

CME 570 HW4

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1. Determine the phase of the binary couple (e.g., both sides are bcc or fcc) and the composition difference between the couple. You should be able to determine the phase and composition range based on the phase diagram of the system. Plot the phase diagram and show that in your report. Also show the phase and composition range in the phase diagram.

I am using the Nickel-based demo databases, with Al-Cr as the binary alloy.

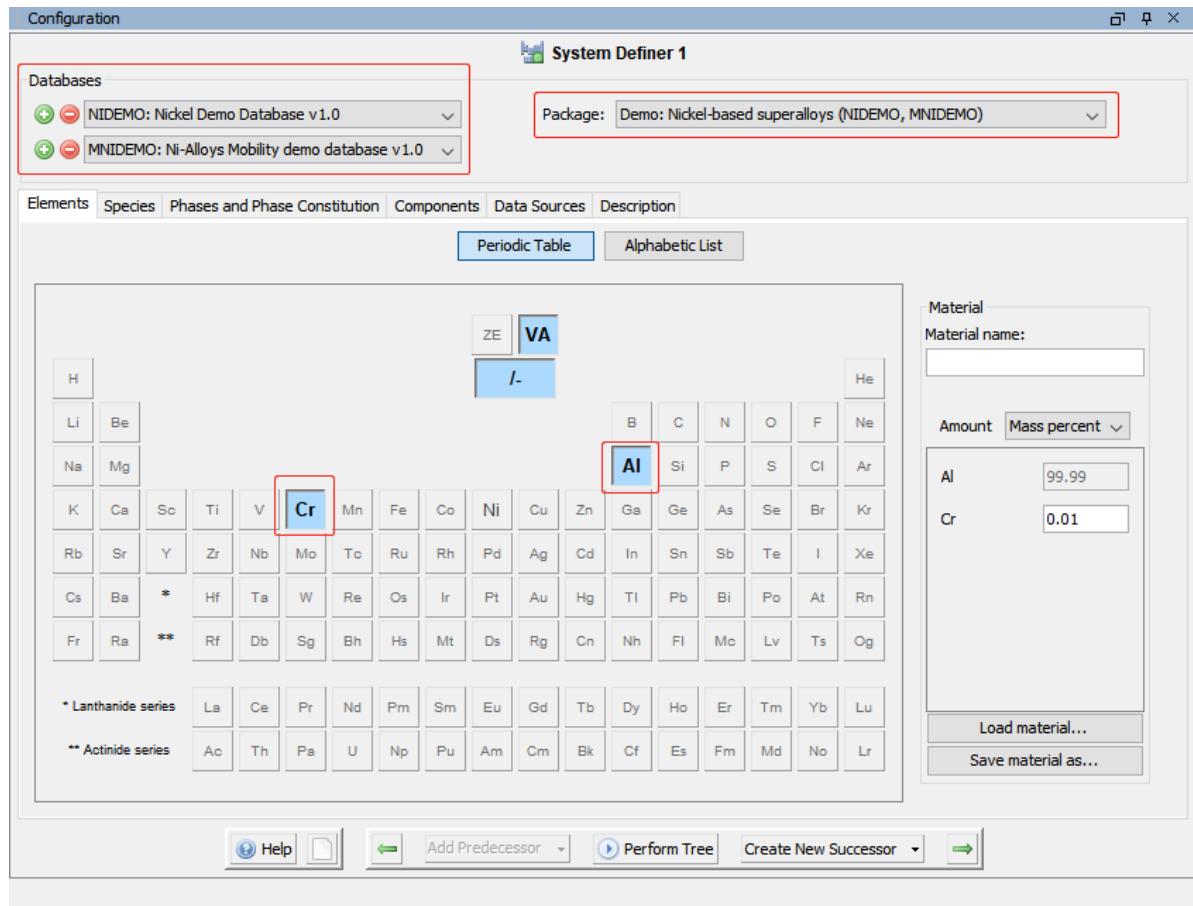


Figure 1: Thermodynamic and Mobility Data Sources

I only included the following phases: BCC_A2, FCC_A1, LIQUID, and GAS(not shown in phase diagram).

Elements	Species	Phases and Phase Constitution	Components	Data Sources	Description
Phases					
Status	Name	NIDEMO	MNIDEMO		
Entered	AL11CR2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL13CR2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL3NI2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL4CR	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL8CR5_H	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL8CR5_L	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL9CR4_H	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL9CR4_L	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	ALCR2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	BCC_A2	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		
Entered	BCC_B2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	BCC_B2#2	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	BCT_D022	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	C14_LAVES	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	CBCC_A12	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	CHI_A12	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	CUB_A13	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	DIAMOND_A4	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	DIS_FCC_A1	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	DIS_MU	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	DIS_SIG	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	FCC_A1	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		
Entered	FCC_L12	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	GAS	<input checked="" type="checkbox"/>	<input type="checkbox"/>		
Entered	HCP_A3	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	LIQUID	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		
Entered	MU_PHASE	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	NI3TA_D0A	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	NI3TI_D024	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	SIGMA	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL3NI1	<input type="checkbox"/>	<input type="checkbox"/>		
Entered	AL3NI5	<input type="checkbox"/>	<input type="checkbox"/>		

Check/uncheck all

Add composition set...

Figure 2: Included Phases in the Calculations

I plotted the phase diagram within $T \in [500 \text{ K}, 1250 \text{ K}]$ and $x_{Al} \in [0.00, 1.00]$.

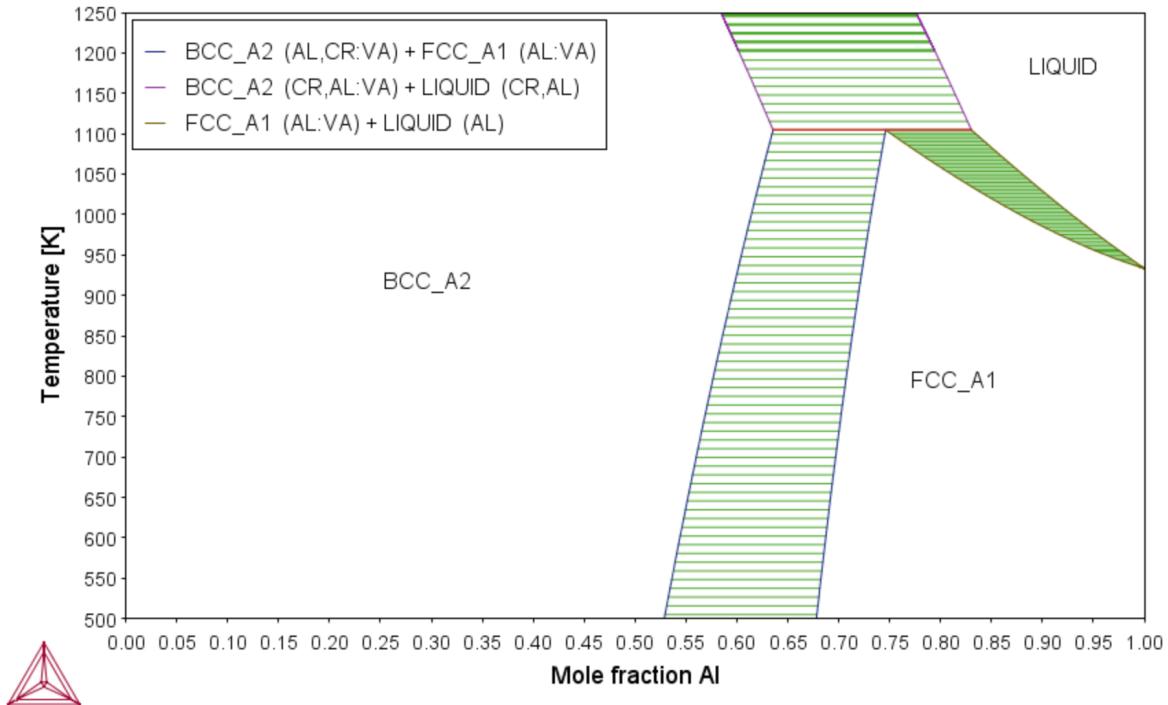


Figure 3: Phase Diagram of Al-Cr

Binary Couple at the following mole fraction compositions:

1. $x_{Al} = 0.8, x_{Cr} = 0.2,$
2. $x_{Al} = 1.0, x_{Cr} = 0.0,$

At $T = 900 \text{ K}$,

in the FCC_A1 region: (AL:VA=1:1).

