1 2	Review Article
2	High resolution separation methods for the determination of intact human erythropoiesis
4	stimulating agents. A review
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22	Keywords:
23	Erythropoietin (EPO)
24	capillary electrophoresis

25	high p	performa	nce liqui	id chroma	atography
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- 26 mass spectrometry
- 27 doping control
- 28 EPO biosimilars
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31	Abbreviations:
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- 32 AGP, Alfa 1-acid glycoprotein
- 33 BGE, Background electrolyte.
- 34 BRP, Biological reference preparation.
- 35 BSA, Bovine serum albumin.
- 36 CE, Capillary electrophoresis
- 37 CERA, Continuous erythropoiesis receptor activator.
- 38 CHO, Chinese hamster ovary
- 39 CIEF, Capillary isoelectric focusing.
- 40 CMC, Carboxymethyl chitosan.
- 41 CZE, Capillary zone electrophoresis.
- 42 DEAE-agarose, Diethylaminoethyl-agarose.
- 43 DPO, Darbepoetin alfa.
- 44 DTT, Dithiothreitol.
- 45 eEPO, Equine erythropoietin.
- 46 ELISA, Enzyme linked immunosorbent assay.
- 47 EOF, Electroosmotic flow.

- 48 EP, European pharmacopoeia.
- 49 EPO, Erythropoietin.
- 50 ESA, Erythropoiesis stimulating agent.
- 51 ESI, Electrospray ionization.
- 52 ESP, Erythropoiesis stimulating protein.
- 53 FOBs, Follow-on biologics.
- 54 FTICR, Fourier transform ion cyclotron resonance.
- 55 HMW, High molecular weight.
- 56 hEPO, Human erythropoietin.
- 57 HILIC, Hydrophilic interaction chromatography
- 58 HPLC, High performance liquid chromatography.
- 59 HRP, Horseradish peroxidase
- 60 HSA, Human serum albumin.
- 61 IAC, Immunoaffinity chromatography.
- 62 IEF, Isoelectric focusing.
- 63 IEF-PAGE, Isoelectric focusing Polyacrylamide gel electrophoresis
- 64 IEX, Ion exchange.
- 65 IT, Ion trap.
- 66 LacNAc, *N*-Acetyllactosamine.
- 67 LC, Liquid chromatography.
- 68 LIF, Laser induced fluorescence
- 69 MS, Mass spectrometry.
- 70 NESP, Novel erythropoiesis stimulating protein.

- 71 NeuNAc, *N*-Acetylneuraminic acid.
- 72 NeuNGc, *N*-Glycolylneuraminic acid.
- 73 PBS, Phosphate buffer saline
- 74 PEG, Polyethyleneglycol.
- 75 PTM, Post-translational modification.
- 76 PVP, Polyvinylpyrrolidone.
- 77 QC, Quality control.
- 78 rhEPO, recombinant human erythropoietin.
- 79 RMT, Relative migration time
- 80 RP, Reversed phase.
- 81 SDS, Sodium dodecyl sulphate.
- 82 SDS-PAGE, Sodium dodecyl sulphate Polyacrylamide gel electrophoresis.
- 83 SE, Size exclusion.
- 84 SEBs, Subsequent entry biologics.
- 85 SPS, Sodium Pentasulfonate.
- 86 TOF, Time of flight.
- 87 uhEPO, Urinary human erythropoietin.
- 88 WHO, World health organization
- 89 WADA, World antidoping agency.

- 91 Abstract
- 92

93 Human erythropoietin (hEPO), a hormone involved in the formation of red blood cells, is 94 a 30 kDa glycoprotein with a high carbohydrate content. The production of recombinant hEPO 95 has made possible its widespread therapeutic use and its banned use in competition sports. 96 Methods to analyze EPO and other erythropoiesis stimulating agents (ESAs) are necessary for 97 the characterization and quality control of these biopharmaceuticals and also for doping control. 98 In this paper, high resolution separation methods, namely high performance liquid 99 chromatography (HPLC) and capillary electrophoresis (CE), with special attention to CE-100 coupled mass spectrometry, are reviewed. The usefulness of these techniques when applied in 101 different modes to separate the glycoprotein isoforms, aggregates or excipients are detailed. In 102 addition, sample preparation methods that have been applied to ESA samples for subsequent 103 determination by HPLC or CE, as well as the potential compatibility of other preparation 104 methods, are discussed. Applications of the HPLC and CE methods regarding regulatory 105 considerations for biopharmaceuticals analysis, with emphasis on biosimilars, and doping control 106 are also included. Finally, limitations of the present methods and their possible solutions are 107 considered.

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120 **1. Introduction**

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122 Human erythropoietin (hEPO) is a glycoprotein hormone produced mainly in the kidneys 123 that regulates the erythropoiesis process, that is, the formation of red blood cells [1]. The large-124 scale production of a recombinant version of human erythropoietin (rhEPO) by biotechnological 125 methods has made possible its widespread therapeutic use as well as its misuse in sport 126 competitions. It has a molecular mass of about 30 kDa, 40% of which corresponds to three N-127 linked and one O-linked carbohydrate chains attached to the polypeptide chain. A representation 128 can be observed in Figure 1 [2]. Glycosylation depends on several factors such as the organism 129 or cell line in which EPO is produced or the culture conditions. This has given rise to the 130 production of several EPO products featuring variations in their glycosylation pattern. In addition, several modifications have been carried out in recombinant products, giving rise to 131 132 different erythropoiesis stimulating agents (ESAs). 133 The nomenclature and main characteristics of EPO and EPO-related substances have 134 been summarized by Jelkmann [3] and by Reichel and Gmeiner [4]. For EPO-type substances, 135 the stem "-poetin" is used according to the International Nonproprietary Names system of the 136 World Health Organization (WHO). The term "epoetin" corresponds to eukaryotic cell-derived 137 rhEPO, with a peptide sequence identical to that of urinary human EPO (uhEPO). A prefix is

used to indicate changes in the peptidic sequence, as in the case of darbepoetin alfa (DPO, also known as novel erythropoiesis stimulating protein or NESP). Greek letters after the name are employed to indicate analogues of a given EPO type substance with an altered glycosylation, as in the case of epoetin- α and epoetin- ω . For other EPO related substances, pegylated epoetin- β is known as CERA (continous erythropoiesis receptor activator), and the name Hematide accounts for a synthetic dimeric EPO mimetic peptide linked to polyethyleneglycol (PEG). Schemes of
EPO, DPO, and CERA are shown in Figure 2 [3].

145 Analysis methods are aimed at the control of the drug for its therapeutic use as well as in 146 situations of doping control. In particular, the increase in biosimilars and uncontrolled drugs 147 claimed to be rhEPO is enlarging the need for these analysis methods. In addition, ESAs are 148 now produced in many countries around the world as exemplified in Table 1, which shows a 149 non-comprehensive list of ESAs available worldwide; further information about ESAs in the 150 world can be found in [5-8]. Depending on the source of information, mainly the web pages of 151 the manufacturers, contradictory data are sometimes found. Efforts are also being made for the 152 in-depth characterization of recombinant EPO [9-12].

When performing EPO determination, it is important to consider that variations in glycosylation result in alternate versions of the same protein called glycoforms. These glycoforms can be separated using different techniques and appear as different peaks in capillary electrophoresis or liquid chromatography, and as different bands or spots in gel electrophoresis, these peaks or bands being known as isoforms.

158 Different strategies can be followed to perform the determination of glycoproteins. Some 159 of them include the controlled hydrolysis of the glycoprotein, resulting in the study of glycans 160 and/or glycopeptides. In other instances, hydrolysis is performed to solely release the glycans, 161 leaving the peptide sequence intact, thus enabling the determination of glycans and/or of the 162 peptide sequence. Yet in other cases, hydrolysis is not carried out and the whole intact 163 glycoprotein is analyzed. This review will focus on the determination of intact EPO and other 164 ESAs, namely, erythropoiesis stimulating proteins (ESPs). The determination of glycoforms of 165 intact glycoproteins presents several advantages in comparison to hydrolyzed glycoproteins, such as saving time, less sample manipulation (less influence of the operator and the sample treatmenton the results) and obtaining a more holistic information about the glycoprotein.

168 A number of analytical methods for EPO and related substances have been recently 169 summarized [4]. In that book chapter chromatography, mass spectrometry, immunoassay, and 170 electrophoresis are considered. Special attention is paid to this last technique by including 171 different methods performed using zone electrophoresis in agarose suspension, isoelectric 172 focusing both in solution and in polyacrylamide gels, sodium dodecyl sulphate-polyacrylamide 173 gel electrophoresis (SDS-PAGE) and bidimensional electrophoresis. In the present review, high 174 resolution separation techniques, namely high performance liquid chromatography (HPLC) and 175 capillary electrophoresis (CE), will be considered. Each of the analytical steps, from sample 176 preparation to detection, with special attention paid to mass spectrometry (MS) coupled to CE, 177 will be discussed. Finally, applications of these techniques for the determination of biosimilars 178 and for doping control will be covered. The review will address the methods published and also 179 potential approaches to solve current limitations. In this regard, methods employed to analyze 180 other glycoproteins or other aims will be discussed focusing on their potential applicability for 181 HPLC or CE determination of ESAs.

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184 **2. Sample preparation**

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186 The process by which a sample is made ready for analysis by HPLC or by CE depends on 187 the type of sample, the aim of the analysis, the HPLC or CE separation mode as well as the detection technique to be employed. In this context, two main groups of samples can beconsidered: pharmaceutical preparations and biological fluids.

190 2.1 Sample preparation methods carried out for HPLC or CE determination of ESAs

191 For pharmaceutical preparations, the ESA concentration is usually high enough to allow 192 its detection by most of the usual methods (e.g., UV and MS). The sample preparation step is 193 aimed at eliminating substances that may interfere during analysis. The need to remove these 194 substances, as can be the case for formulation excipients, in CE analysis depends on the capillary 195 coating and the background electrolyte (BGE) employed. For example, EPO products formulated 196 with human serum albumin (HSA) as excipient can be directly analyzed using a BGE containing 197 nickel chloride and amine-coated capillaries [13]. Alternatively, analysis using widely available 198 and inexpensive bare fused silica capillaries requires prior removal of HSA, which can be 199 efficiently carried out with commercial anti-HSA cartridges. In this case special attention must 200 be paid to avoid the presence of phosphate in the purified rhEPO sample as this prevents its 201 determination by capillary zone electrophoresis (CZE) [14]. Low-molecular weight sample 202 components can be readily removed using centrifugal ultrafiltration devices [15-17]. ESA 203 samples containing several excipients, including polysorbate 80, were prepared using these 204 devices and analyzed by CZE without problem [18]. However, some authors [19] reported, 205 without presenting results that these devices are unable to remove polysorbate 80 and that they 206 preferred the use of a commercial non-ionic detergent trap cartridge prior to the centrifugal 207 filtration to prepare ESA samples containing this excipient. In order to increase EPO recovery 208 passivation of the filtration devices with 5% BRIJ 35 was recommended [14,20]. Similar 209 recoveries were achieved for devices with 10 kDa or 30 kDa cut-off membranes [14]. When the 210 analysis was carried out by capillary isoelectric focusing (CIEF), better results were obtained if

excipients were not completely removed or, for a better control of the salt content, known
amounts of salts were added after excipients were eliminated [15,21]. For characterization by
CE-MS, a high rhEPO concentration (about 3 mg mL⁻¹) is desirable. To increase the EPO
concentration, ultrafiltration through a 10 kDa cut-off membrane device was used, followed by
freeze-drying and subsequent dissolution in a small volume of water [22].

