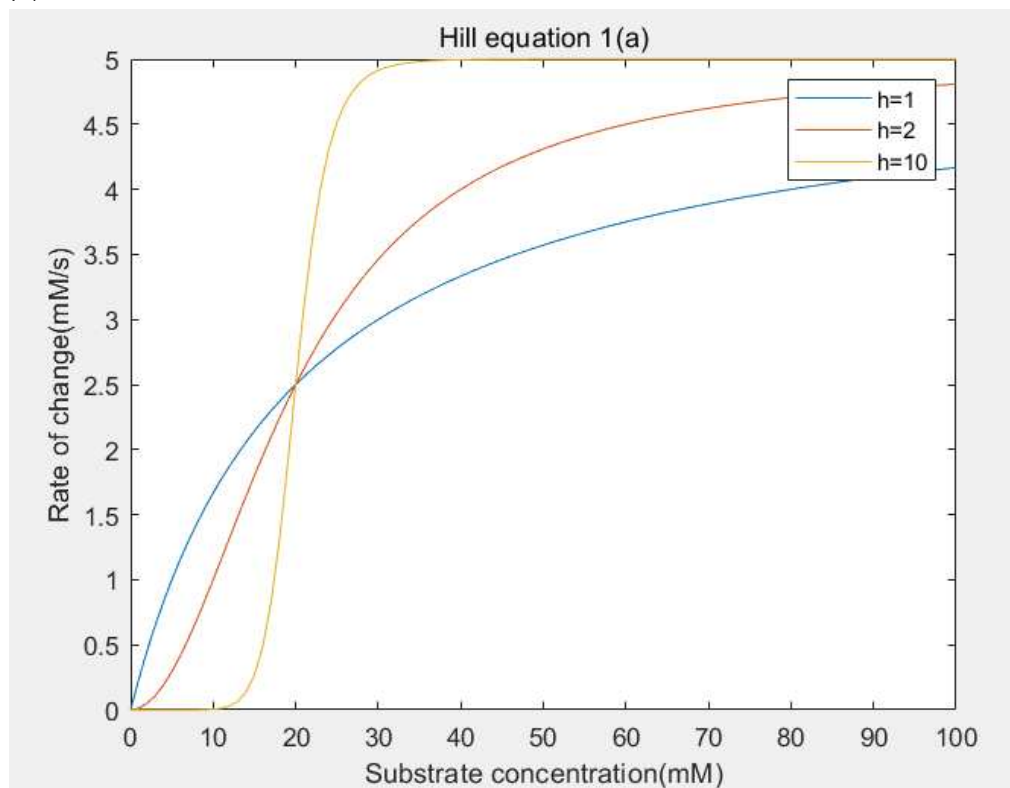


Part A

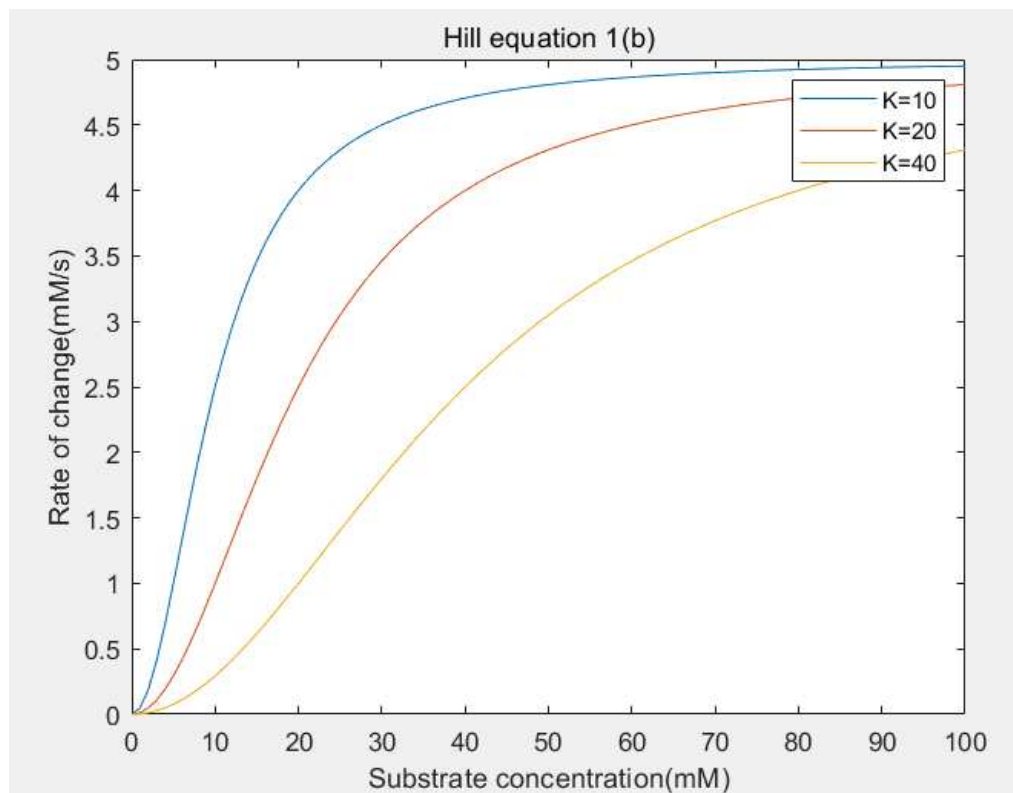
1.