Compositions of elements and vacuum are taken from the ThermoCalc "TCAL5" document[1].

2. Assume that these two materials are joined together in a weldment and then kept at an elevated temperature for a period of time. Assume that the length of each side is 0.5 mm. Plot the following:

- (a) concentration profile
- (b) flux of each species
- (c) activity profile
- (d) chemical diffusion profile

You should be able to determine the simulation temperature based on the phase diagram.

I configured the diffusion simulation as the following:

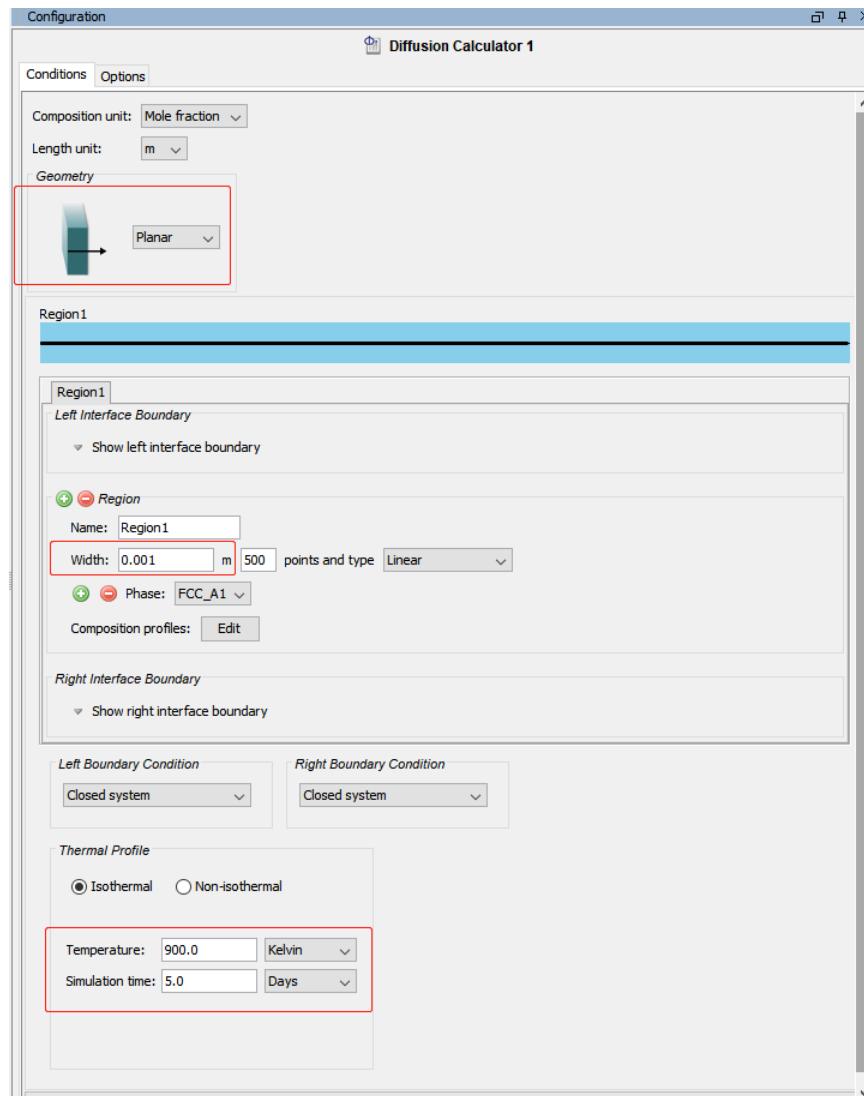


Figure 4: Simulation Configuration: FCC_A1, $T = 900\text{K}$

with the initial concentration profile as:

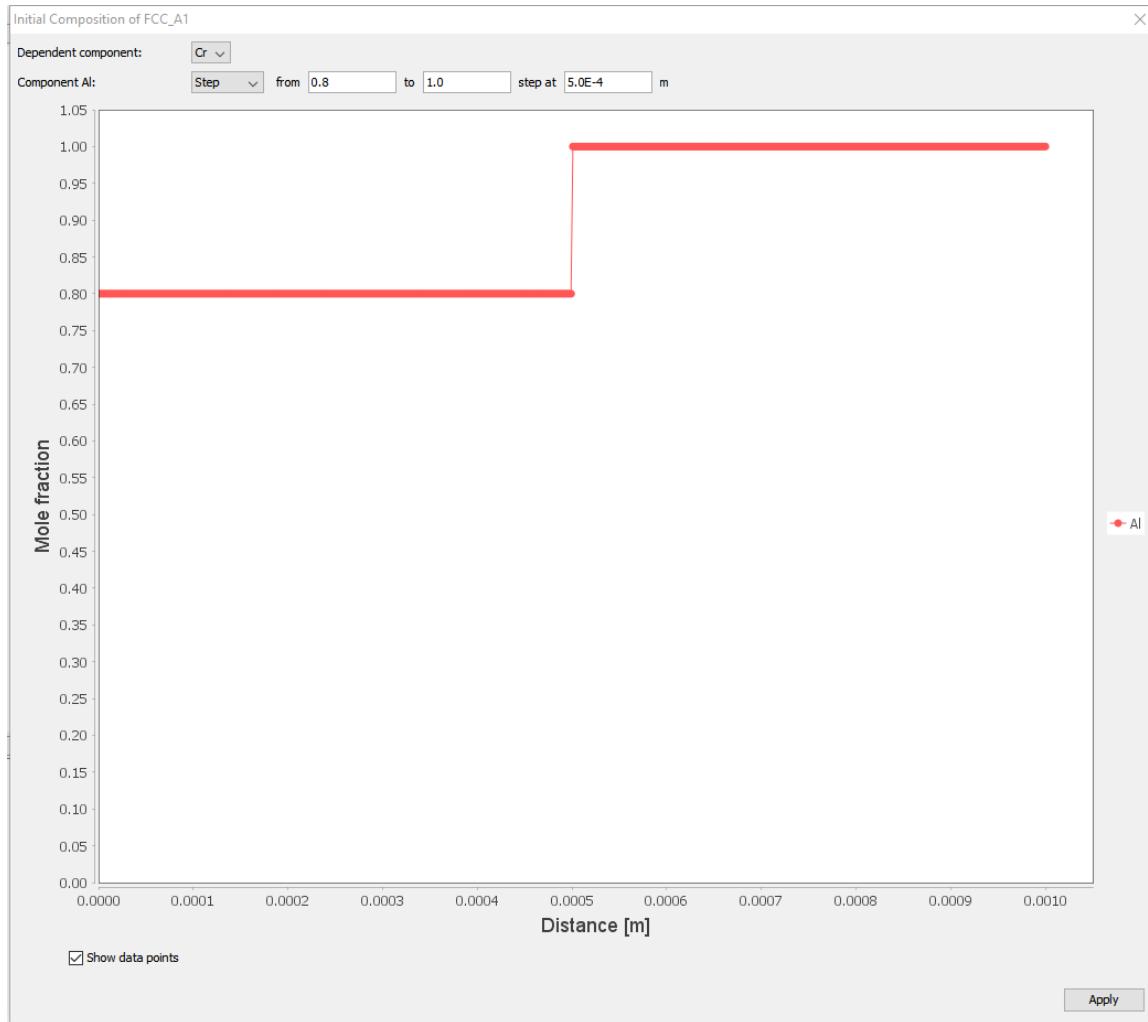


Figure 5: Initial Concentration Profile of Al

I set the solver to "homogenization" in the simulation options:

