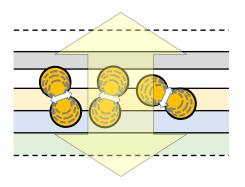
Internal Light Source-Power Density

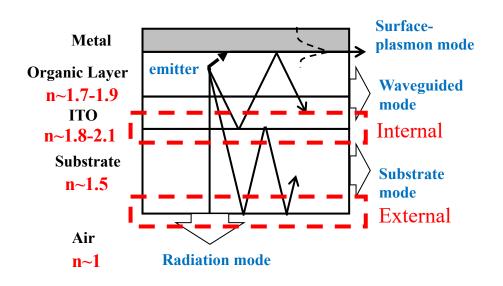
-PowerDensityCmd.pyc

Author: Wei-Kai Lee

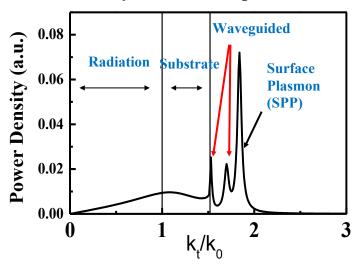


Objective

• To calculate the power density of the device.



Power Density/Power Dissipation



How to execute the calculator

python: windows python3: mac, linux

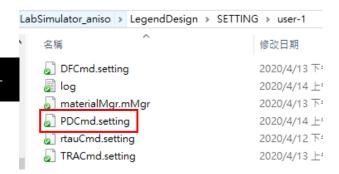
h>python PowerDensityCmd.pyc

Execution file

>>> Please insert username : user-1

Type user name

Power Density Setting is not built in ../../SETTING/user-1 Now saving the default power density setting.



```
>>> <u>?</u>
User Control Command
 --------

    Settting Command:

changeUser
                      exit
Material Manager Command

    Settting Command:

                      saveMgr
printMgr
Structure/Structure List Command
1. Structure List Command:
ReadStructListPath
SaveStructListPath
                     ReadStructListName
                      SaveStructListName
eadStructList
                      saveStructList
Structure Command:
ReadStructPath
                      ReadStructName
                     SaveFileName
deleteStruct
SaveFilePath
-ead$truct
Print Information Command:
                      printStructSettingInfo
print$tructInfo
.
printListInfo
4. Result Command:
                      ResultFileName
ResultFilePath
deleteResult
save_run_time_result_Bool
esetSN
```



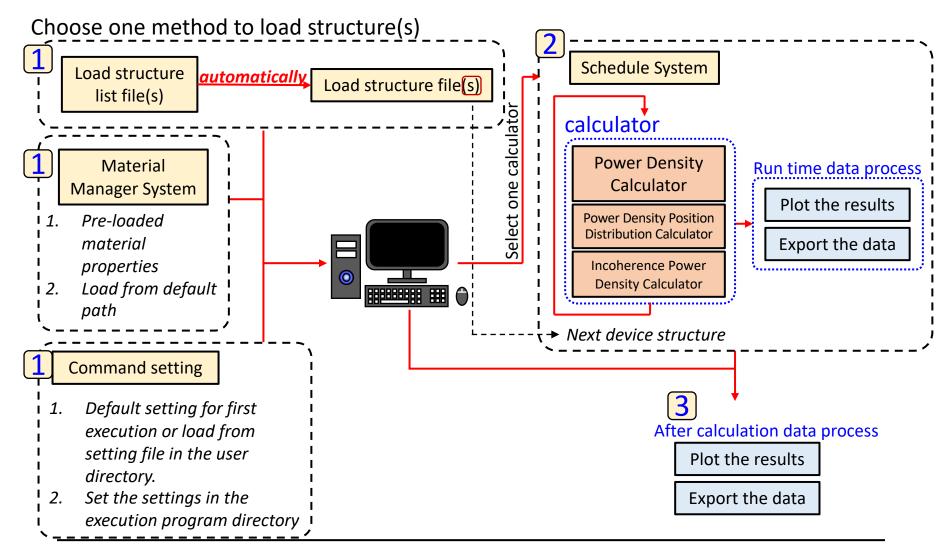
```
ower Density Command
    Notice the mode interference power !!!
 .. Settting Command:
 ettingFilePath
                     SettingFileName
etDefaultSetting
                     printPDInfo
.oadPDSETTING
                     savePDSETTING
Power Density in kxky domain:
                                              -Parameter
                     kyko
unPD_kxky
                     plotPDvskxky
                     writekxkyMatrix
plotPDkxkyContour
Power Density in k polar domain:
                                              -Parameter
                     Thetakt
unPD PolarInK
                     plotPDvskPolar
plotPDPolarContour
                     writekPolarMatrix
4. Power Density z in kxky domain:
                                                                ----Parameter
runPDz_kxky
                     pĺotPDvsZ kxkv
5. Power Density z in k polar domain:
                                                                  --Parameter
                     Thetakt
runPDz PolarInK
                     plotPDvsZ kPolar
6. Incoherence Layer Power Density in kxky domain:
                                          ----Parameter
unInCoPD_kxky
                     pĺotInCoPDvskxky
plotInCoPDkxkyContour writeInCokxkyMatrix
7. Incoherence Layer Power Density in k polar domain:
unInCoPD PolarInK
                     plotInCoPDvskPolar
```



```
Write Bool:
 -- Mode (related to Power Density/ Power Density z) ---
rite_TotalPowerBool
write_ForwardPowerBool
write_M1M3PowerBool
write_M1PowerBool
                                  write_BackwardPowerBool
                                  write_M2M4PowerBool
                                  write_M2PowerBool
write_M3PowerBool
                                  write M4PowerBool
 -- Region (Only related to Power Density) ---
                                 ---at the emission position
---at the position just above the emission position
⊬rite EMLBool
write_EML_zPlusBool
                                  ----at the position just below the emission position
vrite_EML_zMinusBool
write_SEMI_zPlusBool
write_SEMI_zMinusBool
                                  ----at the position just above the 1st interface of the device
                                  ----at the position just below the last interface of the device
9. Write Matrix Bool:
 -- Mode (related to Power Density/ Power Density z/ Incoherence Power Density) ---
writeMatrix_TotalPowerBool
writeMatrix_ForwardPowerBool
                                  writeMatrix_BackwardPowerBool
writeMatrix_M1M3PowerBool
                                  writeMatrix_M2M4PowerBool
                                  writeMatrix_M2PowerBool
vriteMatrix M1PowerBool
writeMatrix_M3PowerBool
                                  writeMatrix_M4PowerBool
 --- Region (Only related to Power Density) ---
                                  ----at the emission position
----at the position just above the emission position
vriteMatrix EMLBool
writeMatrix EML zPlusBool
                                  ----at the position just below the emission position
 riteMatrix_EML_zMinusBool
writeMatrix_SEMI_zPlusBool
                                  ----at the position just above the 1st interface of the device
riteMatrix SEMI zMinusBool
                                  ----at the position just below the last interface of the device
```

```
10. Plot Bool:
  changefigshowBool
     -- Mode (related to Power Density/ Power Density z/ Incoherence Power Density) ---
plot_TotalPowerBool
plot_ForwardPowerBool
                                                                                                                             plot_BackwardPowerBool
  olot M1M3PowerBool
                                                                                                                             plot M2M4PowerBool
   plot_MiPowerBool plot_M
 plot_M1PowerBool
plot M3PowerBool
                                                                                                                          ---at the emission position
---at the position just above the emission position
---at the position just below the emission position
plot EMLBool
 plot EML zPlusBool
plot_EML_zMinusBool
plot_SEMI_zPlusBool
                                                                                                                            ----at the position just above the 1st interface of the device
                                                                                                                             ----at the position just below the last interface of the device
plot_SEMI_zMinusBool
11. Run Time Bool:
 runtime plot
   --- (Only related to Power Density/ Incoherence Power Density)) ---
  runtime_write_matrix runtime_contourplot
```