Let $K(1/2) = K$

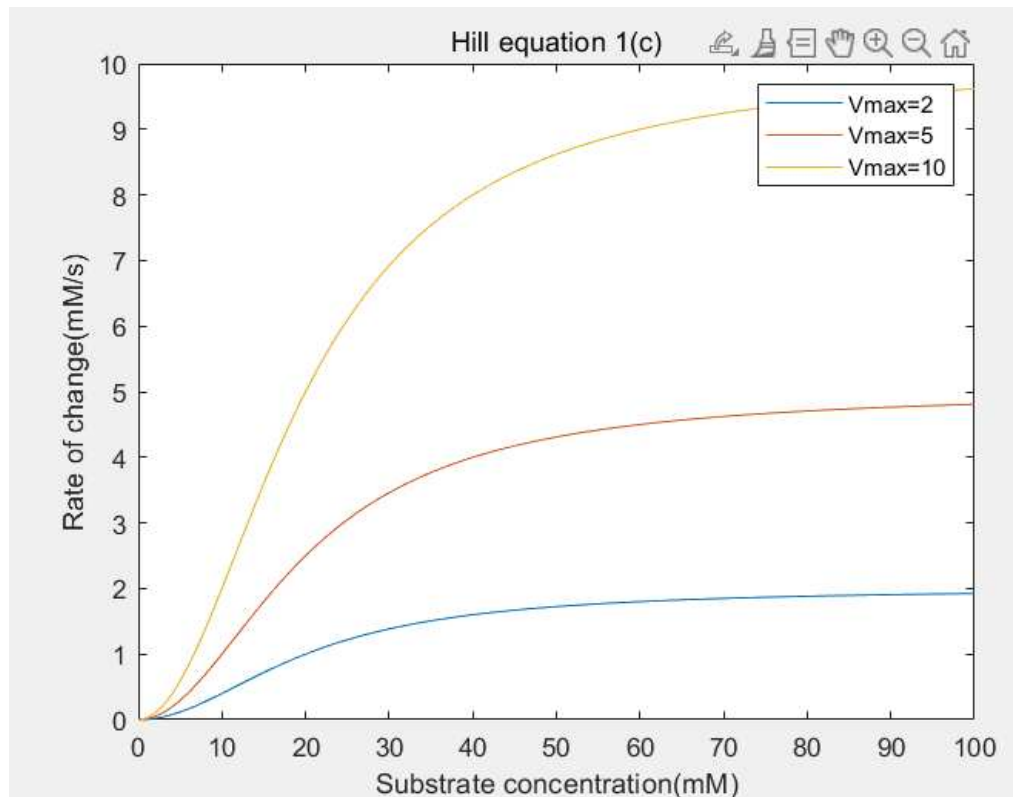
(a)



(b)



(c)



