

These results are in agreement with the possibilities that, on the release of stored amines in brain, total amine levels, but not necessarily free amine levels, are lowered, and that the central effects of reserpine depend on the levels of free amine at the receptor sites (9). Because of the very rapid synthesis of serotonin (10) as compared to nor-epinephrine (11) in brain, the effects of free serotonin may predominate after reserpine action.

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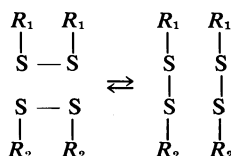
### Disulfide Interchange by Ionizing Radiation

**Abstract.** The irradiation of a solution of two symmetric disulfides produces detectable amounts of the nonsymmetric mixed disulfide. The effect is abolished by the addition of ethylmaleimide. The finding indicates that radiation causes a disulfide opening and recombining process which may be of radiobiological interest.

The sulfur groups present in the structures of the cell have generally been considered some of the loci most vulnerable to radiation. The nature of the changes induced by radiation in these groups is not yet completely understood. In considering the possible extension of studies of simple systems to more complex ones, chemical changes in sulfur compounds of low molecular weight resulting from radiation have been frequently investigated. Researches have been focused on the degradation of these compounds (1) or on the oxidation (2) and the reduction (2, 3) of their sulfur portion. The occurrence of a less destructive action of ionizing radiation on sulfur compounds—that is, disulfide interchange—is reported here; though less destructive, this phenomenon might equally effect a disorganiza-

tion of disulfide-containing substances.

Disulfide interchange was studied in a system composed of cystamine and N-diformylcystine. This system permitted the use of compounds soluble over a large pH range and of paper electrophoresis for the detection of the interchange product. As a matter of fact, cystamine and diformylcystine, having different dissociation, migrate in opposite directions, while the mixed disulfide produced by the interchange



( $R_1 = -CH_2-CH_2-NH_2$ ;

$R_2 = -CH_2-CH(NHCH)-COOH$ )

having an intermediate charge, migrates at an intermediate rate.

Four hundred micromoles of cystamine dihydrochloride (4) were dissolved in 4 ml of the same acetate buffer that was used for electrophoresis; 400  $\mu$ mole of N-diformylcystine was suspended in 4 ml of water and brought into solution by the addition of solid sodium carbonate. The two solutions were mixed (final pH, 3.8 to 4.0) and irradiated for a suitable length of time with a Philips 50-kv x-ray source at a distance of 1 cm. The magnitude of the radiation dose was checked by a ferrous sulfate dosimeter. At any desired time a 0.02-ml sample of the solution was spotted at the center of a strip of Whatman 3-mm filter paper. The paper was then wetted with a 0.45M sodium acetate buffer of pH 3.75 (ionic strength, 0.05). Electrophoresis was performed in the same buffer, with a potential of 300 volts (about 10 volt/cm of paper), for 30 minutes. After the paper was dried the compounds were located by spraying with the Folin-Marenzi reagent, with bisulfite added according to a procedure previously devised for the detection of disulfides (5).

The results are shown in Fig. 1; they indicate that under the action of radiation a new disulfide compound, which in electrophoresis does not migrate, is promptly produced in amounts related to the radiation dose. In order to identify the new compound, larger amounts of it were prepared by large-scale electrophoresis. As was to have been expected from the mixed disulfide formed by interchange between cystamine and diformylcystine, upon oxidation with  $H_2O_2$  and ammonium molybdate (6), followed by hydrolysis in 1N HCL for 2 hours at 100°C, the new compound yielded taurine and cysteic acid, which were detected by paper chromatography.

