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# **AtomNeb**

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## ATOMNEB API FUNCTIONS

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:

```
>> 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
>> citation = atomneb.get_aeff_collection_reference_citation(atom_rc_file,
↪ atom, ion)
>> print(citation)
Davey, A. R., Storey, P. J. and Kisielius, R., Astron.Astrophys.Suppl.,
↪ 142, 85, 2000
```

**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\_PFS12.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_he_ii_PFS12.fits')
>> atom='he'
>> ion='ii' # He I
>> reference='PFS13'
>> citation = atomneb.get_aeff_he_i_pfs12_reference_citation(atom_rc_
↪file, atom, ion, reference=reference)
>> print(citation)
Porter, R. L., Ferland, G. J., Storey, P. J. and Detisch, M. J., MNRAS,
↪ 433L, 89, 2013
```

**Returns** This function returns the Citation.

**Return type** str

**Parameters**

- **atom\_rc\_file** (*str*) – the FITS data file name ('rc\_he\_ii\_PFS12.fits')
- **atom** (*str*) – atom name e.g. 'he'
- **ion** (*str*) – ionic level e.g. 'ii'
- **reference** (*str*, *optional*) – set for the reference e.g. 'PFS13'

atomneb.get\_aeff\_n\_ii\_fsl13\_reference\_citation(atom\_rc\_file, atom, ion, refer-  
ence=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,
↪ A18; 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:

```
>> 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
>> reference='SSB17'
>> citation = atomneb.get_aeff_o_ii_ssb17_reference_citation(atom_rc_file,
↪ atom, ion)
>> print(citation)
Storey, P.J., Sochi, T. and Bastin, R. 2017, MNRAS, 470, 379; VizieR,
↪ On-line Data Catalog: VI/150
```

**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:

```
>> 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'
>> citation = atomneb.get_aeff_ppb91_reference_citation(atom_rc_file,
↪ atom, ion)
>> print(citation)
Pequignot, D., Petitjean, P. and Boisson, C. Astron.Astrophys., 251,
↪ 680, 1991
```

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**Returns** This function returns the Citation.

**Return type** str

**Parameters**

- **atom\_rc\_file**(*str*) – the FITS data file name ('rc\_PP91.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, case1=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, 41S, 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', default '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')

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'
>> 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:

```
>> 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'
>> citation = atomneb.get_omij_reference_citation(atom_omij_file, atom,
↪ion, reference)
>> print(citation)
    Storey, P. J., Sochi, T., and Badnell, N. R. 2014, Astron.Astrophys.,
↪441, 3028
```

**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:

```
>> 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
>> list_cii_aeff_reference = atomneb.list_aeff_collection_references(atom_
↪rc_file, atom, ion)
>> print(list_cii_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\_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\_PFS12.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_he_ii_PFS12.fits')
>> atom='he'
>> ion='ii' # He I
>> list_hei_aeff_reference = atomneb.list_aeff_he_i_pfsd12_
↪references(atom_rc_file, atom, ion)
>> print(list_hei_aeff_reference)
PFS12 PFS13
```

**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\_PFS12.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:

```
>> 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
>> list_nii_aeff_reference = atomneb.list_aeff_n_ii_fsl13_references (atom_
↪rc_file, atom, ion)
>> print(list_nii_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\_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_oi_aeff_reference = atomneb.list_aeff_o_ii_ssb17_references(atom_
↳ rc_file, atom, ion)
>> print(list_oi_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

**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.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:

```
>> 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
>> list_hi_aeff_reference = atomneb.list_aeff_sh95_references(atom_rc_
↪file, atom, ion)
>> print(list_hi_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\_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:

```
>> 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'
>> list_oiii_aij_reference = atomneb.list_aij_references(atom_aij_file,
↪atom, ion)
>> print(list_oiii_aij_reference)
FFT04-SZ00 FFT04 GMZ97-WFD96 SZ00-WFD96
```

**Returns** This function returns the references.

**Return type** an array of data

**Parameters**

- **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_oiij_omij_reference = atomneb.list_omij_references(atom_omij_file,
→ atom, ion)
>> print(list_oiij_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 ( $A_{eff}$ ) from the table extensions of the FITS data file (‘rc\_collection.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_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.
→24310000    -0.88000000
    962.00000    1.0998000    -0.0042000000    -0.027900000    -0.
→22940000    -0.96560000
    ...
```

**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')

`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\_PFS12.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_PFS12.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, 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\_PFS12.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\_PFS12.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\_PFS12.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\_PFS12.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\_PFS12.fits')

`atomneb.read_aeff_n_iii_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').

