact on efficacy
- Isoelectric point (pI) determination and charge heterogeneity comparison
- Bioinformatical and literature data screen and comparison with experimental data
- High resolution separation of peptide charge variants by cIEF

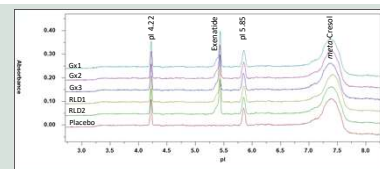


Figure 5. Charge variants determination: native cIEF electropherograms of the Exenatide injection (Gx) and the RLD Byetta® samples

Site Name	Owned by	pI Value
Protein Isoelectric Point Calculator	Lukasz P. Kozłowski	4.55
Protein Calculator	The Scripps Research Institute	4.70
PepCalc	Innovagen AB	4.77
Karger Kidney and Blood Pressure Research	Yi Zhang et al.	4.86
Peptide Calculator	Bachem	4.90
Biomaterials	Jae Y. Kim et al.	4.96

Table 2. List of pI databases and literature sources with theoretical pI values

CONCLUSION

The performed characterization of different batches of the generic Exenatide injection and the RLD Byetta® proved the same primary structure, aggregation propensity and charge heterogeneity. Moreover, similar impurity profiles of Exenatide injection and RLD were confirmed. For impurities which were not detected in RLD batches or levels were higher in generic Exenatide drug, the immunogenicity risk assessment was conducted. It was shown that generic Exenatide injection does not contain impurities which induce greater or distinct stimulation of innate or functional immunoresponse than the RLD Byetta®. Furthermore, comparability of Exenatide's attributes such as physicochemical properties, identity, secondary and tertiary structure, and biological activity were evaluated. It is demonstrated that the generic Exenatide injection drug shows highly similar quality attributes related to safety and efficacy, including immunogenicity, and meets the same standards as the innovator drug.

LITERATURE

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