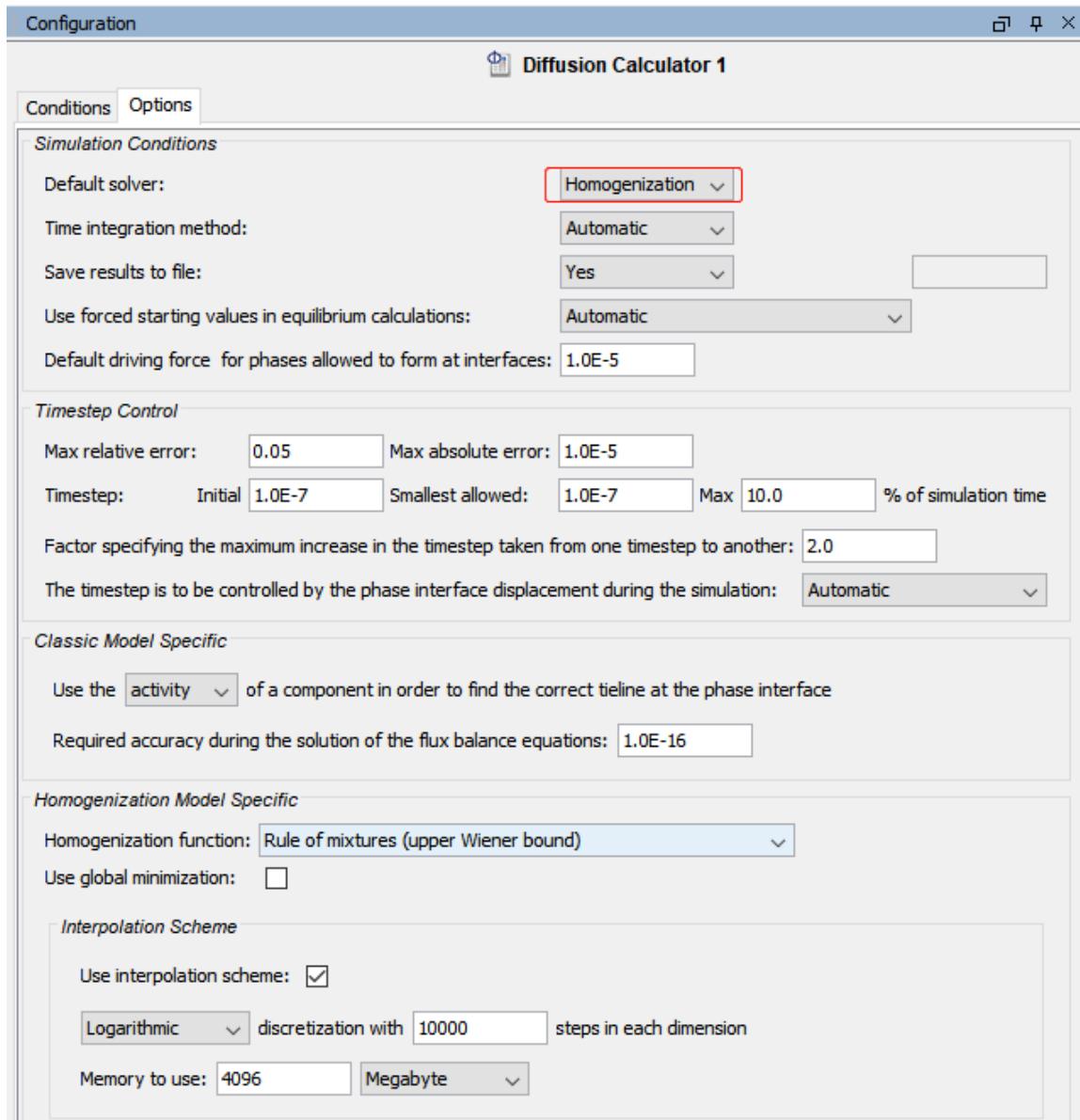


Figure 6: Simulation Options

(a) concentration profile

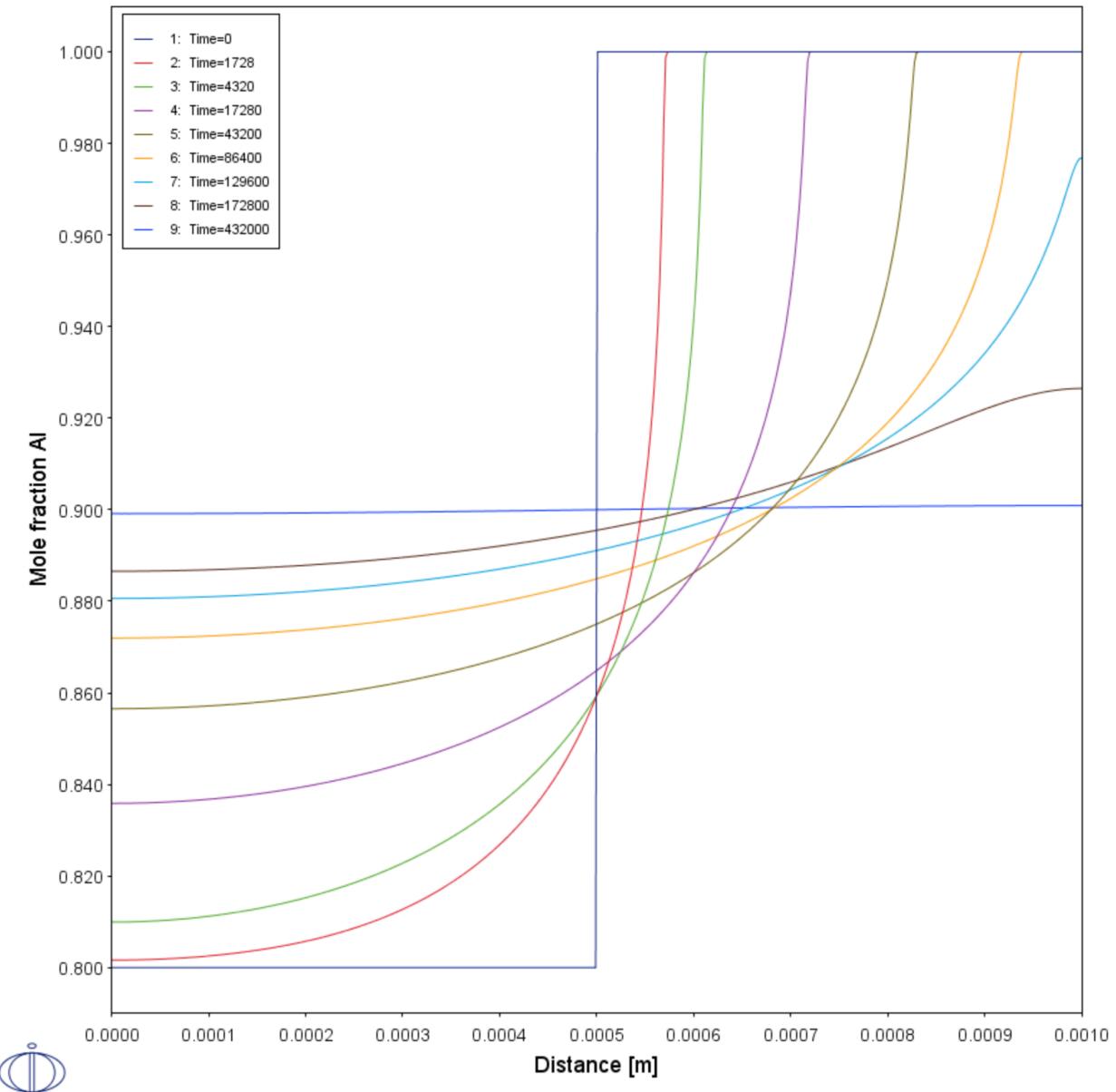
