In [1]:

```
import numpy as np
from scipy.integrate import odeint
from matplotlib import pyplot as plt
plt.rcParams["font.family"] = "Times New Roman"
#plt.rcParams.update({'font.size': 14})
%matplotlib notebook
```

In [2]:

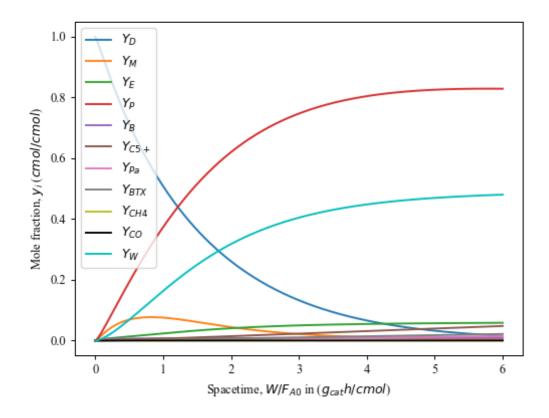
```
### Model parameters:
T0 = 623 \# K
R = 8.206 \# (m^{**}3 \ atm) / (K \ mol)
Ka = 1.27e1 \#1/atm
#Kinetic parameters:
k_01 = 4.99e-2
k_02 = 7.88e1
k_03 = 2.42e-3
k_04 = 2.48e-1
k_05 = 2.54
k_06 = 1.44e-1
k_07 = 3.02e1
k_08 = 2.63
k_09 = 4.24e-1
k_010 = 1.03
k_011 = 1.16e-2
k_012 = 3.10e-1
k_013 = 5.63e-4
f1 = 1.78e-1
f2 = 6.92e-1
k_010 = 1.03
k_010_1 = k_010*f1
k_010_2 = k_010*f2
k_012 = 3.10e-1
k_012_1 = k_012*f1
k_012_2 = k_012*f2
E1 = 4.2e1
E2 = 1.2e1
E3 = 3.4e1
E4 = 1.7e1
E5 = 2.6e1
E6 = 9.8e1
E7 = 1.6e1
E8 = 1.7e1
E9 = 6.9e1
E10 = 2.1e1
E11 = 6e-1
E12 = 2.1e1
E13 = 3.4e1
###
```

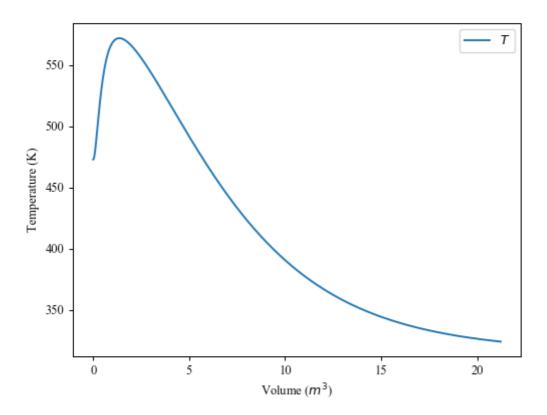
```
### Non-adiabatic non isothermal heat transfer parameters:
rho c = 1790 \#kq/m^{**3}
epsilon = 0.29
rho bed = rho c*(1-epsilon) \#kq/m**3
Tu = 300 #K cooling water constant temperature
U = 750 \ \#(W/m^{**}2 \ K) \ p656 \ Coulson \ @ boiling organics (paraffins) and cooling tower water
# heat transfer area (dependant on tube geometry used!)
a = 1990 #m**2 # solve until the temperature NEVER exceeds 573 K to prevent coking (n
o steam fed)
#max a (at max pressure=20.1 atm) is 2375! remember to check max catalyst!
H1D = -40129 \ \#kJ/kmolD
H2D =1980 #kJ/kmolD
H3M = -31152 \# kJ/kmolM
H4D = -1886 \# kJ/kmolD
H5D = -42734 \ \#kJ/kmolD
H6D = -56355 \#kJ/kmolD
H7M = -12972 \#kJ/kmoLM
H8M = -32454 \# kJ/kmolM
H9M = -39265 \# kJ/kmoLM
H10 1E =-111938.03 #kJ/kmolE
H10P = -109404.8 \#kJ/kmolP
H10_{2B} = -116607.14 \ \#kJ/kmolB
H11_C5_plus =123615 #kJ/kmolC5_plus
H12 1E =-62495 \#kJ/kmolE
H12P = -35298 \# kJ/kmolP
H12\ 2B = -19818\ \#kJ/kmolB
H13D = 2745 \#kJ/kmolD
CpD = 102.8 \#kJ/kmolK
CpM = 50.94 \#kJ/kmolK
CpE =68.73 \#kJ/kmolK
CpP = 104 \#kJ/kmolK
CpB = 142.56 \#kJ/kmolK
Cp_C5_plus =202.25 #kJ/kmolK
CpPa =125.02 \#kJ/kmolK
CpBTX =186.6 #kJ/kmolK
CpCH4 = 65.4 \#kJ/kmolK
CpCO = 30.27 \#kJ/kmolK
CpW = 36.11 \#kJ/kmolK
###
def ri(x):
         pD,pM,pE,pP,pB,pC5_plus,pPa,pBTX,pCH4,pCO,pW, T = x
        K2 = np.exp(-(-9.76 + 3200/T + 1.07*np.log(T) -6.6e-4*T + 4.9e-8*T**2 + 6500/T**2))
  #'dimensionless'
        \theta = 1/(1 + Ka*(pM + pW)) #dimensionless
         k1 = k_01*np.exp((-E1/R)*(1/T - 1/T0)) #molc/(gcat h atm)#I think this should be: m
olc/(gcat h atm)
        k2 = k_02*np.exp((-E2/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)#should be: molc/(gcat h atm**2)
  h atm**2)
         k3 = k_03*np.exp((-E3/R)*(1/T - 1/T0)) #molc/(gcat h atm)
         k4 = k_04*np.exp((-E4/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
         k5 = k_05*np.exp((-E5/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
         k6 = k_06*np.exp((-E6/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
```

