

## STUDIES ON THE AFFINITY OF CHICK EMBRYO DNA LIGASE FOR RIBONUCLEOTIDES AND DEOXYRIBONUCLEOTIDES

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### 1. Introduction

We have recently described a five-step purification and the properties of a soluble chick embryo DNA ligase (EC 6.5.1.1) [1]. This enzyme, like the other previously described enzymes from T<sub>4</sub> and T<sub>7</sub> infected *Escherichia coli* and all the eukaryotic DNA ligases [2–8], has been shown to use ATP as cofactor [1]. Moreover, chick embryo DNA ligase was inhibited, with low  $K_i$  ( $4 \times 10^{-6}$  M), by dATP and this observation has made possible a single step purification of the enzyme [9]. The mechanism of this inhibition and binding is, however, still unclear and we decided to study it using equilibrium dialysis.

### 2. Materials and methods

Electrophoretically pure DNA ligase was obtained as already described, mol. wt 165 000 [1]. Radioactive ribonucleotides, [2-<sup>3</sup>H]adenosine monophosphate (18 Ci mmol<sup>-1</sup>), [2-<sup>3</sup>H]adenosine diphosphate (15 Ci mmol<sup>-1</sup>), and [2-<sup>3</sup>H]adenosine (25 Ci mmol<sup>-1</sup>), were obtained from Amersham Radiochemical Centre (Amersham, England). Radioactive deoxyribonucleotides, [8-<sup>3</sup>H]deoxyadenosine monophosphate (4.6 mCi mmol<sup>-1</sup>) and deoxyadenosine diphosphate (5 mCi mmol<sup>-1</sup>) were from Amersham Radiochemical Centre. [<sup>3</sup>H-G]Deoxyadenosine (20 Ci mmol<sup>-1</sup>), [2,8-<sup>3</sup>H]adenylylimidodiphosphate ( $\alpha,\beta$ ) (7.5 mCi mmol<sup>-1</sup>) were from New England Nuclear Corporation. [2,8-<sup>3</sup>H]Deoxyadenylylimidodiphosphate ( $\alpha,\beta$ ) was a gift from New England Nuclear Corporation

and had a specific activity of 1.1 mCi mmol<sup>-1</sup>. The corresponding non-radioactive compounds were all purchased from Sigma with the exception of the ATP analogs and dATP analogs, which were gifts from Dr Forterre (IRBM, Paris).

In the equilibrium dialysis experiments, the enzyme in 50 mM Tris-HCl buffer, pH 7.6, was dialysed against the ribonucleotides or deoxyribonucleotides at various concentrations in the same buffer. The method used followed that described by Koshland et al. [10]. At the start of dialysis, the first compartment (1 ml) contained  $10^{-6}$  M to  $5 \times 10^{-5}$  M non radioactive nucleotides plus 1 nmol corresponding radioactive compound. The second compartment (1 ml) contained the enzyme at a concentration equivalent to that of the nucleotide in the first compartment.

To establish that equilibrium had been achieved in 24 h at 4°C a control was run with Tris-HCl buffer alone instead of the enzyme solution. After 24 h the radioactivity in both compartments was found to be equal, within the experimental error (5%).

The radioactivity was counted on 100  $\mu$ l aliquots from each compartment. The solutions were spotted onto Millipore filters, completely desiccated and the radioactivity measured in a liquid scintillation counter.

Thin-layer chromatography of the dialyzed material showed that both in the presence and in the absence of enzyme there was no significant degradation of nucleotides and/or ATP or dATP analogs.