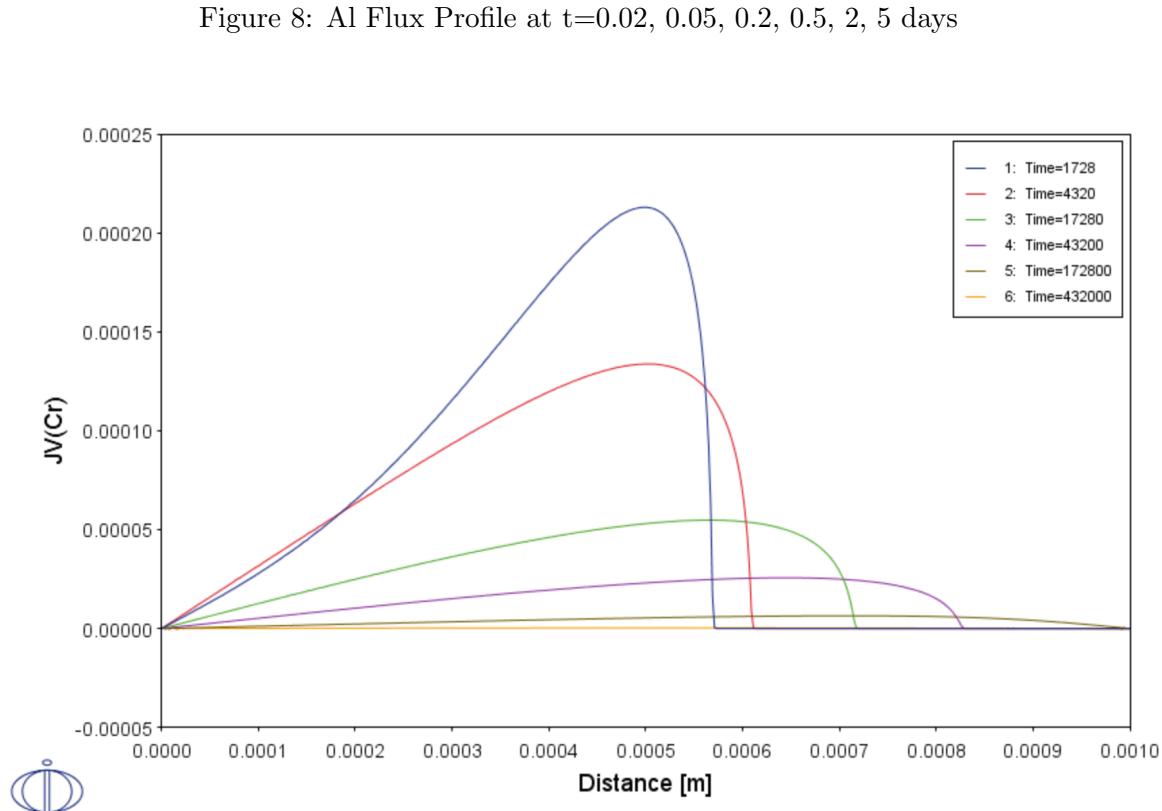
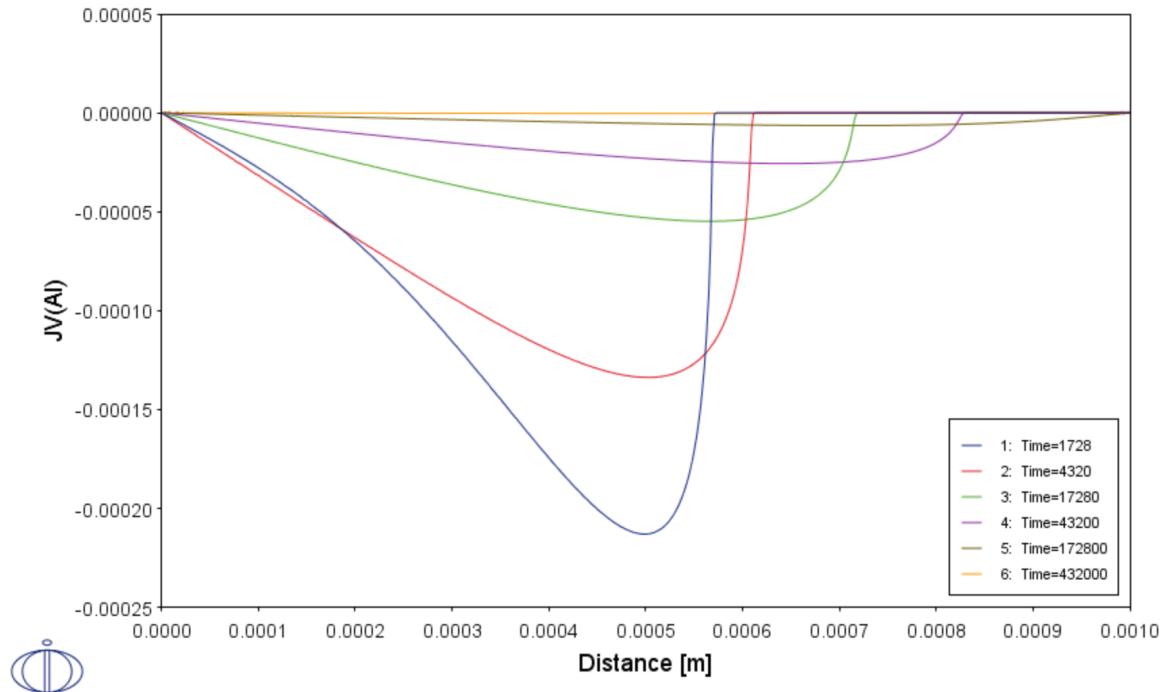


Figure 7: Composition Profile of the system at t=0, 0.02, 0.05, 0.2, 0.5, 1.0, 1.5, 2, 5 days