The radiation-induced disulfide exchange was observed also at neutral pH and with concentrations of the reactants other than those reported above. Be-

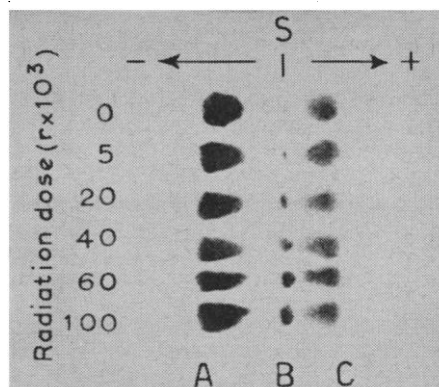


Fig. 1. Disulfide interchange by radiation. Paper electrophoresis of a solution of equimolar amounts of cystamine and diformylcystine after irradiation with increasingly large doses of x-rays. S, starting point; A, cystamine; C, N-diformylcystine; B, mixed disulfide. Spots were developed by spraying with the Folin-Marenzi reagent for disulfides.

cause of the slow spontaneous exchange (7), which tends to mask the results, the described conditions are the most suitable for a clear-cut demonstration of the radiation effect. The presence of an excess of N-ethylmaleimide abolishes the exchange; this indicates that the reaction probably proceeds through the temporary opening of the disulfide bonds with liberation of thiol groups. However, the nitroprusside test for thiol groups, carried out soon after irradiation of the solutions, was found invariably negative.

By slightly modifying the experimental conditions, so as to solubilize cystine (solution of the disulfides in final 0.2N HCl), the exchange can be observed also between cystine and cystamine.

Two conclusions should be drawn from the above results. First, the radiation-induced exchange can be expected to occur also within a single molecular species of disulfide. This effect of radiation, which is not followed by analytical change of the compound, might dissipate radiation energy, thus contributing to the reduction of the damaging action of radiation observed in the presence of disulfides. Second, present results, if extended to proteins, might provide another approach to the understanding of the disorganization of secondary and tertiary structures of disulfide-containing proteins brought about by radiation. In this connection, the disulfide interchange which has been reported to occur in the course of the chemical denaturation of proteins (8) and polypeptides (9) is highly suggestive (10).

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## Rh<sub>0</sub>(D) Genotype and Red Cell Rh<sub>0</sub>(D) Antigen Content

**Abstract.** The Rh<sub>0</sub>(D) content of red cells obtained from different individuals as determined with I<sup>131</sup> anti-Rh<sub>0</sub>(D) showed a bimodal distribution. Family studies indicate that the cells with the lower antigen content represent the heterozygous Rh<sub>0</sub>(D) state and that the cells with 1.6 times more Rh<sub>0</sub>(D) correspond to the homozygous state.

The blood group antigens on the stroma of the red cell represent specific immunochemical molecules whose synthesis is genetically determined (1). If the one gene-one product hypothesis applies to the blood group antigens, the red cell derived from a homozygous individual should have twice the antigen content found on a red cell derived from a heterozygous individual.

The Rh<sub>0</sub>(D) antigen content of the human red cell was determined with incomplete I<sup>131</sup> trace labeled anti-Rh<sub>0</sub>(D) and the methods described previously (2). The method of approach was to determine the frequency distribution of the Rh<sub>0</sub>(D) antigen content in a population of different Caucasian red cells (3). The heterozygous state in this distribution then was identified by study of genetically defined Rh<sub>0</sub>(D) heterozygous cells obtained from family studies. The use of this experimental approach was necessitated because of the inability to differentiate in advance of the experiment the heterozygous from the homozygous Rh<sub>0</sub>(D) red cell by conventional techniques (4).

Red cells were obtained by selection from the donor population entering the Blood Bank (5) and were typed for the ABO and Rh antigens with commercially obtained antisera. Only a few Rh<sub>0</sub>(D) negative red cells were used as controls. The red cells from 199 indi-

viduals were studied, but only the results obtained on rh'(C) negative, Rh<sub>0</sub>(D) positive cells are presented in this report. The results are expressed as quantity of antibody nitrogen bound to 0.01 ml of centrifuged red cells as determined by the microhematocrit technique (6) using the International microhematocrit centrifuge (5 minutes at  $g_{max} = 12,000$ ). The mean number of red cells in 0.01 ml of centrifuged red cells was determined for a number of cell suspensions with the Coulter electronic counter (7, 8) and was found to equal  $9.01 \pm 1.48 \times 10^7$  cells (9).

