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   There is probably very little loss of T<sub>3</sub> from nuclei during the isolation procedure, which takes 1 hour at 0°C. Nuclei isolated by this method stably retain added T<sub>3</sub> for 5 hours at 22°C [R. C. Jaffee and A. R. Means, Endocrinology 101, 447 (1977)]. The scubilized recentor from these nuclei for the second stable of the second Jaffee and A. R. Means, Endocrinology 101, 447 (1977)]. The solubilized receptor from these nuclei retains added T<sub>3</sub> with little loss over a period of 20 hours at 0°C (Fig. 2).
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- The efficiency of extraction was not affected by the ratio of extracting buffer to nuclei. Por-14. tions of nuclei from a single control preparation were extracted with one-half to twice the usual ratio of buffer to tissue weight. The extract was diluted so that all preparations should have had the same receptor concentration if extraction had been equally efficient. The ratios of bound to free [ $^{125}$ I]T<sub>3</sub> in these extracts were similar, indicating that within the range observed with method A, differences in the ratio of extracting buffer to nuclei did not affect the efficiency of traction. Direct measurement of bound/free  $[^{125}I]T_3$  in the more concentrated extracts revealed that when bound/free ratios were high in-creases in them did not fully reflect increases in the estimated receptor content. With method A (extracting volume constant) this could have had a small tendency to exaggerate the difference be tween extracts from control and fasted rats. In contrast, with method B (extracting volume proportional to DNA), the same lack of proportion-ality between the bound/free ratio and the esti-mated receptor content would have diminished the apparent difference between extracts from control and fasted rats. The excellent agreement between the results obtained with methods A and B indicates that the effect of any lack of proportionality between the bound/free ratio and re-15
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- Preliminary studies in the mouse show that fasting sufficient to decrease hepatic nuclear  $T_3$  receptors does not decrease brain  $T_3$  receptor caacity
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- 20. donation of T<sub>3</sub> immunoassay kits

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## **Covalent Immunoglobulin Assembly in vitro: Reactivity of Light** Chain Covalent Dimers (L<sub>2</sub>) and Blocked Light Chain Monomers

Abstract. Covalent light chain dimers  $(L_2)$  and cysteine-blocked L chain monomers readily react with partially reduced heavy (H) chains. A rapid disappearance of these blocked L chain species is followed by the appearance of covalent intermediates HL,  $H_2$ , and  $H_2L$ —leading to fully assembled  $H_2L_2$ . The mechanism of initial disulfide bond formation between heavy and light chains is disulfide interchange.

In a study of the factors controlling antibody assembly, we have investigated in vitro the covalent assembly of a human IgG1 $\kappa$  immunoglobulin (Fro) (1). The quaternary structure of proteins of this immunoglobulin class is maintained by strong noncovalent forces as well as by four covalent interchain disulfide bonds. Two of these link the heavy (H) chains to each other, and two bridge heavy to light (L) chains, utilizing the carboxy terminal light chain residues and residue 220 in each heavy chain (2).

The interchain disulfides can be reduced without detectable reduction of any of the intrachain bonds (3, 4). Our experimental procedure, accordingly, entails selective (partial) reduction of the four interchain disulfide bonds, followed in certain experiments by reoxidation without prior disruption of the noncovalently associated tetramer (3) and in others by reoxidation after the H and L chains are separated and then recombined in varying molar ratios (4). The last type of experiment permits study of the assembly and reoxidation processes when L chains are modified, or present in excess, as is often the case both in normal lymphoid cells (5, 6) and in mouse myeloma tumors or cloned cell lines (7).

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Methods have been devised for the quantitative determination of assembly intermediates during reduction as well as reoxidation (3, 8), so that not only initial and final states, but the kinetics of conversion from one to the other can be analyzed. Moreover, a general theory has been formulated to compare assembly in vitro with reported intracellular pathways and kinetics of assembly (9).

Among the key experimental findings with the IgG1 $\kappa$  studied are (i) that the rates and pathways of covalent assembly are not significantly affected by whether or not the H and L chains are separated from one another prior to reoxidation at neutral pH, and (ii) that large molar excesses of L over H chains do not markedly affect the pathways compared to equimolar reoxidations, although excesses diminish the rate of covalent  $H_2L_2$ assembly. At the same time, excess L chains retard the formation of insoluble H chain aggregates in vitro, and this suggests a comparable role for such excesses in vivo (4).