(b) flux of each species



(c) activity profile

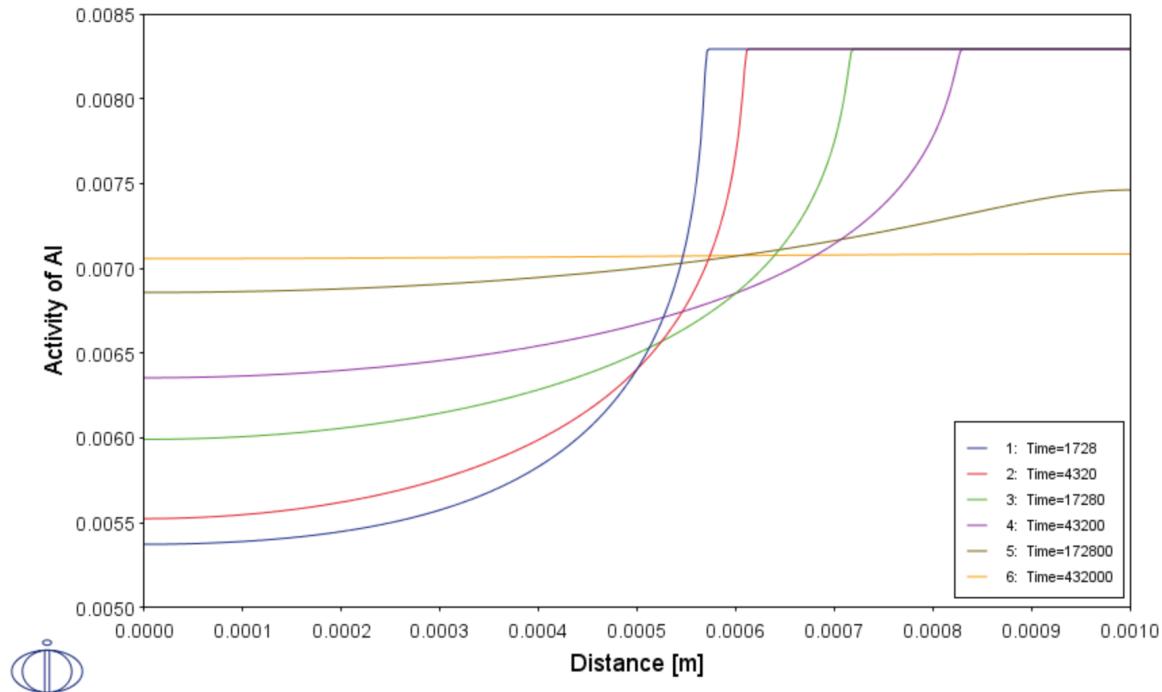


Figure 10: Al Activity Profile at $t=0.02, 0.05, 0.2, 0.5, 2, 5$ days

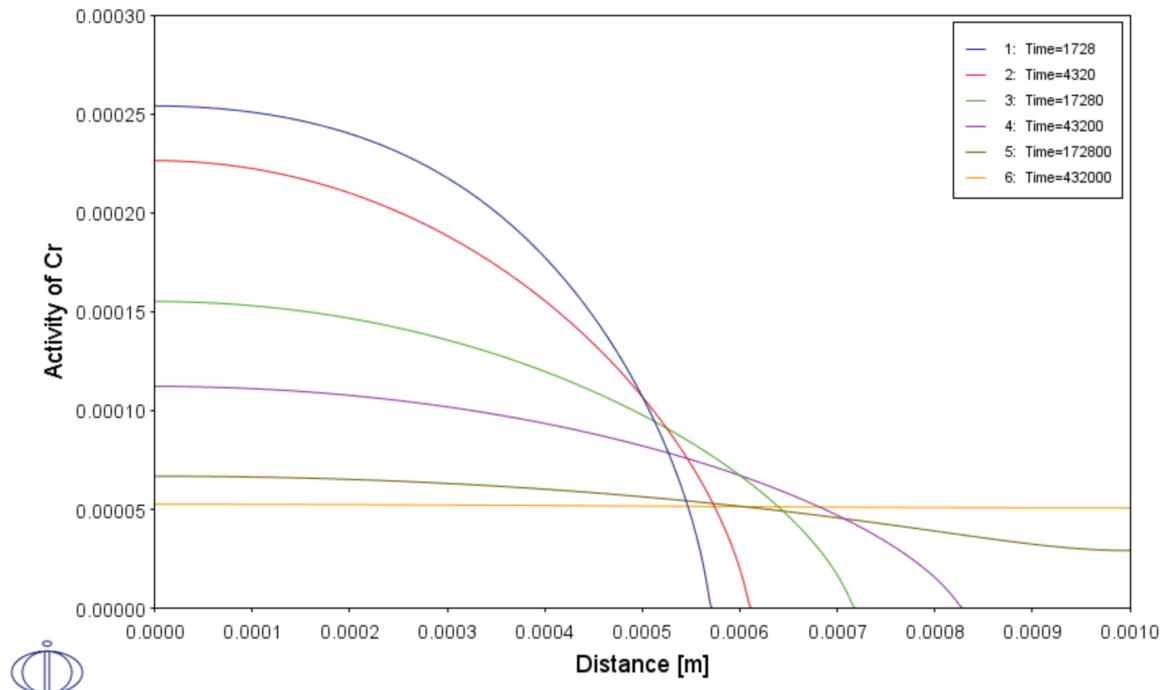


Figure 11: Cr Activity Profile at $t=0.02, 0.05, 0.2, 0.5, 2, 5$ days

(d) chemical diffusion profile

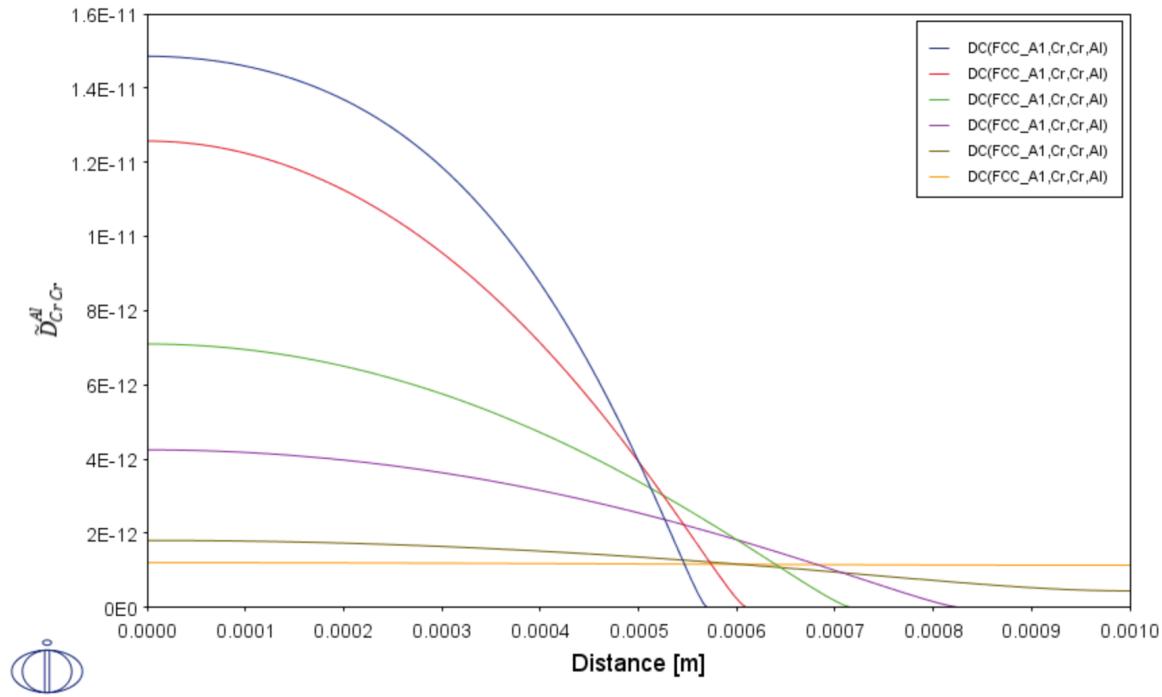


Figure 12: Al Chemical Diffusion Profile at $t=0.02, 0.05, 0.2, 0.5, 2, 5$ days

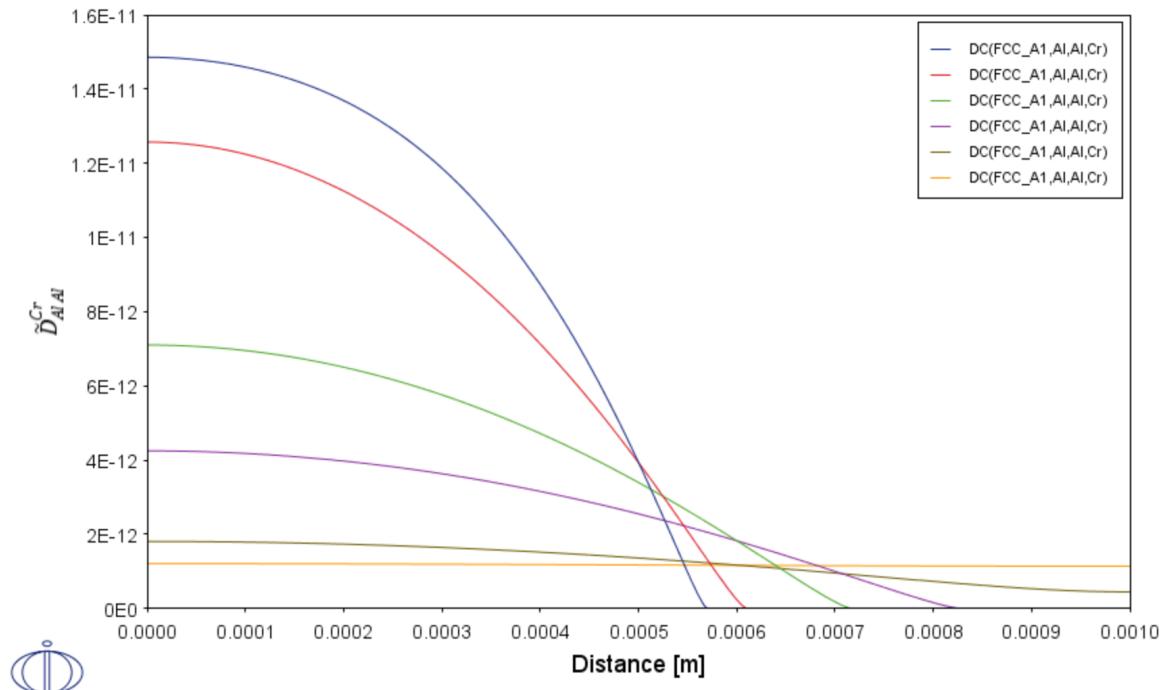


Figure 13: Cr Chemical Diffusion Profile at $t=0.02, 0.05, 0.2, 0.5, 2, 5$ days

3. Having the concentration profile for a specific annealing time, use the Matano-Boltzmann relation to reproduce the chemical diffusivity for different concentration. Compare that with the results of your simulation.

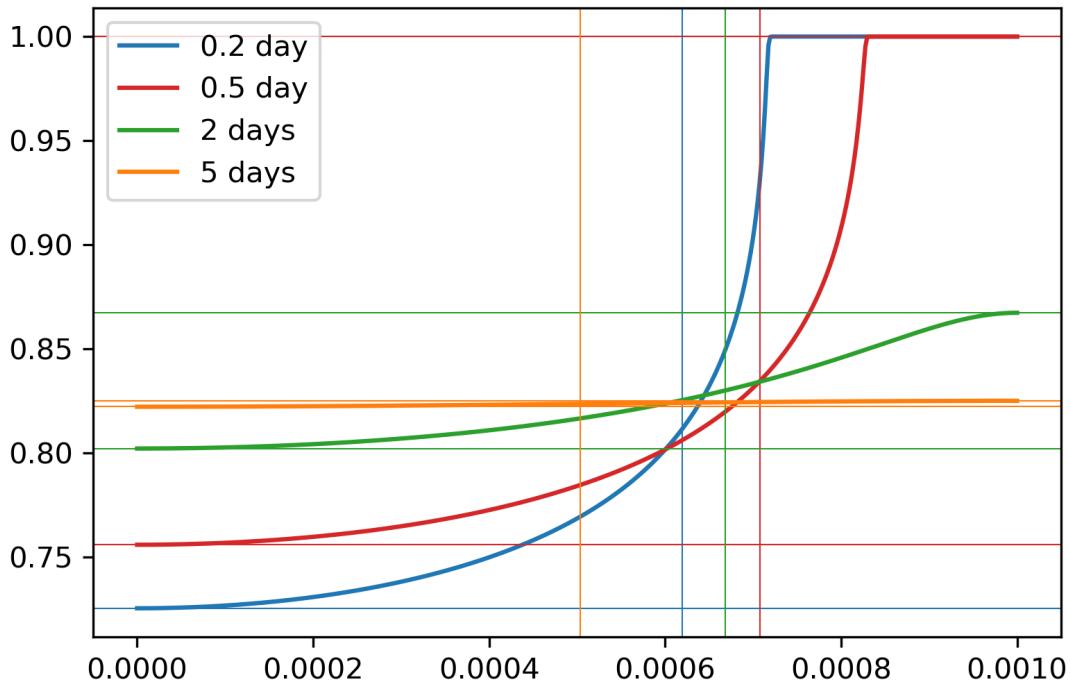


Figure 14: Al Concentration Profile and Matano Plane Positions at $t=0.2, 0.5, 2, 5$ days

At each annealing time: First I used the following formula to find the Matano plane position X_M :

$$X_M = \frac{1}{c(x=0.0) - c(x=0.001)} \int_{c(x=0.001)}^{c(x=0.0)} xdc \quad (1)$$

The code used to find Matano planes:

```
def matano(x, c):
    x = list(x)
    c = list(c)

    dc = [None] * (len(x)-1)

    sum_xdc = 0.0

    for i in range(0, len(dc)):
        dc[i] = c[i] - c[i-1]

    if dc[i] == 0:
        print(i)
        break
```

```

    sum_xdc = sum_xdc - (dc[i] * x[i+1])

    cL = c[0]
    cR = c[-1]

    x_M = sum_xdc / (cL - cR)

    return x_M

fig, ax = plt.subplots()

