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QUANTITATIVE X-RAY MICROANALYSIS OF BIOLOGICAL CRYOSECTIONS DEPENDS ON ICE CRYSTAL DAMAGE

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Résumé - A l'aide de la microanalyse par rayons X sur des cryocoupes d'épaisseur de 100 nm de standards gélatine-glycérol et de foie de rat on constate que le rapport pic sur fond (p/b) décroît avec l'augmentation de la taille des cristaux de glace.

Abstract - It is shown by X-ray microanalysis of 100 nm thick cryosections from glycerol-gelatine standards and rat liver that the measured peak-to-background ratio (p/b) decreases with increasing ice crystal size.

INTRODUCTION

X-ray microanalysis of freeze-dried cryosections is used to measure the distribution of elements in different compartments of biological cells and tissues (1-5). Ice crystal damage of the ultrastructure due to insufficient freezing velocity is a well known preparation artefact of this method (6). However, the displacement of elements, particularly of diffusible ions as sodium, chlorine, potassium and calcium, is assumed to take place in dimensions not larger than one ice crystal diameter. Therefore X-ray microanalysis by scanning an electron beam in an area larger than a few ice crystal diameters in size was expected to be independent on freezing damage. The experiments described in the following contradict to this assumption.

MATERIALS AND METHODS

The cooling chain preparation method as described previously (7, 8) is used for the preparation of the cryosections. Two kinds of specimen were studied:

1. Droplets of 20% glycerol-gelatine and 80% electrolyte solution of known composition, e.g. KCl varying in concentration between 3.1 and 200 mMol/liter. These specimens were used as standards for quantitative X-ray microanalysis (9).

2. Freshly excised pieces of rat liver, about 1 mm in diameter. 100 nm thick cryosections, prepared by means of the Reichert FC4 cryoultramicrotome were transferred to the electron microscope under cold nitrogen gas atmosphere. Freeze-drying of the sections was enabled by evacuating the cryotransfer chamber and loosening the cold contact to the grid holder. X-ray microanalysis was performed in a Siemens Elmiskop ST 100 F, a scanning transmission electron microscope (STEM) with a field emission gun, operated at 100 kV, by means of an energy dispersive Sili-detector (nuclear semiconductor) and a multichannel analyzer (Link Systems). The scanning area is varied from 278 nm x 444 nm to 6940 nm x 11100 nm, analysis time was 100 s.
The sections were kept at 138 K in the electron microscope. For the evaluation of the X-ray peaks, p/b-values were calculated by dividing the peak height (p) through the mean X-ray intensity between 4.5 and 5.5 keV (b).

**RESULTS**

Fig. 1 shows a cryosection of glycerol-gelatine mixed with 200 mMol/l KCl with the mean ice crystal diameter less than 50 nm. A corresponding X-ray spectrum is added (Fig. 2). Fig. 3 was obtained after slower freezing the same specimen type as in Fig. 1. The mean ice crystal diameter is ca. 1/µm. Fig. 4 shows the corresponding X-ray spectrum. p/b-values for sulfur and potassium, derived from X-ray spectra of cryosections of glycerol-gelatine mixed with KCl are sketched in Fig. 5 and 6, respectively. The main result is that the p/b-value decreases with increasing ice crystal diameter. This holds for diffusible ions (e.g. potassium) as well as for bound elements (e.g. sulfur). Similar results are obtained in biological specimens. Fig. 7 shows rat liver with intermediate ice crystal size. The p/b-value of phosphorus depending on the ice crystal size in the cytoplasm and the nucleus is drawn in Fig. 8.
DISCUSSION

These results could be explained as follows: In biological material of low density a 100 keV-electron has a mean free path of about 100 nm. The mean distance between characteristic atoms homogeneously distributed with a volume density of 100 mMol/l = 62000 atoms in a cube of 100 nm length of side is about 2.5 nm. Projected to the section surface the interatomic distance as seen from the impinging electrons is 0.4 nm. After precipitation of ice crystals during cryofixation the density of the remaining material accumulated between the ice crystals is enhanced, and the interatomic distance between the characteristic atoms is reduced as schematically drawn in Fig. 9. Relatively small ice crystals as sketched in Fig. 9b result in small ion displacements,
whereas larger ice crystals compress the ions in material walls of enhanced density (Fig. 9c). As a consequence electrons hitting holes left after ice sublimation penetrate the section without interaction with the specimen. Electrons hitting the material walls are scattered preferably by the atoms in the upper part of the section, and the excitation probability of the atoms below is reduced. This effect results in reduced p/b-values depending on the ice crystal size.

Fig. 9 - Schematic drawing of ion displacements in cryosections due to ice crystal growth as seen from the side. Circles = ions, hatched areas = ice crystals. a) The ion distribution is assumed to be statistically homogeneous in a section without ice crystals. b) Small ice crystals cause small ion displacements. The ions are still homogeneously distributed. c) Large ice crystals accumulate the ions in narrow material walls. The ion distribution becomes inhomogeneous.

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