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_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])
255.000      79.5000      47.3000      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,
↳ case1, wavelength_range)
>> 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', default '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.c[i],
>>           cii_rc_data.d[i], cii_rc_data.br[i], cii_rc_data.q[i],
↵cii_rc_data.y[i])
C2+A      9903.4600      0.69700000      -0.78400000      ...
C2+A      4267.1500      1.0110000      -0.75400000      ...
...
```

**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])
100.00000      500.00000      0.0000000      4.2140000e-27      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', default '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)
0.0000000  2.5960000e-05  3.0300000e-11  2.3220000e-06  0.
↪0000000  0.0021910000
0.0000000  0.0000000  9.6320000e-05  0.0069510000  0.
↪22550000  230.80000
0.0000000  0.0000000  0.0000000  0.020290000  0.
↪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  0.0000000  0.
↪0000000  0.0000000
```

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

**Return type** an array of data

**Parameters**

- `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(np.asarray(oiii_elj_data.j_v))
0.00000    1.00000    2.00000    2.00000    0.00000    2.
↪00000
>> print(np.asarray(oiii_elj_data.ej))
0.0000000    113.17800    306.17400    20273.270    43185.
↪740    60324.790
```

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

**Return type** an array of data

**Parameters**

- `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(np.asarray(oiii_omij_data.level1))
    0      1      1      1      1      2      2      2      3
↳ 3      4
>> print(np.asarray(oiii_omij_data.level2))
    0      2      3      4      5      3      4      5      4
↳ 5      5
>> print(np.asarray(oiii_omij_data.strength)[0])
    100.00000      125.89254      158.48932      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'

**Parameters**

- **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:

```
>> 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
>> list_cii_aeff_data = atomneb.search_aeff_collection(atom_rc_file, atom,
↪ ion)
>> print(list_cii_aeff_data)
c_iii_aeff
```

**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\_PFS12.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_he_ii_PFS12.fits')
>> atom='he'
>> ion='ii' # He I
>> list_hei_aeff_data = atomneb.search_aeff_he_i_pfs12(atom_rc_file,
↪ atom, ion)
>> print(list_hei_aeff_data)
he_ii_aeff_PFS12 he_ii_aeff_PFS13
```

**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\_PFS12.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.

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_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(np.asarray(list_nii_aeff_data.wavelength))
5679.56
>> print(np.asarray(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' # O II
>> case1='B'
>> wavelength=5325.42
>> list_oi_iaeff_data = atomneb.search_aeff_o_ii_ssb17(atom_rc_file, atom,
↪ ion, case1, wavelength)
>> print(np.asarray(list_oi_iaeff_data.wavelength))
5325.42
>> print(np.asarray(list_oi_iaeff_data.iaeff))
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', default '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.

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'
>> list_cii_aeff_data = atomneb.search_aeff_ppb91(atom_rc_file, atom, ion)
>> print(list_cii_aeff_data)
c_iii_aeff
```

**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.

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'
>> list_oiii_aij_data = atomneb.search_aij(atom_aij_file, atom, ion)
>> print(list_oiii_aij_data)
    o_iii_aij_FFT04-SZ00 o_iii_aij_FFT04 o_iii_aij_GMZ97-WFD96 o_iii_aij_
↪SZ00-WFD96
```

**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:

```
>> 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_data = atomneb.search_omij(atom_omij_file, atom, ion)
>> print(list_oiii_omij_data)
    o_iii_omij_AK99 o_iii_omij_LB94 o_iii_omij_Pal12-AK99 o_iii_omij_SSB14
```

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

**Return type** an array of data

**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'

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