Sample preparation for the determination of EPO and other ESAs in culture supernatants and biological fluids is far more complex than in pharmaceutical preparations due to the low analyte concentration and the complexity of the sample matrix. In these samples the preparation step is aimed not only to purify the ESA but also to concentrate it. The determination of ESAs in these complex samples can be of interest from a therapeutic point of view (e.g., culture supernatant control during production, pharmacokinetics studies, etc) as well as for doping control considerations.

223 Purification of rhEPO from culture supernatants for subsequent CZE analysis has been 224 carried out through a series of chromatographic steps including ion-exchange (IEX), reversed-225 phase (RP), and gel filtration or size-exclusion (SE) modes [23]. The protocol followed for the 226 EPO purification is referred to other papers [24-25] and it appears to be based on the seven-step 227 procedure developed by Miyake et al. [26] for urinary EPO. It includes ethanol precipitation, 228 diethylaminoethyl-agarose (DEAE-agarose) fractionation, sulfopropyl-Sephadex 229 chromatography, gel filtration, and hydroxylapatite chromatography. Samples purified from cell 230 cultures in this way were of high purity, thus making CE analysis possible [23]. Although the 231 length of the purification procedure is not indicated, the large number of steps involved implies 232 that it is a laborious and time-consuming method that would not be adequate for doping or drug 233 production control.

The on-line coupling of immunoaffinity with CZE separation has been attempted using rhEPO [27]. Polyclonal anti-EPO antibodies were attached to glutaraldehyde-treated glass beads and packed between two frits in a small Polybrene-coated capillary column, which was then linked to a separation capillary of the same diameter and type. The authors indicated that the capture and release of EPO were observed but the resolution of isoforms was completely lost. 2.2 Suitability of other methods for sample preparation prior to HPLC or CE determination of ESAs

As seen above, only a handful of sample preparation methods have been specifically described for the isolation of intact ESAs for analysis by CE or HPLC. In the following paragraphs, other methods described to purify EPO for determination by other techniques will be considered and their suitability for subsequent determination by CE and HPLC analysis discussed.

246 Immunoaffinity using anti-EPO antibodies has been performed on different platforms 247 such as magnetic beads, ELISA plates, monolith disks and, mainly, chromatographic columns. 248 Monoclonal antibodies covalently linked to magnetic beads were used by Wognum et al. to 249 purify EPO from serum and plasma [28]. The elution buffer included PEG 8000. Polyclonal anti-250 EPO antibodies also attached to magnetic beads were used to purify different ESAs from equine 251 plasma [29-30]. In this case, desorption was carried out using PEG 6000 in phosphate buffer 252 saline (PBS). In both cases, PEG should be eliminated if CE or SE-HPLC is used to analyze 253 intact EPO isoforms. A modification of the method by Wognum et al. that includes elimination 254 of PEG, has proved useful to isolate rhEPO and ESA from equine plasma [31] and DPO from 255 human plasma [32]. An alternative method for EPO elution from magnetic beads derivatized 256 with polyclonal antibody, consisting of 1% sodium dodecyl sulphate (SDS) in PBS under

vigorous shaking and subsequent removal of SDS by dialysis, was effective to isolate EPO fromhuman serum [33].

259 The optimization of EPO purification using monoclonal anti-EPO 9C21D11 on 260 immunoaffinity well plates (similar to the ones used in Enzyme Linked Immunosorbent Assays -261 ELISA) led to around 56% recovery of hEPO in plasma without isoform discrimination [34]. 262 Although the total protein content had been reduced 3000 times, the remaining concentration of 263 interfering proteins from plasma was still too high and would not be compatible with the direct 264 CZE determination on uncoated capillaries or with the SE-HPLC determination of EPO. Figure 3 265 shows the total concentration of proteins (black line, right scale) and that of hEPO (bars, left 266 scale).

267 Gupta et al. [35] prepared three immunocolumns using three different commercial antibodies from different sources to purify EPO and analogues in the development of a method 268 269 aimed at detecting horse doping. No indication about the nature of the packing material or the 270 antibodies characteristics was provided. By using an antibody/antigen molar ratio of about 1 271 million (50 mg antibody for each column), incubating the sample in the column for 4 h, and 272 employing 100 mM glycine pH 2.8 as eluent, equine EPO (eEPO), rhEPO, and DPO were 273 purified with about 87% recovery. Analyses of the intact ESAs in the purified samples were 274 performed by ELISA and gel electrophoresis with either Coomasie or immunological staining. 275 The desorption agent could be easily eliminated with centrifugal filtration devices; however, the 276 amount of EPO purified in this way (about 20 ng) would not be sufficient for determination by 277 CE or SE-HPLC with the currently available detection methods. When this immunoaffinity 278 chromatography (IAC) method was slightly modified by using 0.2 M glycine buffer pH 2.5 as 279 desorption agent and used to isolate rhEPO and darbepoietin from horse plasma, it was observed

280 that albumin co-eluted with the ESPs [36]. Among the different strategies assayed to eliminate 281 co-elution, the combination of lectin (concanavalin A) affinity with IAC provided the best results 282 with 70% recovery of rhEPO or darbepoietin; however, the samples were still contaminated with 283 albumin and other proteins. This procedure would most probably be useful in the determination 284 of epoetins by RP-HPLC or IEX-HPLC but not by CZE on uncoated capillaries, or by SE-HPLC. 285 Another IAC procedure, potentially compatible with further CE or SE-HPLC 286 determination of the purified EPO, was developed by Mi et al. using anti-EPO polyclonal 287 antibodies bound to Sepharose [37]. Recoveries in the range 78-86% were achieved using Gly-288 HCl pH 2.5 as elution buffer. This buffer could be easily removed after the immunopurification 289 step. The binding capacity (2 µg) should be slightly increased to obtain enough purified EPO to 290 be detected by the usual procedures. 291 IAC with columns made of the 9C21D11 anti-EPO antibody bound to a chromatographic 292 support with hydrazide groups was demonstrated to be useful to isolate the glycoprotein from 293 serum and urine prior to Western blot isoelectric focusing analysis [38]. Although the use of 294 bovine serum albumin (BSA) as a component of the elution buffer would make the method

inappropriate for the subsequent determination of EPO by either CZE on uncoated capillaries or
by SE-HPLC, it could probably be compatible with CZE analysis using the Bietlot and Girard

297 method [13] and with HPLC in the RP and IEX modes.

IAC was also demonstrated to be adequate for eliminating interferences caused by unfractionated heparin on IEF-PAGE of EPO [39]. This anticoagulant, used by some athletes during recombinant EPO therapies, forms a heparin gradient on the IEF gel that modifies or even destroys the IEF EPO pattern. Saturation of samples with urea was also shown to eliminate the heparin interferences on IEF gels [39]. Although the effect of heparin on EPO determination by CE has not been studied, it could also affect focusing of EPO in CIEF and potentially modify its charge, which would affect its electrophoretic mobility in CZE. Among the two methods to remove heparin, IAC seems to be more appropriate for CE analysis because it eliminates heparin at the same time as the rest of the interfering compounds from the sample matrix. It should also be noted that a BGE frequently used for CZE of EPO contains 7M urea [15,18,40] and that such a concentration of urea is usually included in the sample mixture for CIEF [15,21], a situation that could help to avoid interactions between heparin and EPO.

310 A commercial kit based on anti-EPO IAC and made of disposable monoliths was shown 311 to be useful to purify different recombinant EPOs from several matrices without isoform 312 discrimination [41-42]. A picture of the device is shown in Figure 4. The usefulness of this 313 technique for subsequent analysis by CZE on uncoated capillaries or SE-HPLC may be limited 314 due to the presence of BSA in the pH-adjustment buffer provided with the kit. The elimination of 315 BSA would be required and the impact of such a modification studied. The same antibody (anti-316 EPO 3F6) had been previously employed in larger columns using a pH 2.2 solution 317 supplemented with an inhibitor of aspartyl proteases (pepstatin) as desorption agent [43]. In that 318 case, nothing was mentioned about the need to use a pH adjustment buffer including BSA. In 319 principle, this IAC process, if followed by removal of the desorption agent, could be compatible 320 with CE or HPLC analysis.

As discussed, both polyclonal and monoclonal antibodies have been used as affinity ligands. When considering their use, they should be carefully chosen to allow EPO capture and elution without crossreaction with other proteins, as would be the case for the AE7A5 clone, which can crossreact with other proteins such as eukaryotic or bacterial, and namely with the urinary proteins Tamm Horsfall glycoprotein, alfa- antichymotrypsin, alfa 2-thiol proteinase
inhibitor, and alfa 2-HS glycoprotein [44-45].

327 Besides antibodies, other affinity ligands have also been used for EPO purification. 328 Among them, the use of synthetic peptides developed by a combinatorial strategy has recently 329 been shown to be effective for isolating rhEPO from Chinese Hamster Ovary (CHO) cell culture supernatant spiked with the recombinant glycoprotein [46]. The selected peptides were 330 331 immobilized on Sepharose and EPO was eluted with 90% recovery and around 96% purity using 332 100 mM sodium acetate buffer, pH 3.0, 0.25 M NaCl. These conditions should be compatible for 333 further CE or HPLC analysis provided that the sample is first cleaned of the elution buffer. 334 Lectins have also been used to purify EPO [47]. However, these affinity ligands, if not combined 335 with another isolation step, may not be appropriate for preparing EPO samples to be analyzed by 336 CE or HPLC as lectins show affinity for a wide range of glycoproteins. 337 Purification of rhEPO from human plasma was also performed by depletion of the seven 338 major plasma proteins followed by reversed phase desalting and purification [48]. This method,

which has been optimized for tandem mass spectrometry analysis of tryptic peptides, leads to
samples containing proteins that would make it unsuitable for CZE analysis of intact EPO on
uncoated capillaries or by SE-HPLC.

