# atomneb Documentation

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# **USER DOCUMENTATION**

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**USER DOCUMENTATION** 

ONE

# INTRODUCTION

**atomneb** is a library written in Python for reading atomic data from a database containing atomic data stored in the Flexible Image Transport System (FITS) file format for *collisionally excited lines* and *recombination lines* typically observed in spectra of ionized gaseous nebulae. The AtomNeb database were generated for use in pyEQUIB, proE-QUIB, and other nebular spectral analysis tools.

# 1.1 Collisional Excitation Unit

AtomNeb for collisionally excited lines contains sets of atomic datasets, which include energy levels  $(E_j)$ , collision strengths  $(\Omega_{ij})$ , and transition probabilities  $(A_{ij})$  of the most ions commonly observed in ionized nebulae.

The atomic datasets for collisionally excited lines are as follows:

- Collection from the National Institute of Standards and Technology (NIST) Atomic Spectra Database, the CHI-ANTI atomic database, and some improved atomic data from Cloudy v13.04 and pyNeb v1.0. This collection was compiled according to the atomic data used in pyNeb v1.0.
- Chianti52 from the CHIANTI atomic database version 5.2. This dataset was compiled according to the atomic data used in MOCASSIN.
- Chianti60 from the CHIANTI atomic database version 6.0. This dataset was compiled according to the atomic
  data used in MOCASSIN.
- Chianti70 from the CHIANTI atomic database version 7.0. This dataset was compiled according to the atomic data used in MOCASSIN.
- Chianti90 from the CHIANTI atomic database version 9.0. This dataset was compiled according to the atomic data used in NEAT.

Each dataset contains the following atomic data FITS files: AtomElj.fits for *Energy Levels*  $(E_j)$ , AtomOmij. fits for *Collision Strengths*  $(\Omega_{ij})$ , and AtomAij.fits for *Transition Probabilities*  $(A_{ij})$ .

# 1.2 Recombination Unit

AtomNeb for recombination lines contains sets of effective recombination coefficients ( $\alpha_{eff}$ ) of recombination lines of H I, He II, C I, C III, C III, C VI, N III, N III, N IV, N VI, N VII, O II, O III, O IV, O V, O VI, O VIII, and Ne II ions typically observed in ionized nebulae, as well as Branching ratios (Br) of O II and N II lines.

The atomic datasets for recombination lines are as follows:

• RC Collection, effective recombination coefficients for C II (Davey et al. 2000), N II (Escalante and Victor 1990), O II (Storey 1994; Liu et al. 1995), and Ne II ions (Kisielius et al. 1998), including Branching ratios (Br) for O II and N II ions. This collection was compiled according to the atomic data used in MOCASSIN.

- SH95 Collection, hydrogenic ions for Z=1 to 8, namely H I, He II, Li III, Be IV, B V, C VI, N VII, and O VIII ions from Storey and Hummer (1995).
- PPB91 Collection, effective recombination coefficients for H, He, C, N, O, Ne ions from Pequignot, Petitjean and Boisson (1991).
- PFSD12 He I data, effective He I recombination coefficients from Porter et al (2012) and (2013a).
- FSL13 N II data, effective N II recombination coefficients (corrigendum) from Fang, Storey and Liu (2011) and (2013b).
- SSB17 O II data, effective O II recombination coefficients of 8889 recombination lines for Cases A, B, and C, and 2433 optical (3500-9000Å) recombination lines for Case B from Storey, Sochi and Bastin (2017).

# **TWO**

# **INSTALLATION**

To install the last version, all you should need to do is:

```
python setup.py install
```

To install the stable version, you can use the preferred installer program (pip):

```
pip install atomneb
```

or you can install it from the cross-platform package manager conda:

```
conda install -c conda-forge atomneb
```

To get this package with all the FITS file, you can simply use git command as follows:

```
git clone https://github.com/atomneb/AtomNeb-py
```

If you plan to use the recent O II recombination coefficients (Storey, Sochi and Bastin 2017), you need to unpack them:

```
cd AtomNeb-py/atomic-data-rc/
tar -xvf *.fits.tar.gz
```

This package requires the following packages:

- NumPy
- Astropy

THREE

# **USAGE**

The Documentation of the functions provides in detail in the *API Documentation* (atomneb.github.io/AtomNeb-py/doc). There are two main categories: *collisionally excited lines (CEL)* and *recombination lines (RC)*.

# 3.1 Collisional Excitation Unit

The atomic data for **collisional excitation unit (CEL)** contain Energy Levels  $(E_j)$ , Collision Strengths  $(\Omega_{ij})$ , and Transition Probabilities  $(A_{ij})$ . We have four atomic datasets for them: collection, chianti52, chianti60, and chianti70.

You need to load the **atomneb** library as follows:

```
import atomneb
```

Also:

```
import atomneb
```

Also:

```
import numpy as np
import os

atom_elj_file = os.path.join(base_dir,data_dir, 'AtomElj.fits')
atom_omij_file = os.path.join(base_dir,data_dir, 'AtomOmij.fits')
atom_aij_file = os.path.join(base_dir,data_dir, 'AtomAij.fits')
elj_data_list = atomneb.read_elj_list(atom_elj_file)
omij_data_list = atomneb.read_omij_list(atom_omij_file)
aij_data_list = atomneb.read_aij_list(atom_aij_file)
```

Now you have access to:

• Energy Levels  $(E_j)$ :

```
atom='o'
ion='iii'
oiii_elj_data = atomneb.read_elj(atom_elj_file, atom, ion, level_num=6)
print(oiii_elj_data['j_v'])
print(oiii_elj_data['ej'])
```