Calculating Workflow



Default Setting

```
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd.setting
kt/ko : 0.00000:0.10000:1.00000
Theta in kt/ko domain : 0.00000:10.00000:360.00000
kx/ko : -1.00000:0.10000:1.00000
ky/ko : -1.00000:0.10000:1.00000
z(nm) : -50.00000:1.00000:50.00000
Save Run Time Result : True
Run Time Write Matrix : False
Run Time Plot : False
Run Time Contour Plot : False
Plot Region Bool:
                                  : True (EMLz+ add EMLz- )
       EML(EMLz+)
                                  : False (at the position just above emitting position)
       EML(EMLz-)
                                    False (at the position just below emitting position)
       +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
+z (M1+M2) : False
               (M3+M4)
                                     False
       -z
M1
M2
M3
M/
                                     False
                                     False
                                     False
                                    False
Write Region Bool:
                                     True (EMLz+ add EMLz- )
                                    True (at the position just above emitting position)
True (at the position just below emitting position)
       EML(EMLz+)
      EML(EMLz-)
       +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
       -Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
       Total (M1+M2+M3+M4)
                                     True
               (M1+M2)
(M3+M4)
       +z
                                     True
       -z
M1
M2
                                     True
                                     True
                                     True
                                     True
WriteMatrix Region Bool:
                                     True (EMLz+ add EMLz- )
       EML(EMLz+)
                                     False (at the position just above emitting position)
       EML(EMLz-) : False (at the position just below emitting position)
+Semi-infinite region: True (at the interface of the 1st semi-infinite region)
       -Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
               (M1+M2)
       +z
                                     False
               (M3+M4)
      -z
M1
M2
                                     False
                                    False
                                  : False
```

Setting

```
kt/ko : 0.00000:0.10000:1.00000
Theta in kt/ko domain : 0.00000:10.00000:360.00000
```

kt/ko and θ_{kt} setting method

```
<PDCmd> help ktko
Set the kt/ko. (Empty for delete setting)
Set the kt/ko. (Empty for delete setting)
[Usage] ktko [ktko] - single value, start:spacing:end, (v1,v2,v3,v4)

<PDCmd> help Thetakt
Set the polar angle in kx ky domain.
[Usage] Thetakt [theta] - single value, start:spacing:end, (v1,v2,v3,v4)
```

Setting

```
kx/ko : -1.00000:0.10000:1.00000
ky/ko : -1.00000:0.10000:1.00000
```

kx/ko and ky/ko setting method

```
<PDCmd> help kxko
Set the kx/ko. (Empty for delete setting)
[Usage] kxko [kxko] - single value, start:spacing:end, (v1,v2,v3,v4)

<PDCmd> help kyko
Set the ky/ko. (Empty for delete setting)
[Usage] kyko [kyko] - single value, start:spacing:end, (v1,v2,v3,v4)
```

```
z(nm): -50.00000:1.00000:50.00000
```

z setting method

```
<PDCmd> help z
Set the simulation position z. (Empty for delete setting)
[Usage] z [z] – single value, start:spacing:end, (v1,v2,v3,v4)
```

Run-Time Setting

Save Run Time Result : True

Whether to save the data in the memory after the calculation. If the user would like to execute plot or other data manipulation commands, the save-run-time-result bool should be "True". However, the user should notice the memory usage when scanning a lot of parameters.

Run Time Write Matrix : False

Whether to write the results when calculation.

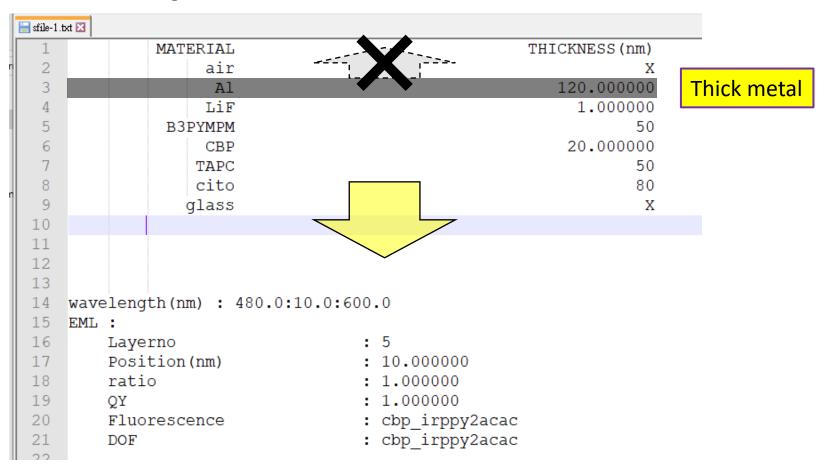
```
Run Time Write Matrix : False
Run Time Plot : False
Run Time Contour Plot : False
```

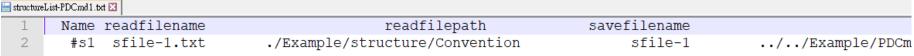
Whether to write the results when calculation.