An alternative sample preparation method for EPO doping control in blood samples has been recently published [49]. The method is based on EPO solubility and stability at acidic pH and consists of sample (serum or plasma) precipitation with perchloric acid, followed by neutralization of the supernatant and further elimination of the precipitate formed during neutralization. Although this method was shown to be valid for performing IEF in gel followed by blotting assay, it may not be adequate for CZE analysis on uncoated capillaries as it was

348	observed when performing sample preparation of another glycoprotein (alfa-1 acid glycoprotein,
349	AGP) from serum using perchloric precipitation. AGP is also a very acidic glycoprotein; and
350	although most of the interferences from the sample were eliminated by acidic precipitation, the
351	remaining ones markedly decreased the electroosmotic flow (EOF). This is most probably due to
352	adsorption to the capillary wall [50]. In all likelihood, for EPO the remaining interferences would
353	also make this preparation method incompatible with SE-HPLC. This isolation method could
354	potentially be appropriate for samples to be analyzed by CE using coated capillaries.
355	2.3 General concerns for sample preparation for ESAs determination
356	Independent of the specific purification method chosen, EPO-containing samples should
357	generally be subject to pre-treatment steps. For example the modification, reduction and even
358	disappearance of the EPO bands in the IEF analysis of urine samples have been related to the
359	enzymatic activity from micro-organisms. Besides the activity from normal flora or other
360	microbial contaminants, concerns arose recently that purposely-added proteolytic enzymes, to
361	avoid doping control, can distort EPO determination results. Physical and chemical methods to
362	stabilize human urine samples for doping control have been summarized elsewhere [51]. The
363	addition of a general protease inhibition cocktail was the method of choice in some protocols,
364	and was considered more efficient in preventing EPO degradation by proteases than was heating
365	or the addition of pepstatin [45]. Coating plastic urine collection containers with a chemical
366	mixture of antibiotics, antimycotics and protease inhibitors was also advised [52]. In addition,
367	when analyzing urine samples, the need to solubilize precipitates prior to the purification step is
368	considered mandatory because the precipitates can contain up to 85% of the total amount of EPO
369	in the sample [53]. This is particularly relevant when analyzing frozen samples.
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372 3. High performance liquid chromatography for the determination of intact EPO and other
373 ESAs

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375 High performance liquid chromatography (HPLC) methods have been reported for the 376 determination of intact EPO and other ESP preparations in a variety of situations that included 377 identification, purity assessment and assay of the active ingredient. Of the several HPLC modes 378 SE-HPLC, IEX-HPLC and RP-HPLC have been successfully applied; a summary can be 379 observed in Table 2. As its name implies SE-HPLC separates compounds according to their size 380 (or hydrodynamic volume), which can, in turn, be approximated to the molecular weight. SE-381 HPLC columns are made of porous packing materials that allow compounds to interact with the 382 pores according to their size. Thus, in the absence of non-specific interactions, larger molecules 383 interact less with the stationary phase and elute ahead of smaller molecules. Depending on the 384 column porosity different ranges of molecular size can be adequately separated. For RP-HPLC, 385 separation is obtained based on differences in hydrophobicity between the compounds, with the 386 more hydrophobic species showing greater retention. RP-HPLC columns vary in their 387 hydrophobicity depending on the bonded phase used. Typically, C4 and C8 bonded phase columns with wide pores (300 Å) have been used for protein separations. In IEX-HPLC, 388 389 separation involves coulombic (also called electrostatic or ionic) interactions between the 390 components and ionic functional groups on the stationary phase. Both cationic and anionic 391 groups have been used for the separation of proteins.

392 The formation of dimers and high molecular weight (HMW) aggregates constitutes one393 of the main degradation pathways of EPO. Their presence in pharmaceutical proteins has been a

major concern to manufacturers and regulatory authorities alike due to their immunogenic
 potential. As such, the monitoring and control of these species in EPO and other ESAs have been
 the subject of several studies using chromatographic techniques.

397 Early reports have shown that SE-HPLC conditions could be used to study dimer and 398 HMW aggregate formation in EPO samples placed under stress conditions [54-55]. Silica-based 399 columns providing separation in the molecular weight range of 10– 500 kDa (medium-range) 400 were used. The detection of dimers and oligomers was found to be adequate by monitoring with 401 UV at either 214 nm or 280 nm. However, the use of a low angle laser light scattering detector 402 provided increased sensitivity for the detection of very large aggregates [54]. These two studies 403 showed that heat-stressed EPO samples led mainly to the formation of HMW aggregates. 404 Dimerization was favored when samples were placed under denaturing and reducing conditions 405 in the presence of dithiothreitol (DTT), presumably through the cleavage of disulfide bridges and 406 the subsequent random re-oxidation of the free thiols [55]. Similar SE-HPLC conditions were 407 later used to monitor dimers and HMW aggregates in formulation studies of EPO [56] and DPO 408 [57]. RP-HPLC conditions have also been shown to separate EPO dimers and HMW aggregates 409 [55]. In this case, both dimers and HMW aggregates showed increased hydrophobicity compared 410 to monomeric EPO.

A quantitative test for dimers and HMW aggregates has been prescribed in the European Pharmacopoeia (EP, also abbreviated as Ph.Eur.) monograph for EPO concentrated solution since the first publication of the monograph [58]. It is based on SE-HPLC with UV detection at 214 nm and applies to unformulated bulk preparations. A limit for total aggregates, including dimers and HMW aggregates, of less than 2% is specified. The method has been successfully 416 applied in collaborative studies for the establishment of the EP biological reference preparations417 for EPO [59-60].

418 The applicability of SE-HPLC for monitoring dimers and HMW aggregates in final 419 products has been complicated by the presence of excipients. For instance, formulations 420 containing HSA as a stabilizer are not amenable to determination by SE-HPLC due to the 421 interference of HSA with the detection of the EPO monomer. In most formulations the amount of HSA is much larger than the amount of EPO (2500 μ g mL⁻¹ of HSA vs 30-100 μ g mL⁻¹ of EPO). 422 423 In formulations where polysorbate 80 is present instead of HSA, there have been reports of peaks 424 due to the polysorbate interfering with the detection of EPO dimers and oligomers [61-63]. To 425 circumvent this problem detection by fluorescence was used [63-64]. Interestingly, other studies 426 have not reported the interference of polysorbate peaks on the detection of dimers and aggregates 427 [65-66] despite using separation conditions that do not appear too dissimilar, that is, phosphate-428 containing mobile phases and silica-based gel columns. It is not clear at this time on what basis 429 the latter conditions enabled to differentiate the polysorbate 80 peaks from the EPO dimers and 430 aggregates without further evaluation.

431 Methods based on RP-HPLC have also been used for the analysis of HSA-containing and 432 polysorbate-containing EPO formulations. RP-HPLC conditions using wide-pore, mildly 433 hydrophobic, C4 bonded (as opposed to the strongly hydrophobic, C18 bonded phase) reversed-434 phase columns were first reported by De Paolis et al. [55] and later used to provide adequate 435 separation between HSA and EPO [67]. This was also the case with strongly hydrophobic, C18-436 bonded phase columns that enabled the separation of HSA, EPO and aggregates [68]. Notably, 437 the latter conditions enabled the partial separation of the Ser-125 unglycosylated variant that was 438 detected in EPO-omega formulations. Similarly, underglycosylated EPO obtained from

transgenic pigs was also separated from CHO-derived EPO by RP-HPLC conditions [69]. In
both cases, the underglycosylated EPO was found to elute after EPO as a consequence of its
increased hydrophobicity.

A method based on anion-exchange HPLC with fluorescence detection was reported for the analysis of HSA-containing formulations [70]. Fluorescence detection was found to be more sensitive than UV at either 214 or 280 nm. In addition, the method provided selectivity between EPO-alfa and EPO-beta. IEX-HPLC conditions were also reported for the fractionation of EPO glycoforms. As expected, the gradient elution of EPO on anion-exchange columns using increasing salt concentrations resulted in fractions containing isoforms with decreasing average pI [71-72].

The EPO content measured by SE-HPLC and RP-HPLC has been correlated to the
biological activity in pharmaceutical finished products using the EP biological reference
preparation (BRP) for EPO as reference [66-67]. The average potency determined by RP-HPLC
was 11% higher than by the bioassay, whereas it was 2.5% higher by SE-HPLC. The authors
suggested that the latter method could be used as an alternative to the bioassay for the potency
determination of EPO preparations.

An attempt at the identification of intact EPO-alfa, EPO-beta and DPO by RP-HPLC coupled on-line to electro-spray ionization mass spectrometry (ESI-MS) has been described [73]. While the chromatographic step enabled the three components from a mixture to be resolved, the mass spectra of each of the three ESAs were of poor quality in terms of their mass resolution and signal intensity. The authors hypothesized that this was likely due to the presence of clusters of multicharged species deriving from the various isoforms.

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463 4. Separation and detection of intact EPO using capillary electrophoresis methodology
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465	Capillary electrophoresis (CE) modes of free zone (CZE) and isoelectric focusing (CIEF)
466	are well suited for the isoform separation of glycoproteins [74-75]. In particular, both
467	pharmaceutical preparations and natural samples of EPO have been analyzed using CE
468	[15,40,76]. It should be noted that the EP prescribes CZE as an identification method in its
469	monograph for Erythropoietin Concentrated Solution [58,77]. Most of the work on CE analysis
470	has been carried out using UV detection (at 214 nm) which has a limited sensitivity (in the range
471	of 10 ⁻⁶ M). Other detection techniques such as MS have been used and will be reviewed in this
472	paper.
473	In this section a general overview of the different CE methodologies for the
474	determination of intact EPO will be presented with a particular emphasis on papers published
475	during the period from 2006 until January 2011, which have been summarized in Table 2. Works
476	performed earlier have been already reviewed by us [74]. Considerations regarding different
477	possibilities to improve detection will also be discussed. The reader may also refer to review
478	papers of protein separation using CE that contain a summary of EPO (intact and digested)
479	studies [75,78].
480	4.1 Separation of intact EPO using CZE
481	Most of the recent work carried out by CE for the determination of isoforms of intact
482	EPO has been based on the method prescribed by the EP [58,77]. This method uses CZE for

483 identification of EPO (assay B). It uses uncoated fused silica capillaries and a separation buffer

484 composed of 0.01 M tricine, 0.01 M NaCl, 7 M urea and 2.5 mM putrescine. This diamine is

485 employed to avoid adsorption of the protein on the capillary surface and to control the EOF. 486 Separation is carried out at 35°C and UV detection at 214 nm is used. A critical point in the use 487 of this CE method is the capillary preconditioning. The monograph recommends rinsing new 488 capillaries with 0.1 M NaOH (60 min) and with the separation buffer (60 min), the latter applied 489 using 20 kV (for 12 h). Between runs, the monograph recommends rinsing the capillary 490 successively with water (10 min), 0.1 M NaOH (5 min) and separation buffer (10 min). With 491 these precautions the method provides the separation of 8 isoforms in 60 min with good 492 repeatability for the EP BRP standard.