### 3. Results and discussion

A typical Scatchard plot is shown in fig.1. The

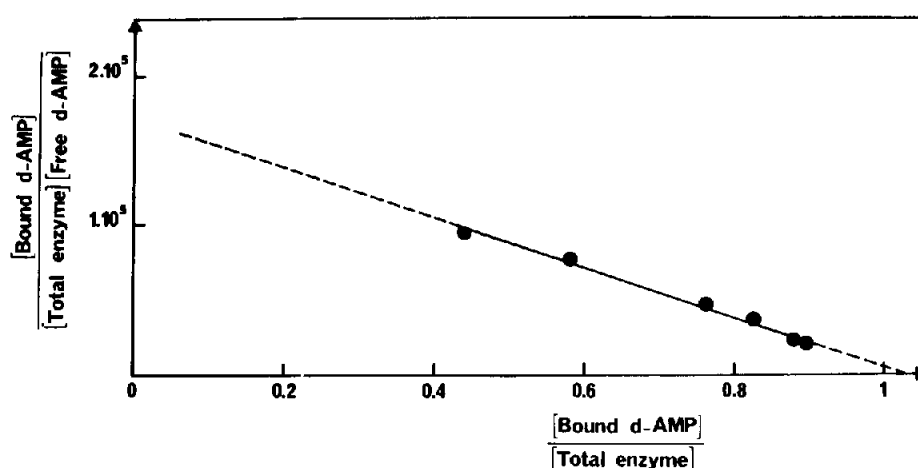


Fig.1. Scatchard plot of equilibrium dialysis data at 4°C for [<sup>3</sup>H]deoxyadenosine monophosphate-DNA ligase system. The medium contained 50 mM Tris-HCl buffer, pH 7.6;  $5 \times 10^{-6}$  M DNA ligase,  $10^{-6}$  M to  $5 \times 10^{-5}$  deoxyadenosine monophosphate. Represented in ordinate is the ratio of bound dAMP to total enzyme multiplied by free dAMP. Represented in abscissa is the ratio of bound dAMP to total enzyme concentration.

intercept on the abscissa gives the number of binding sites.

In this experiment the enzyme concentration was  $5 \times 10^{-6}$  M and the intercept corresponds to one binding site per enzyme molecule. From the slope of the Scatchard plot, the dissociation constant of the DNA ligase-dAMP complex was calculated to be  $7 \times 10^{-6}$  M. Similar experiments were done with other ribonucleotides, deoxyribonucleotides and analogs. The results are shown in table 1. For all the compounds studied only one binding site was found per enzyme molecule. For the ribonucleotides the  $K_d$  values ranges from  $9 \times 10^{-5}$  M to  $4 \times 10^{-5}$  M as

the number of phosphate groups in the ligand increased. For the deoxyribonucleotides the  $K_d$  values were from  $8 \times 10^{-6}$  M to  $4.7 \times 10^{-6}$  M as the number of phosphates increased. The corresponding  $\Delta G'_0$  values do not differ greatly with increasing numbers of phosphate residues.

To elucidate the role of the phosphate moiety in the binding equilibrium dialysis was also done in the presence of pyrophosphate. As shown in fig.2, there was no change in the apparent number of binding sites in the presence of  $10^{-5}$  M,  $10^{-4}$  M or  $10^{-3}$  M sodium pyrophosphate. Only a slight increase in the  $K_d$  value (from  $8 \times 10^{-6}$  M to  $1.2 \times 10^{-5}$  M for

Table 1  
Dissociation constants and  $\Delta G'_0$  values of DNA ligase for nucleotides and deoxynucleotides

Nucleotides	$K_d$	$\Delta G'_0$ cal mol <sup>-1</sup>	Deoxynucleotides	$K_d$	$\Delta G'_0$ cal.mol <sup>-1</sup>
Adenosine	$9 \times 10^{-5}$ M	14 636	d-Adenosine	$0.8 \times 10^{-5}$ M	6755
Adenosine monophosphate	$7.5 \times 10^{-5}$ M	14 036	d-Adenosine monophosphate	$0.7 \times 10^{-5}$ M	6305
Adenosine diphosphate	$5 \times 10^{-5}$ M	12 685	d-Adenosine diphosphate	$0.6 \times 10^{-5}$ M	5779
Adenylyl $\alpha,\beta$ imidodiphosphate	$4 \times 10^{-5}$ M	12 009	d-Adenylyl $\alpha,\beta$ imidodiphosphate	$0.47 \times 10^{-5}$ M	5253