Plot Setting

```
Plot Region Bool:
                           : True (EMLz+ add EMLz- )
     EML(EMLz+)
                            False (at the position just above emitting position)
                           : False (at the position just below emitting position)
     FMI(FMI z-)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
     Total (M1+M2+M3+M4)
                           · True
           (M1+M2)
     +Z
                             False
           (M3+M4)
                             False
     -z
M1
M2
M3
M4
                             False
                             False
                             False
                             False
Write Region Bool:
                             True (FMlz+ add FMlz-)
     \mathsf{EML}
     FMI (FMI 2+)
                             True (at the position just above emitting position)
                             True (at the position just below emitting position)
     FMI(FMI z-)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
     Total (M1+M2+M3+M4)
                             True
           (M1+M2)
                             True
     +Z
           (M3+M4)
     -z
M1
M2
M3
                              True
                             True
                             True
                             True
                             True
WriteMatrix Region Bool:
                           : True (EMLz+ add EMLz- )
     \mathsf{EML}
     EML(EMLz+)
                           : False (at the position just above emitting position)
                           : False (at the position just below emitting position)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
     Total (M1+M2+M3+M4)
                           : True
           (M1+M2)
                             False
     +Z
           (M3+M4)
                             False
     -z
M1
M2
                             False
                             False
```

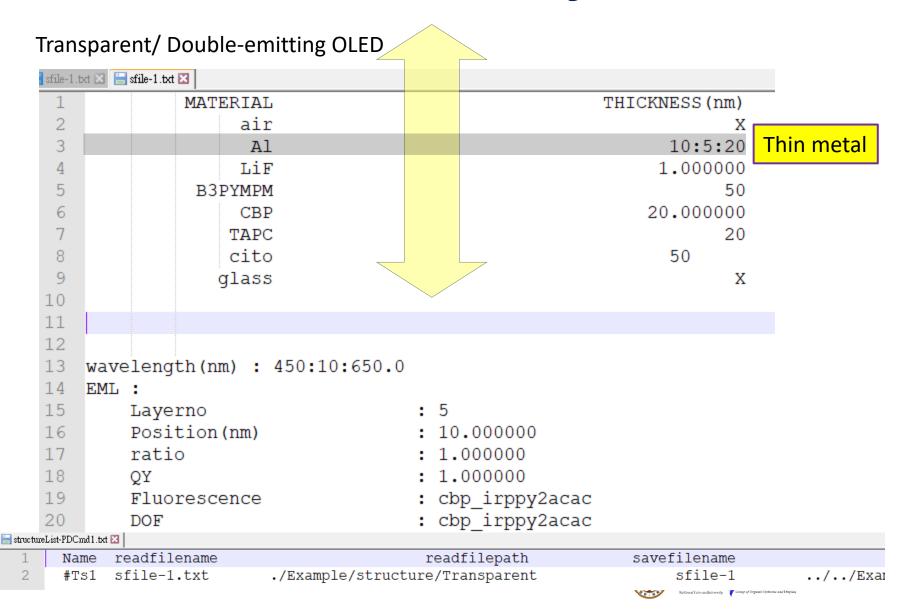
Bottom-emitting OLED











```
>>> changefigshowBool F
>>> save run time result Bool F
>>> ReadStructListPath ./Example/structure/Convention
>>> readStructList
Now reading structure list file ./Example/structure/Convention\structureList-PDCmd1.txt
                           savefilename CommandID Check
                                                                     readfilepath
              filename
 No./Name
           sfile-1.txt
                               sfile-1
                                            0.0
                                                   X ./Example/structure/Convention ../../Example/PDCmd-1
     #s1
Structure file reading...
Now reading structure file ./Example/structure/Convention\sfile-1.txt
>>> readStructList
Now reading structure list file ./Example/structure/Transparent\structureList-PDCmd1.txt
 No./Name
                           savefilename CommandID Check
                                                                      readfilepath
           sfile-1.txt
                               sfile-1
                                            0.0
                                                   X ./Example/structure/Transparent ../../Example/PDCmd-
     #Ts1
Structure file reading...
Now reading structure file ./Example/structure/Transparent\sfile-1.txt
```

```
>>> printStructInfo
*******<del>****</del>
     Material
                         Thickness(nm)
                        X
120.0
     B3PYMPM
                         50.0
     CBP
     TAPC
     cito
     glass
wavelength(nm) : 480.00000:10.00000:600.00000
                                   Position(nm)
                                                 PLQY
     Fluorescence
                                                                Ratio
[5] cbp_irppy2acac cbp_irppy2acac 10.0
                                                 1.0
                                                                1.0
Device number : 1
***********
Name: #Ts1
                         Thickness(nm)
     Material
                         10.00000:5.00000:20.00000
     B3PYMPM
     CBP
     TAPC
     cito
     glass
wavelength(nm) : 450.00000:10.00000:650.00000
                                   Position(nm)
                                                 PLQY
     Fluorescence
                                                                Ratio
[5] cbp_irppy2acac cbp_irppy2acac 10.0
                                                 1.0
                                                                1.0
Device number : 3
```

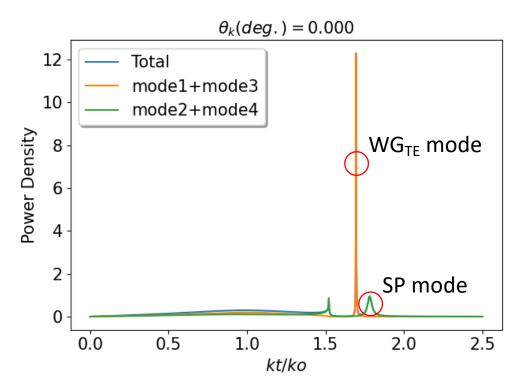
```
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd setting
kt/ko : 0.00000:0.00100:2.50000
Theta in kt/ko domain : 0.0
kx/ko : -1.00000:0.10000:1.00000
ky/ko : -1.00000:0.10000:1.00000
z(nm) : -50.00000:1.00000:50.00000
Save Run Time Result : False
Run Time Plot
Run Time Contour Plot : False
Plot Region Bool:
                                : True (EMLz+ add EMLz- )
      EML
      EML(EMLz+)
                                : False (at the position just above emitting position)
EML(EMLz-) : False (at the position just below emitting position)
+Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
      Total (M1+M2+M3+M4)
                               : True
             (M1+M2)
(M3+M4)
                                   False
      +Z
                                   False
                                   False
                                   False
                                   False
                                   False
Write Region Bool:
                                 : True (EMLz+ add EMLz- )
      EML(EMLz+)
                                  True (at the position just above emitting position)
                                 : True (at the position just below emitting position)
      +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
      -Semi-infinite region: True (at the interface of the last semi-infinite region)
```