493 However, some researchers have claimed a lack of reproducibility with the EP method. 494 For instance, Zhang et al. [19] have found that the causes of the poor reproducibility of the 495 method are: i) inefficient elimination of the excipient (e.g., polysorbate 80 in EPO-alfa) in the 496 sample and ii) inadequate capillary conditioning procedure. As mentioned in Section 2, removal 497 of polysorbate 80 from formulated EPO-alfa can be achieved using a commercial non-ionic, 498 detergent trap cartridge. The EPO recovered from the cartridge is then desalted using the method 499 proposed in the EP monograph prior to analysis by CE. For the capillary conditioning, the 500 authors highlighted the critical role of putrescine in the separation method. According to them, a 501 portion of the protein anionic isoforms is associated via electrostatic interactions with the 502 putrescine anchored on the silica surface through hydrogen bonding, ionic, and/or hydrophobic 503 interactions. The putrescine-isoform association reduces the net electrophoretic migration of the 504 isoforms, resulting in a modification of the migration velocity. The authors suggest that the rinse 505 of the capillary is a critical point for obtaining good repeatability of the migration time for the 506 EPO isoforms. This step is necessary to completely remove the putrescine and re-establish the 507 pH on the silica surface. Consequently, the authors proposed a rinsing procedure consisting of a

508 combination of base, acid, and water rinses over an extended period of time (more than 260 509 min). Although the recommended rinsing procedure to be done after every 10 injections is long, 510 its efficiency was demonstrated by the good reproducibility of the absolute migration time (for 511 peak number 2) of the EPO samples (RSD 1.8%, for 3 different samples with n=3 for each 512 sample) and even better for the relative migration time to the water peak (RMT) (RSD 0.2%, 513 under the same conditions). For peak areas, the inter-run RSD values (for 4 runs with a total 514 n=10) ranged from 0.2 to 0.7% (for the more abundant ones) and from 3.3 to 6.8% for isoforms 515 3 and 8 (those being the less abundant isoforms). As expected, better RSD values (ranging from 516 0.0 to 6.8 %) for areas were obtained in these experiments when intra-run values were 517 considered. The use of relative migration time to the EOF as (indicated by the water peak) 518 instead of the absolute migration time has been also advised by other authors. This is even in 519 cases where good repeatability of the absolute migration time (RSD < 0.6%, n= 6) has been 520 achieved for the separation of EPO isoforms [18]. The experience of our group is also that an 521 appropriate rinsing of the capillary is crucial for obtaining good repeatability among injections, 522 when using a procedure for sample preparation and a CE buffer similar to that of the EP [18]. 523 With a shorter capillary conditioning step (35 min between consecutive analysis) than the one 524 proposed by Zhang et al. (20 min between consecutive analyses and 260 min every 10 analysis) 525 [19], the repeatability of the migration time for the peaks of the isoforms of the EP-BRP EPO 526 sample was in the range of 0.29-0.56% (RSD, n=6) for analyses in the same day and in the range 527 0.98-1.72% (RSD, n=18) for analyses carried out in three different days with two different 528 instruments.

529 Other authors proposed the use of commercial polymeric coatings, such as hexadimetrine 530 bromide (Polybrene) or Ultratrol LN [79] to avoid the lack of reproducibility that had previously been observed for uncoated capillaries [80]. Using volatile buffers compatible with MS detection 6 isoforms were detected for the EP BRP standard when using the Polybrene coating, and more than 7 peaks were observed when Ultratrol LN was used. However, a lower number of peaks was obtained in the CE separations in comparison to those obtained using non volatile BGEs containing diamine additive, which could complicate the interpretation of the MS spectra (vide infra).

537 Another commercial polymeric cationic coating, carboxymethyl chitosan (CMC), has 538 shown promising results for rhEPO isoforms separation [81]. New capillaries were rinsed with 1 539 M NaOH (30 min) followed by 0.1 M NaOH (10 min) and water (10 min). Then, capillaries were 540 dynamically coated by rinsing them with 0.6% (w/v) of commercial CMC (MW 29.000) solution 541 in 0.1M HCl (30 min). Finally, the capillaries were left to stand (40 min) to allow CMC to 542 adsorb on the fused silica wall and followed by a 10 min purge with air. Between runs the 543 capillaries were flushed with the separation buffer (2 min). Using a volatile buffer of 100 mM 544 acetate (pH 4.5) and -30 kV for separation voltage, 9 isoforms of EPO (NCPC Gene Tech 545 Biotechnology Development, China) were separated in 9 min. Reproducibility of the migration 546 times for the isoforms ranged from 0.75 to 1.29 (RSD (%), n=5). Although, base-line resolution 547 was not achieved for the 9 isoforms, the CMC coating showed good potential for rhEPO 548 determination using both CZE and CZE-MS.

549 Amine-coated capillaries (eCAP) and a separation buffer containing 8 M urea and 300 550 mM phosphate were used for QC of rhEPO produced by several companies from Korea, China, 551 and India and compared to the one produced by Amgen (Epogen) in terms of the isoform 552 distribution [82], as mentioned in Section 6.1 of this paper. Information about capillary 553 conditioning and repeatability of the separation method was not provided.

554 A similar, but not identical, CE method to the one prescribed by the EP has been 555 developed for quantification of the 8 isoforms in samples of Epoetin beta [83]. The method used 556 70 mM MES buffer instead of 0.01 M tricine and 0.01 M sodium acetate employed by the EP 557 method. Forty production batches of this pharmaceutical protein were selected to cover adequate 558 range of precisely established potency values. The percentage of each isoform in the set of 559 samples was determined using CE analysis. The sialylation degree of each batch was also 560 measured. Several relationships between biological activity and chemical parameters were 561 statistically evaluated. As a consequence, a suitable model for correlating physico-chemical 562 parameters (percentage of each isoform and sialic acid content) and biological activity of the 563 EPO samples was established. The results obtained using the model were equivalent to those 564 obtained using the established in vivo bioassay in terms of accuracy, but they were superior in 565 terms of precision. The CE method was also proposed as an alternative to animal 566 experimentation in Quality Control (QC) of EPO.

567 Recently [23] the CZE method of the EP was used for determination of the sialic acid 568 content per molecule of different rhEPO subfractions; these subfractions were obtained using 569 chromatography from a conditioned medium of CHO cells expressing rhEPO. The sialic acid per 570 molecule for each subfraction was calculated using a first order polynomial equation of the 571 relative proportion (expressed as %) of each of the isoforms separated by CZE. The sialic acid 572 per molecule content from each subfraction was correlated to the in vivo bioactivity of the 573 fraction determined on the basis of increased number of reticulocytes following each EPO 574 fraction injection into a murine model. The authors observed a low correlation coefficient (0.76)575 among these two parameters, suggesting the possibility that other glycan moieties structures 576 besides sialic acid residues may have a significant impact on *in vivo* bioactivity of rhEPO

577 subfractions, a suggestion in accordance with the known influence of branching of the N-linked

- 578 glycans on the EPO *in vivo* activity [25].
- 579 4.2 Separation of intact EPO using CIEF

580 Since EPO isoforms can be separated based on differences in their respective isoelectric 581 point [4], CIEF is also well suited for isoform determination. However, due to difficulties in 582 achieving reproducible separations by CIEF only one report has been published during the period 583 reviewed [84]. In this article, CIEF was carried out with a commercial instrument (iCE 280 from 584 Convergent Bioscience) that used a separation cartridge which contains an internally 585 fluorocarbon-coated fused silica capillary (i.d. $100 \ \mu m$) with integrated buffer reservoirs. The 586 detection was carried out by whole column (50 mm) imaging using UV light at 280 nm and a 587 CCD camera that takes images every 30 sec during the separation process. The sample was prepared by mixing an aliquot of the EPO stock solution (typically 1 mg mL⁻¹ after desalting) 588 589 with 2% (v/v) ampholytes (Fluka pH 3-10), 1% (v/v) polyvinylpyrrolidone (PVP) and 4 M urea. 590 The anolyte was 100 mM phosphoric acid with 1% (w/v) PVP and the catholyte was 100 mM 591 NaOH also containing the same amount of PVP. The polymer was used to increase the viscosity 592 of the sample and to decrease the effect of the EOF in the separation. When the electric field (1.5 593 kV) was applied, the focalization of the EPO isoforms and the detection took place 594 simultaneously. With this procedure, isoforms 1-9 were separated in different EPO preparations 595 in 5 min (see Figure 5) with good reproducibility (RSD values for migration time relative to the 596 pI standard of 0.3% and 0.9% for run-to-run and day-to-day, respectively). The sensitivity of the technique with UV detection was estimated to be in the range of 10^{-6} M; at such a level, it is of 597 598 interest for pharmaceutical preparations analysis, but it would be of limited value for some 599 applications such as doping control.

600 4.3 Improving detection sensitivity in CE for intact EPO

601 Due to its favorable resolution, the potential of CZE for doping control is good [85]. 602 CIEF and gel CE could potentially be valid for this purpose taking into consideration the 603 experience acquired using IEF in gel [86] and SDS-PAGE [87-88]. However, in our opinion, the 604 major limitation for the development of a doping control method based on CE techniques is 605 detection. It is tempting to speculate that the use of laser induced fluorescence (LIF) detection, 606 which is around 1000 times more sensitive than UV, could help for these applications. 607 Nonetheless, some difficulties have to be overcome before achieving the monitoring of EPO 608 isoforms using LIF detection. These problems are analyzed and potential solutions based on the 609 literature are presented in the next few paragraphs.

610 LIF detection is one of the most sensitive techniques for CE [89]. However, few analytes 611 have native fluorescence when excited by visible laser light. For this reason, covalent 612 derivatization and laser with emission in the visible region of the spectrum must be used to get 613 detection sensitivity in the low nM range. For LIF detection of proteins non-covalent and 614 covalent derivatization procedures have been proposed [90]. Non-covalent derivatization relies 615 on the adsorption of a dye, which is dissolved in the separation buffer, on the hydrophobic 616 patches of the proteins of the sample. The adsorption takes place inside the column once the 617 sample has been injected and the dye wavelength and emission intensity increase with respect to 618 unbound dye. In this way the protein-dye complex emits fluorescence when it passes through the 619 laser beam at the detection point. However, hydrophobic interaction of such substances with 620 glycoproteins is generally very small or prevented. In fact, to our knowledge, no substance has 621 been reported to be useful for the non-covalent derivatization of EPO isoforms. Although several 622 methods for covalent derivatization through amino group of Lys residues have been proposed for

623 protein derivatization, multiple derivatization products (leading to multiple peaks or broad peaks 624 in CE) and slow reaction (when the protein is at low concentration ($< 10^{-9}$ M) in the sample) 625 limit covalent derivatization of proteins. The use of on-column derivatization using submicellar 626 concentration of detergent (SDS or sodium pentasulfonate - SPS) in the separation buffer have 627 been used successfully in many cases [91]. However, the use of an ionic detergent in the 628 separation buffer may also prevent the separation of isoforms of glycoproteins [92]. Preliminary 629 results recently obtained by our group for another glycoprotein (prostate-specific antigen) have 630 shown the feasibility of separating isoforms by CZE of on-column fluorescently-labeled 631 glycoprotein [93].

632 Covalent derivatization through thiol group offers better perspective for avoiding 633 multiple derivatization products since Cys amino acids are less abundant than Lys in proteins. 634 This strategy has proved effective for separation of isoforms of some glycoproteins [94] and its 635 applicability to the determination of EPO isoforms is currently under study in our group. 636 It should be mentioned that native fluorescence of proteins has been probed [95] using 637 laser light at 275 nm that is able to excite the fluorescence of the tryptophan residues of the 638 proteins. However, this approach could be difficult to carry out for proteins in most natural 639 samples due to impurities that are always present in natural samples which fluoresce at the same 640 wavelength. This can lead to a strong fluorescence background and a poor sensitivity for LIF 641 detection in this type of samples.