# which gives:

0.00000	1.00000	2.00000	2.00000	0.00000	2.00000
0.00000	113.200	306.200	20273.30	43185.69	60324.80

• Collision Strengths  $(\Omega_{ij})$ :

```
atom='o'
ion='iii'
oiii_omij_data = atomneb.read_omij(atom_omij_file, atom, ion)
print(oiii_omij_data['level1'])
print(oiii_omij_data['level2'])
print(oiii_omij_data['strength'][0])
```

## which gives:

```
    0
    1
    1
    1
    ...

    0
    2
    3
    4
    5
    ...

    100.0
    158.50
    251.20
    398.10
    631.0
    ...
```

• Transition Probabilities  $(A_{ij})$ :

```
atom='o'
ion='iii'
oiii_aij_data = atomneb.read_aij(atom_aij_file, atom, ion)
print(oiii_aij_data['aij'][0])
```

## which gives:

```
0.0000 2.5969e-05 0.0000 2.3220e-06 ...
```

# 3.2 Recombination Unit

The atomic data for **recombination unit (RC)** contain effective recombination coefficients ( $\alpha_{eff}$ ) of emission lines from different collections: RC Collection, SH95 Collection, PPB91 Collection, PFSD12 He I data, FSL13 N II data, and SSB17 O II data.

You need to load the atomneb libary:

```
import atomneb
```

Also:

```
import numpy as np import os
```

Now you have access to effective recombination coefficients ( $\alpha_{eff}$ ) of the following collections:

• RC Collection:

which gives:

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```
914.00000
             0.69280000
                           0.021400000
                                         -0.016300000
→24310000
            -0.88000000
             1.0998000
                        -0.0042000000
962.00000
                                         -0.027900000
                                                         -0.
→22940000
             -0.96560000
997.00000
            0.78210000
                          -0.36840000 0.00030000000
                                                         -0.
→12170000
             -0.78740000
. . .
```

#### • SH95 Collection:

```
atom_rc_file = os.path.join(base_dir,data_dir, 'rc_SH95.fits')
atom='h'
ion='ii'
hi_rc_data = atomneb.read_aeff_sh95(atom_rc_file, atom, ion)
print(hi_rc_data['aeff'][0])
```

## which gives:

```
100.00000 500.00000 0.0000000 4.2140000e-27 1.7560000e-

→27 1.0350000e-27 ...
```

#### • PPB91 Collection:

## which gives:

```
C2+A 9903.4600 0.69700000 −0.78400000 4.2050000

→ 0.72000000 1.0000000 1.6210000

C2+A 4267.1500 1.0110000 −0.75400000 2.5870000

→ 0.71900000 0.95000000 2.7950000

...
```

#### • PFSD12 He I data:

```
atom_rc_file = os.path.join(base_dir,data_dir, 'rc_he_ii_PFSD12.fits')
atom='he'
ion='ii'
hei_rc_data = atomneb.read_aeff_he_i_pfsd12(atom_rc_file, atom, ion)
hei_rc_data_wave = atomneb.read_aeff_he_i_pfsd12(atom_rc_file, atom, ion,
    wavelength=True)
print(hei_rc_data['aeff'][0])
```

# which gives:

```
5000.0000 10.000000 −25.379540 −25.058970 −25.

→948440 −24.651820 −25.637660

...
```

• FSL13 N II data:

#### which gives:

```
255.000
            79.5000
                         47.3000
                                      12.5000
                                                   6.20000
                                                                4.00000
      2.86000
258.000
                         29.7000
            54.4000
                                      7.92000
                                                   4.11000
                                                                2.72000_
      2.00000
310.000
            48.1000
                         23.7000
                                      5.19000
                                                   2.55000
                                                                1.65000_
     1.21000
434.000 50.3000
                         23.2000
                                      4.71000
                                                   2.26000
                                                                1.45000
     1.05000
6413.23 6g - 4f2p6g G[9/2]o4 - 2p4f F[7/2]e3
6556.32 6g - 4f2p6g G[9/2]o5 - 2p4f G[7/2]e4
6456.97 6g - 4f2p6g G[9/2]o5 - 2p4f F[7/2]e4
6446.53 6g - 4f2p6g F[7/2]o3 - 2p4f D[5/2]e2
6445.34 6g - 4f2p6g F[7/2]o4 - 2p4f D[5/2]e3
```

• SSB17 O II data: You first need to unpack rc\_o\_iii\_SSB17\_orl\_case\_b.fits.tar.gz, e.g.:

```
tar -xvf rc_o_iii_SSB17_orl_case_b.fits.tar.gz
```

If you need to have access to the full dataset (entire wavelengths, case A and B):

```
tar -xvf rc_o_iii_SSB17.fits.tar.gz
```

Please note that using the entire atomic data will make your program very slow and you may need to have a higher memory on your system. Without the above comment, as default, the cose uses rc\_o\_iii\_SSB17\_orl\_case\_b.fits:

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### which gives:

```
1.64100e-30 1.60000e-30 1.56400e-30 1.54100e-30 1.52100e-30 1.

