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# **CRRLpy Documentation**

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## CRRLPY

Tools for processing Carbon **R**adio **R**ecombination **L**ine spectra.

The models are not shipped with the modules and scripts.

The documentation can be found in: <http://astrofle.github.io/CRRLpy/>





## CRRLPY.CRRLS MODULE

`crrlpy.crrls.alphanum_key(s)`

Turn a string into a list of string and number chunks.

**Parameters** *s* – String

**Returns** List with strings and integers.

**Return type** list

**Example**

```
>>> alphanum_key("z23a")
["z", 23, "a"]
```

`crrlpy.crrls.average(data, axis, n)`

Averages data along the given axis by combining n adjacent values.

**Parameters**

- **data** (*numpy array*) – Data to average along a given axis.
- **axis** (*int*) – Axis along which to average.
- **n** (*int*) – Factor by which to average.

**Returns** Data decimated by a factor n along the given axis.

**Return type** numpy array

`crrlpy.crrls.best_match_idx(value, array, tol)`

Searchs for the best match to a value inside an array given a tolerance.

**Parameters**

- **value** (*float*) – Value to find inside the array.
- **tol** (*float*) – Tolerance for match.
- **array** (*numpy.array*) – List to search for the given value.

**Returns** Best match for val inside array.

**Return type** float

`crrlpy.crrls.best_match_idx2(value, array)`

Searchs for the index of the closest entry to value inside an array.

**Parameters**

- **value** (*float*) – Value to find inside the array.
- **array** (*list or numpy.array*) – List to search for the given value.

**Returns** Best match index for the value inside array.

**Return type** float

**Example**

```
>>> a = [1, 2, 3, 4]
>>> best_match_idx2(3, a)
2
```

`crrlpy.crrls.best_match_value(value, array)`

Searchs for the closest occurrence of value in array.

**Parameters**

- **value** (*float*) – Value to find inside the array.
- **array** (*list or numpy.array*) – List to search for the given value.

**Returns** Best match for the value inside array.

**Return type** float.

**Example**

```
>>> a = [1, 2, 3, 4]
>>> best_match_value(3.5, a)
3
```

`crrlpy.crrls.blank_lines(freq, tau, reffreqs, v0, dv)`

Blanks the lines in a spectra.

**Parameters**

- **freq** (*array, MHz*) – Frequency axis of the spectra.
- **tau** (*array*) – Optical depth axis of the spectra.
- **reffreqs** (*list*) – List with the reference frequency of the lines. Should be the rest frequency.
- **v0** (*float*) – Velocity shift to apply to the lines defined by *reffreq*. (km/s)
- **dv** (*float*) – Velocity range to blank around the lines. (km/s)

`crrlpy.crrls.blank_lines2(freq, tau, reffreqs, dv)`

Blanks the lines in a spectra.

**Parameters**

- **freq** (*array*) – Frequency axis of the spectra. (MHz)
- **tau** (*array*) – Optical depth axis of the spectra.
- **reffreqs** (*list*) – List with the reference frequency of the lines. Should be the rest frequency.
- **dv** (*float*) – Velocity range to blank around the lines. (km/s)

`crrlpy.crrls.df2dv(f0, df)`

Convert a frequency delta to a velocity delta given a central frequency.

**Parameters**

- **f0** (*float*) – Rest frequency. (Hz)
- **df** (*float*) – Frequency delta. (Hz)

**Returns** The equivalent velocity delta for the given frequency delta.

**Return type** float in Hz

`crllpy.crlls.dv2df(f0, dv)`

Convert a velocity delta to a frequency delta given a central frequency.

**Parameters**

- **f0** (*float*) – Rest frequency. (Hz)
- **dv** (*float*) – Velocity delta. (m/s)

**Returns** The equivalent frequency delta for the given velocity delta.

**Return type** float in  $\text{m s}^{-1}$

`crllpy.crlls.dv_minus_doppler(dV, ddV, dD, ddD)`

Returns the Lorentzian contribution to the line width assuming that the line has a Voigt profile.

**Parameters**

- **dV** (*float*) – Total line width
- **ddV** (*float*) – Uncertainty in the total line width.
- **dD** (*float*) – Doppler contribution to the line width.
- **ddD** – Uncertainty in the Doppler contribution to the line width.

**Returns** The Lorentz contribution to the total line width.

**Return type** float

`crllpy.crlls.dv_minus_doppler2(dV, ddV, dD, ddD)`

Returns the Lorentzian contribution to the line width assuming that the line has a Voigt profile.

**Parameters**

- **dV** (*float*) – Total line width
- **ddV** (*float*) – Uncertainty in the total line width.
- **dD** (*float*) – Doppler contribution to the line width.
- **ddD** – Uncertainty in the Doppler contribution to the line width.

