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**pmcpy**  
*Release 0.0.2*

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## OVERVIEW

<b>1</b>	<b>About</b>	<b>3</b>
1.1	Introduction . . . . .	3
1.2	Relevant Publications . . . . .	3
<b>2</b>	<b>Installation</b>	<b>5</b>
<b>3</b>	<b>pmcpy quickstart</b>	<b>7</b>
3.1	load the raw output file from PartMC . . . . .	7
3.2	number concentration . . . . .	7
3.3	aerosol mass concentration of selected species . . . . .	7
3.4	overall mass concentration . . . . .	8
3.5	aerosol density . . . . .	8
3.6	gas mixing ratio . . . . .	9
3.7	mixing state index calculation . . . . .	9
3.8	particle size distribution visualization (number concentration v.s. diameter) . . . . .	9
3.9	black carbon fraction in dry particles . . . . .	11
3.10	black carbon core volume . . . . .	12
<b>4</b>	<b>mixing state calculator</b>	<b>13</b>
<b>5</b>	<b>benchmarking</b>	<b>15</b>
<b>6</b>	<b>format of PartMC raw output</b>	<b>19</b>
<b>7</b>	<b>particles visualization</b>	<b>21</b>
<b>8</b>	<b>How to ask for help</b>	<b>23</b>



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## 1.1 Introduction

This package is used to post-process [PartMC](#) data (see the [reference](#) here). Additionally, a [mixing state calculator](#) can be used to calculate the mixing state index.

You can play with the mixing state calculator [HERE](#)

## 1.2 Relevant Publications

- Zheng, Z., Curtis, J. H., Yao, Y., Gasparik, J. T., Anantharaj, V. G., Zhao, L., et al. (2021). Estimating submicron aerosol mixing state at the global scale with machine learning and Earth system modeling. *Earth and Space Science*, 8, e2020EA001500. <https://doi.org/10.1029/2020EA001500>
- Zheng, Z., West, M., Zhao, L., Ma, P.-L., Liu, X., and Riemer, N.: Quantifying the structural uncertainty of the aerosol mixing state representation in a modal model, *Atmos. Chem. Phys.*, 21, 17727–17741, <https://doi.org/10.5194/acp-21-17727-2021>, 2021.
- Riemer, N. and West, M.: Quantifying aerosol mixing state with entropy and diversity measures, *Atmos. Chem. Phys.*, 13, 11423–11439, <https://doi.org/10.5194/acp-13-11423-2013>, 2013.





## INSTALLATION

Step 1: create a conda environment

```
$ conda create -n pmcpy python=3.8
$ conda activate pmcpy
$ conda install -c conda-forge numpy pandas xarray netcdf4
```

Step 2: install using pip

```
$ pip install pmcpy
```

(optional) install from source::

```
$ git clone https://github.com/zhonghua-zheng/pmcpy.git
$ cd pmcpy
$ python setup.py install
```



## PMCPY QUICKSTART

```
[1]: # import necessary package
import pmcpy
import xarray as xr
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

### 3.1 load the raw output file from PartMC

```
[2]: # define the path to data
p = "../data/urban_plume_0001_00000002.nc"
pmc = pmcpy.load_pmc(p)
```

### 3.2 number concentration

```
[3]: print("overall number conc.:", pmc.get_num_conc(), "# m^{-3}")
print("diameter<=2.5um:", pmc.get_num_conc(pmc.get_aero_particle_diameter()<=2.5e-6), "# m^{-3}")
print("diameter>=2.5um:", pmc.get_num_conc(pmc.get_aero_particle_diameter()>=2.5e-6), "# m^{-3}")
```

```
overall number conc.: 8854807162.459946 # m^{-3}
diameter<=2.5um: 8854806282.274422 # m^{-3}
diameter>=2.5um: 880.1855249722205 # m^{-3}
```

### 3.3 aerosol mass concentration of selected species

```
[4]: # consider BC and OC only
aero_species_ls = ["BC","OC"]
print("BC and OC mass conc.:",
      pmc.get_aero_mass_conc(aero_species_ls),
      "kg m^{-3}")
```

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```
# consider all species and dry species
all_aero_species = pmc.aero_species.copy()
dry_aero_species = all_aero_species.copy()
dry_aero_species.remove("H2O")

print("overall mass conc.:",
      pmc.get_aero_mass_conc(all_aero_species).sum(),
      "kg m^{-3}")
print("overall dry mass conc.:",
      pmc.get_aero_mass_conc(dry_aero_species).sum(),
      "kg m^{-3}")
print("PM2.5 mass conc. (with water):",
      pmc.get_aero_mass_conc(all_aero_species,
                             part_cond=(pmc.get_aero_particle_diameter()<=2.5e-6)).sum(),
      "kg m^{-3}")
```