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k7 = k_07*np.exp((-E7/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
    k8 = k_08*np.exp((-E8/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
    k9 = k_0^2 + np.exp((-E9/R)*(1/T - 1/T0)) #molc/(gcat h atm**2)
    k10 = k_010*np.exp((-E10/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k10_1 = k_010_1*np.exp((-E10/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k10_2 = k_010_2*np.exp((-E10/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k11 = k_011*np.exp((-E11/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k12 = k_012*np.exp((-E12/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k12_1 = k_012_1*np.exp((-E12/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k12_2 = k_012_2*np.exp((-E12/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    k13 = k_013*np.exp((-E13/R)*(1/T - 1/T0)) #molc/(gcat h atm)
    r1 = k1*pD*\theta #because units were fixed this is: molc/(gcat h)
    r2 = k2*(pD*pW - (pM**2)/K2)*\theta #reversible reaction
    r3 = k3*pM*\theta
    r4 = k4*pD*pE*\theta
    r5 = k5*pD*pP*\theta
    r6 = k6*pD*pB*\theta
    r7 = k7*pM*pE*\theta
    r8 = k8*pM*pP*\theta
    r9 = k9*pM*pB*\theta
    r10 = k10*pP*\theta
    r10_1 = k10_1*pE*\theta
    r10_2 = k10_2*pB*\theta
    r11 = k11*pC5_plus*\theta
    r12 = k12*pP*\theta
    r12_1 = k12_1*pE*\theta
    r12 \ 2 = k12 \ 2*pB*\theta
    r13 = k13*pD*\theta
                            4 5 6 7 8 9
                                                                                      12
                      3
                                                           10
                                                                10_1 10_2
       12_1
                12_2
                        13
    rD = -6*r1-2*r2
                        -2*r4-2*r5-2*r6
                      -6*r13
               2*r2-3*r3
    rM =
                                          -r7 -r8 -r9 #molMc/ (gcat h)
    rE = 2*r1
                   + r3 + 2*r4
                                         + r7
                                                             - r10 1
  -2.5*r12_1 #molEc/ (gcat h)
                            +2*r5
    rP = 2*r1
                                              + r8
                                                        -r10
                                                                                    -2.5*r
                   + r3
12 #molPc/ (gcat h)
    rB = 2*r1
                   + r3
                                   +2*r6
                                                    + r9
                                                                    - r10 2
            -2.5*r12_2
    rC5_plus =
                                                          r10 +r10_1 +r10_2-2.5*r11 #molC
5_plusc/ (gcat h)
    rPa =
                                                                                 r11 + r
12 + r12 1+
                r12 2
    rBTX =
                                                                             1.5*r11+1.5*r
12+1.5*r12 1+1.5*r12 2
    rCH4 =
                      4*r13
    rCO =
                      2*r13
           r1 - r2 + r3 + r4 + r5 + r6 + r7 + r8 + r9 #molWc/ (qcat h)
    rW =
    return [[rD,rM,rE,rP,rB,rC5_plus,rPa,rBTX,rCH4,rC0,rW, T],
            [r1,r2,r3,r4,r5,r6,r7,r8,r9,r10,r10_1,r10_2,r11,r12,r12_1,r12_2,r13]]
def cmolbalance(variables, catalyst):
    rD,rM,rE,rP,rB,rC5 plus,rPa,rBTX,rCH4,rCO,rW, T = ri(variables)[0]
    r1,r2,r3,r4,r5,r6,r7,r8,r9,r10,r10_1,r10_2,r11,r12,r12_1,r12_2,r13 = ri(variables)
[1]
    FD, FM, FE, FP, FB, F_C5_plus, FPa, FBTX, FCH4, FCO, FW, placehold =
```