Values are for 4°C

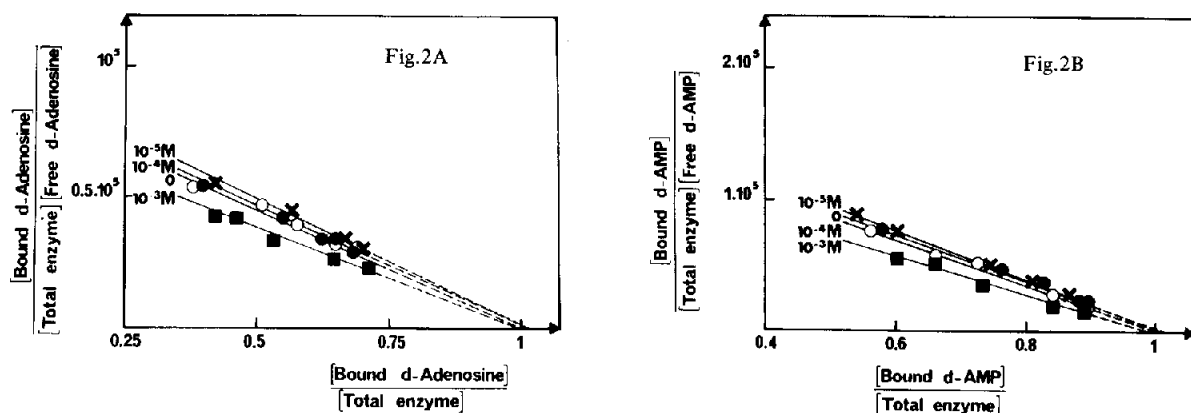


Fig.2A. Scatchard plot of equilibrium dialysis data at 4°C for [ $^3\text{H}$ ]deoxyadenosine–DNA ligase system. The medium contained 50 mM Tris–HCl buffer pH 7.6;  $10^{-5}$  M DNA ligase and  $10^{-6}$  M to  $5 \times 10^{-5}$  M deoxyadenosine. Represented in ordinate is the ratio of bound d-adenosine to total enzyme multiplied by free d-adenosine. Represented in abscissa is the ratio of bound d-adenosine to total enzyme concentration. This experiment in the presence of  $10^{-5}$  M,  $10^{-4}$  M and  $10^{-3}$  M pyrophosphate. Fig.2B. Scatchard plot of equilibrium dialysis data at 4°C for [ $^3\text{H}$ ]deoxyadenosine monophosphate–DNA ligase system. The medium contained 50 mM Tris–HCl buffer, pH 7.6;  $5 \times 10^{-6}$  M DNA ligase and  $10^{-6}$  M to  $5 \times 10^{-5}$  M deoxyadenosine monophosphate. Represented in ordinate is the ratio of bound dAMP to total enzyme multiplied by free dAMP. Represented in abscissa is the ratio of bound dAMP to total enzyme concentration. This experiment in the presence of  $10^{-5}$  M,  $10^{-4}$  M and  $10^{-3}$  M pyrophosphate.

deoxyadenosine and  $7 \times 10^{-6}$  M to  $10^{-5}$  M for dAMP) was observed at  $10^{-3}$  M pyrophosphate. This result is in good agreement with the observed inhibition of chick embryo DNA ligase activity (30% decrease) in the presence of  $10^{-3}$  M sodium pyrophosphate [1]. Similar results were obtained with dAMP in the presence of sodium pyrophosphate. These results are consistent with the hypothesis that the ligase binds to the adenosine moiety. This is similar to the conclusions from the observations made with phage T<sub>4</sub> ligase by Harvey et al. [11]. Further, the existence of only one binding site and the very small increase in affinity of the enzyme for the mono-, di- and triphosphonucleotides indicate that the enzyme reacts with the nucleoside part of the molecule.

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