The theoretical analysis revealed that the reoxidation process is not random; the four interchain disulfide bonds form with unequal probability throughout the course of the reaction. Initially, HL bonds form about twice as rapidly as HH bonds, but the formation of the second HL bond in the molecule is not favored over HH bond formation. In other words, the existence of one HL bond in a noncovalently assembled tetramer slows the rate at which the second forms.

Preliminary results have also established that covalent L<sub>2</sub> dimers rapidly react with reduced H chains or H<sub>2</sub> dimers and generate intermediates that assemble normally into H<sub>2</sub>L<sub>2</sub> tetramers, a finding unanticipated by any previous studies on immunoglobulins (10, 11). Our report further documents this result and demonstrates also the reaction of partially reduced H chains with a stable L chain monomer whose carboxy terminal cysteine residue is in mixed disulfide linkage with the amino acid L-cysteine (12).

These L chain forms are derived from the urinary Bence Jones protein. Chromatography on Sephadex G-100 yields two fractions, a covalent L chain dimer and a stable L chain monomer. These forms are not interconvertible, and neither contains detectable free sulfhydryl (12). Procedures for preparation and purification of the plasma monoclonal protein (Fro) and of the partially reduced H chains have been published (3, 4), as have all of the other analytical procedures employed in this study (3, 4, 11). For recombination experiments, reduced H chains and unreduced dimeric or

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blocked monomeric L chains were mixed under nitrogen in equimolar quantities at pH 3.2, in 10 mM acetic acid. Reoxidation conditions were established by exposing the mixture to air, and raising the pH to 7.5 and the ionic strength to 0.14. At various times during the reoxidation, portions were removed and reacted with excess iodoacetamide. These alkylated samples were then subjected to electrophoresis on polyacrylamide gel in the presence of sodium dodecyl sulfate (SDS). Reactants (H,L,L<sub>2</sub>), product  $(H_2L_2)$ , and the three covalent intermediates-HL, H<sub>2</sub>, H<sub>2</sub>L-were identified by their migration distances relative to the unreduced protein and partially reduced H and L chains (3, 4).

Figure 1 shows the gels from an experiment in which partially reduced H chains were reacted with the Bence Jones covalent dimer containing a small amount (less than 10 percent) of blocked L monomer (well 2). By the first reoxidation time point (well 3)  $L_2$  disappeared and intermediates leading to H<sub>2</sub>L<sub>2</sub> were observed, with noticeable increases in L, HL, and H<sub>2</sub>L. Control experiments, aimed at eliminating the possibility that thiols of low molecular weight were involved, established that the isolated  $L_2$ fraction, when maintained under reaction conditions for 1 hour or longer. ran true on gels and molecular sieve columns, as well as in sedimentation experiments. Moreover, Gally and Edelman (10), in an early investigation of L chain dimers, showed that alkylated heavy chains are not reactive with  $L_2$ .

These intermediates, therefore, arise by two disulfide interchange reactions:

$$H + L_2 \rightarrow HL + L \tag{1}$$

$$H_2 + L_2 \rightarrow H_2 L + L$$
 (2)

After these initial interchange reactions, the reoxidation closely resembles other experiments with unblocked, reduced L chain preparations. Quantitative densitometry of the gels (4) showed that the pathways are comparable, as inferred from the maximum intensities achieved by the intermediates, and from the positions of gel band maxima on an abscissa representing the average number of disulfide bonds formed at any point during the reoxidation (3). In addition, the kinetics of the assembly process agree with those in other 1:1 recombinations, as judged by the half-times of sulfhydryl disappearance.

In other experiments with covalent  $L_2$  dimers, the H chain fraction was largely in the form of  $H_2$ . In such cases, reaction 2 predominates, as indicated by a high initial level of  $H_2L$ .