**Returns** The Lorentz contribution to the total line width.

**Return type** float

`crllpy.crlls.f2n(f, line, n_max=1500)`

Converts a given frequency to a principal quantum number  $n$  for a given line.

**Parameters**

- **f** (*array*) – Frequency to convert. (MHz)
- **line** (*string*) – The equivalent  $n$  will be referenced to this line.
- **n\_max** (*int*) – Maximum  $n$  number to include in the search. (optional, Default 1)

**Returns** Corresponding  $n$  for a given frequency and line. If the frequency is not an exact match, then it will return an empty array.

**Return type** array

`crllpy.crlls.find_lines_sb(freq, line, z=0, verbose=False)`

Finds if there are any lines of a given type in the frequency range. The line frequencies are corrected for redshift.

**Parameters**

- **freq** (*array*) – Frequency axis in which to search for lines.
- **line** (*string*) – Line type to search for.
- **z** (*float*) – Redshift to apply to the rest frequencies.
- **verbose** (*bool*) – Verbose output?

**Returns** Lists with the principal quantum number and the reference frequency of the line. The frequencies are redshift corrected in MHz.

**Return type** array.

**Example**

```
>>> freq = [10, 11]
>>> ns, rf = crrls.find_lines_sb(freq, 'CIalpha')
>>> ns
array([ 843.,  844.,  845.,  846.,  847.,  848.,  849.,  850.,  851.,
        852.,  853.,  854.,  855.,  856.,  857.,  858.,  859.,  860.,
        861.,  862.,  863.,  864.,  865.,  866.,  867.,  868.,  869.] )
```

`crrlpy.crlls.fit_continuum(x, y, degree, p0)`

Divide tb by given a model and starting parameters p0. Returns: tb/model - 1

`crrlpy.crlls.fit_model(x, y, model, p0, wy=None, mask=None)`

Fits a model to the data defined by x and y. It uses p0 as starting values.

**Parameters**

- **x** (*array*) – Abscissa values of the data to be fit.
- **y** (*array*) – Ordinate values of the data to be fit.
- **model** (*callable*) – Model to be fit.
- **p0** (*dict*) – Dictionary with the starting values for the fit.
- **wy** (*array*) – Weights of the ordinate values. (Optional)
- **mask** (*array*) – Mask to apply to the x and y values. (Optional)

**Returns** An object containing the results of the fit.

**Return type** `lmfit.model.ModelResult`

`crrlpy.crlls.freq2vel(f0, f)`

Convert a frequency axis to a velocity axis given a central frequency. Uses the radio definition of velocity.

**Parameters**

- **f0** (*float*) – Rest frequency for the conversion. (Hz)
- **f** (*numpy array*) – Frequencies to be converted to velocity. (Hz)

**Returns** f converted to velocity given a rest frequency  $f_0$ .

**Return type** numpy array

`crrlpy.crlls.fwhm2sigma(fwhm)`

Converts a FWHM to the standard deviation,  $\sigma$  of a Gaussian distribution.

**Parameters** **fwhm** (*array*) – FWHM of the Gaussian.

**Returns** Equivalent standard deviation of a Gaussian with a Full Width at Half Maximum *fwhm*.

**Return type** array

**Example**

```
>>> 1/fwhm2sigma(1)
2.3548200450309493
```

`crrlpy.crrls.gauss_area(amplitude, sigma)`

Returns the area under a Gaussian of a given amplitude and sigma.

**Parameters**

- **amplitude** – Amplitude of the Gaussian,  $A$ .
- **sigma** (*array*) – Standard deviation fo the Gaussian,  $\sigma$ .

**Returns** The area under a Gaussian of a given amplitude and standard deviation.

**Return type** array

`crrlpy.crrls.gauss_area_err(amplitude, amplitude_err, sigma, sigma_err)`

Returns the error on the area of a Gaussian of a given *amplitude* and *sigma* with their corresponding errors. It assumes no correlation between *amplitude* and *sigma*.

**Parameters**

- **amplitude** (*array*) – Amplitude of the Gaussian.
- **amplitude\_err** (*array*) – Error on the amplitude.
- **sigma** – Standard deviation of the Gaussian.
- **sigma\_err** – Error on sigma.

**Returns** The error on the area.

**Return type** array

`crrlpy.crrls.gaussian(x, sigma, center, amplitude)`

Gaussian function in one dimension.

**Parameters**

- **x** (*array*) – x values for which to evaluate the Gaussian.
- **sigma** (*float*) – Standard deviation of the Gaussian.
- **center** (*float*) – Center of the Gaussian.
- **amplitude** (*float*) – Amplitude of the Gaussian.