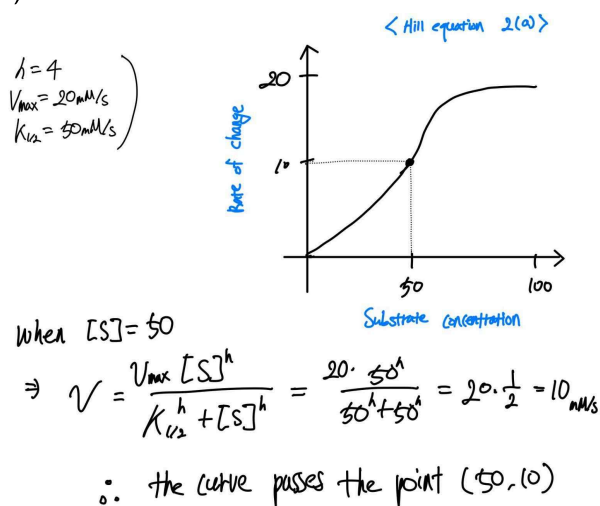
(d)

The curves have the same starting point, but because the variables are different, the curve's model and the value it converges are different.

From (a), I could see that if $[S]=K(1/2)$, the rate of change has $(V_{max}/2)$. Also, as the Hill coefficient increases, the graph converges very rapidly to the 'rate of change'= V_{max} line. From (b), it was confirmed that the graph slowly converges to 'rate of change'= V_{max} line as the value of $K(1/2)$ increases. And from (c), It was confirmed that 'rate of change'= V_{max} line is the point where the curves approach

2.

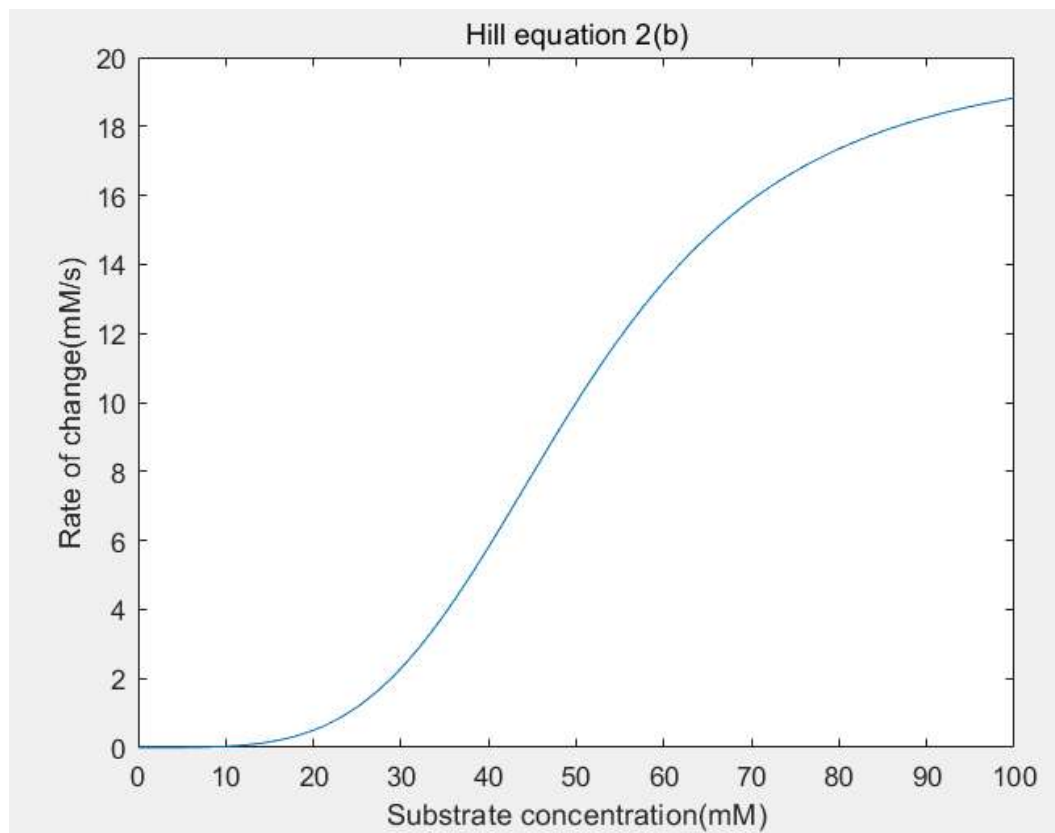
(a)



I expected when $[S]$ is small, the rate of change increases little by little because the influence of $K(1/2)$ is large and as $[S]$ increases, the influence of $K(1/2)$ becomes negligible and eventually converges to the maximum speed, V_{max} .