```
variables*FA0/Pin
    #Energy balance:
    \#dTdW FA0 = T
    dTdW_FA0 = (FA0*((U*a/rho_bed)*(Tu - T)+(6*r1*-H1D))
                                              + 2*r2*-H2D
                                              + 3*r3*-H3M
                                              + 2*r4*-H4D
                                              + 2*r5*-H5D
                                              + 2*r6*-H6D
                                              + r7*-H7M
                                              + r8*-H8M
                                              + r9*-H9M
                                              + r10*-H10P
                                              + r10_1*-H10_1E
                                              + r10_2*-H10_2B
                                              + 2.5*r11*-H11 C5 plus
                                              + 2.5*r12*-H12P
                                              + 2.5*r12_1*-H12_1E
                                              + 2.5*r12_2*-H12_2B
                                              + 6*r13*-H13D))/
              (FD*CpD
             + FM*CpM
             + FE*CpE
             + FP*CpP
             + FB*CpB
             + F_C5_plus*Cp_C5_plus
             + FPa*CpPa
             + FBTX*CpBTX
             + FCH4*CpCH4
             + FCO*CpCO
             + FW*CpW))
    #Pressure drop:
    #Mole balances:
    dyDdW_FA0 = rD #molDc/ (gcat h)
    dyMdW FA0 = rM
    dyEdW_FA0 = rE
    dyPdW FA0 = rP
    dyBdW_FA0 = rB
    dyC5_plusdW_FA0 = rC5_plus
    dyPadW FA0 = rPa
    dyBTXdW FA0 = rBTX
    dyCH4dW FA0 = rCH4
    dyCOdW_FA0 = rCO
    dyWdW FA0 = rW
    return [dyDdW_FA0,dyMdW_FA0,dyEdW_FA0,dyPdW_FA0,dyBdW_FA0,dyC5_plusdW_FA0,
            dyPadW FA0,dyBTXdW FA0,dyCH4dW FA0,dyCOdW FA0,dyWdW FA0, dTdW FA0]
#Scale up for reactor:
cmolfactor = 2 #cmol DME/mol DME
FA0 = 2247.84*cmolfactor*1000 #cmol DME feed (to reactor inlet)
Pin = 16.91 #atm #use P=16.91 atm for max catalyst(WHSV) AND check max temperature> ch
ange 'a'
Tin= 473 #K or 200C #also MAX P = 20.1 atm for WHSV of 4 then see 'a' for increase HX
variables = [Pin, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, Tin]
spacetime = np.linspace(0,6,10000) # gcat h/ cmol
```

```
solve = odeint(cmolbalance, variables, spacetime)
Ptotal = Pin #atm
plt.figure(1)
plt.plot(spacetime, solve[:,0]/Pin, label='$Y D$')
plt.plot(spacetime, solve[:,1]/Pin, label='$Y_M$')
plt.plot(spacetime, solve[:,2]/Pin, label='$Y_E$')
plt.plot(spacetime, solve[:,3]/Pin, label='$Y_P$')
plt.plot(spacetime, solve[:,4]/Pin, label='$Y B$')
plt.plot(spacetime, solve[:,5]/Pin, label='$Y_{C5+}$')
plt.plot(spacetime, solve[:,6]/Pin, label='$Y_{Pa}$')
plt.plot(spacetime, solve[:,7]/Pin, label='$Y_{BTX}$')
plt.plot(spacetime, solve[:,8]/Pin, label='$Y_{CH4}$')
plt.plot(spacetime, solve[:,9]/Pin, label='$Y_{CO}$', color='black')
plt.plot(spacetime, solve[:,10]/Pin, label='$Y_W$', color='c')
plt.xlabel('Spacetime, $W/F_{A0}$ in ($g_{cat} h/ cmol$)')
plt.ylabel('Mole fraction, $y_i$ ($cmol/cmol$)')
plt.legend(loc='best')
plt.show()
plt.savefig("figure1_Model2.png")
moleflow = solve*FA0/cmolfactor/Ptotal/1000 #kmol i/h
catalyst_scaled = spacetime*FA0/1000 # kg
Volume = catalyst_scaled/rho_bed
plt.figure(2)
plt.plot(Volume, solve[:,11], label='$T$')
plt.xlabel('Volume ($m^3$)')
plt.ylabel('Temperature (K)')
plt.legend(loc='best')
plt.show()
plt.savefig("figure2_Model2.png")
print ('Maximum temperature =', max(solve[:,11]), 'K')
print ('Total catalyst =', catalyst_scaled[-1], 'kg')
print ('Catalyst to use @ max Propylene =', catalyst_scaled[moleflow[:,3].argmax()], 'k
g')
print ('Volume of packed bed reactor =', Volume[-1], 'm^3')
```





Maximum temperature = 572.094394116 K Total catalyst = 26974.08 kg Catalyst to use @ max Propylene = 25892.3112151 kg Volume of packed bed reactor = 21.224392163 m^3

| In []: | | |
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