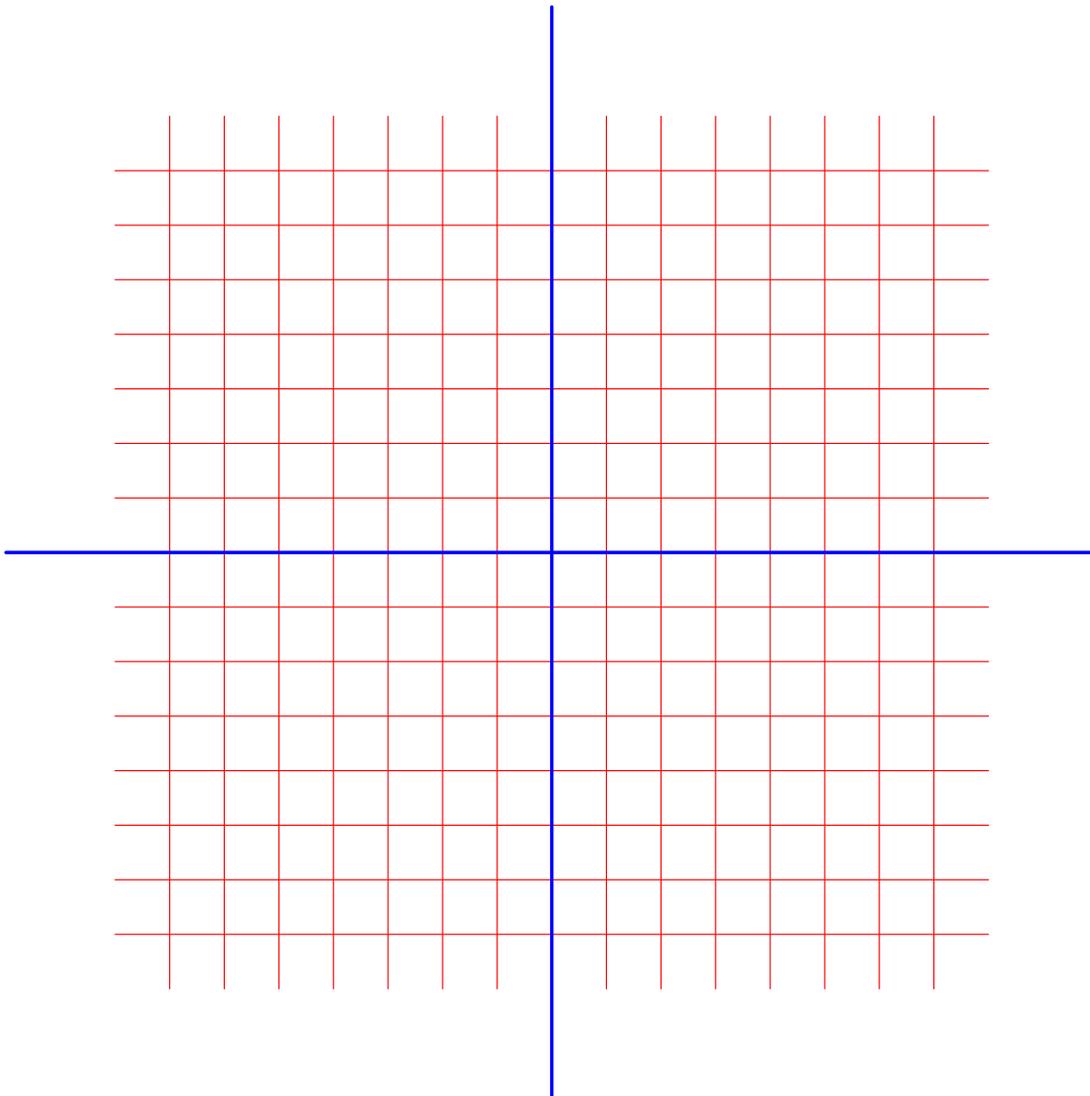


The **threefoil knot** is the space curve

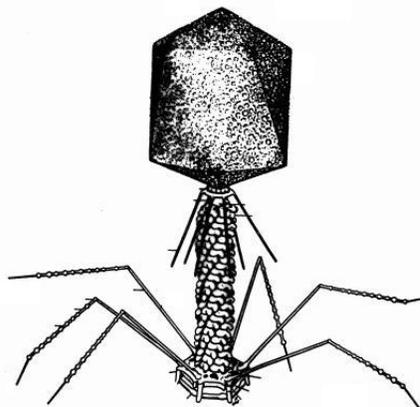
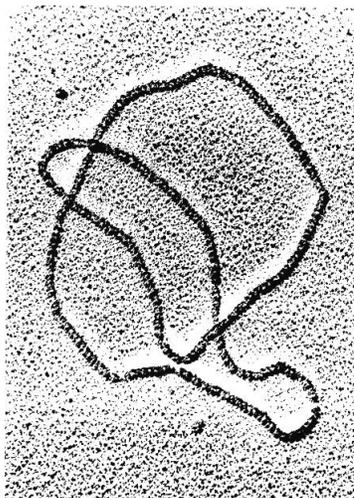
$$r(t) = (2 + \cos(3t/2) \cos(t), (2 + \cos(3t/2)) \sin(t), \sin(3t/2))$$