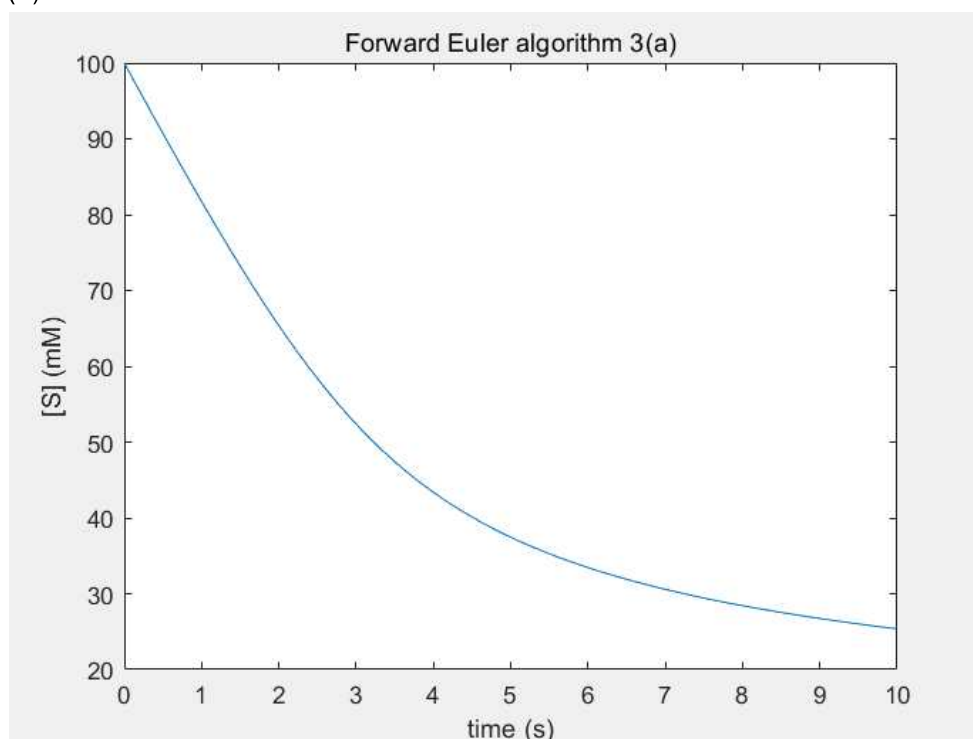
(b)

Before $[S]$ became 50 I thought the speed would increase slowly, but due to the influence of the hill coefficient, I found that even a small increase in $[S]$ increases the speed significantly.



3. Let $K(1/2) = K$ and $S = [S]$

(a)



(b)

2(b) represents the change in V according to $[S]$ through hill equation, and 3(a) is different because it is a graph that predicts how $[S]$ changes according to the change in time for 10 seconds from the initial point using that V we got in 2(b).

However, Since they both use the same Hill equation with the same parameters, we can see that depending on the V value at the specific value of $[S]$ in 2(a) graph, the rate of change of the value of $[S]$ in 3(a) graph becomes large or small.