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5. Determination of intact EPO and other ESAs by capillary electrophoresis-mass
spectrometry

647	Owing to the structural information that MS detectors can provide, this technique has
648	been coupled online to CE (CE-MS) for the determination of different compounds including
649	glycoproteins [74]. In particular, several papers were published in 2005-2006 on the
650	determination and characterization of intact EPO and DPO by CE-MS [79,96-98]. Intact standard
651	EPOs (BRP EPO, epoetin alfa, and epoetin beta) were fairly well, though not completely,
652	characterized by CE-MS. After a thorough analysis of the deconvoluted mass spectra for each
653	electrophoretic peak, about 64 glycoforms of EPO differing in their glycan moieties were
654	characterized by CE-time of flight (TOF)-MS [79]. Minor modifications of these 64 glycoforms,
655	such as acetylation and exchange of N-acetylneuraminic acid (NeuNAc) with N-
656	glycolylneuraminic acid (NeuNGc), were also observed. Detection of NeuNGc in the glycan
657	moiety of EPO is especially relevant as an indicator of misuse of EPO (doping) because glycans
658	containing NeuNGc are produced by CHO cells [99] but not by human cells [10,100-101]. In
659	spite of the extensive work carried out, some post-translational modifications (PTMs) involving
660	shifts of 1 or 2 Da such as deamidation ($\Delta = 1$ Da), addition of 2 fucose molecules <i>vs</i> addition of
661	1 molecule of sialic acid to the glycan ($\Delta = 1$ Da), cleavage of disulfide bridge ($\Delta = 2$ Da),
662	hydroxylation vs methylation (Δ = 2 Da), and combinations of oxidation and water loss (Δ = 2
663	Da), in a 30 kDa (approx.) protein could not be assessed due to performance limitations of the
664	MS equipment employed [96]. From 2007 to the present, only a few papers dealing with the CE-
665	MS determination of intact EPO have been published [22,102].
666	When CE-MS determination of proteins or glycoproteins is carried out, several practical
667	problems should be considered. From a separation point of view, as it is known, proteins can
668	undergo non-specific adsorption to the capillary wall, and/or lack of reproducible EOF can

669 occur. Highly alkaline BGE (to ensure the electrostatic repulsion of analytes from the capillary 670 wall) or coated capillaries (covalent or non-covalent – either dynamic or semi-permanent – 671 coatings) have been used to prevent adsorption. Between these different options, the most 672 common approach for CE-MS of proteins is the use of capillaries with non-covalent, semi-673 permanent coatings. For instance, polybrene [79,96,98], different ionene polymers [97], and N-674 substituted acrylamide copolymers (UltraTrolTM) with high reversed EOF (UltraTrolTM HR) 675 [102] or almost suppressed EOF (UltraTrol[™] LN) [22,79,102], have been used for separation of 676 glycoforms of EPO by CE-MS. The efficiency of the CE-MS determination of glycoprotein 677 glycoforms depends on the partial separation obtained in the CE analysis, the ionization 678 efficiency and the performance of the MS employed for the detection; the best results were 679 obtained using UltraTrol[™] LN as coating. When comparing the two UltraTrol coatings [102], 680 UltraTrolTM LN provided better results in terms of the determination of the glycoforms of EPO. 681 but UltraTrolTM HR showed higher stability and less leakage, which, from the MS detection 682 point of view, is a positive characteristic. With UltraTrol[™] LN better glycoform resolution was 683 achieved for EPO than for DPO due to the higher microheterogeneity of the latter glycoprotein 684 [102]. In addition, better separation of EPO glycoforms was obtained with UltraTrol[™] LN in 685 comparison to the polybrene coating [79] and with either of these two coatings than with ionene 686 polymers [97]. In Figure 6, the electropherogram of EPO BRP analyzed by CE-MS using an 687 UltraTrol[™] LN coated capillary is shown.

In all of these papers dealing with CE-MS of EPO, which are collected in Table 2, the use of a volatile BGE was compulsory due to the MS detection. Acetic acid – ammonium acetate buffers [97], or acetic acid (1M or 2M) [22,79,96,98,102], with [79,98] or without [22,79,96-97,102] methanol, were used as BGE for the CE separation. 692 In the above-mentioned papers, the coupling of the CE instrument to the MS detector, 693 operating in electrospray ionization (ESI) mode, was through a sheathflow interface. In most 694 instances [22,79,96,98,102], the composition of the make-up liquid was a mixture of water and 695 isopropanol (1:1, v/v) containing 1% (v/v) acetic acid at a flow rate in the range of $3.3 - 4 \mu L$ 696 min⁻¹ operating in the positive ion detection mode. For the negative ion detection mode [97] a 697 completely different make-up liquid composition was required and consisted of 5 mM ammonium acetate in 50 % v/v water/acetonitrile at a flow rate of 10 µL min⁻¹. However, in the 698 699 latter case the performance of the analysis of EPO glycoforms was poor. Although some peaks 700 assigned to EPO glycoforms by the authors were resolved by CE-MS, their MS spectra was very 701 weak and without the typical MS peaks pattern of multiply-charged glycoproteins. This is likely 702 due to a combination of the separation conditions, the ionization settings and the detector 703 employed. For instance, the negative ionization mode was used, though positive ionization mode 704 tends to be the best option for the determination of proteins as ammonia seems to give rise to 705 inefficient ionization of proteins. In fact, Neususs et al. [96] were unsuccessful in detecting EPO 706 glycoforms in the negative ionization mode, while, as mentioned above, succeeded using 707 positive ionization mode. Besides, the dilution of the eluent from the CE column was larger as they used a higher make-up liquid flowrate (10 μ L min⁻¹ vs 3.3-4 μ L min⁻¹). 708 709 Different MS instruments with different MS analyzers such as ion trap (IT) [97,102], 710 time of flight (TOF) [79,96,98] and extra-long flight path TOF [22] have been used for the

determination of EPO glycoforms. Two equipments with IT analyzer (Agilent [97] and Bruker
[102]) have been used. The two instruments led to different results in terms of identification of
EPO glycoforms, probably because different separation conditions and ionization settings were
used. The Bruker MicrOTOFTM has been the only "normal" TOF analyzer used for the CE-MS

715 determination of EPO glycoforms and provided better results than IT analyzers. This was 716 expected as, in general, TOF analyzers show better performance than IT analyzers when there is 717 a high number of partially co-migrating glycoforms over a broad mass/charge range. Reasons 718 include better suitability for larger m/z values (quadrupole based instruments including ITs have 719 limited mass range), better de-clustering due to higher vacuum and shorter or no storage time in 720 TOF instruments. For similar separation conditions (same BGE and capillary coating) and 721 ionization settings (sheathflow composition) 64 glycoforms of EPO were detected with a TOF 722 analyzer [79] while only 17 were detected with an IT analyzer [102]. Recently, a new study [22] 723 was published dealing with improvement of the determination of EPO by CE-MS where an 724 extralong ion flight path TOF (5 m flight path, maXis quadrupole time-of-flight mass 725 spectrometer also called maxis 4G UHR-TOF from Bruker) was used as opposed to the "normal" 726 TOF analyzer (2 m flight path, Bruker MicrOTOFTM), also called benchtop TOF-MS. In this 727 work EPO was presented as a proof-of-concept for resolution improvement in MS spectra 728 achieved for large proteins (Figure 6). With the extralong flight path TOF analyzer, isotopically 729 resolved MS spectra can be obtained leading to the accurate determination of the empirical 730 formula of the protein and, as a consequence, to the determination of the accurate structure of the 731 protein having PTMs involving mass shifts as small as 1 Da (such as deamidation). The authors 732 determined the accurate empirical formula of the protein by comparison between the isotopically 733 resolved, deconvoluted MS spectra and the theoretical isotopic MS spectra of the protein. 734 However, the authors recognize that even with the very high resolution provided by the extra-735 long flight path TOF analyzer, the unambiguous identification of PTMs involving very small 736 mass shift (1 Da) is not always accomplished due to the comigration of glycoforms and the 737 overlap of the isotopic pattern. However, in view of the technical improvement provided by the

extra-long flight path TOF, the publication of the CE-MS determination of the glycoforms ofEPO with this new instrumentation in the near future can be expected.

740 There are other MS instruments with high mass resolution and accuracy such as orbitrap MS and

741 Fourier Transform Ion Cyclotron Resonance (FTICR)-MS, but their features -resolution and

accuracy- depend on the measuring time. As a consequence, in order to obtain the best

performance when FTICR-MS is coupled to CE separation, the measured mass range is reduced

in order to devote the required time to measure each m/z ion. Although CE-FTICR-MS has been

visual relation visual relation of used to analyze protein glycoforms [103], it has not been applied yet to the characterization of

746 EPO.

747 As mentioned in the Introduction section, the determination of glycoforms of intact 748 glycoproteins presents several advantages in comparison to hydrolyzed glycoproteins. This 749 approach permits obtaining the whole glycoform profile without further "reconstruction" of the 750 glycoprotein structure. However, the observed microheterogeneity cannot always be 751 unequivocally assigned to either the glycan structure or the protein backbone. For instance, a 16 752 Da mass shift can be assigned to several pathways: hydroxylation, exchange of NeuNAc with 753 NeuNGc or oxidation at either methionine, histidine (to oxo-histidine), cysteine (to sulfenic acid) 754 or proline. Although it may be possible to ascertain the nature of such a PTM from the MS 755 results and the electrophoretic behavior of the glycoform (as the PTM may have an effect on the 756 migration of the glycoprotein), a combination of the CE-MS determination of the intact 757 glycoprotein with other techniques is very useful to characterize the structure of glycoproteins. 758 Such approaches could involve selective digestion processes (e.g., deglycosylation, proteolytic 759 digestion) and the determination of the different fractions (glycans, glycopeptides, and peptides) 760 by different techniques such as HPLC (IEX, RP, or HILIC), CE, MS, and hyphenations CE-MS

761 and HPLC-MS. CE-MS analysis of an intact glycoprotein can provide the glycoforms profile of 762 the target glycoprotein and, as a consequence, the global composition of its glycan moiety, but 763 the exact structure of the glycans cannot be completely determined. In addition, when different 764 glycosylation sites are present in the glycoprotein, as it is the case for EPO, it is not possible to 765 determine the glycan composition of each glycosylation site individually. Besides, there are 766 several possible isomeric structures associated with a given monosaccharide composition that 767 cannot be distinguished by simple mass spectrometric measurements without fragmentation. For 768 instance, a tetra-antennary structure is equivalent to one tri-antennary glycan containing one N-769 acetyllactosamine (LacNAc) repeat. Moreover, information relative to the interglycosidic 770 linkages cannot be obtained using this methodology [79]. Thus, in general, the above-mentioned 771 approaches complementary to CE-MS of the intact analyte are useful for a deeper 772 characterization of the glycoprotein.