BC and OC mass conc.: [6.53030409e-10 5.73984931e-09] kg m^{-3}  
 overall mass conc.: 2.2186891469474523e-08 kg m^{-3}  
 overall dry mass conc.: 1.316284475347949e-08 kg m^{-3}  
 PM2.5 mass conc. (with water): 2.2164289553077785e-08 kg m^{-3}

### 3.4 overall mass concentration

```
[5]: print("overall mass conc.:",
          pmc.get_mass_conc(dry=False), "kg m^{-3}")
      print("overall dry mass conc.:",
            pmc.get_mass_conc(dry=True), "kg m^{-3}")
      print("PM2.5 mass conc. (with water):",
            pmc.get_mass_conc(dry=False, part_cond=pmc.get_aero_particle_diameter()<=2.5e-6),
            "kg m^{-3}")
```

overall mass conc.: 2.2186891469474523e-08 kg m^{-3}  
 overall dry mass conc.: 1.316284475347949e-08 kg m^{-3}  
 PM2.5 mass conc. (with water): 2.2164289553077805e-08 kg m^{-3}

### 3.5 aerosol density

```
[6]: print("overall density:",
          pmc.get_aero_density(dry=False), "kg m^{-3}")
      print("overall dry density:",
            pmc.get_aero_density(dry=True), "kg m^{-3}")
      print("PM2.5 density (with water):",
            pmc.get_aero_density(dry=False, part_cond=pmc.get_aero_particle_diameter()<=2.5e-6),
            "kg m^{-3}")
```

overall density.: 1179.020687087021 kg m^{-3}  
 overall dry density: 2265.3505453667535 kg m^{-3}  
 PM2.5 density (with water): 1178.3768174927013 kg m^{-3}

### 3.6 gas mixing ratio

```
[7]: gas_list = ['CO','O3']
      pmc.get_gas_mixing_ratio(gas_list)

[7]: <xarray.DataArray 'gas_mixing_ratio' (gas_species: 2)>
      array([354.311384,  43.069374])
      Coordinates:
        * gas_species  (gas_species) int32 17 11
      Attributes:
        unit:          ppb
        long_name:      mixing ratios of gas species
```

### 3.7 mixing state index calculation

based on group\_list

```
[8]: group_list = [['SO4','Cl','ARO1','ARO2','ALK1','OLE1',
                   'API1','API2','LIM1','LIM2','Na'],
                  ['BC','OC','OIN']]
      pmc.get_mixing_state_index(group_list=group_list, diversity=False)

[8]: 0.838104186813985
```

based on all species (without water)

```
[9]: pmc.get_mixing_state_index(drop_list=["H2O"], diversity=False)

[9]: 0.6471920046172539
```

specific range of diameters and display diversity ( $D_\alpha$ ,  $D_\gamma$ ,  $\chi$ )

```
[10]: pmc.get_mixing_state_index(drop_list=["H2O"],
                                   part_cond=pmc.get_aero_particle_diameter()<=2.5e-6,
                                   diversity=True)

[10]: (3.1876143541221267, 4.355134759609094, 0.6520198176412457)
```