**Returns** Gaussian function of the given amplitude and standard deviation evaluateated at x.

**Return type** array

`crrlpy.crrls.get_axis(header, axis)`

Constructs a cube axis.

**Parameters**

- **header** (*pyfits header*) – Fits cube header.
- **axis** (*int*) – Axis to reconstruct.

**Returns** cube axis

**Return type** numpy array

`crrlpy.crrls.get_line_mask(freq, reffreq, v0, dv)`

Return a mask with ranges where a line is expected in the given frequency range for a line with a given reference frequency at expected velocity  $v_0$  and line width  $dv_0$ .

**Parameters**

- **freq** (*numpy array or list*) – Frequency axis where the line is located.
- **reffreq** (*float*) – Reference frequency for the line.
- **v0** (*float, km/s*) – Velocity of the line.
- **dv** (*float, km/s*) – Velocity range to mask.

**Returns** Mask centered at the line center and width  $dv_0$  referenced to the input *freq*.

`crrlpy.crrls.get_line_mask2(freq, reffreq, dv)`

Return a mask with ranges where a line is expected in the given frequency range for a line with a given reference frequency and line width  $dv$ .

**Parameters**

- **freq** (*numpy array or list*) – Frequency axis where the line is located.
- **reffreq** (*float*) – Reference frequency for the line.
- **dv** (*float, km/s*) – Velocity range to mask.

**Returns** Mask centered at the line center and width  $dv_0$  referenced to the input *freq*.

`crrlpy.crrls.get_min_sep(array)`

Get the minimum element separation in an array.

**Parameters** **array** (*array*) – Array where the minimum separation is wanted.

**Returns** The minimum separation between the elements in *array*.

**Return type** *float*

`crrlpy.crrls.get_rchi2(x_obs, x_mod, y_obs, y_mod, dy_obs, dof)`

Computes the reduced  $\chi$  squared,  $\chi^2_\nu = \chi^2/dof$ .

**Parameters**

- **x\_obs** (*array*) – Abscissa values of the observations.
- **x\_mod** (*array*) – Abscissa values of the model.
- **y\_obs** (*array*) – Ordinate values of the observations.
- **y\_mod** (*array*) – Ordinate values of the model.
- **dy\_obs** (*array*) – Error on the ordinate values of the observations.
- **dof** (*float*) – Degrees of freedom.

`crrlpy.crrls.get_rms(data, axis=None)`

Computes the rms of the given data.

**Parameters**

- **data** (*numpy array or list*) – Array with values where to compute the rms.
- **axis** (*int*) – Axis over which to compute the rms. Default: None

**Returns** The rms of data.

$$\text{rms} = \sqrt{\langle \text{data} \rangle^2 + V[\text{data}]}$$

where  $V$  is the variance of the data.

`crrlpy.crpls.is_number(str)`

Checks whether a string is a number or not.

**Parameters** `str` (*string*) – String.

**Returns** True if `str` can be converted to a float.

**Return type** `bool`

**Example**

```
>>> is_number('10')
True
```

`crrlpy.crpls.linear(x, a, b)`

Linear model.

**Parameters**

- `x` (*array*) – x values where to evaluate the line.
- `a` (*float*) – Slope of the line.
- `b` (*float*) – y value for x equals 0.

**Returns** A line defined by  $ax + b$ .

**Return type** `array`

`crrlpy.crpls.load_model(prop, specie, temp, dens, other=None)`

Loads a model for the CRRL emission.

`crrlpy.crpls.load_ref(specie, trans)`

Loads the reference spectrum for the specified atomic specie and transition. Available species and transitions: CI alpha CI beta CI delta CI gamma CI13 alpha HeI alpha HeI beta HI alpha HI beta SI alpha SI beta

`crrlpy.crpls.load_ref2(transition)`

Loads the reference spectrum for the specified atomic specie and transition. Available transitions: CIalpha CIbeta CIDelta CIGamma CI13alpha HeIalpha HeIbeta HIalpha HIBeta SIalpha SIBeta

`crrlpy.crpls.lookup_freq(n, specie, trans)`

Returns the frequency of a given transition.

`crrlpy.crpls.lorentz_width(n, ne, Te, Tr, W, dn=1)`

Gives the Lorentzian line width due to a combination of radiation and collisional broadening. The width is the FWHM in Hz. It uses the models of Salgado et al. (2015).

**Parameters**

- `n` (*array*) – Principal quantum number for which to evaluate the Lorentz widths.
- `ne` (*float*) – Electron density to use in the collisional broadening term.
- `Te` (*float*) – Electron temperature to use in the collisional broadening term.
- `Tr` (*float*) – Radiation field temperature.
- `W` (*float*) – Cloud covering factor used in the radiation broadening term.
- `dn` (*int*) – Separation between the levels of the transition. e.g.,  $dn=1$  for CIalpha.