773 Finally, one of the problems of the CE-MS determination of glycoforms of glycoproteins 774 is its lack of sensitivity. For instance, the concentration of the standard EPO analyzed in the articles cited above was in the range of $1.5 - 15 \text{ mg mL}^{-1}$. While these concentrations may be 775 776 appropriate for characterizing pharmaceutical preparations they would not be suitable for EPO 777 concentration expected in human biological fluids. To our best knowledge, there is only one 778 published article [97] where intact standard urinary EPO was analyzed by CE-MS and the MS 779 results were not very useful for the comparison of the glycoform profile between recombinant 780 and urinary EPO due to lack of quality in the MS spectra as it was mentioned before. 781 Improvements in the purification and concentration methods for EPO determination should be 782 developed, as discussed in Section 2, enabling future comparisons of the glycoform pattern of 783 recombinant and endogenous EPO. The knowledge of these differences could be of interest both

784	in pharmaceutical drug development, in order to decrease chances of immunological reactions,
785	and for doping control.
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788	6. Applications
789	
790	6.1 Regulatory considerations and Biosimilars.
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792	EPO was one of the first recombinant protein pharmaceuticals to reach the market at the
793	end of the 1980s and, just as for any other therapeutic entity, market authorization was granted
794	on the demonstration of safety, quality and efficacy. These are the three pillars on which the
795	regulatory approval process is based in most jurisdictions and they have been the focus of
796	extensive harmonization efforts that culminated in the development of the International
797	Conference on Harmonization (ICH) guidelines that are now widely accepted by regulatory
798	agencies and manufacturers [104]. In addition, in several jurisdictions manufacturers must
799	comply with pharmacopoeial requirements for the establishment of quality attributes for the drug
800	substance. Altogether, these regulatory considerations have led to the development and
801	widespread adoption of high resolution separation techniques and methods for the
802	characterization of biopharmaceuticals due in large part to their increased precision and
803	quantitative nature. This has been particularly the case for EPO.
804	As mentioned in Section 4, capillary electrophoresis has been prescribed as an
805	identification test since the publication of the first EPO monograph of the European
806	Pharmacopoeia (EP) [58,77]. The test is based on the separation of intact EPO isoforms by CZE

and applies to the analysis of the drug substance. As such, a sample treatment is required to
eliminate excipients that may be present for stability purposes. The isoform profile of the test
sample is compared to that of the EPO BRP for the identification of the specific isoforms. The
percentage content of each isoform in the test sample is then calculated and must be within a
specified range.

The quantitative determination of dimers and other high molecular mass aggregates by SE-HPLC has also been a prescribed test in the EP monograph for EPO since its initial publication [58,77] (see section 3).

815 As indicated above, EPO was one of the first recombinant protein pharmaceuticals to 816 reach the market and it has been one of the most successful, with worldwide annual sales 817 exceeding 10 billion US dollars in 2010. Soon after the patent expiry of the two innovator 818 products, EPO-alfa (Eprex and Epogen) and EPO-beta (Neorecormon), in Europe and the USA 819 many biopharmaceutical companies indicated their intention to bring their own EPO version to 820 the market. This led regulatory authorities to look into the concept of biosimilars, which was 821 formally adopted in Europe in 2005 [105]. It differs significantly from the concept of generics 822 used for small molecule pharmaceuticals. In the standard generic approach the demonstration of 823 bioequivalence is made with a reference product through appropriate bioavailability studies. This 824 approach is not scientifically feasible for biopharmaceuticals due to their complex nature. 825 Instead, a comparability exercise is recommended and it is the degree of similarity that 826 ultimately is evaluated. In Canada, the terminology "subsequent entry biologics" (SEBs) has 827 been adopted and a pathway similar to that of Europe has been recommended [106]. In the 828 United States, where the terminology of follow-on biologics (FOBs) is used, no guidelines have 829 been issued as of the beginning of 2011.

830 The first EPO biosimilars were approved in 2007 in Europe (under the names Binocrit, 831 Epoetin alfa Hexal and Abseamed) using the EPO-alfa innovator product, Eprex, as comparator. 832 Since then a few others have been marketed. Among the contentious issues to surface in the 833 debate between innovators and biosimilar manufacturers was the lack of availability of the 834 comparator's bulk drug substance for biophysical and chemical comparability testing. In such a 835 case, a biosimilar manufacturer must rely on obtaining it through purification from the 836 comparator's finished drug product. This has led to the publication of studies on the biophysical 837 comparability of EPO purified from different formulations [107-108]. 838 Using a wide range of techniques that included CE and HPLC, Deechongkit et al. [107] 839 were the first to compare EPO-alfa isolated from two commercially-available formulations, 840 namely Eprex and Epogen. EPO was purified from formulations by IEX -HPLC. They used CZE 841 to examine the isoform profiles of the two purified EPO products and found them to be similar. 842 On the other hand, determination of the dimers and HMW aggregates by SE-HPLC showed that 843 the purification procedure gave rise to a slight increase in the aggregate content for one of the 844 two products. Ultimately, significant differences were observed in the tertiary structure and they 845 concluded that the two proteins were not structurally identical. Heavner et al [108] conducted a 846 follow-up study and made a counter-argument to the approach of Deechongkit et al. by showing 847 that the purification procedure used irreversibly altered the physical structure of the purified 848 EPO. 849 EPO products obtained from several countries around the world, although not strictly 850 considered biosimilars, have been compared to innovator products such as Eprex [109] and 851 Epogen [82]. In the latter study, the determination of EPO isoforms was carried out by CZE and

showed differences in isoform profiles between several EPO products obtained from Asia. Both

studies showed wide variability in terms of glycoform profiles, content and biological activity,
and even batch-to-batch consistency. Isoform profile determination by CZE was also used in the
comparative assessment of the approved biosimilar, Binocrit, to the innovator product, Eprex.
Binocrit was found to be comparable, of high quality, safe and efficacious [110]. A similar
conclusion was reached by another group on the quality assessment of two biosimilars, Binocrit
and Retacrit, and two innovator products, Eprex and Dynepo [111]. Again, CZE determination of
isoform profiles was used along with SE-HPLC.

A recent article featuring the ESA product, Aranesp, has brought forward into the public domain another contentious issue hotly debated for some time between innovators and biosimilar manufacturers. It relates to the degree with which differences in quality attributes are acceptable [17]. Among several other techniques, they used CZE to demonstrate significant differences in the isoform distribution profile between different batches of the product and concluded that these variations had been deemed acceptable by health authorities since the batches had been marketed without any label alterations.

867

868 6.2 Doping control

869

EPO and the rest of the ESAs are included in the List of Prohibited substances by the
World Anti-Doping Agency (WADA); they are part of the substances classified as prohibited at
all times (in- and out-of-competition) [112].

For epoetin alfa, epoetin beta (both produced in CHO cell lines) and epoetin omega
(produced in hamster kidney cells) the basis for detecting rhEPO doping relies on the different
glycosylation of the protein as a function of the organism or cell line in which it is produced. As

876	a result of glycosylation changes, the isoelectric points and molecular weights of recombinant
877	EPOs are higher than in uhEPO. The isoforms in darbepoetin are more acidic than those in
878	uhEPO due to its increased number of sialic acid residues from the higher number of
879	glycosylation positions. CERA can be distinguished from endogenous EPO due to its higher
880	molecular weight. Epoetin delta (produced in human fibrosarcoma cell line) in comparison to the
881	endogenous EPO shows more acidic isoforms on IEF, but these differences are not sufficient to
882	differentiate them according to the WADA's criteria; doping with epoetin delta is detected by
883	differences observed by SDS-PAGE. A summary of ESA doping strategies and their abuse
884	detection methods was published by Jelkmann [113].
885	None of the methods approved for EPO-related doping control is based on HPLC or CE

886 techniques. The method presently used for EPO and other ESAs doping control is based on 887 Western blot after IEF and, if additional scientific evidence is needed, SDS-PAGE or an 888 equivalent procedure can be used [114]. IEF allows partial discrimination of isoforms of 889 endogenous (either urinary, plasma or serum) EPO from those of different recombinant epoetins. 890 Identification windows for uhEPO and ESAs are established by WADA [114]. Detection is 891 based on a double blotting procedure followed by chemiluminescent detection, which provides 892 high selectivity and sensitivity [86]. In addition to these gel electrophoretic methods, ELISA-893 based methods (to measure EPO and CERA) are among the techniques used at present by 894 WADA-accredited laboratories to detect ESAs [115-116].

A CE competitive immunoassay which, according to the authors, could be potentially useful for EPO doping control has been developed using silica dioxide nanoparticles as pseudo stationary phase and chemiluminescence detection. The assay is based on separation of EPO labeled with horseradish peroxidase (EPO-HRP) from the immunocomplex-HRP. High sensitivity (0.9 ng mL⁻¹ = 3×10^{-11} M) and high correlation with commercial immunoassay kit are achieved. However, the assay does not separate EPO isoforms and cannot differentiate endogenous from exogenous EPO; thus doping detection should be based on total EPO concentration [117].

903 Taking into account the separation mechanisms involved in differentiation of exogenous 904 from endogenous EPO by IEF or SDS electrophoresis performed in conventional gels, EPO 905 doping control by CIEF, CZE, or gel CE should potentially allow distinguishing EPO from both 906 origins. However, doping control based on separation of isoforms of intact EPO (or related 907 molecules) either by HPLC or by CE has not been carried out. Similarly, gel CE has not been 908 applied for EPO doping control. The main limitations are the low concentration of endogenous 909 and exogenous EPO or EPO-related substances in biological fluids (in the range of mIU mL⁻¹, this is about $pg mL^{-1}$) and the complexity of the matrices which include a high number of 910 911 proteins in a broad concentration range. To solve these limitations, sample preparation methods 912 aimed to isolate and concentrate EPO substances, and detection methods providing enhanced 913 sensitivity while preserving isoform resolution, are needed.

914 The CE or HPLC published papers dealing with differentiation of endogenous from 915 exogenous EPO substances based on isoform separation have focused on the development of 916 methods using standard glycoproteins of both origins. In our laboratory, a preliminary study 917 performed by CZE with UV detection showed different isoform profiles for human EPO 918 standards from recombinant and urinary origin. uhEPO showed some isoforms migrating slower 919 than the isoforms in rhEPO (see Figure 7) [85]. If an almost constant size is assumed for all 920 isoforms, the slower isoforms in uhEPO would correspond to more acidic forms than in rhEPO, 921 a result that is in accordance with those obtained by conventional electrophoresis by Wide et al.

[118] and by Lasne and de Ceaurriz [86]. Although this result was encouraging, the sensitivity
obtained was not enough to analyze endogenous and exogenous EPO at the levels existing in
natural samples.

Differentiation of urinary and recombinant human EPO standards has been attempted by coupling CE-ESI-MS [97], as mentioned above. Although the authors indicated that a unique m/z value was observed for rhEPO and a different one for uhEPO, identification of those ions was not performed and the low sensitivity of this CE-MS method prevented its use for doping control.

An attempt to increase sensitivity while maintaining CZE resolution of EPO isoforms has been performed using standard rhEPO. Combination of large volume sample stacking and reversed pH junction allowed about 50 times on-line sample preconcentration at the cost of reduced resolution [119]. However, this increase in sensitivity was insufficient for doping control.

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936

937 7. Concluding remarks and future perspectives

938

At the present time HPLC and CE methods developed for the determination of EPO and
other ESAs enable most of the challenges presented by such complex proteins to be addressed.
Sufficient isoform resolution can be achieved using CE methods, particularly in CZE and CIEF
modes, although some isoforms can also be separated by RP-HPLC and IEX-HPLC. These two
HPLC modes have also been shown to be adequate for the separation of ESAs from excipients.