### 3.8 particle size distribution visualization (number concentration v.s. diameter)

overall number concentration

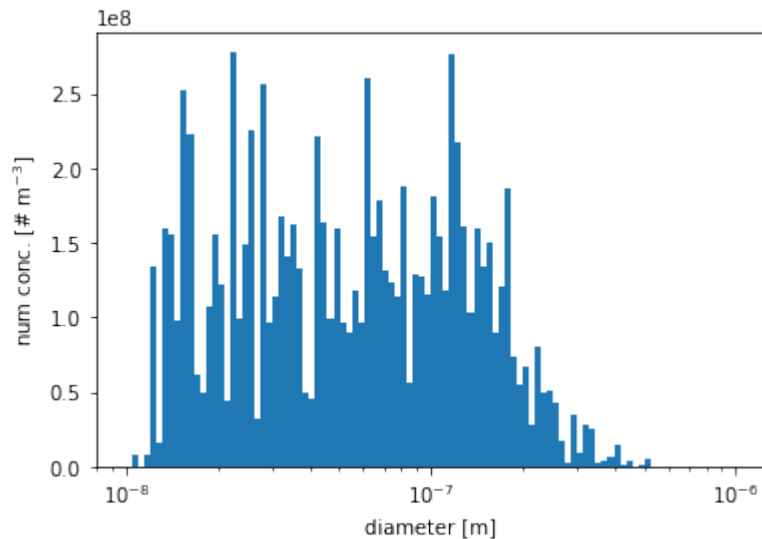
```
[11]: # get number distribution
      aero_diameter = pmc.get_aero_particle_diameter()
      num_conc_per_particle = pmc.ds["aero_num_conc"]

      # setup the 111 bins ranged from 10^-8 to 10^-6
      bins = np.logspace(-8,-6,2*50+1)
```

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```
# plot the number distribution
plt.hist(aero_diameter, bins=bins, weights=num_conc_per_particle)
plt.xscale('log')
plt.xlabel('diameter [m]')
plt.ylabel(r'num conc. [# m$^{-3}$]')
plt.show()
```



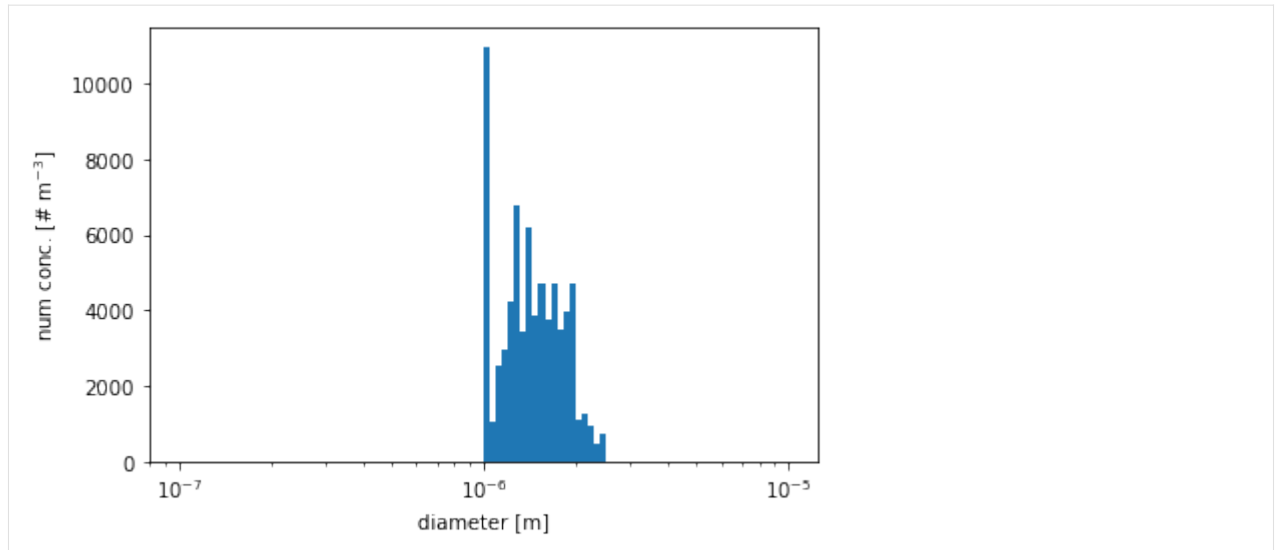
number concentration for particles with diameters between  $1\mu\text{m}$  and  $2.5\mu\text{m}$

```
[12]: # get the diameters of all particles
aero_diameter = pmc.get_aero_particle_diameter()
# select particles with diameters between 1um and 2.5um
part_cond = (aero_diameter<=2.5e-6) & (aero_diameter>=1.0e-6)

aero_diameter_cond = aero_diameter[part_cond]
num_conc_per_particle_cond = pmc.ds["aero_num_conc"][part_cond]

# setup the 111 bins ranged from 10^-8 to 10^-6
bins = np.logspace(-7,-5,2*50+1)

# plot the number distribution
plt.hist(aero_diameter_cond, bins=bins, weights=num_conc_per_particle_cond)
plt.xscale('log')
plt.xlabel('diameter [m]')
plt.ylabel(r'num conc. [# m$^{-3}$]')
plt.show()
```



### 3.9 black carbon fraction in dry particles