The over-all precision of the technique was evaluated by the following experiment. Red cells from a given individual were stored in Alsever's solution at 4°C and on subsequent days an aliquot was reconstituted and reacted with the I<sup>131</sup> anti-Rh<sub>0</sub>(D). A fresh specimen of cells obtained from the same individual was run in parallel with the stored cells. The mean and standard error of the mean for repeated determinations on the stored cells over a period of 11 days was  $1.85 \pm 0.13$ ,  $10^{-2}$   $\mu$ g of N per 0.01 ml of red blood cells. The corresponding value for fresh unstored red cells was  $1.85 \pm 0.15$ ,  $10^{-2}$   $\mu$ g of N. There was no significant difference between these two values ( $t = 0.18$ ,  $n = 14$ ). Rh<sub>0</sub>(D) negative red cells took up less than 3 to 7 percent of the I<sup>131</sup> bound to Rh<sub>0</sub>(D) positive red cells. The amount of nitrogen bound was independent of both the Rh and ABO phenotype of the Rh<sub>0</sub>(D) negative cell. All the red cell suspensions were reacted in an excess of antibody in order to obtain maximum saturation of the Rh<sub>0</sub>(D) antigen sites. Antibody excess was determined by demonstrating free I<sup>131</sup> anti-Rh<sub>0</sub>(D) in the supernatants after reaction and by re-reacting the I<sup>131</sup>-sensitized red cells with additional I<sup>131</sup> anti-Rh<sub>0</sub>(D) to show that the available antigen sites were saturated. There was less than 10 percent increase in the amount of antibody bound to the sensitized red cell after a second 60-minute incubation with the I<sup>131</sup> anti-Rh<sub>0</sub>(D).

The absolute frequency of the antibody nitrogen bound to 47 Rh<sub>0</sub>(D) positive, rh'(C) negative cells is shown in Fig. 1. It can be seen that the cells segregate into two groups, one with a mean of 2.44 and the other with a mean of 3.95 ( $10^{-2}$   $\mu$ g of N per 0.01 ml of red blood cells). The cells in the higher nitrogen value group take up 1.62 times more antibody nitrogen than do the cells in the lower group. Known heterozygous Rh<sub>0</sub>(D) positive red cells obtained from families with a history of Rh hemolytic diseases of the newborn (10) were employed to identify the heterozygous value in this bimodal distribution. Eight genetically determined

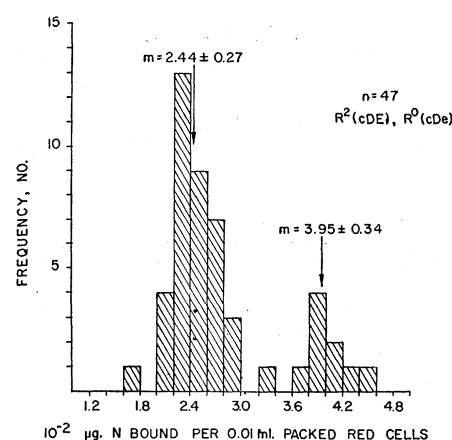


Fig. 1. Distribution of I<sup>131</sup> anti-Rh<sub>0</sub>(D) nitrogen bound to 47 different Rh<sub>0</sub>(D) positive, rh'(C) negative red cells. The values shown are in units of  $10^{-2}$   $\mu$ g of nitrogen per 0.01 ml of centrifuged red cells with the mean and standard error of the mean shown for each group.

heterozygous Rh<sub>0</sub>(D) positive red cells had values which ranged from 1.47 to 2.48 ( $10^{-2}$   $\mu$ g of N per 0.01 ml of red blood cells). These results indicate that the 2.44 antibody nitrogen peak represents the heterozygous state [Rh<sub>0</sub>(D), rh(d)] and that the 3.95 value corresponds to the homozygous state [Rh<sub>0</sub>(D), Rh<sub>0</sub>(D)]. If certain assumptions are granted, namely uniformity of iodination of the globulins in the gamma globulin fraction and a molecular weight of  $1.6 \times 10^5$  for the anti-Rh<sub>0</sub>(D), then the heterozygous Rh<sub>0</sub>(D) red cell contains about 6400 Rh<sub>0</sub>(D) antigen sites per cell and the homozygous cell about 10,300 per cell (11).

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9. Mean and standard error of the mean are used throughout this report.
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