- 1) Find an interval  $[a, b]$  on which  $r(t)$  parameterizes a closed curve in space. A closed curve in space is called a **knot**. These objects are not only interesting in mathematics or physics. For example, DNA of bacteria and some proteins form knots (see back of the page).
- 2) (compare problem 34 on page 711) Verify that the projection of this curve onto the  $xy$ -plane is given in polar coordinates by  $r(t) = 2 + \cos(3t/2), \theta(t) = t$ .
- 3) Sketch this curve.



Indicate at the crossings, which part of the curve is above the other.

- 4) Calculate the velocity vector of  $r(t)$  at  $t = 0$ . If this vector is  $v = (a, b, c)$ , then  $(a, b)$  is its projection on the  $xy$  plane. Draw this vector  $(a, b)$  in the above  $xy$ -projection of the knot.



Communications to the Editor

A Real Knot in Protein

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It is well-known that circular DNAs exhibit a rich variety of knotted structures.<sup>1</sup> Recent surveys of the X-ray structures deposited in the Brookhaven Protein Data Bank have revealed the presence of pseudolinks and pseudoknots in protein structures caused by formation of disulfide bonds and metal coordination bonds.<sup>2</sup> However, there has been no report so far of knots in native proteins or polypeptides.<sup>2</sup> We now report our finding of a linear knot in the structure of (S)-adenosylmethionine synthetase (MAT) recently determined in our laboratory.<sup>3</sup>

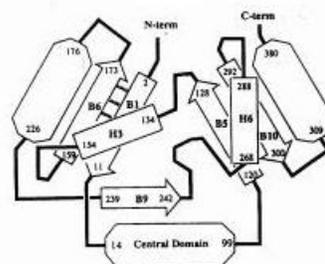


Figure 1. Schematic drawing of the unusual knot structure of the polypeptide chain of MAT. The knot is formed by passage of the B9  $\beta$ -strand leading to the C-terminus through a loop formed by the sequence B1  $\rightarrow$  [central domain]  $\rightarrow$  B5  $\rightarrow$  H3  $\rightarrow$  B6. The rectangles, arrows, and elongated octagons represent  $\alpha$ -helices,  $\beta$ -strands, and portion of domains, respectively. The numbers at both ends are the start and end of the amino acid residue numbers. Hydrogen bonds between B1 and B6  $\beta$ -strands are shown by dotted lines.



Chart 1. Ribbon Presentation of the MAT Subunit\*

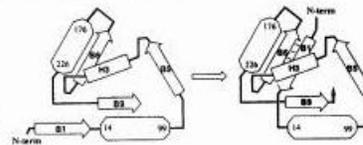
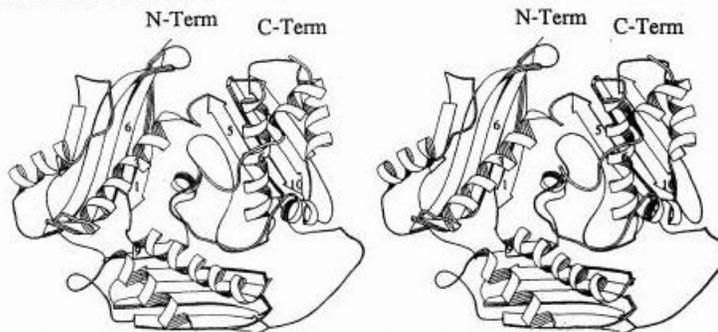


Figure 2. Hypothesis for biosynthetic knot formation. The N-terminal region (residues 1-11) locates near the central domain until after the synthesis of the B9  $\beta$ -strand (residue 242). After synthesis of the B9  $\beta$ -strand region, the N-terminal region moves into the closed loop, and the B1  $\beta$ -strand (residues 2-11) forms antiparallel  $\beta$ -sheet hydrogen bonds with the B6  $\beta$ -strand shown by dotted lines.



For more information on DNA knots, see the MSRI talk by DeWitt Summers:  
<http://www.msri.org/publications/ln/msri/2000/molbio/summers/1/>