$\infty$50900e-30 \\
\therefore \text{5327.17 2s22p2(1s) 3p 2Po} \\
5325.42 2s22p2(1s) 3p 2Po \\
5327.18 2s22p2(1D) 3d 2Ge \\
5326.84 2s22p2(1D) 3d 2Ge \\
\therefore \text{...}
```

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# **FOUR**

# **REFERENCES**

- Danehkar, A. (2020). AtomNeb Python Package, an addendum to AtomNeb: IDL Library for Atomic Data of Ionized Nebulae. *J. Open Source Softw.*, **5**, 2797. doi:10.21105/joss.02797 ads:2020JOSS....5.2797D.
- Danehkar, A. (2019). AtomNeb: IDL Library for Atomic Data of Ionized Nebulae. *J. Open Source Softw.*, **4**, 898. doi:10.21105/joss.00898 ads:2019JOSS....4..898D.

**FIVE** 

# ATOMNEB.MAIN PACKAGE

# 5.1 atomneb main module

This module contains functions for Atomic Data of Ionized Nebulae

```
atomneb.get_aeff_collection_reference_citation(atom_rc_file, atom, ion, br=None, reference=None)
```

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_collection.fits').

#### For example:

**Returns** This function returns the Citation.

# Return type str

### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_collection.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'
- **br** (boolean, optional) set for the branching ratios (Br)
- reference (str, optional) set for the reference

```
atomneb.get_aeff_he_i_pfsd12_reference_citation(atom_rc_file, atom, ion, reference=None)
```

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_he\_ii\_PFSD12.fits').

#### For example:

**Returns** This function returns the Citation.

#### Return type str

### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_he\_ii\_PFSD12.fits')
- atom (str) atom name e.g. 'he'
- ion (str) ionic level e.g 'ii'
- reference (str, optional) set for the reference e.g. 'PFSD13'

```
atomneb.get_aeff_n_ii_fsl13_reference_citation(atom_rc_file, atom, ion, reference=None)
```

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_n\_iii\_FSL13.fits').

# For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = ['atomic-data-rc']
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_n_iii_FSL13.fits')
>> atom='n'
>> ion='iii' # N II
>> reference='FSL13'
>> citation = atomneb.get_aeff_n_ii_fsl13_reference_citation(atom_rc_file, atom, ion)
>> print(citation)
    Fang X., Storey P.J., and Liu X.-W., R. 2011, Astron.Astrophys. 530, Als; 2013, Astron.Astrophys. 550, C2
```

**Returns** This function returns the Citation.

### Return type str

# **Parameters**

• atom rc file (str) - the FITS data file name ('rc n iii FSL13.fits')

```
• atom (str) - atom name e.g. 'n'
```

- ion (str) ionic level e.g 'iii'
- reference (str, optional) set for the reference e.g. 'FSL13

atomneb.get\_aeff\_o\_ii\_ssb17\_reference\_citation(atom\_rc\_file, atom, ion, reference=None)

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_o\_iii\_SSB17.fits').

#### For example:

**Returns** This function returns the Citation.

# Return type str

## **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_o\_iii\_SSB17.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- reference (str, optional) set for the reference e.g. 'SSB17

atomneb.get\_aeff\_ppb91\_reference\_citation(atom\_rc\_file, atom, ion, reference=None)

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_PPB91.fits').

### For example:

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```
>> print(citation)
Pequignot, D., Petitjean, P. and Boisson, C. Astron.Astrophys., 251, 4680, 1991
```

**Returns** This function returns the Citation.

**Return type** str

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_PPB91.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'
- reference (str, optional) set for the reference

atomneb.get\_aeff\_sh95\_reference\_citation(atom\_rc\_file, atom, ion, reference=None, caseI=None)

This function returns the reference citation for a recombination coefficient (Aeff) from the 2nd binary table extension of the FITS data file ('rc SH95.fits').

#### For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = ['atomic-data-rc']
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_SH95.fits')
>> atom='h'
>> ion='ii' # H I
>> citation = atomneb.get_aeff_sh95_reference_citation(atom_rc_file, atom, ion)
>> print(citation)
Storey, P. J. and Hummer, D. G., MNRAS, 272, 415, 1995
```

**Returns** This function returns the Citation.

# Return type str

### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_SH95.fits')
- atom (str) atom name e.g. 'h'
- ion (str) ionic level e.g 'ii'
- reference (str, optional) set for the reference
- case1 (str, optional) set for the case 'a' or 'b', defualt 'b'

# atomneb.get\_aij\_reference\_citation(atom\_aij\_file, atom, ion, reference)

This function returns the reference citation for a transition probability (Aij) from the 2nd binary table extension of the FITS data file ('AtoAij.fits')

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..','atomic-data', 'collection')
>> atom_aij_file = os.path.join(base_dir,data_dir, 'AtomAij.fits')
>> atom='o'
>> ion='iii'
>> reference='FFT04'
>> citation = atomneb.get_aij_reference_citation(atom_aij_file, atom, ion, reference)
>> print(citation)
Froese Fischer et al 2004, ADNDT 87, 1
```

**Returns** This function returns the Citation.

#### Return type str

#### **Parameters**

- **Atom\_Aij\_file** (str) the FITS data file name ('AtoAij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- **reference** (str) set for the reference e.g. 'FFT04'

```
atomneb.get_elj_reference_citation(atom_elj_file, reference)
```

This function returns the reference citation for energy levels (Ej) from the 2nd binary table extension of the FITS data file ('AtomElj.fits').

# For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..','atomic-data', 'collection')
>> atom_elj_file = os.path.join(base_dir,data_dir, 'AtomElj.fits')
>> reference='L7288'
>> citation=atomneb.get_elj_reference_citation(atom_elj_file, reference)
>> print(citation)
    C. E. Moore, in CRC Series in Evaluated Data in Atomic Physics, 339 pp.

-- (CRC Press, Boca Raton, FL, 1993)
```

**Returns** This function returns the Citation.

#### Return type str

# **Parameters**

- atom\_elj\_file (str) the FITS data file name ('AtomElj.fits')
- **reference** (*str*) set for the reference e.g. 'L7288'

```
atomneb.get_omij_reference_citation(atom_omij_file, atom, ion, reference)
```

This function returns the reference citation for collision strengths (Omega\_ij) from the 2nd binary table extension of the FITS data file ('AtomOmij.fits').