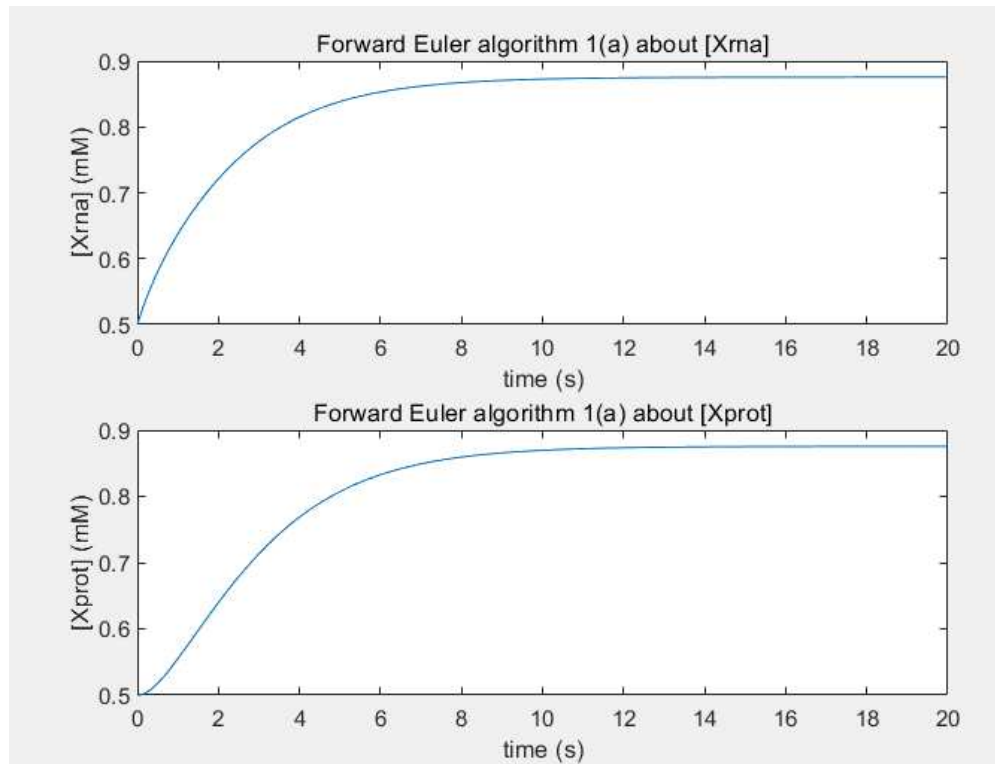
Part B

1.

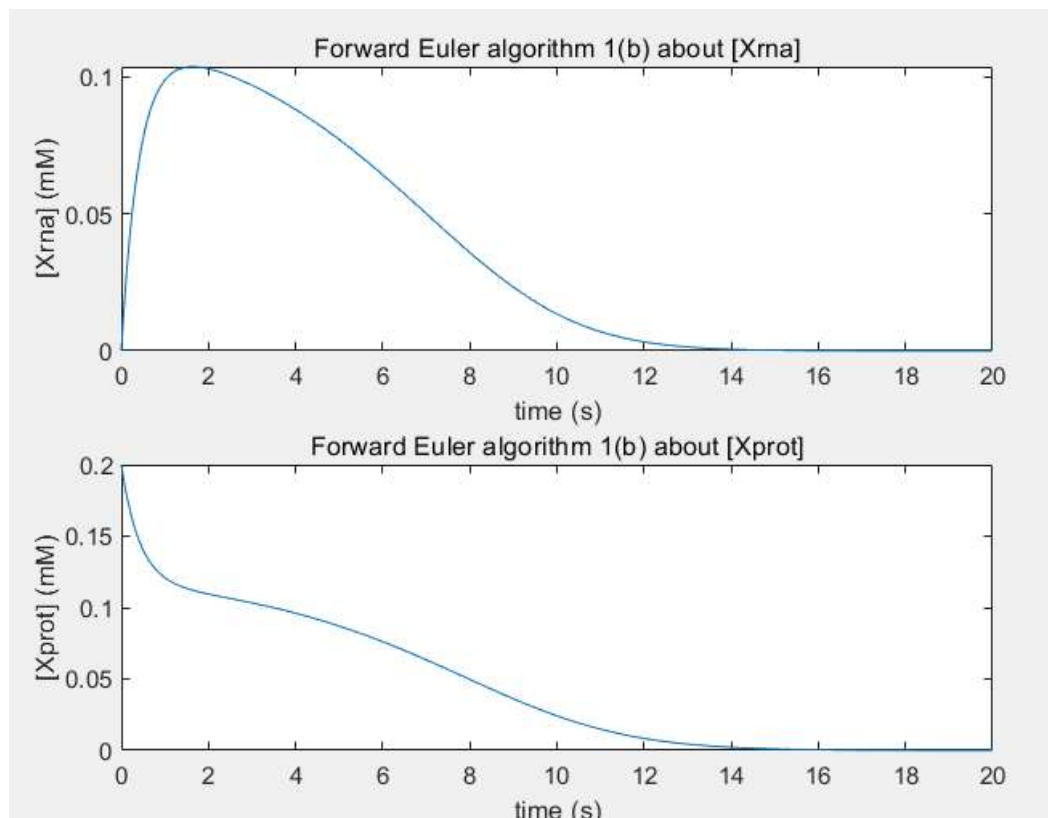
Terms X_{prot} & X_{rna} in this coding mean the concentration of X_{prot} and X_{rna}