al = pd.read_csv("al_17280.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=17280'])
ax.plot(x, c, label="0.2 day", color='tab:blue')
x_M = matano(x, c)
ax.axvline(x=x_M, linewidth=0.5, color='tab:blue')
ax.axhline(y=c[0], linewidth=0.5, color='tab:blue')
ax.axhline(y=c[-1], linewidth=0.5, color='tab:blue')

al = pd.read_csv("al_43200.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=43200'])
ax.plot(x, c, label="0.5 day", color='tab:red')
x_M = matano(x, c)
ax.axvline(x=x_M, linewidth=0.5, color='tab:red')
ax.axhline(y=c[0], linewidth=0.5, color='tab:red')
ax.axhline(y=c[-1], linewidth=0.5, color='tab:red')

al = pd.read_csv("al_172800.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=172800'])
ax.plot(x, c, label="2 days", color='tab:green')
x_M = matano(x, c)
ax.axvline(x=x_M, linewidth=0.5, color='tab:green')
ax.axhline(y=c[0], linewidth=0.5, color='tab:green')
ax.axhline(y=c[-1], linewidth=0.5, color='tab:green')

al = pd.read_csv("al_432000.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=432000'])
ax.plot(x, c, label="5 days", color='tab:orange')
x_M = matano(x, c)
ax.axvline(x=x_M, linewidth=0.5, color='tab:orange')
ax.axhline(y=c[0], linewidth=0.5, color='tab:orange')
ax.axhline(y=c[-1], linewidth=0.5, color='tab:orange')

#ax.grid()
ax.legend()
fig.savefig("matano_planes.png", dpi=300)
plt.show()

```

Then I used the following formula to calculate the diffusivity:

$$D(c') = -\frac{1}{2t} \frac{\partial x}{\partial c} \Big|_{c'} \int_{c_0}^{c'} x dc \quad (2)$$

where c_0 is the concentration at $x = 0.001$ m.