```
[13]: # get dry species per particle
all_aero_species = pmc.aero_species.copy()
dry_aero_species = all_aero_species.copy()
dry_aero_species.remove("H2O")
mass_per_dry_particle = pmc.ds["aero_particle_mass"]\
    .sel(aero_species = pmc.aero_species_to_idx(dry_aero_
    ↪ species))\
    .sum(dim="aero_species").values
# get black carbon mass per particle
bc_per_particle = pmc.ds["aero_particle_mass"]\
    .sel(aero_species=pmc.aero_species_to_idx(["BC"])).values.reshape(-
    ↪ 1)
bc_per_particle

[13]: array([0.00000000e+00, 0.00000000e+00, 0.00000000e+00, ...,
        5.78304546e-20, 0.00000000e+00, 0.00000000e+00])

[14]: bc_frac_per_dry_particle = bc_per_particle / mass_per_dry_particle
(bc_frac_per_dry_particle > 0.01).sum()

[14]: 269
```

## 3.10 black carbon core volume

```
[15]: BC_density = pmc.ds["aero_density"].sel(aero_species = pmc.aero_species_to_idx(["BC"])).  
      ↪ values  
      BC_density
```

```
[15]: array([1800.])
```

```
[16]: # volume = mass / density  
      pmc.ds["aero_particle_mass"].sel(aero_species = pmc.aero_species_to_idx(["BC"])).values /  
      ↪ BC_density
```

```
[16]: array([[0.00000000e+00, 0.00000000e+00, 0.00000000e+00, ...,  
            3.21280303e-23, 0.00000000e+00, 0.00000000e+00]])
```



## MIXING STATE CALCULATOR

This script is used for calculating [mixing state index](#)

**Step 0:** click the link [here](#) to launch a jupyter notebook

**Step 1:** upload your own csv file (each row is a particle, each column is the mass of a species)

`|upload\_csv|`

**Step 2:** edit the `file_name` and `group_list` below

Below is an example with three surrogate species

The definition of surrogate species is [here](#) (Figure 2)

- surrogate species 1: "BC" and "POM"
- surrogate species 2: "DUST"
- surrogate species 3: "SS"

**Step 3:** click Run button on the top of your jupyter notebook

`|run\_cell|`

Please edit the "file\_name" and "group\_list", then run the code below

```
[1]: file_name = "sample_data.csv"
      group_list = [["BC", "POM"],
                    ["DUST"],
                    ["SS"]]

      # ===== please don't change =====
      import numpy as np
      import pandas as pd
      import pmcpy

      # load data
      df = pd.read_csv(file_name)
      # get matrix for calculation
      da = np.concatenate([df[group].values.sum(axis=1).reshape(-1,1) for group in group_list],
                           ↪axis=1)
      # calculate mixing state index
      D_alpha, D_gamma, chi = pmcpy.get_chi(da)
      print("mixing state index:", chi)
```

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```
print("average particle (alpha) species diversity:", D_alpha)
print("bulk population (gamma) species diversity:", D_gamma)
```

```
mixing state index: 0.9408323695414793
average particle (alpha) species diversity: 2.838147712915256
bulk population (gamma) species diversity: 2.9537462489849164
```

## BENCHMARKING

This script is used to compare the pmcpy results with urban\_plume\_process.F90

```
[1]: # import necessary package
import pmcpy
import xarray as xr
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt

# define the path to data
p = "../data/"
```

results from ``pmcpy``

```
[2]: # define a dictionary to save the results
d = {"tot_num_conc": [], "tot_mass_conc": [],
     "chi": [], "d_alpha": [], "d_gamma": [],
     "chi_a": [], "d_alpha_a": [], "d_gamma_a": []}

# define the surrogate groups for mixing state calculation
group_list = [["OC", "BC"],
               ["API1", "API2", "LIM1", "LIM2"],
               ["SO4", "NO3", "NH4"]]

# loop across scenarios using pmcpy
for i in range(1, 26):
    pmc = pmcpy.load_pmc(p+"/urban_plume_0001_0000000"+str(i).zfill(2)+".nc")
    d["tot_num_conc"].append(pmc.get_num_conc())
    d["tot_mass_conc"].append(pmc.get_mass_conc(dry=False))

    # calculate mixing state for grouped species with water
    D_alpha, D_gamma, chi = pmc.get_mixing_state_index(group_list, diversity=True)
    d["d_alpha"].append(D_alpha)
    d["d_gamma"].append(D_gamma)
    d["chi"].append(chi)

    # calculate mixing state for all species (without water)
    D_alpha_a, D_gamma_a, chi_a = pmc.get_mixing_state_index(drop_list=["H2O"],
    diversity=True)
    d["d_alpha_a"].append(D_alpha_a)
```