Terms CX_{prot} & CX_{rna} in this coding mean X_{prot} and X_{rna} in the system of ODEs

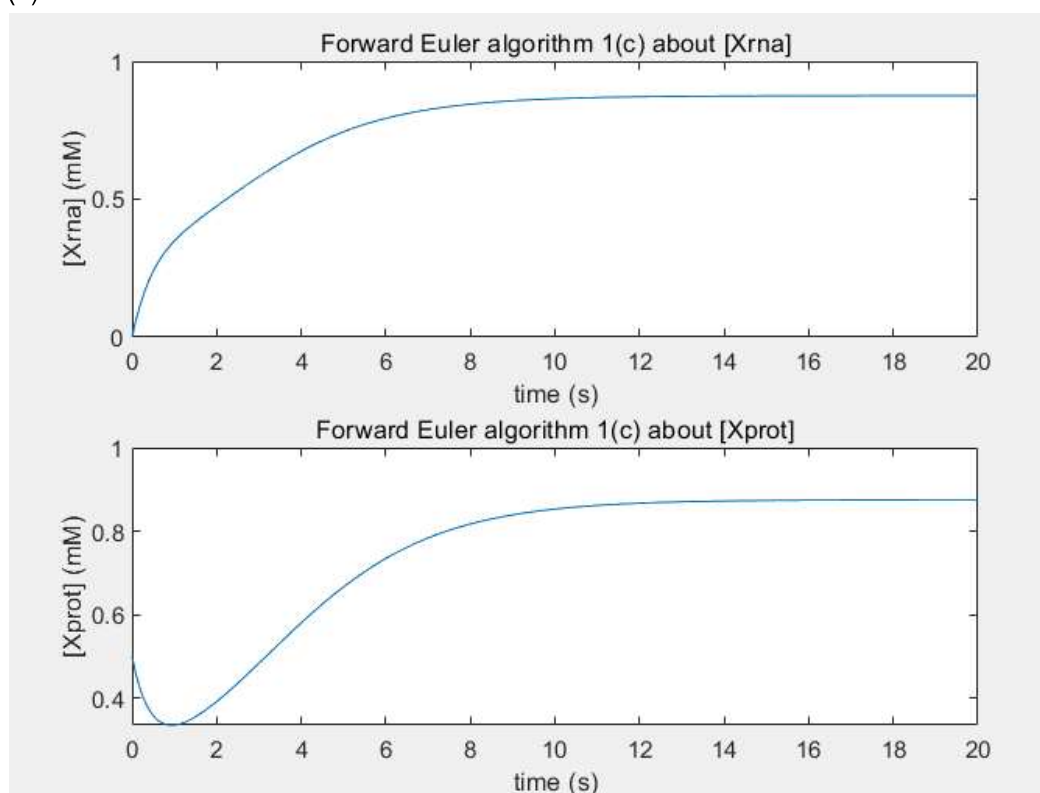
(a)



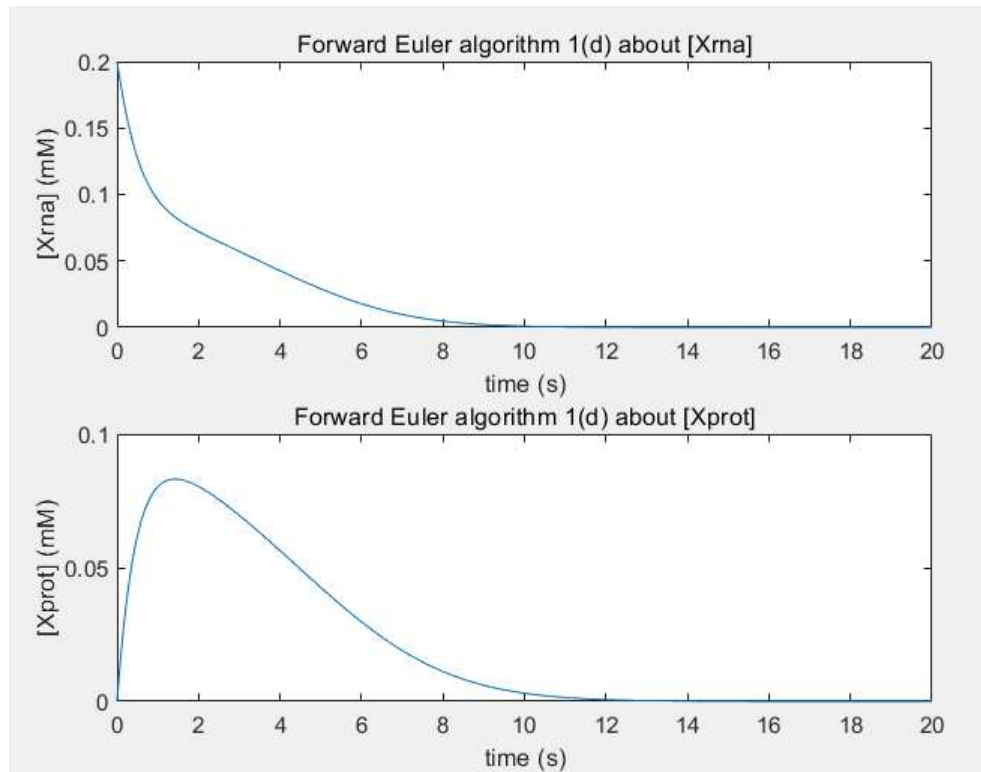
(b)