Python code used to plot and analyze:

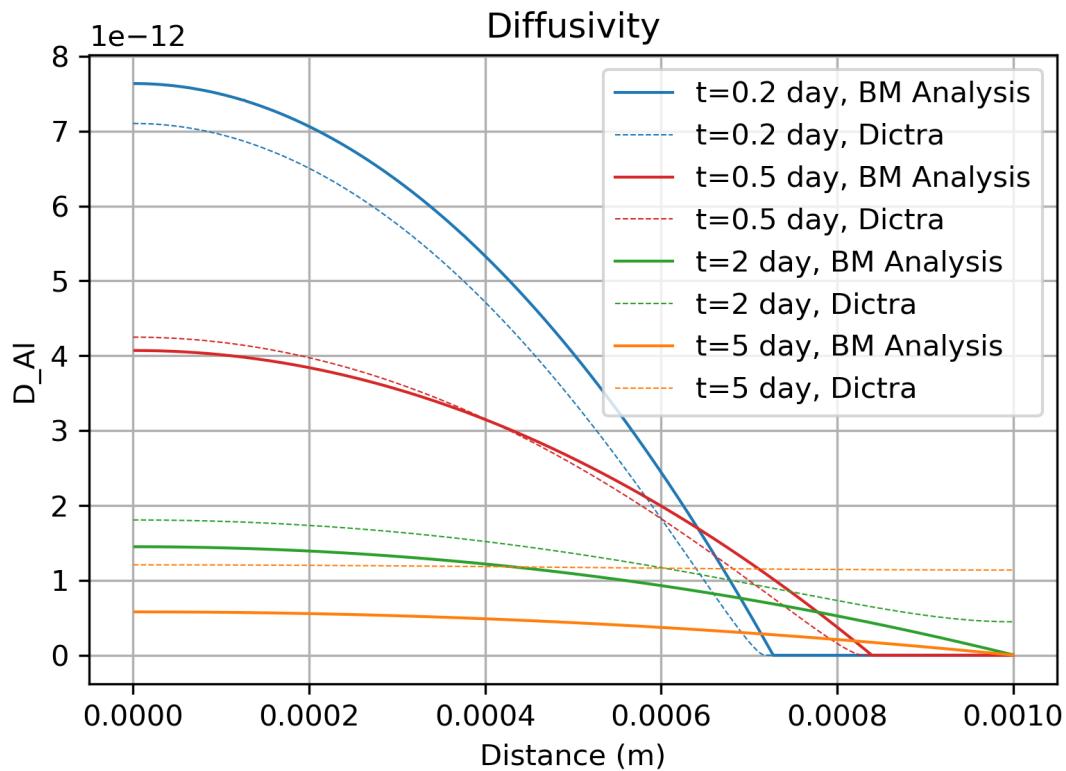


Figure 15: Al Chemical Diffusion Profile Comparison at $t=0.2, 0.5, 2, 5$ days

```

import pandas as pd
import numpy as np
import matplotlib
import matplotlib.pyplot as plt
%matplotlib inline

def dc_dx(x, c):
    x = list(x)
    c = list(c)
    dx = [None] * (len(x)-1)
    dc = [None] * (len(x)-1)
    cp = [None] * (len(x)-1)
    for i in range(0, len(x) - 1):
        dx[i] = x[i+1] - x[i]
        dc[i] = c[i+1] - c[i]
        cp[i] = dc[i] / dx[i]
    return x, c, cp

def int_x_dc(x, c, end_index):
    x = list(x)
    c = list(c)

```

```

dc = [None] * (len(x)-1)
int_x = [None] * (len(x))

int_x[0] = 0.0
max_index = end_index + 1
for i in range(1, len(x)):
    int_x[i] = 0.0
    if i > end_index:
        int_x[i] = 0.0
    else:
        for j in range(i, end_index + 1):
            dc[i] = c[i] - c[i-1]
            int_x[i] = int_x[i] - (dc[i] * x[j])
return int_x

fig, ax = plt.subplots()

t = 17280.0
al = pd.read_csv("al_17280.csv")

x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=17280'])
cd = [None] * (len(x) - 1)
end_index=len(c)-1
for j in range(0, len(c)):
    if c[j] == 1.0:
        end_index = j
        #print(j)
        break
int_x = int_x_dc(al['Distance [m]'], al['1: Time=17280'], end_index)
for i in range(0, (len(x) - 1)):
    if cp[i] == 0.0:
        cd[i] = 0.0
    else:
        cd[i] = -int_x[i+1] / cp[i] / (2.0 * t)

ax.plot(x[1:], cd, label="t=0.2 day, BM Analysis", linewidth=1.0, color='tab:blue')
diff = pd.read_csv("diff17280.csv")
ax.plot(x, list(diff['1: Time=17280']), '--', label="t=0.2 day, Dictra", linewidth=0.5,
        color='tab:blue')

t = 43200.0
al = pd.read_csv("al_43200.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=43200'])
end_index=len(c)-1
for j in range(0, len(c)):
    if c[j] == 1.0:
        end_index = j
        #print(j)
        break
int_x = int_x_dc(al['Distance [m]'], al['1: Time=43200'], end_index)
for i in range(0, (len(x) - 1)):
    if cp[i] == 0.0:
        cd[i] = 0.0

```

```

else:
    cd[i] = -int_x[i+1] / cp[i] / (2.0 * t)
ax.plot(x[1:], cd, label="t=0.5 day, BM Analysis", linewidth=1.0, color='tab:red')
diff = pd.read_csv("diff43200.csv")
ax.plot(x, list(diff['1: Time=43200']), '--', label="t=0.5 day, Dictra", linewidth=0.5,
        color='tab:red')

t = 172800.0
al= pd.read_csv("al_172800.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=172800'])
end_index=len(c)-1
for j in range(0, len(c)):
    if c[j] == 1.0:
        end_index = j
        #print(j)
        break
int_x = int_x_dc(al['Distance [m]'], al['1: Time=172800'], end_index)
for i in range(0, (len(x) - 1)):
    if cp[i] == 0.0:
        cd[i] = 0.0
    else:
        cd[i] = -int_x[i+1] / cp[i] / (2.0 * t)
ax.plot(x[1:], cd, label="t=2 day, BM Analysis", linewidth=1.0, color='tab:green')
diff = pd.read_csv("diff172800.csv")
ax.plot(x, list(diff['1: Time=172800']), '--', label="t=2 day, Dictra", linewidth=0.5,
        color='tab:green')

t = 432000.0
al = pd.read_csv("al_432000.csv")
x, c, cp = dc_dx(al['Distance [m]'], al['1: Time=432000'])
end_index=len(c)-1
for j in range(0, len(c)):
    if c[j] == 1.0:
        end_index = j
        #print(j)
        break
int_x = int_x_dc(al['Distance [m]'], al['1: Time=432000'], end_index)
for i in range(0, (len(x) - 1)):
    if cp[i] == 0.0:
        cd[i] = 0.0
    else:
        cd[i] = -int_x[i+1] / cp[i] / (2.0 * t)
ax.plot(x[1:], cd, label="t=5 day, BM Analysis", linewidth=1.0, color='tab:orange')
diff = pd.read_csv("diff432000.csv")
ax.plot(x, list(diff['1: Time=432000']), '--', label="t=5 day, Dictra", linewidth=0.5,
        color='tab:orange')

ax.set(xlabel='Distance (m)', ylabel='D_Al', title='Diffusivity')
ax.grid()
ax.legend()
fig.savefig("D_Al.png", dpi=300)
plt.show()