#### For example:

**Returns** This function returns the Citation.

# Return type str

### **Parameters**

- atom\_omij\_file (str) the FITS data file name ('AtomOmij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- **reference** (*str*) set for the reference e.g. 'SSB14'

#### atomneb.list aeff collection references (atom rc file, atom, ion, br=None)

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc\_collection.fits').

### For example:

**Returns** This function returns the references.

**Return type** an array of strings

- atom\_rc\_file (str) the FITS data file name ('rc\_collection.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'

• **br** (boolean, optional) – set for the branching ratios (Br)

#### atomneb.list\_aeff\_he\_i\_pfsd12\_references(atom\_rc\_file, atom, ion)

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc\_he\_ii\_PFSD12.fits').

For example:

**Returns** This function returns the references.

Return type an array of strings

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_he\_ii\_PFSD12.fits')
- atom (str) atom name e.g. 'he'
- ion (str) ionic level e.g 'ii'

# atomneb.list\_aeff\_n\_ii\_fsl13\_references(atom\_rc\_file, atom, ion)

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc\_n\_iii\_FSL13.fits').

For example:

**Returns** This function returns the references.

Return type an array of strings

- atom\_rc\_file (str) the FITS data file name ('rc\_n\_iii\_FSL13.fits')
- atom (str) atom name e.g. 'n'

• ion (str) – ionic level e.g 'iii'

```
atomneb.list_aeff_o_ii_ssb17_references(atom_rc_file, atom, ion)
```

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc o iii SSB17.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = ['atomic-data-rc']
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_o_iii_SSB17_orl_
--case_b.fits')
>> atom='o'
>> ion='iii' # O II
>> list_oii_aeff_reference = atomneb.list_aeff_o_ii_ssb17_references(atom_---rc_file, atom, ion)
>> print(list_oii_aeff_reference)
```

**Returns** This function returns the references.

Return type an array of strings

### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_o\_iii\_SSB17.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'

## atomneb.list\_aeff\_ppb91\_references(atom\_rc\_file, atom, ion)

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc\_PPB91.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = ['atomic-data-rc']
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_PPB91.fits')
>> atom='c'
>> ion='iii'
>> list_cii_aeff_reference = atomneb.list_aeff_ppb91_references(atom_rc_
--file, atom, ion)
>> print(list_cii_aeff_reference)
```

**Returns** This function returns the references.

Return type an array of strings

- atom\_rc\_file (str) the FITS data file name ('rc\_PPB91.fits')
- atom (str) atom name e.g. 'c'

• ion (str) – ionic level e.g 'iii'

```
atomneb.list_aeff_sh95_references(atom_rc_file, atom, ion)
```

This function returns a list for all references of recombination coefficients (Aeff) for given element and ionic level from the FITS data file ('rc SH95.fits').

For example:

**Returns** This function returns the references.

Return type an array of strings

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_SH95.fits')
- atom (str) atom name e.g. 'h'
- ion (str) ionic level e.g 'ii'

### atomneb.list\_aij\_references(atom\_aij\_file, atom, ion)

This function returns a list for all references of transition probabilities (Aij) for given element and ionic level from the FITS data file ('AtoAij.fits').

For example:

**Returns** This function returns the references.

Return type an array of data

- atom\_rc\_file (str) the FITS data file name ('AtoAij.fits')
- atom (str) atom name e.g. 'o'

• ion (str) – ionic level e.g 'iii'

#### atomneb.list\_omij\_references(atom\_omij\_file, atom, ion)

This function returns a list for all references of collision strengths (Omega\_ij) for given element and ionic level from the FITS data file ('AtomOmij.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..', 'atomic-data', 'collection')
>> atom_omij_file = os.path.join(base_dir, data_dir, 'AtomOmij.fits')
>> atom='o'
>> ion='iii'
>> list_oiii_omij_reference = atomneb.list_omij_references(atom_omij_file, atom, ion)
>> print(list_oiii_omij_reference)
AK99 LB94 Pal12-AK99 SSB14
```

**Returns** This function returns the references.

Return type str

#### **Parameters**

- atom\_omij\_file (str) the FITS data file name ('AtomOmij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'

atomneb.read\_aeff\_collection(atom\_rc\_file, atom, ion, br=None, reference=None)

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc\_collection.fits').

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_collection.fits')
>> atom='c'
>> ion='iii' # C III
>> cii_rc_data = atomneb.read_aeff_collection(atom_rc_file, atom, ion)
>> n_line = len(cii_rc_data['wavelength'])
>> for i in range(0, n_line):
        print(cii_rc_data['wavelength'][i], cii_rc_data['a'][i],
>>
            cii_rc_data['b'][i], cii_rc_data['c'][i],
>>
>>
             cii_rc_data['d'][i], cii_rc_data['f'][i])
  914.00000 0.69280000
                               0.021400000 -0.016300000
                                                                -0.
\rightarrow 24310000 -0.88000000
  962.00000
               1.0998000 -0.0042000000
                                               -0.027900000
                                                                -0.
              -0.96560000
-22940000
```

**Returns** This function returns the effective recombination coefficients. aeff\_data (c\_iii\_aeff) { Wavelength:0.0,a: 0.0, b: 0.0, c: 0.0, d: 0.0, f: 0.0}, aeff\_data (n\_iii\_aeff) { a: 0.0, b: 0.0, c: 0.0}, aeff\_data (n\_iii\_br) {Wavelength: 0.0, BR: 0.0, g1:0, g2:0, Mult1:", LowerTerm:", UpperTerm:" }, aeff\_data (o\_iii\_aeff) {Term: ", Case1: ", a2: 0.0, a4: 0.0, a5: 0.0, a6: 0.0, b: 0.0, c: 0.0, d: 0.0}, aeff\_data (o\_iii\_br) {Wavelength:double(0.0), Br\_A: 0.0, Br\_B: 0.0, Br\_C: 0.0, g1: 0, g2: 0, Mult1: ", LowerTerm: ", UpperTerm: "}, aeff\_data (ne\_iii\_aeff) {Wavelength:0.0, a: 0.0, b: 0.0, c: 0.0, d: 0.0, f: 0.0, br: 0.0},