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```
d["d_gamma_a"].append(D_gamma_a)
d["chi_a"].append(chi_a)
```

compare the results from Fortran postprocessing ``urban\_plume\_process.F90``

```
[3]: ds_b = xr.open_dataset(p+"urban_plume_process.nc")

# ===== comparison =====
for k in d:

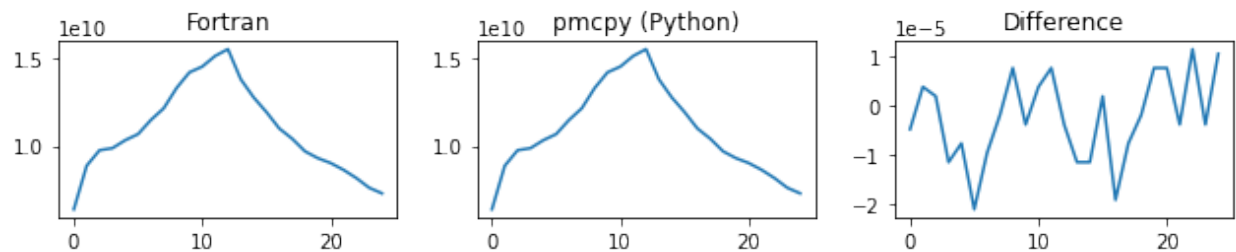
    print("#####",k,"#####")
    fig = plt.figure(figsize=(9,2))
    ax1 = fig.add_subplot(131)
    pd.Series(ds_b[k].values).plot(ax=ax1)
    ax1.set_title("Fortran")

    ax2 = fig.add_subplot(132)
    pd.Series(d[k]).plot(ax=ax2)
    ax2.set_title("pmcpy (Python)")

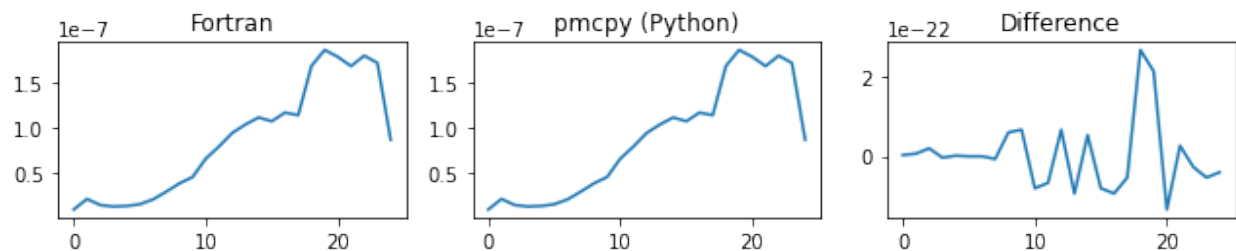
    ax3 = fig.add_subplot(133)
    pd.Series(ds_b[k].values-np.array(d[k])).plot(ax=ax3)
    ax3.set_title("Difference")