```

4. Examine Darken's equation based in the calculated tracer diffusivity, thermodynamic factor, and chemical diffusivity.

$$\tilde{D} = (x_B D_A^{Tr} + x_A D_B^{Tr})\varphi \quad (3)$$

x_A, x_B are the mole fractions of atom types A and B; D_A^{Tr}, D_B^{Tr} are the tracer diffusivities of atom types A and B; φ is the thermodynamic factor.

During the calculation, I am using the plots at same simulation condition and annealing time at 2 days.

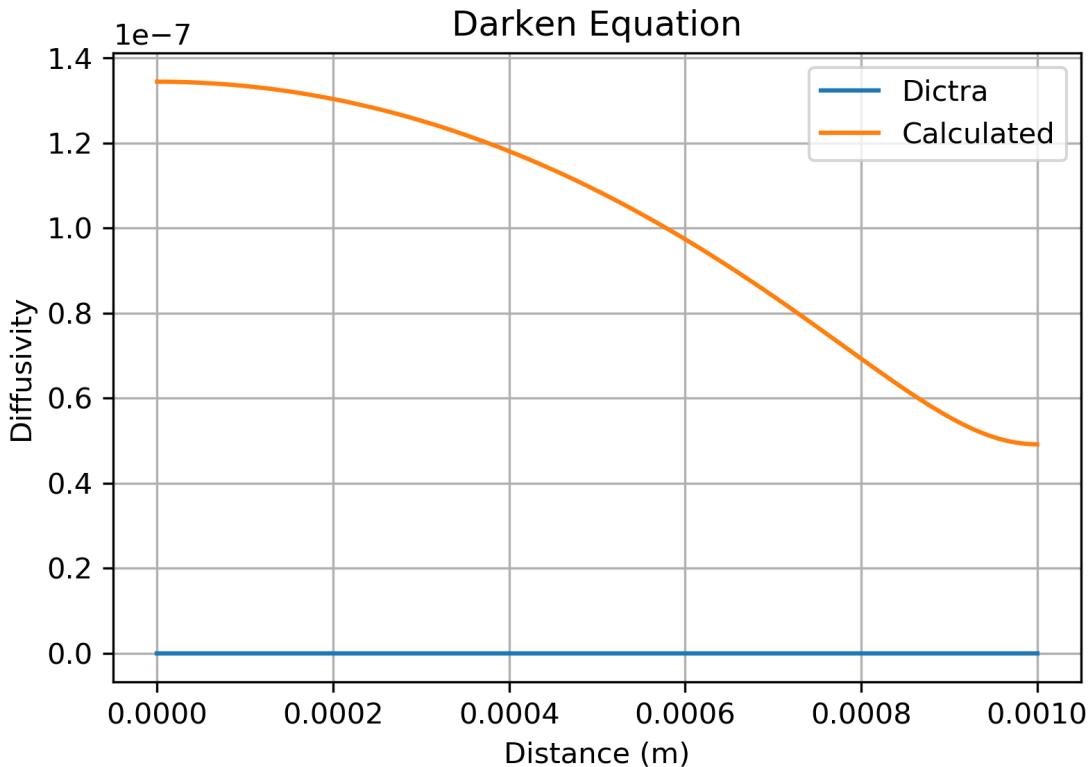


Figure 16: Dictra Calculated Diffusivity and Darken Calculated Difusivity

The plots are not in the same order of magnitude, but after I removed the thermodynamic factor,

$$\tilde{D} = x_B D_A^{Tr} + x_A D_B^{Tr} \quad (4)$$

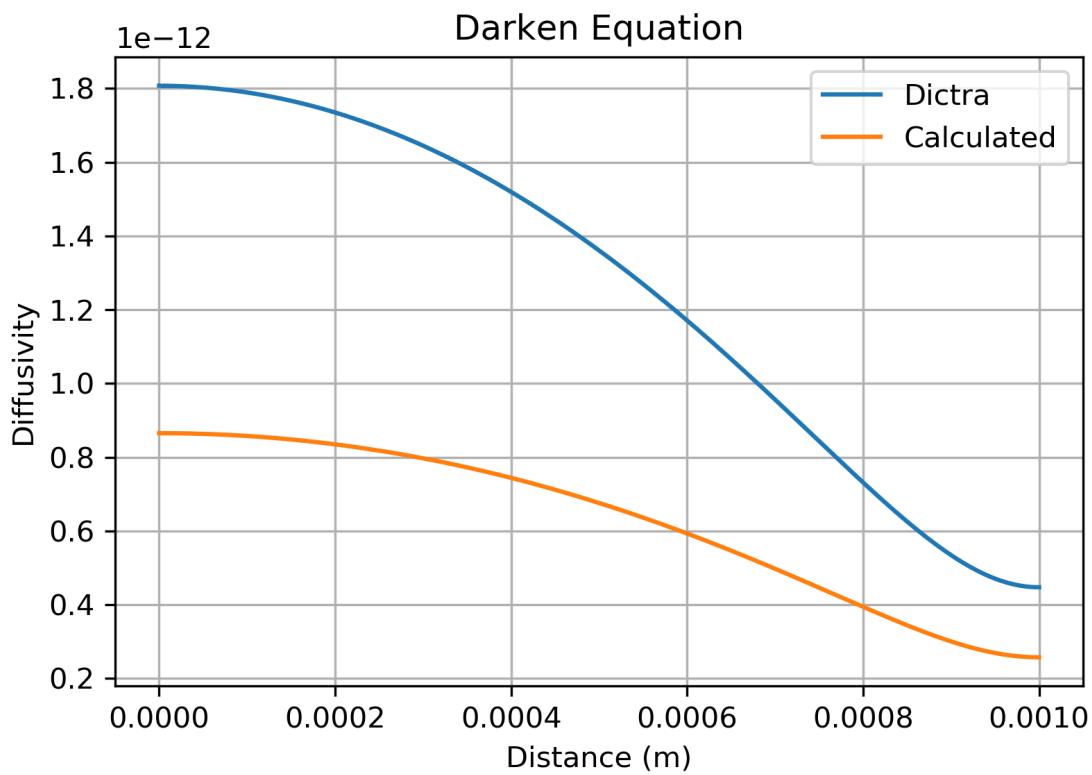


Figure 17: Dictra Calculated Diffusivity and Darken Calculated Difusivity

References

- [1] Thermocalc.com. [online] Available at:
https://www.thermocalc.com/media/19849/tcal5_extended_info.pdf [Accessed 9 Nov. 2019].