**Return type** an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_collection.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'
- **br** (boolean, optional) set for the branching ratios (Br)
- reference (str, optional) set for the reference

```
atomneb.read_aeff_collection_list(atom_rc_file)
```

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc\_collection.fits')

**Returns** This function returns the aeff\_data\_list: { Aeff\_Data:", Extension:0.0}

Return type an array of data

**Parameters atom\_rc\_file** (str) – the FITS data file name ('rc\_collection.fits')

```
atomneb.read_aeff_collection_references(atom_rc_file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc collection.fits').

**Returns** This function returns the aeff\_data\_reference: { Reference:", Citation:"}

**Return type** an array of data

**Parameters atom rc file** (str) – the FITS data file name ('rc collection.fits')

 $\verb|atomneb.read_aeff_he_i_pfsd12| (atom_rc\_file, atom, ion, wavelength=None, reference=None)|$ 

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc he ii PFSD12.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir, data_dir, 'rc_he_ii_PFSD12.fits')
>> atom='he'
>> ion='ii' # He I
>> hei_rc_data = atomneb.read_aeff_he_i_pfsd12(atom_rc_file, atom, ion)
>> hei_rc_data_wave = atomneb.read_aeff_he_i_pfsd12(atom_rc_file, atom, ion)
wion, wavelength=True)
```

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```
>> print(hei_rc_data['aeff'][0])
  5000.0000
                  10.000000
                                  -25.379540
                                                  -25.058970
                                                                  -25.
→948440
>> n_line = len(hei_rc_data_wave['wavelength'])
>> for i in range(0, n_line):
      print(hei_rc_data_wave['wavelength'][i],
>>
            hei_rc_data_wave['lowerterm'][i], hei_rc_data_wave['upperterm
→'][i])
  2945.00005p^{3}P2s^{3}S
  3188.00004p^{3}P2s^{3}S
  3614.00005p^{1}P2s^{1}S
```

**Returns** This function returns the effective recombination coefficients.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_he\_ii\_PFSD12.fits')
- atom (str) atom name e.g. 'he'
- ion (str) ionic level e.g 'ii'
- wavelength (boolean, optional) set for returning the wavelengths
- reference (str, optional) set for the reference

```
atomneb.read_aeff_he_i_pfsd12_list(atom_rc_file)
```

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc\_he\_ii\_PFSD12.fits')

**Returns** This function returns the aeff\_data\_list: { Aeff\_Data:", Extension:0.0}

**Return type** an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_he\_ii\_PFSD12.fits')

```
atomneb.read_aeff_he_i_pfsd12_references(atom_rc_file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc he ii PFSD12.fits').

**Returns** This function returns the aeff\_data\_reference:{ Reference:", Citation:"}

Return type an array of data

**Parameters atom\_rc\_file** (str) – the FITS data file name ('rc\_he\_ii\_PFSD12.fits')

atomneb.read\_aeff\_n\_ii\_fsl13 (atom\_rc\_file, atom, ion, wavelength\_range, wavelength=None, reference=None)

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc\_n\_iii\_FSL13.fits').

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_n_iii_FSL13.fits')
>> atom='n'
>> ion='iii' # N II
>> wavelength_range=[4400.0, 7100.0]
>> nii_rc_data = atomneb.read_aeff_n_ii_fsl13(atom_rc_file, atom, ion,...
→wavelength_range)
>> nii_rc_data_wave = atomneb.read_aeff_n_ii_fsl13(atom_rc_file, atom,_
⇒ion, wavelength_range, wavelength=True)
>> print(nii_rc_data['aeff'][0])
                         47.3000
               79.5000
  255.000
                                         12.5000
>> n_line = len(nii_rc_data_wave['wavelength'])
>> for i in range(0, n_line):
      print(nii_rc_data_wave['wavelength'][i], nii_rc_data_wave['tr'][i],
→ nii_rc_data_wave['trans'][i])
  6413.236g - 4f2p6g G[9/2]o4 - 2p4f F[7/2]e3
  6556.326g - 4f2p6g G[9/2]o5 - 2p4f G[7/2]e4
  6456.976g - 4f2p6g G[9/2]o5 - 2p4f F[7/2]e4
```

**Returns** This function returns the effective recombination coefficients.

Return type an array of data

#### **Parameters**

- atom rc file (str) the FITS data file name ('rc n iii FSL13.fits')
- atom (str) atom name e.g. 'n'
- ion (str) ionic level e.g 'iii'
- wavelength (boolean, optional) set for returning the wavelengths
- reference (str, optional) set for the reference

```
atomneb.read_aeff_n_ii_fsl13_list(atom_rc_file)
```

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc\_n\_iii\_FSL13.fits')

**Returns** This function returns the aeff\_data\_list: {Aeff\_Data:", Extension:0, IND:long(0), Wavelength: float(0.0), Tr:", Trans: ", T\_X: "}

Return type an array of data

Parameters atom rc file (str) - the FITS data file name ('rc n iii FSL13.fits')

```
atomneb.read_aeff_n_ii_fsl13_references(atom_rc_file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_n\_iii\_FSL13.fits').