In addition, monitoring of the formation of dimers and aggregates can be achieved by RP-HPLCand especially by SE-HPLC.

HPLC and CE separation techniques are very useful for the analysis of pharmaceutical
products. In fact, SE-HPLC and CE tests are prescribed by the European Pharmacopoeia for
quantitation of EPO dimers and aggregates and for EPO identification through the separation of
isoforms, respectively. The importance of ESA determination is increasing enormously with the
continuing expansion of biosimilars.

951 The main limitation of HPLC and CE for the determination of ESAs is the insufficient 952 sensitivity provided by UV or MS detection. This fact, together with the difficulty in analyzing 953 ESAs in samples of high complexity, limits their application for pharmacokinetics studies and 954 doping control. Two approaches can be followed to overcome these difficulties. Purification and 955 concentration can be achieved by applying sample preparation methods aimed at isolating the 956 target ESA from its complex matrix. Different sample preparation methods based on 957 immunorecognition have been successfully applied for performing doping control following the 958 IEF WADA accepted method. These results could be adapted to prepare ESA samples to be 959 analyzed by CE or HPLC. Alternative ligands, such as aptamers, whose affinity for EPO has 960 been already demonstrated [120-121] could be tested for purification and concentration 961 applications. The aptamer approach would require more sensitive detection compatible with the 962 separation technique. In this regard, LIF is known to provide higher sensitivity than UV 963 detection and, together with fluorescent derivatization through the amino or thiol groups, could 964 still allow CZE separation of isoforms, as was recently developed [93-94]. Another promising 965 strategy has been developed for LIF detection in CIEF [122]. LOD in the high femtomolar range 966 has been achieved by employing photo-bleached ampholytes, derivatization through the thiol

groups using Chromeo P450 as fluorescent derivatization reagent, and a 532 nm laser for
fluorescent excitation. These methods or similar ones could be assayed for CE-LIF of EPO
isoforms.

In order to automate and simplify the analysis, sample preparation, CE or HPLC
separation, derivatization, and detection could be applied on-line either by coupling HPLC
columns or by performing all the steps in the same CE column. Furthermore, miniaturization via
the analysis in microchip format would decrease the analysis time, sample volumes and reagent
consumption, and provide disposable platforms for analysis, thus avoiding cross-contamination
between samples.

976

977 Figure legends.

978

710	
979	Figure 1. Molecular model of erythropoietin with complex N-linked glycans at sites N24, N38
980	and N83. O-linked glycans are not represented. The structure shown is just one possible
981	conformation. Reprinted permission from Macmillan Publishers Ltd. (C. Sheridan, Nature
982	Biotech., 25 145-146) copyright (2007) [2]. The picture was a courtesy of M.R. Wormald, R.A.
983	Dwek and P.M. Rudd.
984	
985	Figure 2. Schemes of epoetin, darbepoetin, and CERA: glycosylation and pegylation sites (for
986	CERA). Reprinted from W. Jelkmann, "Recombinant EPO production-points the nephrologist
987	should know" Nephrol. Dial. Transplant. (2007) 22, 2749-2753 by permission of Oxford
988	University Press [3].
989	
990	Figure 3. Total protein content (black line and right scale) and hEPO (bars and left scale) present
991	in each fraction through the immunopurification procedure using immunoaffinity well plates:
992	initial plasma sample, unbound fraction, first wash, second wash, and elution fraction.
993	Reprinted from J. Chromatogr. B 878, J. Mallorqui, E. Llop, C. de Bolos, R. Gutierrez-Gallego,
994	J. Segura, J.A. Pascual "Purification of erythropoietin from human plasma samples using an
995	immunoaffinity well plate" 2117-2122, Copyright (2010) with permission from Elsevier [34].
996	
997	Figure 4. Immunoaffinity purification set-up. The anti-EPO antibody is covalently bound to the
998	monolith disk which is held inside the plastic housing. The device is used in a multi-channel

999 vacuum manifold. Reprinted from J. Chromatogr. A 1217, M. Lonnberg, Y. Dehnes, M. Drevin,

1000	M. Garle, S. Lamon, N. Leuenberger, T. Quach, J. Carlsson, "Rapid affinity purification of
1001	erythropietin from biological samples using disposable monoliths" 7031-7037, Copyright (2010)
1002	with permission from Elsevier [41].
1003	
1004	Figure 5. CIEF comparison of different EPO samples using a microchip-like separation device
1005	with internally fluorocarbon-coated fused silica microchannel. EPO samples (1 mg mL ^{-1}
1006	containcontaining 4 M urea, 1% PVP and 2 % ampholytes pH 3-10). Reprinted from J. Chrom. A
1007	1190, P. Dou, Z. Liu, J.G. He, J.J. Xu, H.Y. Chen, "Rapid and high-resolution glycoform
1008	profiling of recombinant human erythropoietin by capillary isoelectric focusing with whole
1009	column imaging detection" 372-376, Copyright (2008) with permission from Elsevier [84].
1010	
1011	Figure 6. Electropherogram of an EPO BRP PE sample (ca. 120 ng injected): CE-MS analysis
1012	using a soluble linear polyacrylamide coated capillary and 1 M acetic acid as BGE. All the traces
1013	are the extracted ion electropherograms (EIEs) of the average mass of the most abundant charge
1014	state. Traces of same color represent EIEs of glycoforms showing the same number of sialic acid
1015	(SA) residues. Within these groups the glycoforms are separated by the number of hexose-N-
1016	acetyl-hexosamine (HexHexNAc) units. Reprinted from J. Proteom. 74, A. Taichrib, M. Pelzing,
1017	C. Pellegrino, M. Rossi, C. Neususs, "High resolution TOF MS coupled to CE for the analysis
1018	of isotopically resolved intact proteins" 958-966, Copyright (2011) with permission from
1019	Elsevier [22].
1020	

Figure 7. CZE electropherograms of A) uhEPO, B) rEPO to which mannitol had been added andthen removed, and C) rhEPO. Electropherograms are aligned according to the retention time of

- 1023 the EOF (Negative peak not shown). Modified from M. de Frutos, A. Cifuentes, J.C. Diez-Masa,
- 1024 "Differences in capillary electrophoresis profiles of urinary and recombinant erythropoietin",
- 1025 Electrophoresis with permission from John Wiley and Sons [85].

1026

1027 Acknowledgments

- 1028 Financial support from the Spanish Ministry of Science and Innovation (projects PCI2006-A7-
- 1029 0646 and CTQ2009-09399) is acknowledged. Angel Puerta acknowledges the Spanish National
- 1030 Research Council (CSIC) for a contract in the JAEdoc program.
- 1031

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ESA	Manufacturer	Geographical area of origin
Abseamed	Medice Arzneimittel Pütter	Germany
Aranesp	Amgen	USA
Binocrit	Sandoz	Germany
Bioetin	Nanogenpharma	Vietnam
Biopoin	CT Arzneimittel	Germany
Bioyetin	Probiomed	Mexico
Ceriton	Rambaxy	India
Dynepo	Shire	Ireland
EPIAO	3SBio(Shenyang Sunshine Pharmaceuticals)	China
Epocim	CIMEQ	Cuba
Epocrin	Sotex	Russia
Epoetal	Pliva	Croatia
Epoetin Alfa Hexal	Hexal Biotech Forschungs	Germany
Epofer	Emcure	India
Epofit	INTAS Pharmaceuticals	India
Epoform	Eipico	Egypt
Epogen	Amgen	USA
Epogin	Chugai Pharmaceutical Company	Japan
Epokine	CJ Corporation	South Korea
Eporatio	Ratiopharm	Germany
Eporise	Zuventus	India
Eporon	Dong-A Pharmaceutical Company	South Korea
Eposino	Shandong Kexing	China
Epostim	Reliance Genemedix	UK
Epotin	Claris Lifesciences	India
Epotrex-NP	Novell Pharmaceutical	Indonesia
Epoyet / Epoimmun	Bio Sidus	Argentina
/ Hemax / Hypercrit		1 ii goinnia
/ Zyrop		
Eprex	Janssen – Ortho Biotech	USA
Eritina	Chalver Laboratories	Columbia
Eritrogen	Bioprofarma	Argentina
Eritromax	Blausiegel	Brazil
Erlan	Landsteiner Scientific	Mexico
Erykine	INTAS Pharmaceuticals	India
Erypo	Janssen-Cilag	Europe
Erypoietin	Amoun Pharmaceuticals	Egypt
Erypro	Biocon Pharmaceuticals	India
Erytrex	Cinnagen / Zhavari	India / Iran
Espo	Kirin Brewery Company, Pharmaceutical Division	Japan
Espogen	LG Life Sciences	South Korea
Exetin	Pisa Farmaceutica	Mexico
Gerepo	North China Pharmaceutical Group Corporation	China
Нетаро	Innogene Kalbiotech	Singapore
-	Bergamo	Brazil
Hemoprex	Derganno	DIAZII
	-	

Table 1. Non – comprehensive list of Erythropoiesis Stimulating Agents (ESAs) around the world and the geographical areas of production *

Hemotin	Itaca Laboratorios	Brazil	
Huan'er Bo	Beijing Four Rings Pharmaceuticals	China	
JR-013	Kissei / JCR Pharmaceuticals	Japan	
Mircera	Roche	Switzerland	
Negortire	Armstrong Laboratorios de Mexico	Mexico	
Nephrodil	Feron	India	
NeoRecormon	Roche	Switzerland	
Nespo	Dompe Biotec (withdrawn 2008)	Italy	
PDpoetin	Pooyesh Darou Pharmaceuticals	Iran	
Procrit	Amgen (for Centocor Ortho Biotech)	USA	
Pronivel	Elea	Argentina	
Relipoietin	Reliance Life Sciences	India	
Renogen	Biomedics	Philippines	
Repotin	BioClones	South Africa	
Retacrit	Hospira/STADA	Germany	
Shanpoietin	Shanta Biotechnics	India	
Silapo	Bioceuticals Arzneimittel AG	Germany	
Tinax	Eurofarma	Brazil	
Vintor	Emcure	India	
Wepox	Wockhardt	India	
Zyrop (see Epoyet)	Zydus Biogen	India	
* Adapted from [5.7-8].			

* Adapted from [5,7-8].