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Partially reduced H chain was also combined with the Bence Jones mixed disulfide monomer, as shown in Fig. 2. At the first time point (well 4) there was an increase in HL, in accord with release of cysteine from the mixed disulfide:

# H + L-S-S-cysteine $\rightarrow HL + cysteine$ (3)

Again, gel analysis revealed that once HL is formed, the assembly proceeds with the same pathways. A distinctive feature of this reoxidation, however, is that at the final time point (well 11) the  $H_2L$  level is still high. Moreover, the rate of sulfhydryl disappearance is about 20 percent slower than in experiments using free, reduced L chain.

Intracellular covalent  $L_2$  dimers are observed in mouse myeloma tumors which synthesize excess L chains (5). Although it has been argued that  $L_2$  is unlikely to be an assembly intermediate (13), the present experiments demonstrate that these forms of L chain, as well as monomers in mixed disulfide linkage

Fig. 1 (top). The reoxidation of partially reduced heavy (H) chain with light chain covalent dimer  $(L_2)$  shown on SDSpolyacrylamide gel. Wells 1 and 2 represent the purified H chain and Bence Jones (BJ) covalent L chain dimer, respectively, after Bio-Gel P2 chromatography. The H and L<sub>2</sub> fractions were combined under N<sub>2</sub> at an equimolar ratio in 10 mM acetic acid (final ratio of total L chain to H chain = 1.36). The pH of the mixture was then raised to 7.5 and the mixture was exposed to air. The final concentration of  $H_2L_2$  was 1.6  $\mu M$ . Wells 3 to 11 represent portions from the reoxidation mixture which were immediately alkylated with 1.0M iodoacetamide at the times indicated. All gel samples contained 1 percent SDS and iodoaceta-Fig. 2 (bottom). The mide. reoxidation of partially reduced H chain with cysteineblocked light chain monomer shown on SDS-poly-(L)acrylamide gel. Wells 1 and 2 represent the purified H chain and Bence Jones monomer fractions which were combined at an equimolar ratio as in Fig. 1. However, here the final ratio of L chain to H chain was 1.14, and the concentration of  $H_2L_2$  was 2.0  $\mu M$ . Well 3 represents a portion from the mixture that was alkylated just prior to the pH being raised above 7.5. Wells 4 to 11 represent reoxidation samples that were alkylated at the times indicated.

with cysteine (or, possibly, other lowmolecular-weight thiols), are reactive with H chain or H<sub>2</sub> dimers, and lead to productive intermediates for covalent tetramer assembly. Furthermore, the initial interchange reactions are rapid compared to the remaining covalent bond formations and do not depress the overall assembly rates, except in the case of the blocked monomer. Kazin (12) has shown that the interchange reactions actually occur in at least two stages. An initial rapid reaction results in the disappearance of about one-half of  $L_2$  dimer or blocked L monomer within 1.5 minutes. A second stage leading to 90 percent or greater disappearance requires 30 to 40 minutes for the dimer and as long as 90 minutes for the blocked L chain monomer.

More detailed studies of both noncovalent and covalent aspects of these interactions are required before it can be decided what role disulfide interchange may play in the intracellular assembly process. There have been suggestions





that enzymes may be involved in intracellular disulfide interchange or in catalyzing formation of correct disulfide bonds in immunoglobulins, but there is no conclusive evidence of such intracellular activity (14). It seems clear, however, that the blocked L forms discussed in this work, and possibly a variety of other nonproductive intermediates in assembly, can be readily converted to molecular species on the main pathways without enzymatic participation. The apparently small kinetic barriers thus may allow thermodynamic control over the final covalent assembly steps.

In this last connection, the blocked L monomer experiment is interesting and exceptional in that an incompletely assembled intermediate, H<sub>2</sub>L, remains in high excess after very long times. A possible explanation is that the released cysteine forms a stable mixed disulfide with  $H_2L$ . It is more likely, however, that H<sub>o</sub>L is not as reactive as H with L-cvsteine. This provides an analogy with the reoxidation system using free L chains in which, as noted above, the second HL bond is formed more slowly than the first. In this view, the driving force for these reactions is the establishment of forms which yield maximal concentrations of tetramer, even if not all disulfide bonds are formed within the tetramer, and the last bond to form, in this case between H<sub>2</sub>L and L, may add only marginal stability. The somewhat diminished rate of sulfhydryl disappearance in this experiment is then probably due to the slow rate of oxidation of the cysteine released in reaction 3.