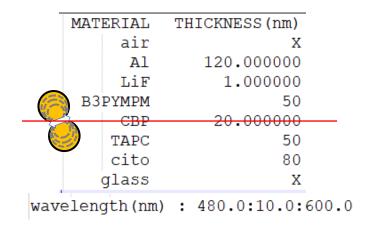
```
Write Mode Bool: (should be cosidered with pl<u>ot region bool)</u>
      Total (M1+M2+M3+M4)
                                 True
             (M1+M2)
                                 True
      +z
     -z
M1
M2
M3
             (M3+M4)
                                 True
                                 True
                                 True
                                 True
                                 True
WriteMatrix Region Bool:
                               : True (EMLz+ add EMLz- )
                                False (at the position just above emitting position)
False (at the position just below emitting position)
     EML(EMLz+)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
             (M1+M2)
      +z
                                 False
             (M3+M4)
                                 False
     -z
M1
M2
M3
M4
                                 False
                                 False
                                 False
                                 False
Figure Show Bool : False
>>> runPD_PolarInK
Now running structure (#1/#2) #s1 ...
|Device-#13/#13|(100%)>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/0min(s)/26sec(s)
Now running structure (#2/#2) #Ts1 ...
|Device-#63/#63|(100%)>>>>>>>>>>>>|Elapsed:0dav(s)/0hr(s)/2min(s)/3sec(s)
```

Data at EML

Bottom-emitting OLED

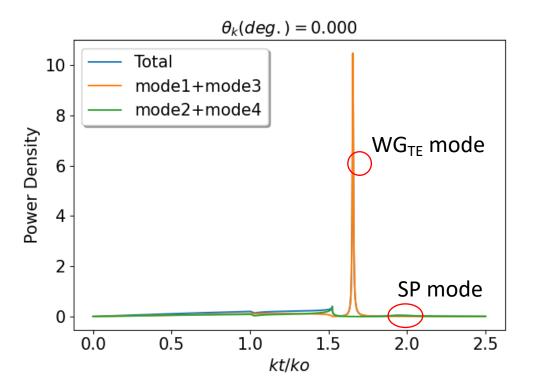
480 nm

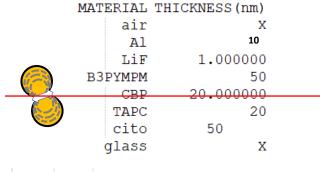




Data at EML

Transparent/ Double-emitting OLED 480 nm





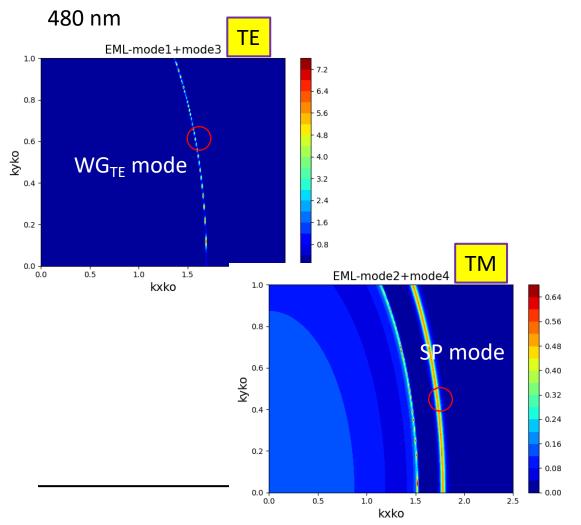
wavelength(nm): 480.0:10.0:600.0

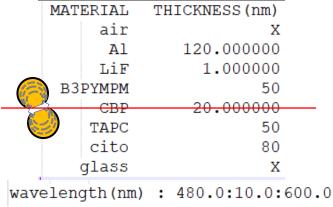
```
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd.setting
kt/ko : 0.00000:0.00100:2.50000
Theta in kt/ko domain · A A
kx/ko : 0.00000:0.01000:2.50000
kv/ko : 0.00000:0.01000:1.00000
z(nm) : -50.00000:1.00000:50.00000
Save Run Time Result : False
Run Time Write Matrix : True
Run Time Plot
                          False
Run Time Contour Plot : True
Plot Region Bool:
                              True (EMLz+ add EMLz- )
                            : False (at the position just above emitting position)
     EML(EMLz+)
                            : False (at the position just below emitting position)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
+z (M1+M2) : False
            (M3+M4)
     -z
M1
M2
M3
                              False
                              False
                              False
                              False
Write Region Bool:
     EML EMLz+)
                            : True (EMLz+ add EMLz- )
                              True (at the position just above emitting position)
                            : True (at the position just below emitting position)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
            (M1+M2)
                               True
     -z
M1
M2
M3
            (M3+M4)
                               True
                               True
                               True
                               True
                               True
```

```
WriteMatrix Region Bool:
                               True (EMLz+ add EMLz- )
                               False (at the position just above emitting position) False (at the position just below emitting position)
     EML(EMLz+)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
     Total (M1+M2+M3+M4)
                               True
            (M1+M2)
                                False
     -z
M1
M2
M3
            (M3+M4)
                                False
                               False
                                False
                               False
                                False
Figure Show Bool : False
>>> runPD_kxky
Now running structure (#1/#2) #s1 ...
|Device-#13/#13|(100%)>>>>>>>>>>>>|Elapsed:Oday(s)/Ohr(s)/3min(s)/13sec(s)
Now running structure (#2/#2) #Ts1 ...
|Device-#63/#63|(100%)>>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/15min(s)/49sec(s)
>>> deleteStruct
```

