8.1

(1)
$$d[E]/dt = -k_1[E][S] + k_2[ES] + k_3[ES]$$

(2)
$$d[S]/dt = -k_1[E][S] + k_2[ES]$$

(3)
$$d[ES]/dt = k_1[E][S] - k_2[ES] - k_3[ES]$$

(4)
$$d[P]/dt = k_3[ES]$$

8.2

To use the Runge-Kutta method, the number of variables need to be reduced. The concentration of free enzyme can be expressed as (total enzyme – ES complex). [E] in equations 2 and 3 can be substituted with

([E₀] – [ES]) which is [1 – ES]. Also,
$$k_1 = \frac{100/\mu M}{60s} = \frac{5}{3}\mu M/s$$
, $k_2 = \frac{600}{min} = \frac{10}{s}$, $k_3 = \frac{150}{min} = \frac{2.5}{s}$.

Therefore, equation 2 can be expressed as $d[S]/dt = -(5/3)(1 - [ES])[S] + 10[ES] \rightarrow f(t, x, y)$

Equation 3 can be expressed as d[ES]/dt = $(5/3)(1 - [ES])[S] - 12.5[ES] \rightarrow g(t, x, y)$

These are 2 simultaneous equations, with dependent variables [S] and [ES], and independent variable t. According to 4^{th} order Runge Kutta method, [S] = [S]₀ + k while [ES] = [ES]₀ + l. k and l represents $\frac{1}{6}(k_1+2k_2+2k_3+k_4)$ and $\frac{1}{6}(l_1+2l_2+2l_3+l_4)$ respectively, and the values of k_{1-4} and l_{1-4} can be derived by:

$$k_1 = h*f(t, x, y)$$

$$I_1 = h*g(t, x, y)$$

$$k_2 = h*f(t + 0.5*h, x + 0.5*k_1, y + 0.5*l_1)$$

$$I_2 = h*g(t + 0.5*h, x + 0.5*k_1, y + 0.5*I_1)$$

$$k_3 = h*f(t + 0.5*h, x + 0.5*k_2, y + 0.5*l_2)$$

$$I_3 = h*g(t + 0.5*h, x + 0.5*k_2, y + 0.5*I_2)$$

$$k_4 = h*f(t + h, x + k_3, y + l_3)$$

$$I_4 = h*g(t + h, x + k_3, y + I_3)$$

By letting step size be 0.02 and maximum time be 30s, the value of [S] and [ES] every 0.02 seconds up till 30s can be calculated.

As mentioned earlier, $[E] = [E_0] - [ES]$, so the value of [E] at every time point can be derived from 1 - [ES] at corresponding time points.

Considering that $[S] = [S_0] - [P] - [ES]$, [P] can be expressed as $[S_0] - [S] - [ES]$. Thus, value of [P] at every time point can be derived from 10 - [S] - [ES] at corresponding time points.

Values of [E], [S], [ES] and [P] every 0.02s can be stored in arrays, and these concentration values can be plotted against an array of time values.

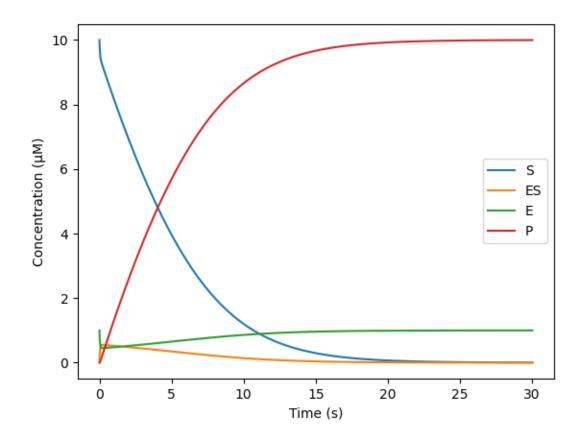
The code and graph output will be:

```
import matplotlib.pyplot as plt
def dydt(t, x, y):
x0 = 10
y0 = 0
t0 = 0
    k1 = h*dxdt(t, x, y)
    k2 = h*dxdt(t + 0.5*h, x + 0.5*k1, y + 0.5*11)
   14 = h*dydt(t + h, x + k3, y + 13)
   x += (1/6)*(k1 + 2*k2 + 2*k3 + k4)
   y += (1/6)*(11 + 2*12 + 2*13 + 14)
x list = [x0]
   x list.append(x)
    y list.append(y)
prod list = [0]
    prod list.append(10 - x list[i] - y list[i])
x arr = np.array(x list)
y arr = np.array(y list)
t_arr = np.array(t list)
enz arr = 1 - y arr
prod arr = np.array(prod list)
plt.plot(t arr,x arr)
plt.plot(t arr,y arr)
plt.plot(t arr,enz arr)
plt.plot(t arr,prod arr)
plt.xlabel('Time (s)')
```

```
plt.legend(['S','ES','E','P'])
plt.show()

velo = 150*y_arr
plateau = np.max(velo)
for i in range(len(velo)-1):
    if velo[i] != plateau:
        velo[i] = plateau
    elif velo[i] == plateau:
        break

plt.plot(x_arr,velo)
plt.ylabel('Rate of change of P (min)')
plt.xlabel('Substrate concentration (µM)')
plt.show()
```



<u>8.3</u>

Since velocity is rate of change of P, v = d[p]/dt = 150[ES]. Previously, arrays containing values for [ES] and [S] every 0.02s were created. This graph will plot 150[ES] against [S] of corresponding time points. However, the graph shows a downwards trend immediately after reaching V_m , as this corresponds to the rapid increase in [ES] at high initial substrate concentration at early time points in part 8.2. The correct trend should show a saturation of velocity at V_m . Thus, in the array for 150[ES], all elements with positions before the maximum value (and also with smaller value) are assigned the value of V_m .

From the plot, value of V_m is **82.6**.