**Returns** This function returns the aeff\_data\_reference: { Reference:", Citation:"}

**Return type** an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_n\_iii\_FSL13.fits')

atomneb.read\_aeff\_o\_ii\_ssb17 (atom\_rc\_file, atom, ion, case1, wavelength\_range, wavelength=None, reference=None)

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc o iii SSB17.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_o_iii_SSB17_orl_
⇒case_b.fits')
>> atom='o'
>> ion='iii' # O II
>> case1='B'
>> wavelength_range=[5320.0, 5330.0]
>> oii_rc_data = atomneb.read_aeff_o_ii_ssb17(atom_rc_file, atom, ion,_
>> oii_rc_data_wave = atomneb.read_aeff_o_ii_ssb17(atom_rc_file, atom,_
⇒ion.
>>
                                                  case1, wavelength_
→range, wavelength=True)
>> print(oii_rc_data['aeff'][0])
  1.64100e-30 1.60000e-30 1.56400e-30 1.54100e-30 ...
>> n_line = len(oii_rc_data_wave['wavelength'])
>> for i in range(0, n_line):
       print(oii_rc_data_wave['wavelength'][i], oii_rc_data_wave['lower_
→term'][i], oii_rc_data_wave['upper_term'][i])
  5327.172s22p2(1S)3p 2Po
  5325.422s22p2(1S)3p 2Po
  5327.182s22p2(1D)3d 2Ge
```

**Returns** This function returns the effective recombination coefficients.

**Return type** an array of data

## **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_o\_iii\_SSB17.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- case1 (str) set for the case 'a' or 'b', defualt 'b'
- wavelength\_range (array) wavelength range e.g. [5320.0, 5330.0]
- wavelength (boolean, optional) set for returning the wavelengths
- reference (string, optional) set for the reference

```
atomneb.read_aeff_o_ii_ssb17_list(atom_rc_file)
```

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc\_o\_iii\_SSB17.fits')

**Returns** This function returns the aeff\_data\_list: {Aeff\_Data:", Extension:0, IND:long(0), Wavelength: float(0.0), Case1:", lower\_term: '', upper\_term: ''}

Return type an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_o\_iii\_SSB17.fits')

```
atomneb.read aeff o ii ssb17 references (atom rc file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_o\_iii\_SSB17.fits').

**Returns** This function returns the aeff\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_o\_iii\_SSB17.fits')

```
atomneb.read_aeff_ppb91 (atom_rc_file, atom, ion, reference=None)
```

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc\_PPB91.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_PPB91.fits')
>> atom='c'
>> ion='iii' # C II
>> cii_rc_data = atomneb.read_aeff_ppb91(atom_rc_file, atom, ion)
>> n_line = len(cii_rc_data['wavelength'])
>> for i in range(0, n_line):
>>
             print(cii_rc_data['ion'][i], cii_rc_data['case1'][i], cii_rc_
→data['wavelength'][i],
                   cii_rc_data['a'][i], cii_rc_data['b'][i], cii_rc_data[
\hookrightarrow 'C'][i],
                   cii_rc_data['d'][i], cii_rc_data['br'][i], cii_rc_data[
\rightarrow'q'][i], cii_rc_data['y'][i])
  C2+A 9903.4600 0.69700000
                                             -0.78400000
              4267.1500
                              1.0110000
                                             -0.75400000
   C2+A
   . . .
```

**Returns** This function returns the effective recombination coefficients: {Ion: ', Case1:", Wavelength:0.0, a: 0.0, b: 0.0, c: 0.0, d: 0.0, br: 0.0, y: 0.0}

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_PPB91.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'
- reference (str, optional) set for the reference

atomneb.read\_aeff\_ppb91\_list(atom\_rc\_file)

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc PPB91.fits')

**Returns** This function returns the aeff\_data\_list: { Aeff\_Data:", Extension:0.0}

Return type an array of data

**Parameters atom\_rc\_file** (str) – the FITS data file name ('rc\_PPB91.fits')

```
atomneb.read_aeff_ppb91_references(atom_rc_file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_PPB91.fits').

**Returns** This function returns the aeff\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_PPB91.fits')

```
atomneb.read aeff sh95 (atom rc file, atom, ion, reference=None, case1=None)
```

This function returns the effective recombination coefficients (Aeff) from the table extensions of the FITS data file ('rc\_SH95.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_SH95.fits')
>> atom='h'
>> ion='ii' # H I
>> hi_rc_data = atomneb.read_aeff_sh95(atom_rc_file, atom, ion)
>> print(hi_rc_data['aeff'][0])
                                 0.0000000 4.2140000e-27
   100.00000
                  500.00000
                                                                1.
→7560000e-27 ...
   . . .
```

**Returns** This function returns the effective recombination coefficients.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_SH95.fits')
- atom (str) atom name e.g. 'h'
- ion (str) ionic level e.g 'ii'
- reference (str, optional) set for the reference
- case1 (boolean, optional) set for the case 'a' or 'b', defualt 'b'

```
atomneb.read_aeff_sh95_list(atom_rc_file)
```

This function returns the list of effective recombination coefficients (Aeff) from the 1st binary table extension of the FITS data file ('rc\_SH95.fits')

**Returns** This function returns the aeff\_data\_list: { Aeff\_Data:", Extension:0.0}

Return type an array of data

**Parameters atom\_rc\_file** (str) - the FITS data file name ('rc\_SH95.fits')