Data at EML

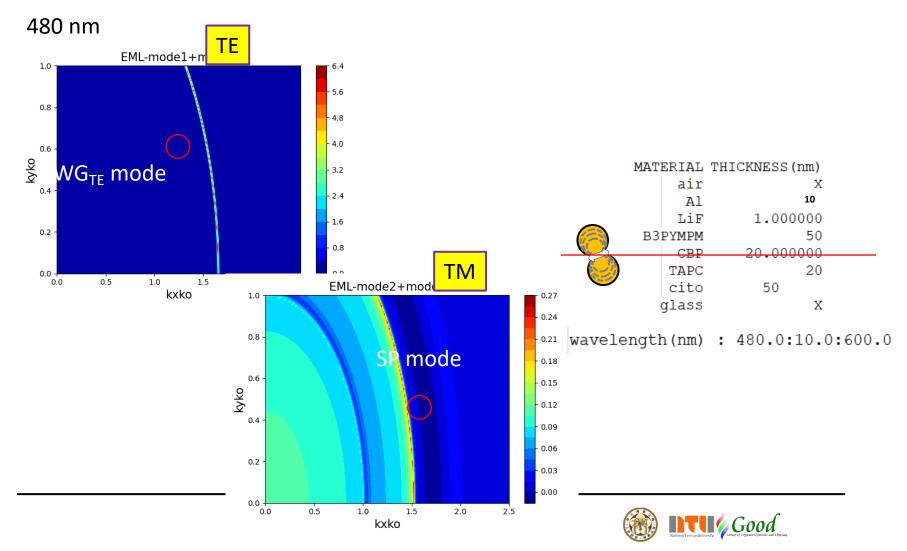
Bottom-emitting OLED



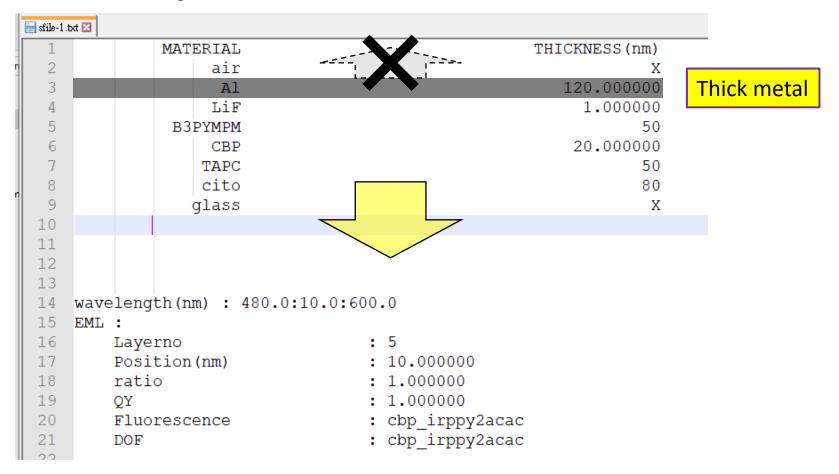


Data at EML

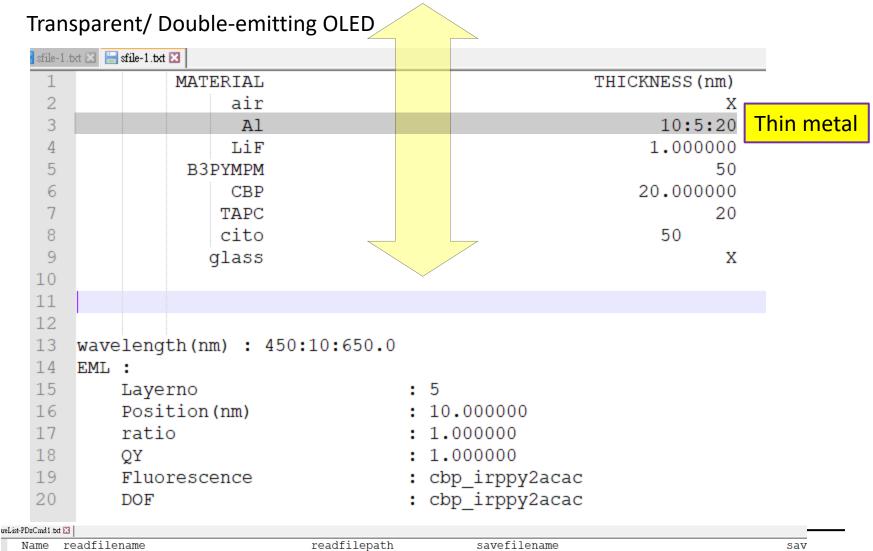
Transparent/ Double-emitting OLED



Bottom-emitting OLED



eList-PDzCmd1.txt 🔀



sfile-1

```
>>> changefigshowBool F
>>> save run time result Bool F
>>> ReadStructListPath ./Example/structure/Convention
>>> read$tructList
Now reading structure list file ./Example/structure/Convention\structureList-PDzCmd1.txt
                           savefilename CommandID Check
                                                                     readfilepath
 No./Name
                                            0.0
           sfile-1.txt
                               sfile-1
                                                   X ./Example/structure/Convention ../../Example/PDzCmd-1/Conven
     #s1
Structure file reading...
Now reading structure file ./Example/structure/Convention\sfile-1.txt
>>> readStructList
Now reading structure list file ./Example/structure/Transparent\structureList-PDzCmd1.txt
                           savefilename CommandID Check
 No./Name
     #Ts1
           sfile-1.txt
                               sfile-1
                                            0.0
                                                   X ./Example/structure/Transparent ../../Example/PDzCmd-1/Trans
Structure file reading...
Now reading structure file ./Example/structure/Transparent\sfile-1.txt
```

```
Structure file reading...
Now reading structure file ./Example/structure/Transparent\sfile-1.txt
>>> printStructInfo
     ************
     Material
                         Thickness(nm)
                         X
120.0
     air
     ВЭРУМРМ
     TAPC
     cito
     glass
wavelength(nm) : 480.00000:10.00000:600.00000
     Fluorescence
                                   Position(nm)
                                                  PLOY
                                                                 Ratio
                                                  1.0
   cbp_irppy2acac cbp_irppy2acac 10.0
                                                                 1.0
Device number : 1
***********
Name: #Ts1
     Material
                         Thickness(nm)
                         10.00000:5.00000:20.00000
     LiF
     ВЭРУМРМ
     TAPC
     cito
     glass
wavelength(nm) : 450.00000:10.00000:650.00000
                                   Position(nm)
                                                  PLOY
     Fluorescence
                                                                 Ratio
     cbp_irppy2acac cbp_irppy2acac 10.0
                                                  1.0
                                                                 1.0
Device number : 3
```

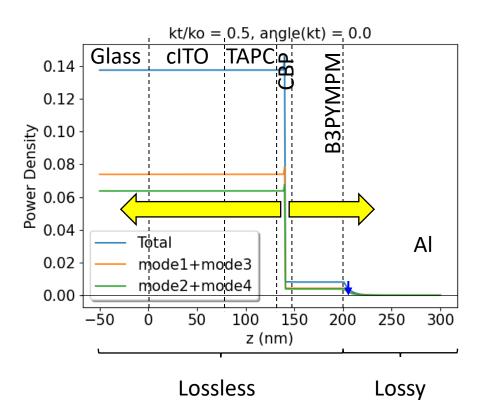
```
>>> ktko (0.5,1,1.5)
>>> Thetakt 0
>>> z -50:1:300
>>> runtime_plot T
```