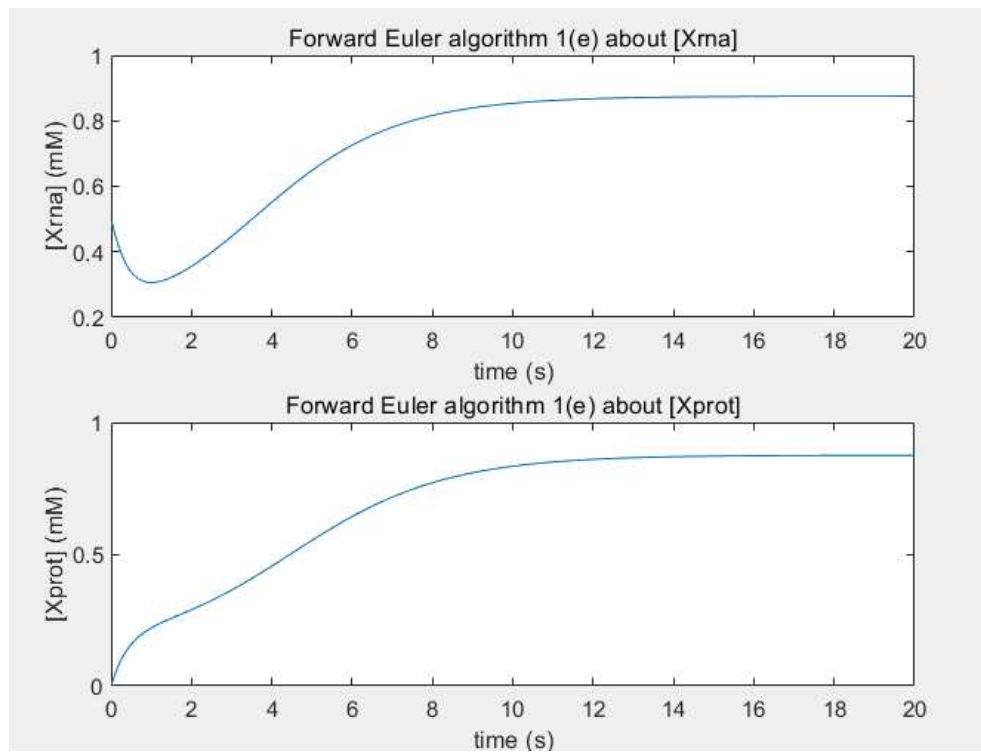
(c)



(d)



(e)