ANALYTE	ANALYSIS TECHNIQUE AND MODE / DETECTION	SPECIES DETECTED	REFERENCE
rhEPO	SE-HPLC / UV	Monomer, dimers and HMW aggregates	[54] [55] [56] [58] [59] [60] [61] [62] [65] [66] [107] [111]
rhEPO	SE-HPLC / Fluorescence	Monomer, dimers and HMW aggregates	[63] [64] [111]
rhEPO	RP-HPLC / UV	Monomer, aggregates and HSA	[55] [67]
rhEPO	RP-HPLC / UV	Monomer, aggregates, HSA and underglycosylated EPO	[68]
rhEPO	RP-HPLC / UV	EPO and underglycosylated EPO	[69]
rhEPO	RP-HPLC / TOF- MS	EPO	[73]
rhEPO	IEX-HPLC / UV	Isoforms	[108]

Table 2. Determination of intact ESAs by high resolution separation methods*

rhEPO	IEX-HPLC / Fluorescence	Isoforms	[70] [71] [72]
rhEPO	CZE / UV	Isoforms	[19] [23] [81] [82] [83] [107] [110] [111] [119]
rhEPO	CIEF / UV	Isoforms	[84]
rhEPO	CZE / TOF-MS	Glycoforms	[22] [79] [96] [98]
rhEPO	CZE / IT-MS	Glycoforms	[102]
rhEPO	CZE / IT-MS	Isoforms. MS spectra did not allow glycoforms characterization	[97]
rhEPO	CE immunoassay / Chemiluminescence	EPO concentration	[117]
DPO	SE-HPLC / UV	Monomer, dimers and HMW aggregates	[57]
DPO	RP-HPLC / TOF- MS	DPO	[73]
DPO	CZE / UV	Isoforms	[17]

DPO	CZE / IT-MS	Isoforms. Glycoforms were not characterized.	[102]
uhEPO	CZE / IT-MS	Isoforms. None of them was characterized based on MS spectra.	[97]
Endogenous EPO in human sera	CE immunoassay / Chemiluminescence	EPO concentration	[117]

* CE methods published before 2006 were reviewed on [74].

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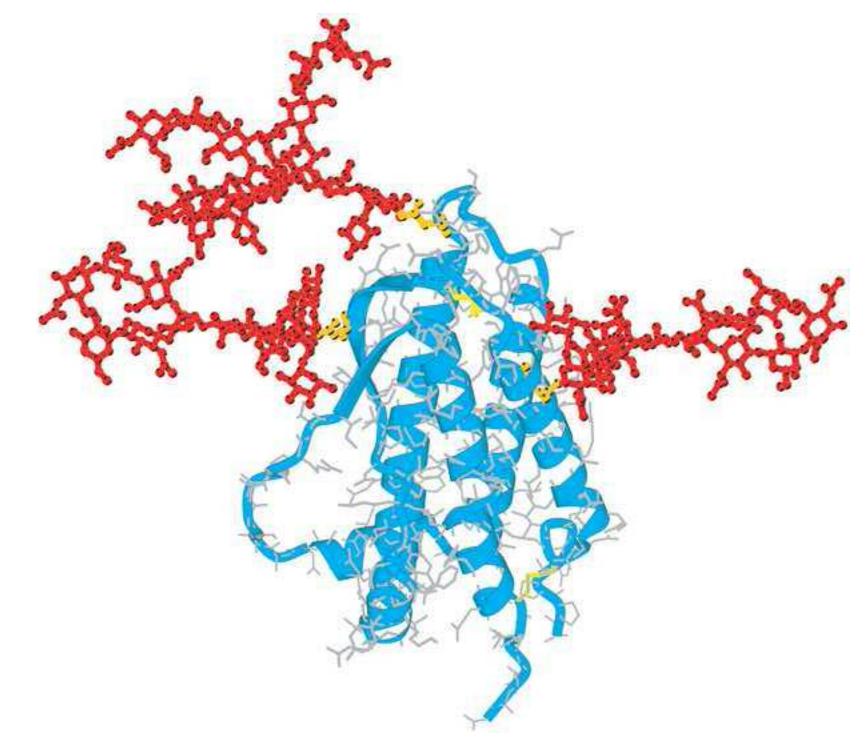
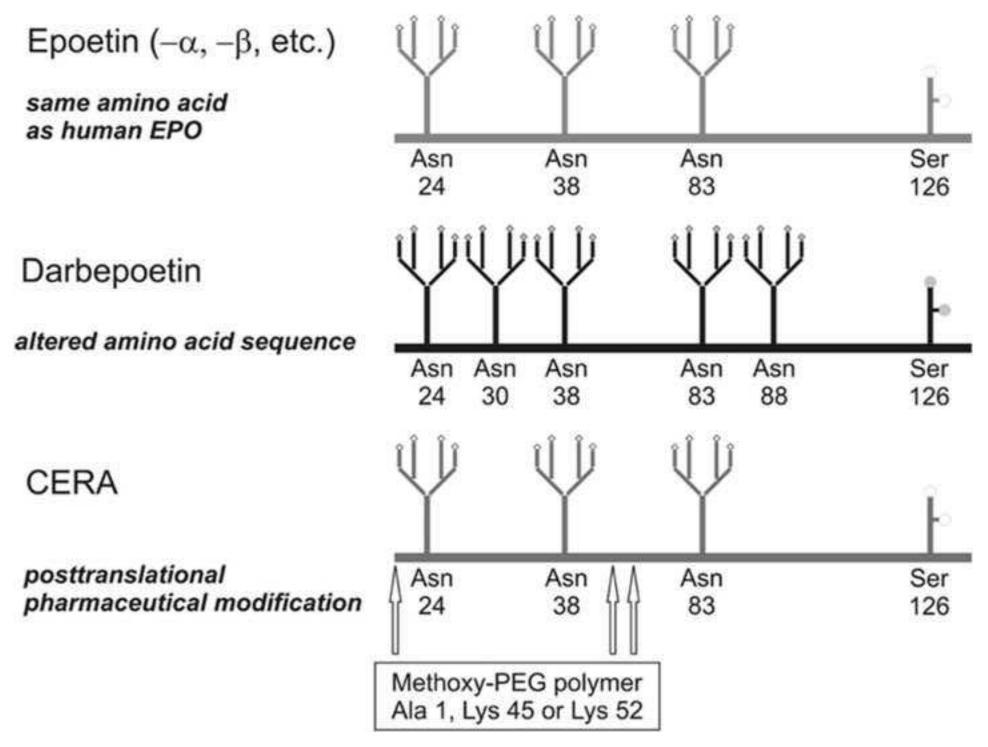
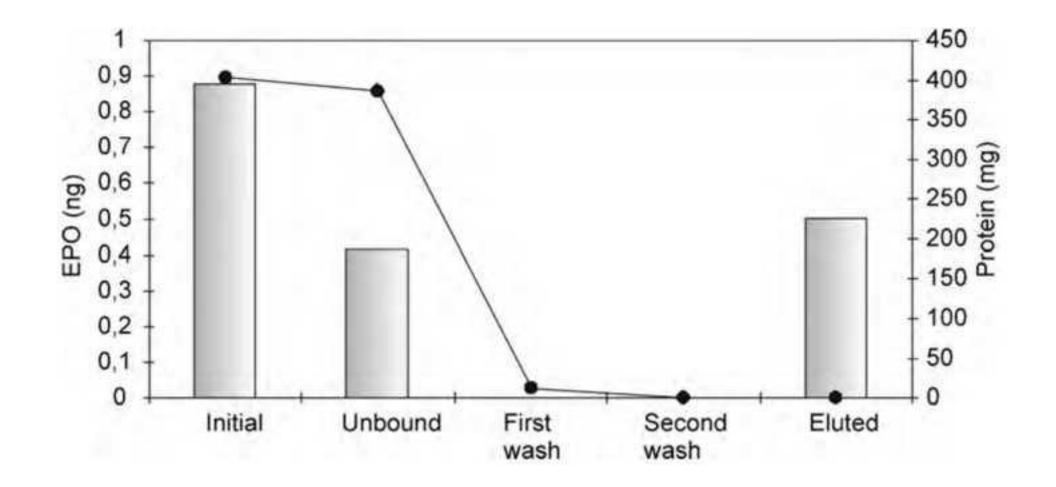


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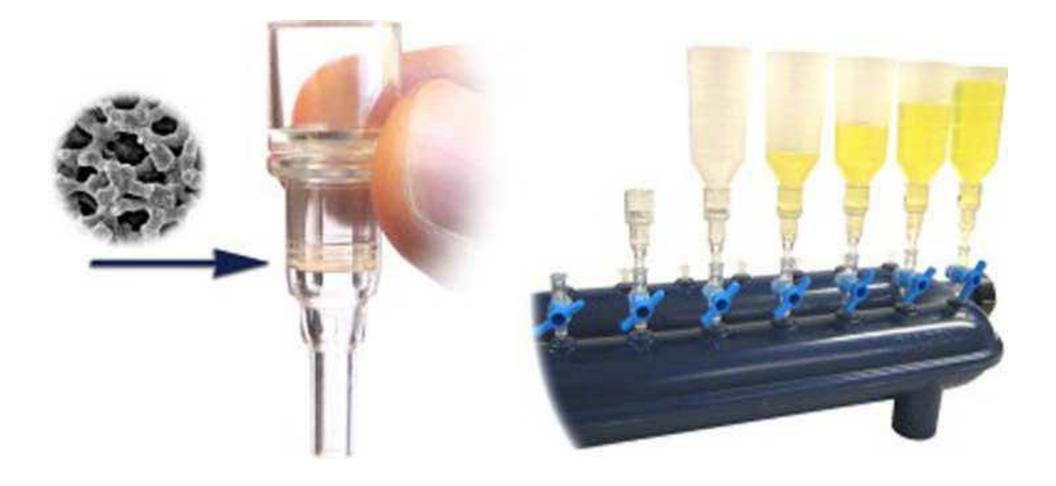


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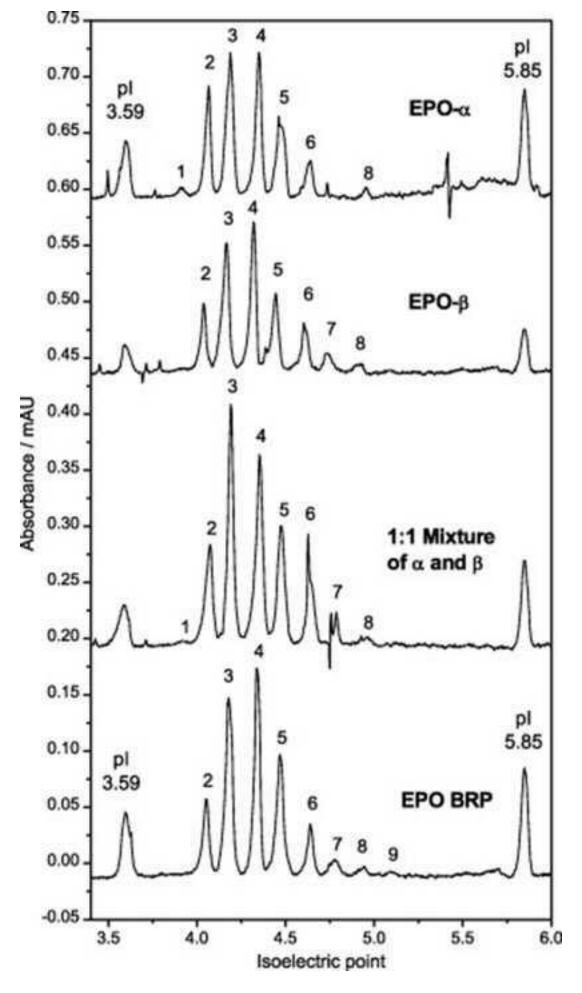


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