```
atomneb.read_aeff_sh95_references(atom_rc_file)
```

This function returns the reference list of recombination coefficients (Aeff) from the 2nd binary table extension of the FITS data file ('rc\_SH95.fits').

**Returns** This function returns the aeff\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters** atom\_rc\_file (str) - the FITS data file name ('rc\_SH95.fits')

atomneb.read\_aij (atom\_aij\_file, atom, ion, reference=None, level\_num=None)

This function returns the transition probabilities (Aij) from the table extensions of the FITS data file ('AtomAij.fits').

## For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data', 'collection')
>> atom_aij_file = os.path.join(base_dir,data_dir, 'AtomAij.fits')
>> atom='o'
>> ion='iii'
>> reference='FFT04'
>> oiii_aij_data = atomneb.read_aij(atom_aij_file, atom, ion, reference)
>> print(oiii_aij_data['aij'][0])
   0.0000000 2.5960000e-05
                              3.0300000e-11 2.3220000e-06
                                                                  0.
→0000000 0.0021910000
  0.0000000
                 0.0000000 9.6320000e-05
                                               0.0069510000
                                                                 0.
               230.80000
\rightarrow 22550000
  0.0000000
                                  0.0000000
                                               0.020290000
                 0.0000000
→00069980000
                  576.50000
  0.0000000
                 0.0000000
                                  0.0000000
                                                  0.0000000
                                                                  1.

→6850000 0.0057770000

  0.0000000
                  0.0000000
                                  0.0000000
                                                  0.0000000
                                                                  0.
→0000000 3.7600000e-11
  0.0000000
                                  0.0000000
                                                  0.0000000
                                                                  0.
              0.0000000
→0000000
               0.0000000
```

**Returns** This function returns the aij\_data: { Aij:dblarr(n\_level,n\_level) }.

Return type an array of data

- atom\_rc\_file (str) the FITS data file name ('AtoAij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- reference (str, optional) set for the reference
- level\_num (str, optional) set for the maximum level number

```
atomneb.read_aij_list(atom_aij_file)
```

This function returns the list of transition probabilities (Aij) from the 1st binary table extension of the FITS data file ('AtomAij.fits').

**Returns** This function returns the aij\_data\_list: { Aij\_Data:", Extension:0.0}

Return type an array of data

**Parameters Atom Aij file** (str) – the FITS data file name ('AtomAij.fits')

```
atomneb.read_aij_references(atom_aij_file)
```

This function returns the reference list of transition probabilities (Aij) from the 1nd binary table extension of the FITS data file ('AtomAij.fits').

**Returns** This function returns the aij\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters Atom\_Aij\_file** (str) – the FITS data file name ('AtomAij.fits')

```
atomneb.read_elj(atom_elj_file, atom, ion, level_num=None)
```

This function returns the energy levels (Ej) from the table extensions of the FITS data file ('AtomElj.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data', 'collection')
>> atom_elj_file = os.path.join(base_dir,data_dir, 'AtomElj.fits')
>> atom='o'
>> ion='iii'
>> oiii_elj_data=atomneb.read_elj(atom_elj_file, atom, ion, level_num=6)
>> print(oiii_elj_data['j_v'])
   0.00000
               1.00000
                             2.00000
                                           2.00000
                                                        0.00000
                                                                     2.
→00000
>> print(oiii_elj_data['ej'])
   0.0000000
                  113.17800
                                    306.17400
                                                    20273.270
                                                                    43185.
            60324.790
→740
```

**Returns** This function returns the elj\_data:{ Configuration:", Term:", J:", J\_v:0.0, Ej:0.0, Reference:"}.

Return type an array of data

- atom\_elj\_file (str) the FITS data file name ('AtomElj.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- **level\_num** (*int*, *optional*) set for the maximum level number.

```
atomneb.read_elj_list(atom_elj_file)
```

This function returns the list of energy levels (Ej) from the 1st binary table extension of the FITS data file ('AtomElj.fits')

**Returns** This function returns the elj\_data\_list: { Elj\_Data:", Extension:0.0}

Return type an array of data

**Parameters atom elj file** (str) – the FITS data file name ('AtomElj.fits')

```
atomneb.read_elj_references(atom_elj_file)
```

This function returns the reference list of energy levels (Ej) from the 2nd binary table extension of the FITS data file ('AtomElj.fits').

**Returns** This function returns the aij\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters** atom\_elj\_file (str) - the FITS data file name ('AtomElj.fits')

```
atomneb.read omij (atom omij file, atom, ion, reference=None, level num=None)
```

This function returns the collision strengths (omega\_ij) from the table extensions of the FITS data file ('AtomOmij.fits').

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..', 'atomic-data', 'collection')
>> atom_omij_file = os.path.join(base_dir,data_dir, 'AtomOmij.fits')
>> atom='o'
>> ion='iii'
>> reference='SSB14'
>> oiii_omij_data=atomneb.read_omij(atom_omij_file, atom, ion,...
→reference=reference)
>> print(oiii_omij_data['level1'])
  0 1
                 1
                         1
                                 1
         4
>> print(oiii_omij_data['level1'])
  0
        2 3
>> print(oiii_omij_data['strength'][0])
                                 158.48932
   100.00000
                  125.89254
                                                 199.52623
                                                                 251.
→18864
```

**Returns** This function returns the omij\_data: { level1:0, level2:0, strength:array(temp\_steps)}.

Return type an array of data

## Params

```
atom_omij_file [in, required, type=string] the FITS data file name ('AtomOmij.fits')
atom [in, required, type=string] atom name e.g. 'o'
ion [in, required, type=string] ionic level e.g 'iii'
```

- reference (str, optional) set for the reference e.g. 'SSB14'
- level num(int, optional) set for the maximum level number.

```
atomneb.read_omij_list(atom_omij_file)
```

This function returns the list of collision strengths (omega\_ij) from the 1st binary table extension of the FITS data file ('AtomOmij.fits').