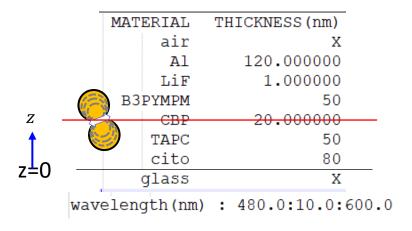
```
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd.setting
kt/ko : 0.50000:0.50000:1.50000
Theta in kt/ko domain : 0.0
kx/ko : 0.00000:0.01000:2.50000
ku/ko · A AAAAA · A1AAA · 1 AAAAA
z(nm) : -50.00000:1.00000:300.00000
Save Run lime Result : False
Run Time Write Matrix : True
Run Time Plot
                         True
Run Time Contour Plot : True
Plot Region Bool:
                           : True (EMLz+ add EMLz- )
     EML(EMLz+)
                             False (at the position just above emitting position)
     EML(EMLz-)
                             False (at the position just below emitting position)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
     Total (M1+M2+M3+M4)
                           : True
           (M1+M2)
(M3+M4)
                             False
     +Z
                              False
     -z
M1
M2
M3
M4
                            : False
                              False
                             False
                            : False
Write Region Bool:
                           : True (EMLz+ add EMLz- )
     EML(EMLz+)
                            : True (at the position just above emitting position)
                            : True (at the position just below emitting position)
     EML(EMLz-)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
            (M1+M2)
                             True
     -z
M1
M2
M3
            (M3+M4)
                             True
                              True
                             True
                             True
                              True
```

```
WriteMatrix Region Bool:
                                         True (EMLz+ add EMLz- )
       \mathsf{EML}
                                         False (at the position just above emitting position)
       EML(EMLz-) : False (at the position just below emitting position) +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
       -Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
                (M1+M2)
                                         False
       -z
M1
M2
                (M3+M4)
                                         False
                                         False
                                         False
                                         False
                                         False
Figure Show Bool : False
>>> runPDz_PolarInK
Now running structure (#1/#2) #s1 ...
|Device-#13/#13|(100%)>>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/0min(s)/12sec(s)
Now running structure (#2/#2) #Ts1 ...
|Device-#63/#63|(100%)>>>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/1min(s)/2sec(s)
>>> deleteStruct
```

Data at EML

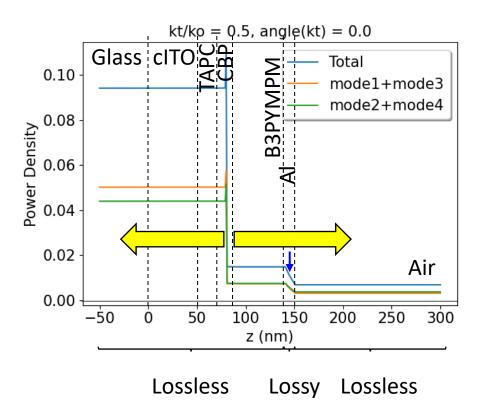
Bottom-emitting OLED 480 nm

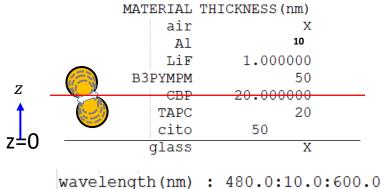




Data at EML

Transparent/ Double-emitting OLED 480 nm





```
>>> changefigshowBool F
>>> save_run_time_result_Bool F
>>> ReadStructListPath ./Example/structure/Convention
>>> readStructList
Now reading structure list file ./Example/structure/Convention\structureList-PDIncohCmd1.txt
                               savefilename CommandID Check
                                                        X ./Example/structure/Convention ../../Example/PDIncohCmd-1/Convent
      #s1 sfile-1-air.txt
                                   sfile-1
                                                 0.0
Structure file reading...
Now reading structure file ./Example/structure/Convention\sfile-1-air.txt
>>> ReadStructListPath ./Example/structure/Transparent
>>> readStructList
Now reading structure list file ./Example/structure/Transparent\structureList-PDIncohCmd1.txt
                               savefilename CommandID Check
 No./Name
                                                0.0
                                                        X ./Example/structure/Transparent ../../Example/PDIncohCmd-1/Transparent
    #Tsal sfile-1-air.txt
                                sfile-1-air
Structure file reading...
Now reading structure file ./Example/structure/Transparent\sfile-1-air.txt
```

```
🔚 sfile-1-air.txt 🔀
               MATERIAL
                                                          THICKNESS (nm)
                     air
                      A 1
                                                              120.000000
  4
                     LiF
                                                                1.000000
                B3PYMPM
                                                                       50
                                                               20.000000
                     CBP
                    TAPC
                                                                       50
                    cito
                                                                       80
  9
                   glass
 10
                     air
                                                                        Χ
 11
 12
 13
 14
     wavelength(nm): 480.0:10.0:600.0
 15
     EML :
 16
                                       : 5
          Layerno
 17
          Position (nm)
                                       : 10.000000
 18
          ratio
                                       : 1.000000
                                       : 1.000000
 19
          ΟY
 20
          Fluorescence
                                       : cbp irppy2acac
 21
                                       : cbp irppy2acac
          DOF
```

```
📑 sfile-1-air.txt 🗵 📙 sfile-1-air.txt 🗵
                                                            THICKNESS (nm)
               MATERIAL
                     air
                      Αl
                                                                   10:5:20
                     LiF
  4
                                                                  1.000000
  5
                                                                         50
                B3PYMPM
  6
                                                                 20.000000
                     CBP
                                                                         20
                    TAPC
                    cito
 8
                                                                   50
  9
                   glass
                                                                          Χ
                                                                          Χ
10
                     air
11
12
     wavelength(nm): 450:10:650.0
14
     EML:
15
                                        : 5
         Layerno
         Position (nm)
16
                                        : 10.000000
17
         ratio
                                        : 1.000000
18
                                        : 1.000000
         QΥ
19
                                        : cbp irppy2acac
         Fluorescence
20
                                        : cbp irppy2acac
         DOF
```