It is hoped that experiments of this kind, coupled with studies of incomplete assembly with the variable half of the L chain  $(V_L)$  instead of whole L chain (15), will lead to a more complete understanding of assembly defects and may also allow controlled preparation of various intra- and interspecies hybrids.

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- The nomenclature used throughout is that recommended by the World Health Organization.
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### **DNA Synthesis in Cartilage Cells Is Stimulated** by Oscillating Electric Fields

Abstract. External oscillating electric fields (1166 volts per centimeter, 5 hertz) enhanced the incorporation of  $[^{3}H]$ thymidine into the DNA of chondrocytes isolated from the proliferative layer of embryonic (16 days) chick epiphysis. Verapamil or tetrodotoxin at 10<sup>-6</sup>M concentrations completely blocked the electric field effect. Tetracaine reduced the incorporation of [<sup>3</sup>H]thymidine in both control and electrically stimulated cells. The findings support the hypothesis that Na<sup>+</sup> and Ca<sup>2+</sup> fluxes generated by the electrical perturbation trigger DNA synthesis in these cells.

Electrical stimuli, applied in various ways, promote appositional and longitudinal bone growth in birds (1), limb regeneration in amphibia (2), and fracture repair in dogs, rabbits, and humans (3). Electric fields (EF's) stimulate proline uptake in cultured fibroblasts (4) and cause dedifferentiation in amphibian erythrocytes (5). Investigation of the mechanism of these effects can contribute to the understanding of epigenetic control of growth and differentiation. In this study we report that oscillating EF's stimulate the incorporation of [<sup>3</sup>H]thymidine into the DNA of cartilage cells in suspension, and that this effect can be blocked by inhibitors of calcium or sodium fluxes.

Cartilage cells were isolated from the proliferative zone of tibia epiphyses from 16-day chick embryos (6). The cells were incubated for 6 hours in a defined medium at 37°C in the presence of [<sup>3</sup>H]thymidine and were exposed to a pulsed d-c external EF of 1166 volt/cm, oscillating at 5 hertz. The EF was chosen on the basis of its inhibitory effect on adenosine 3',5'-monophosphate (cyclic AMP) accumulation (7), a condition conducive to cell proliferation in certain systems (8). Under these conditions the EF invariably stimulated [3H]thymidine incorporation, by  $53 \pm 13$  percent (mean ± standard error, 72 experiments, 216 samples and matched controls, range: 24 to 120 percent). The EF-enhanced incorporation of [3H]thymidine occurred in material that could be digested with deoxyribonuclease, whereas the labeling of the material that could be extracted with trichloroacetic acid was unaffected. The EF effect was first detected after 6 hours of incubation. The lag period suggests that the EF acts on a regulatory step in the cell cycle which precedes DNA synthesis. We also found an increase in DNA synthesis 4 to 6 hours after a brief electrical stimulation (15 minutes).

Skin fibroblasts obtained by collagenase digestion from 10-day-old chick embryos and treated in the same way as the chondrocytes were not stimulated by the EF. Under the same experimental conditions, the EF failed to affect [3H]thymidine incorporation into lymphocytes from rat spleens. However, in bone cells isolated from calvaria of 19-day rat embryos (9). EF's enhanced [3H]thymidine incorporation by  $27 \pm 1.2$ percent (three experiments, 15 samples). Therefore, under the experimental conditions used, the effect appears to be tissue specific.

In formulating a hypothesis for the mechanism of this phenomenon, we took into consideration the similarity of the effects of mechanical and electrical stimuli on bone growth (1, 3) and on cellular cyclic AMP (6, 7, 10) and current theories on the control of cell proliferation (11). Previously, we showed that mechanical stimuli capable of influencing bone growth reduced cellular cyclic AMP in epiphyseal chondrocytes through enhancement of calcium uptake and inhibition of adenyl cyclase (7). Modulation of ion fluxes and cyclic AMP levels are assumed (12) to be general modes of membrane-mediated cellular communication, which reached evolu-

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