    plt.tight_layout()
    plt.show()
```

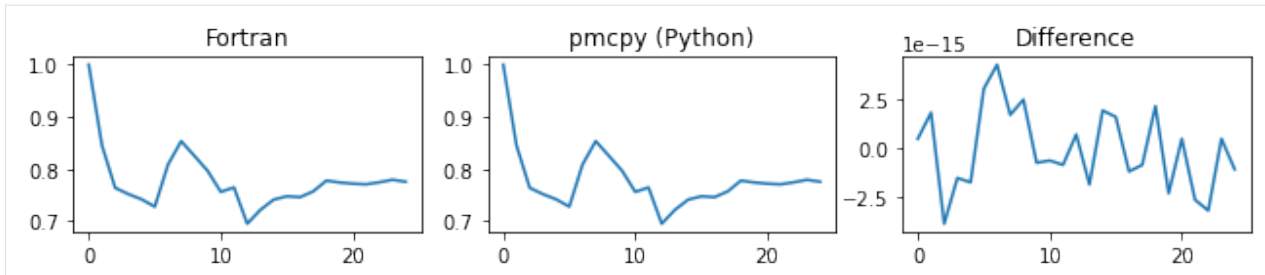
##### tot\_num\_conc #####



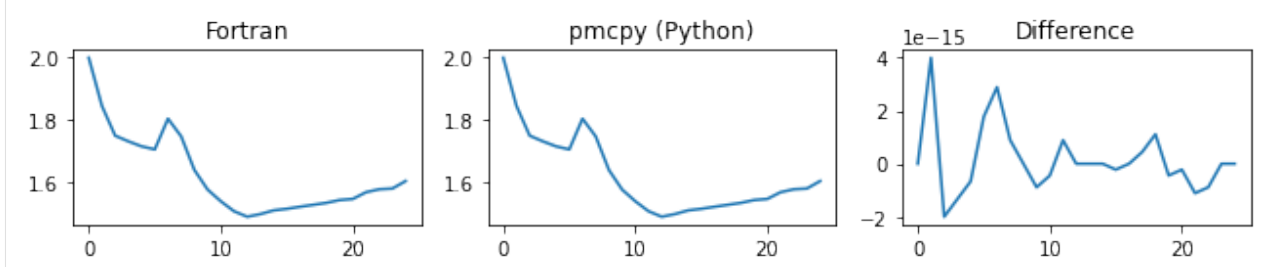
##### tot\_mass\_conc #####



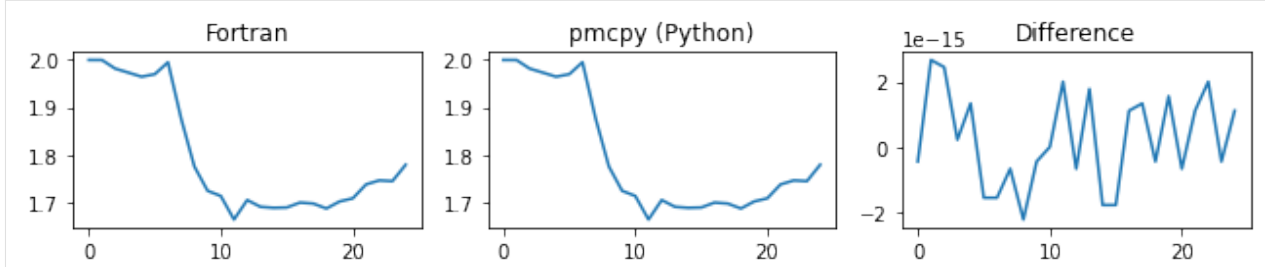
##### chi #####



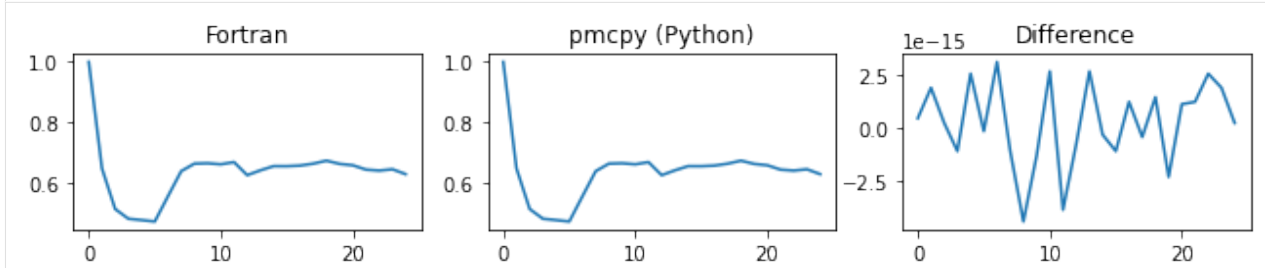
#####  $d_\alpha$  #####



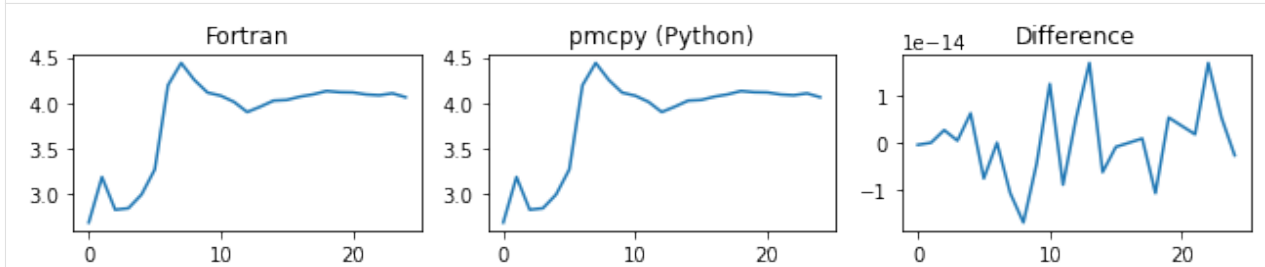
#####  $d_\gamma$  #####



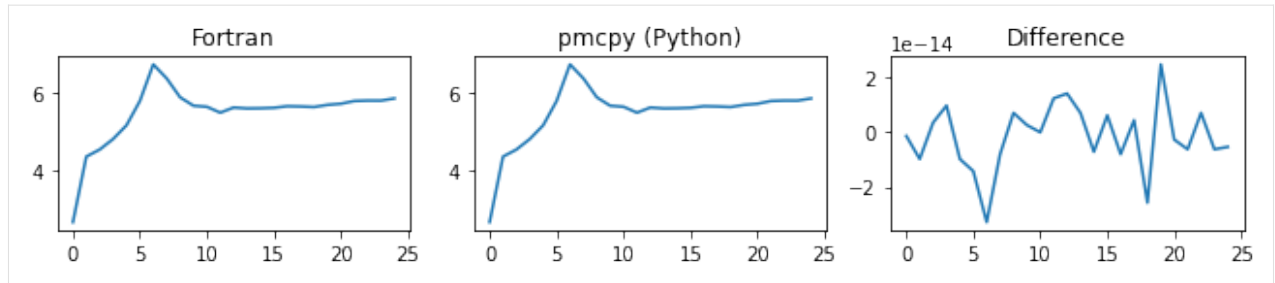
#####  $\chi_a$  #####



#####  $d_{\alpha_a}$  #####



#####  $d_{\gamma_a}$  #####



## FORMAT OF PARTMC RAW OUTPUT

Here is a sample of PartMC raw output

