



DETERMINATION OF **PROTEIN PURITY** AND **HETEROGENEITY** BY CAPILLARY GEL ELECTROPHORESIS AND CAPILLARY ISOELECTRIC FOCUSING

INTRODUCTION

Antibodies, protein-based therapeutics, and other recombinant proteins are final products in biotechnological and pharmaceutical industry. Determination of their purity, stability, and heterogeneity is of utmost importance since post-translational modifications as well as degradation processes can change drastically the biological activity of these proteins.

MEASUREMENT METHOD

To determine the purity and heterogeneity of protein samples, two capillary-based methods are proposed – capillary gel electrophoresis (CGE) and capillary isoelectric focusing (cIEF). The former is applied when proteins differing in molecular mass must be resolved. The latter is applied when proteins or protein isoforms exhibit almost identical molecular masses, but differ in net charges.

In CGE, capillary is filled with a buffered polymer solution. Prior to analysis, protein sample is denatured in the presence of SDS by heating either under reducing (with reducing agent added) or non-reducing conditions. After rapid cooling the sample is injected, high voltage is applied, and proteins start migrating through polymer solution. Similar to SDS-gel electrophoresis in a slab, proteins are separated in a capillary according to their molecular masses due to a sieving effect of the capillary polymer solution. Proteins differing by as little as 4 % in molecular mass can be resolved.

In **cIEF**, the entire capillary is filled with a polymer-based ampholyte-sample mixture. At the first stage, named focusing, high voltage is applied, which results in a pH gradient formation and focusing of proteins in narrow zones according to their pl values (where their net charges are equal to zero). At the end of this stage there is almost no any movement in the capillary and current is about zero. During the second stage, named mobilization, the outlet vial is replaced with mobilization solution, high voltage is applied again, and focused zones start moving toward detector point. Depending on selected ampholytes, proteins and protein isoforms differing by as little as 0.04 pl unit can be resolved and quantified.

For both methods a linear regression exists, reflecting the dependence of log Mw of the proteins (for CGE) or their pl values (for cIEF) upon migration time. This is used to determine both parameters of the unknown proteins by adding the corresponding markers to the sample solution.

EQUIPMENT AND REAGENTS

The CAPEL capillary electrophoresis system is used in measurements. Data acquisition, collection, processing and output are performed using a personal computer running under WINDOWS® XP/7/8/10 operating system with installed dedicated software package ELFORUN.

All reagents must be of analytical grade or better. Appropriate kits of reagents for either CGE or cIEF are available and supplied according to the specified task.





Method: BGE: Capillary: Voltage: Temperature: Detection:	capillary gel electrophoresis special for proteins L _{eff} /L _{tot} 32/42 cm; ID 75 μm + 20 kV 25 °C 220 nm	- 38 mAU
Sample:	four different IgG preparations, analysed under reduced conditions	
Results:	peaks stemming from LC and HC are seen. Small differences in Mw can be detected: 2 kDa only in case of LC (bottom trace) and 5 kDa in case of HC (upper trace)	1 I
Method: BGE: Capillary: Voltage: Detection:	capillary isoelectric focusing special for cIEF coated L _{eff} /L _{tot} 32/42 cm; ID 50 μm + 25 kV 280 nm	pl 10.0 -24 mAU pl 9.5 8. 8.98
Sample:	recombinant IgG preparation	9.04
Results:	four protein isoforms with slightly different pl values were resolved. pl markers, added to the sample (shown in red), help to calculate pl values of protein isoforms (shown in blue)	15 20



Inter-day reproducibility Method: capillary isoelectric focusing

- **Sample:** protein-based therapeutic
- Results: Several isoforms, indicating charge heterogeneity were revealed from the sample, which had earlier appeared to be homogeneous on a slab gel. Sequence of runs was carried out within several days to evaluate reproducibility, which appeared to be on a highest level. Run #5 (shown in blue) was performed two days after Run #1 (shown in red).

-14 mAU run # 1 25 30 35 migration time, min

30

35 migration time, min

25

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