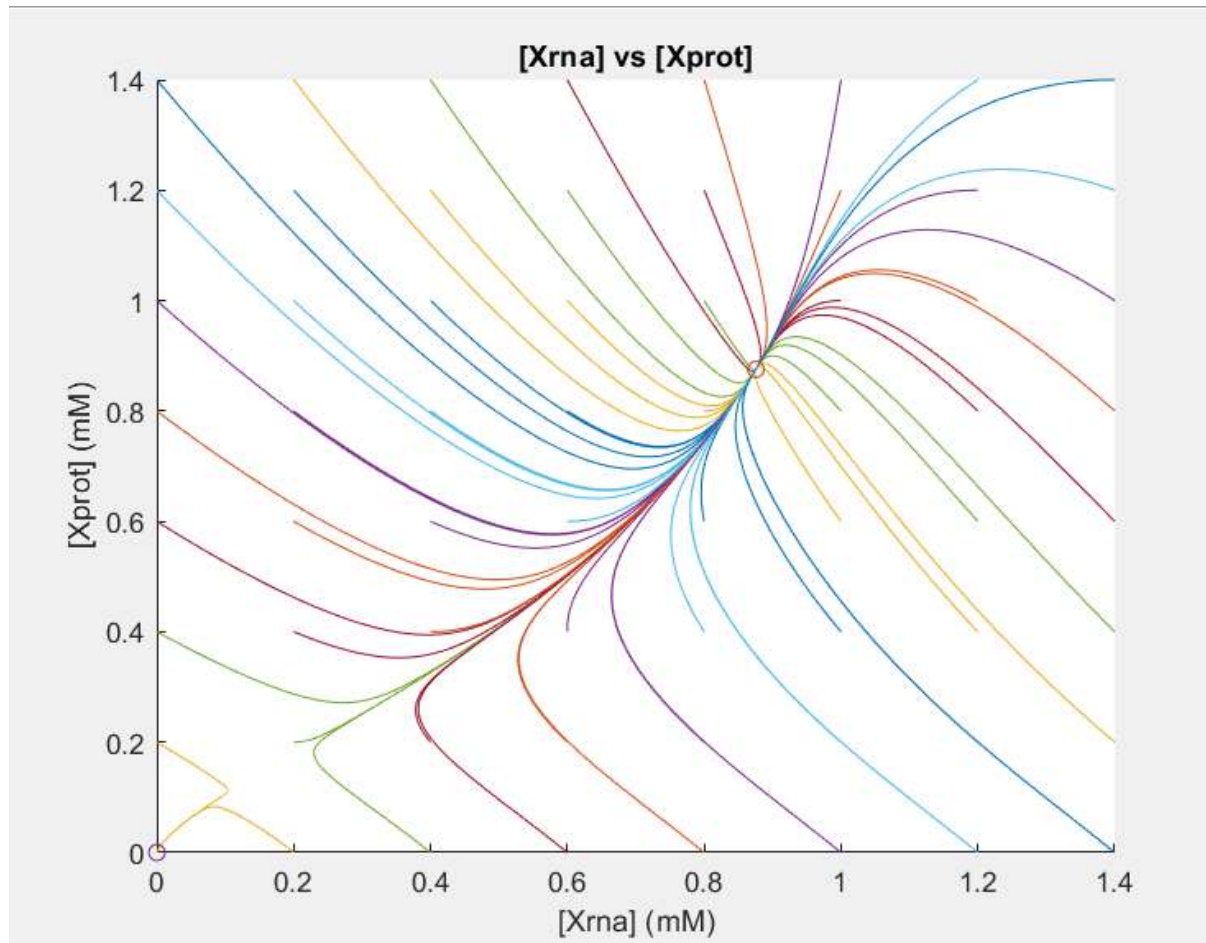


(f)

I found that the result values([Xprot]&[Xrna]) vary greatly depending on the initial value even if they are within the same condition of the system of ODEs and parameters.

2.

(a)



(c)

Each line has a different initial value. The lines indicate where the values of [Xprot] and [Xrna] go, depending on the initial value. Since it is not a graph about time, it is necessary to analyze it by thinking that lines start from the initial value of each graph. In particular, it can be seen that when the initial values of [Xprot] and [Xrna] are less than 1, the end values tend to increase, and when the initial values of [Xprot] and [Xrna] are greater than 1, the end values tend to decrease.

3.

This system is a model for a simple autoregulatory gene. During this process of making a protein, transcription and translation were represented as a system of ODEs. From graph 2(b), I came to know that if [Xprot] or [Xrna] is excessive, it will be reduced through this system, and if too little, it will be increased. This is consistent with the fact that the gradation rate in ODEs is

proportional to $[X_{\text{prot}}]$ and $[X_{\text{rna}}]$. Through this, I found that biologically, when we are in a strange situation, auto-regulatory genes tend to converge back to stable states.