```
[1]: import xarray as xr
p = "../data/urban_plume_0001_000000002.nc"
ds = xr.open_dataset(p)
ds
```

```
[1]: <xarray.Dataset>
Dimensions:                (gas_species: 77, aero_species: 20,
                             aero_source: 8, aero_weight_group: 2,
                             aero_weight_class: 8, aero_particle: 1246,
                             aero_removed: 4636)

Coordinates:
  * gas_species              (gas_species) int32 1 2 3 4 5 ... 73 74 75 76 77
  * aero_species             (aero_species) int32 1 2 3 4 5 ... 17 18 19 20
  * aero_source              (aero_source) int32 1 2 3 4 5 6 7 8
  * aero_weight_group        (aero_weight_group) int32 1 2
  * aero_weight_class        (aero_weight_class) int32 1 2 3 4 5 6 7 8
  * aero_particle            (aero_particle) int32 1 2 3 4 ... 1244 1245 1246
  * aero_removed             (aero_removed) int32 1 2 3 4 ... 4634 4635 4636

Data variables: (12/48)
  time                      float64 ...
  timestep                  float64 ...
  timestep_index            int32 ...
  repeat                    int32 ...
  temperature               float64 ...
  relative_humidity         float64 ...
  ...                       ...
  aero_refract_core_real    (aero_particle) float64 ...
  aero_refract_core_imag    (aero_particle) float64 ...
  aero_core_vol             (aero_particle) float64 ...
  aero_removed_id           (aero_removed) int32 ...
  aero_removed_action       (aero_removed) int32 ...
  aero_removed_other_id     (aero_removed) int32 ...