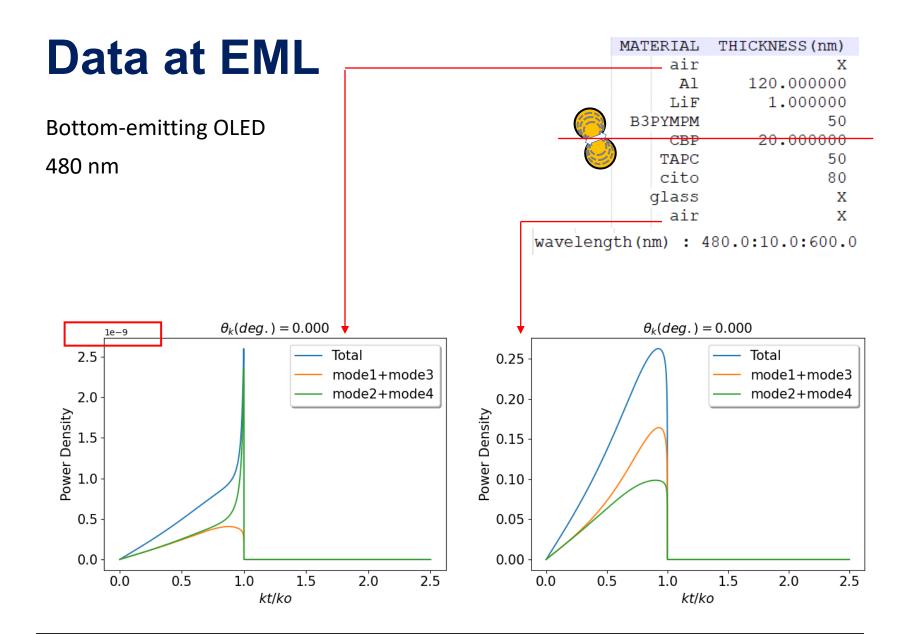
```
>>> printStructInfo
      Material
                            Thickness(nm)
                            X
120.0
      air
      A1
      LiF
                            1.0
      ВЗРУМРМ
      TAPC
                            50.0
      cito
                            80.0
      glass
[9]
      air
wavelength(nm) : 480.00000:10.00000:600.00000
                                        Position(nm)
                                                         PLQY
                                                                          Ratio
      Fluorescence
                       DOF
[5] cbp_irppy2acac cbp_irppy2acac 10.0
Device number : 1
                                                         1.0
                                                                          1.0
Name: #Tsa1
                            Thickness(nm)
      Material
      air
                            10.00000:5.00000:20.00000
      B3PYMPM
      TAPC
      glass
[9]
      air
wavelength(nm) : 450.00000:10.00000:650.00000
                                        Position(nm)
                                                         PLOY
      Fluorescence
                                                                          Ratio
     cbp_irppy2acac cbp_irppy2acac 10.0
                                                         1.0
                                                                          1.0
Device number : 3
```

```
>>> ktko
               0:0.001:2.5
>>> Thetakt 0
>>> runtime write matrix F
>>> runtime plot
>>> runtime contourplot F
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd.setting
kt/ko : 0.00000:0.00100:2.50000
Theta in kt/ko domain : 0.0
kx/ko : 0.00000:0.01000:2.50000
ky/ko : 0.00000:0.01000:1.00000
z(nm) : -50.00000:1.00000:300.00000
Save Run Time Result : False
Run Time Write Matrix : False
Run Time Plot
Run Time Contour Plot : False
Plot Region Bool:
                              True (EMLz+ add EMLz- )
                            : False (at the position just above emitting position)
: False (at the position just below emitting position)
     EML(EMLz+)
     EML(EMLz-)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
            (M1+M2)
                              False
     +z
            (M3+M4)
                              False
                              False
                              False
                              False
                              False
Write Region Bool:
                              True (EMLz+ add EMLz- )
     EML(EMLz+)
                              True (at the position just above emitting position)
                              True (at the position just below emitting position)
     EML(EMLz-)
     +Semi-infinite region: True (at the interface of the 1st semi-infinite region)
     -Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
            (M1+M2)
                               True
     +Z
     -z
M1
M2
            (M3+M4)
                              True
                               True
                               True
```

```
Write Mode Bool: (should be cosidered with plot region bool)
       Total (M1+M2+M3+M4)
+z (M1+M2)
-z (M3+M4)
                                        True
       -z
M1
M2
M3
                                         True
                                         Irue
                                         True
                                        True
WriteMatrix Region Bool:
                                      : True (EMLz+ add EMLz- )
                                        False (at the position just above emitting position)
       EML(EMLz-) : False (at the position just above emitting position)
+Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
       Total (M1+M2+M3+M4)
                (M1+M2)
(M3+M4)
                                        False
                                        False
                                        False
                                        False
                                        False
Figure Show Bool : False
```

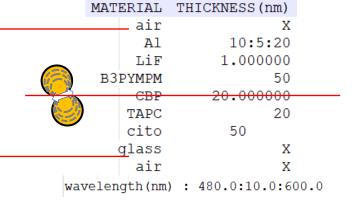
```
>>> runInCoPD_PolarInK

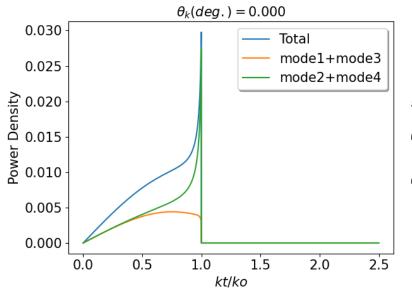
Now running structure (#1/#2) #s1 ...
|Device-#13/#13|(100%)>>>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/0min(s)/45sec(s)
Now running structure (#2/#2) #Tsa1 ...
|Device-#63/#63|(100%)>>>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/3min(s)/37sec(s)
```