**Returns** This function returns the omij\_data\_list: { Omij\_Data:", Extension:0.0}

**Return type** an array of data

**Parameters** atom\_omij\_file (str) - the FITS data file name ('AtomOmij.fits')

```
atomneb.read_omij_references(atom_omij_file)
```

This function returns the reference list of collision strengths (omega\_ij) from the 2nd binary table extension of the FITS data file ('AtomOmij.fits').

**Returns** his function returns the aij\_data\_reference: { Reference:", Citation:"}

Return type an array of data

**Parameters** atom\_omij\_file (str) - the FITS data file name ('AtomOmij.fits')

```
atomneb.search_aeff_collection(atom_rc_file, atom, ion, br=None)
```

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rc collection.fits'), and returns the data entry.

For example:

**Returns** This function returns the Aeff\_Data.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_collection.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'
- **br** (boolean, optional) set for the branching ratios (Br), may not necessary

atomneb.search\_aeff\_he\_i\_pfsd12(atom\_rc\_file, atom, ion)

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rec\_he\_ii\_PFSD12.fits'), and returns the data entry.

### For example:

**Returns** This function returns the Aeff\_Data.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_he\_ii\_PFSD12.fits')
- atom (str) atom name e.g. 'he'
- ion (str) ionic level e.g 'ii'

atomneb.search\_aeff\_n\_ii\_fsl13 (atom\_rc\_file, atom, ion, wavelength)

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rc\_n\_iii\_FSL13.fits'), and returns the data entry.

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_n_iii_FSL13.fits')
>> atom='n'
>> ion='iii' # N II
>> wavelength=5679.56
>> list_nii_aeff_data = atomneb.search_aeff_n_ii_fsl13(atom_rc_file, atom,
→ ion, wavelength)
>> print(list_nii_aeff_data['wavelength'])
  5679.56
>> print(list_nii_aeff_data['aeff'])
  7810.00 1780.00 850.000
                                       151.000
                                                   74.4000
                                                                53.
→1000 47.4000
  7370.00 1700.00
                         886.000
                                       206.000
                                                   110.000
                                                                80.
→1000 70.8000
  7730.00 1680.00
                           900.000
                                       239.000
                                                   138.000
                                                                103.
→000 92.9000
   8520.00 1710.00
                           905.000
                                        244.000
                                                    142.000
                                                                107.
→000 97.0000
```

**Returns** This function returns the Aeff Data.

Return type an array of data

# **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_n\_iii\_FSL13.fits')
- atom (str) atom name e.g. 'n'
- ion (str) ionic level e.g 'iii'
- wavelength (int) set the wavelengths
- reference (str, optional) set for the reference

```
atomneb.search_aeff_o_ii_ssb17 (atom_rc_file, atom, ion, case1, wavelength)
```

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rc\_o\_iii\_SSB17.fits'), and returns the data entry.

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('...', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir,data_dir, 'rc_o_iii_SSB17_orl_
⇒case b.fits')
>> atom='o'
>> ion='iii' # 0 II
>> case1='B'
>> wavelength=5325.42
>> list_oii_aeff_data = atomneb.search_aeff_o_ii_ssb17(atom_rc_file, atom,
→ ion, case1, wavelength)
>> print(list_oii_aeff_data['wavelength'])
  5325.42
>> print(list_oii_aeff_data['aeff'])
  3.41800e-32 3.33300e-32 3.25700e-32 3.20900e-32 3.16800e-32 ...
```

**Returns** This function returns the Aeff Data.

**Return type** an array of data

# **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_o\_iii\_SSB17.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'
- case1 (str) set for the case 'a' or 'b', defualt 'b'
- wavelength (float) set the wavelengths

```
atomneb.search_aeff_ppb91(atom_rc_file, atom, ion)
```

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rec\_PPB91.fits'), and returns the data entry.

Returns This function returns the Aeff Data.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_PPB91.fits')
- atom (str) atom name e.g. 'c'
- ion (str) ionic level e.g 'iii'

```
atomneb.search_aeff_sh95 (atom_rc_file, atom, ion)
```

This function searches effective recombination coefficients (Aeff) for given element and ionic levels in the FITS data file ('rec\_SH95.fits'), and returns the data entry.

For example:

```
>> import atomneb
>> import numpy as np
>> import os
>> base_dir = os.getcwd()
>> data_dir = os.path.join('..', 'atomic-data-rc')
>> atom_rc_file = os.path.join(base_dir, data_dir, 'rc_SH95.fits')
>> atom='h'
>> ion='ii' # H I
>> list_hi_aeff_data = atomneb.search_aeff_sh95(atom_rc_file, atom, ion)
>> print(list_hi_aeff_data)
h_ii_aeff_a h_ii_aeff_b
```

**Returns** This function returns the Aeff\_Data.

Return type an array of data

#### **Parameters**

- atom\_rc\_file (str) the FITS data file name ('rc\_SH95.fits')
- atom (str) atom name e.g. 'h'
- ion (str) ionic level e.g 'ii'

```
atomneb.search_aij (atom_aij_file, atom, ion)
```

This function searches transition probabilities (Aij) for given element and ionic levels in the FITS data file ('AtomAij.fits'), and returns the data entry.

**Returns** This function returns the Aij\_Data.

Return type an array of data

#### **Parameters**

- Atom\_Aij\_file (str) the FITS data file name ('AtomAij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii

atomneb.search\_omij (atom\_omij\_file, atom, ion)

This function searches collision strengths (omega\_ij) for given element and ionic levels in the FITS data file ('AtomOmij.fits'), and returns the data entry.

For example:

**Returns** This function returns the Omij\_Data.

Return type an array of data

- $atom\_omij\_file$  (str) the FITS data file name ('AtomOmij.fits')
- atom (str) atom name e.g. 'o'
- ion (str) ionic level e.g 'iii'

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