Attributes:
  title:                    PartMC version 2.5.0 output file
  source:                   PartMC version 2.5.0
  UUID:                     A96876F0-F62E-45FB-AC47-2FB221548641
  history:                  2021-09-22T15:54:26.773-05:00 created by PartMC version 2.5.0
  Conventions:              CF-1.4
```





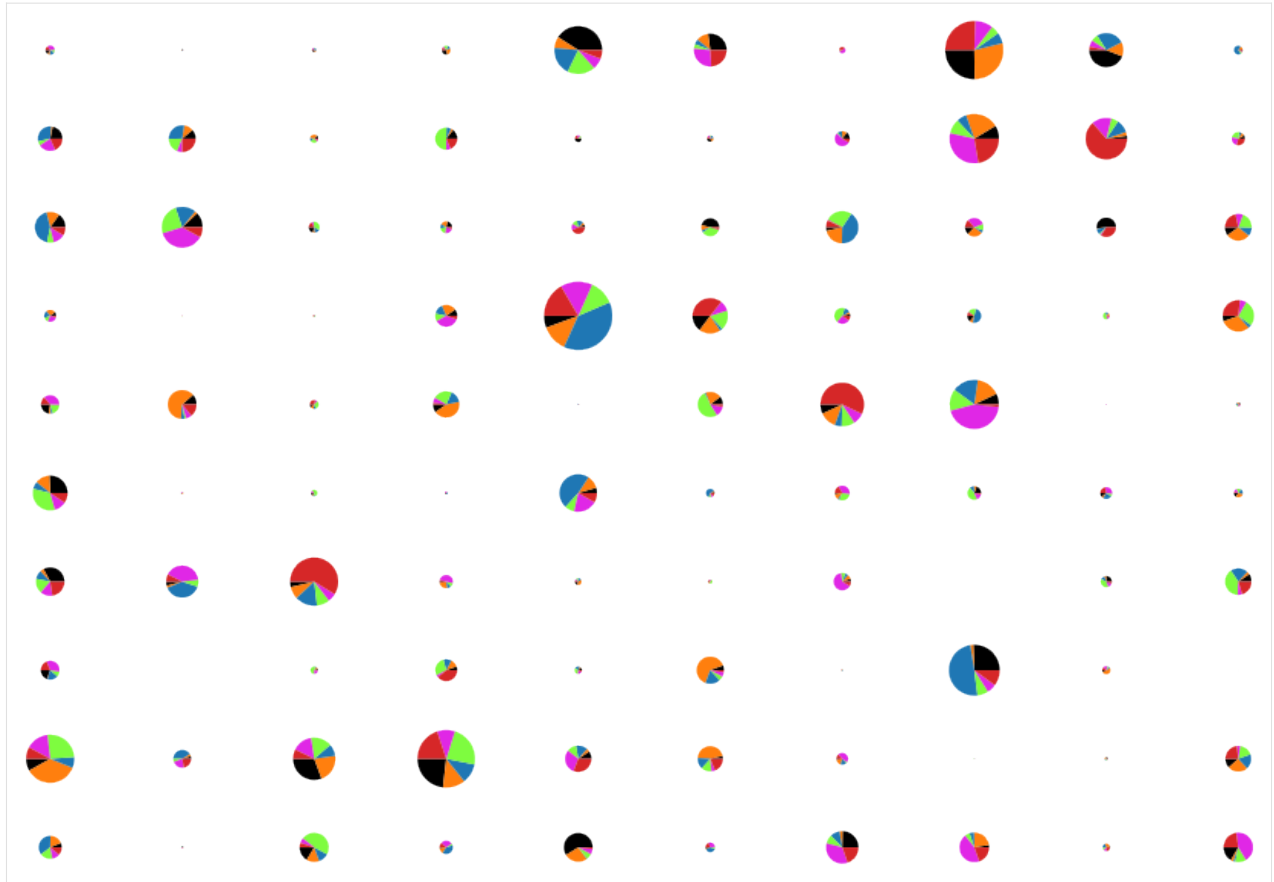
## PARTICLES VISUALIZATION

This script is used for generating and plotting particles

```
[1]: import matplotlib.pyplot as plt
import numpy as np
```

```
[2]: mat = np.random.lognormal(size=(100,6))
radius = np.random.normal(size=(100,1))
colors = ["#000000", "#FF7F0E", "#1F77B4", "#7efc3f", "#e027e6", "#D62728"]
```

```
[3]: fig, axes = plt.subplots(10, 10, figsize=(12,8))
axes=axes.ravel()
# plot each pie chart in a separate subplot
for i in range(100):
    axes[i].pie(mat[i,],radius=radius[i], colors=colors)
plt.tight_layout()
plt.show()
```



## HOW TO ASK FOR HELP

The [GitHub issue tracker](#) is the primary place for bug reports.