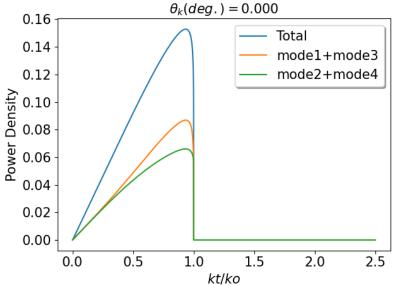


Data at EML

Transparent/ Double-emitting OLED 480 nm



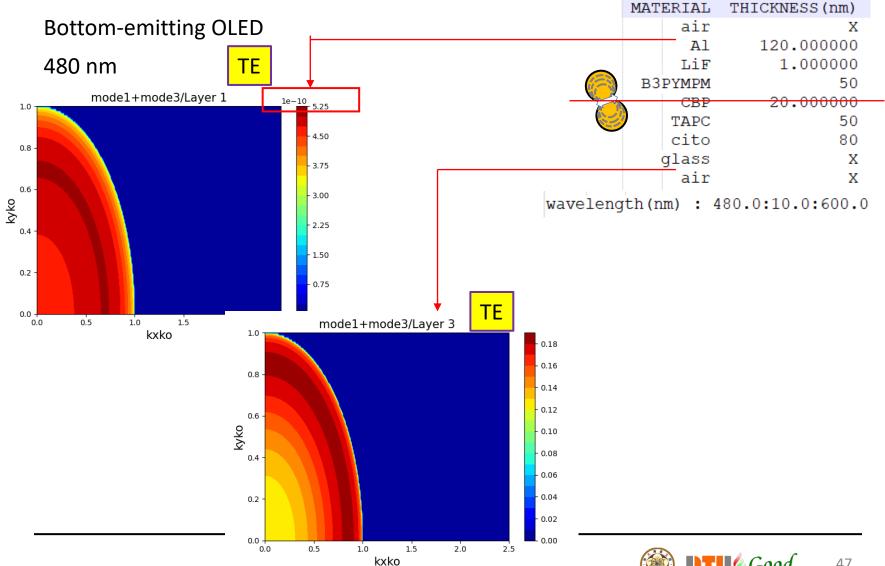


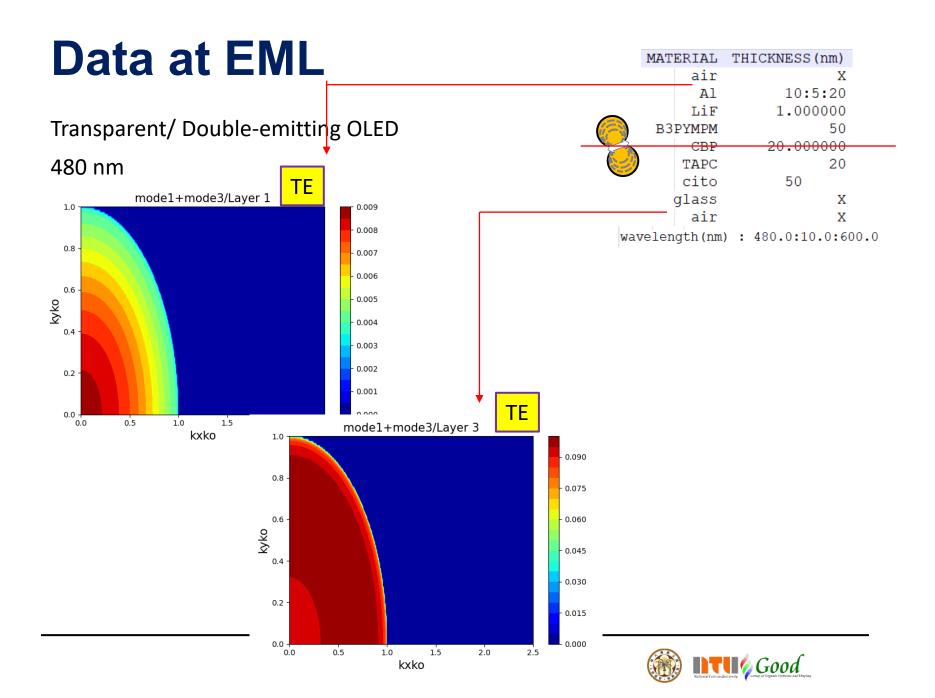


```
>>> kxko
                     0:0.01:2.5
>>> kuko
                     0:0.01:1
>>> runtime_write_matrix T
>>> runtime_plot
>>> runtime_contourplot T
>>> printPDInfo
Setting file path : ../../SETTING/user-1
Setting file name : PDCmd.setting
kt/ko : 0.00000:0.00100:2.50000
Theta in kt/ko domain : 0.0
kx/ko : 0.00000:0.01000:2.50000
ky/ko : 0.00000:0.01000:1.00000
z(nm) : -50.00000:1.00000:300.00000
Save Run Time Result : False
Run Time Write Matrix : True
Run Time Plot
                                    False
Run Time Contour Plot : True
Plot Region Bool:
                                        : True (EMLz+ add EMLz- )
       EML(EMLz+) : False (at the position just above emitting position)
EML(EMLz-) : False (at the position just below emitting position)
+Semi-infinite region: Irue (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
Plot Mode Bool: (should be cosidered with plot region bool)
Total (M1+M2+M3+M4) : True
+z (M1+M2) : False
-z (M3+M4) : False
        -z
M1
M2
M3
M4
                                           False
                                           False
                                          False
                                           False
```

```
Write Region Bool:
                                    True (EMLz+ add EMLz- )
                                    True (at the position just above emitting position)
      EML(EMLz+)
                                    True (at the position just below emitting position)
      EML(EMLz-)
+Semi-infinite region: True (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
Write Mode Bool: (should be cosidered with plot region bool)
      Total (M1+M2+M3+M4)
                                    True
              (M1+M2)
(M3+M4)
      +Z
                                    True
                                    True
                                     True
                                     True
                                    True
WriteMatrix Region Bool:
                                  : True (EMLz+ add EMLz- )
      EML(EMLz+)
                                  : False (at the position just above emitting position)
      EML(EMLz-) : False (at the position just below emitting position) +Semi-infinite region: Irue (at the interface of the 1st semi-infinite region)
-Semi-infinite region: True (at the interface of the last semi-infinite region)
WriteMatrix Mode Bool: (should be cosidered with plot region bool)
      Total (M1+M2+M3+M4)
              (M1+M2)
(M3+M4)
                                    False
                                    False
                                    False
                                    False
                                    False
                                   False
Figure Show Bool : False
>>> runInCoPD_kxky
Now running structure (#1/#2) #s1 ...
|Device-#13/#13|(100%)>>>>>>>>>>>|Elapsed:0day(s)/0hr(s)/6min(s)/43sec(s)
Now running structure (#2/#2) #Tsa1 ...
|Device=#63/#63|(100%)>>>>>>>>>>>>|Elapsed:0dav(s)/0hr(s)/32min(s)/9sec(s)
```

Data at EML





Calculate Device Distribution

Exit the material manager system.

*** The material manager system would be automatically saved into the user's setting directory.

>>> exit

End running GOODLAB anisotropic simulator ver1.0 Tue Apr 14 10:56:58 2020 Elapsed time : 0 day(s)/ 1 hr(s)/ 6 min(s)/ 3.5559067726135254 sec(s)