**Returns** The Lorentz width of a line due to radiation and collisional broadening.

**Return type** `array`

`crrlpy.crrls.mask_outliers(data, m=2)`

Masks values larger than  $m$  times the data median. This is similar to sigma clipping.

**Parameters** `data` (`array`) – Data to mask.

**Returns** An array of the same shape as `data` with `True` where the data should be flagged.

**Return type** `array`

**Example**

```
>>> data = [1,2,3,4,5,6]
>>> mask_outliers(data, m=1)
array([ True, False, False, False, False,  True], dtype=bool)
```

`crrlpy.crrls.n2f(n, line, n_min=1, n_max=1500, unitless=True)`

Converts a given principal quantum number  $n$  to the frequency of a given line.

`crrlpy.crrls.natural_sort(list)`

Sort the given list in the way that humans expect. Sorting is done in place.

**Parameters** `list` (`list`) – List to sort.

**Example**

```
>>> my_list = ['spec_3', 'spec_4', 'spec_1']
>>> natural_sort(my_list)
>>> my_list
['spec_1', 'spec_3', 'spec_4']
```

`crrlpy.crrls.plot_fit(fig, x, y, fit, params, vparams, sparams, rms, x0, refs, refs_cb=None, refs_cd=None, refs_cg=None)`

`crrlpy.crrls.plot_fit_single(fig, x, y, fit, params, rms, x0, refs, refs_cb=None, refs_cd=None, refs_cg=None)`

`crrlpy.crrls.plot_model(x, y, xm, ym, out)`

`crrlpy.crrls.plot_spec_vel(out, x, y, fit, A, Aerr, x0, x0err, sx, sxerr)`

`crrlpy.crrls.pressure_broad(n, Te, ne)`

Pressure induced broadening in Hz. Shaver (1975)

`crrlpy.crrls.pressure_broad_coefs(Te)`

Defines the values of the constants  $a$  and  $\gamma$  that go into the collisional broadening formula of Salgado et al. (2015).

**Parameters** `Te` (`float`) – Electron temperature.

**Returns** The values of  $a$  and  $\gamma$ .

**Return type** `list`

`crrlpy.crrls.pressure_broad_salgado(n, Te, ne, dn=1)`

Pressure induced broadening in Hz. This gives the FWHM of a Lorentzian line. Salgado et al. (2015)

**Parameters**

- `n` (`float` or `array`) – Principal quantum number for which to compute the line broadening.
- `Te` (`float`) – Electron temperature to use when computing the collisional line width.
- `ne` (`float`) – Electron density to use when computing the collisional line width.



- **dn** (*int*) – Difference between the upper and lower level for which the line width is computed. (default 1)

**Returns** The collisional broadening FWHM in Hz using Salgado et al. (2015) formulas.

**Return type** float or array

`crllpy.crlls.radiation_broad(n, W, Tr)`

Radiation induced broadening in Hz.

`crllpy.crlls.radiation_broad_salgado(n, W, Tr)`

Radiation induced broadening in Hz. This gives the FWHM of a Lorentzian line. Salgado et al. (2015)

`crllpy.crlls.radiation_broad_salgado_general(n, W, Tr, nu0, alpha)`

Radiation induced broadening in Hz. This gives the FWHM of a Lorentzian line. The expression is valid for power law like radiation fields. Salgado et al. (2015)

`crllpy.crlls.sigma2fwhm(sigma)`

Converts the  $\sigma$  parameter of a Gaussian distribution to its FWHM.

**Parameters** **sigma** (*float*) –  $\sigma$  value of the Gaussian distribution.

**Returns** The FWHM of a Gaussian with a standard deviation  $\sigma$ .

**Return type** float

`crllpy.crlls.sigma2fwhm_err(dsigma)`

Converts the error on the sigma parameter of a Gaussian distribution to the error on the FWHM.

**Parameters** **dsigma** – Error on sigma of the Gaussian distribution.

**Returns** The error on the FWHM of a Gaussian with a standard deviation  $\sigma$ .

**Return type** float

`crllpy.crlls.stack_interpol(spectra, vmin, vmax, dv, show=True, rmsvec=False)`

`crllpy.crlls.stack_irregular(lines, window='', **kargs)`

Stacks spectra by adding them together and then convolving with a window to reduce the noise. Available window functions: Gaussian, Savitzky-Golay and Wiener.

`crllpy.crlls.temp2tau(x, y, model, p0, wy=None, mask=None)`

Converts a temperature to optical depth. It will fit the continuum using model and then subtract it and divide by it.

**Parameters**

- **x** (*array*) – x values.
- **y** (*array*) – y values to be converted into optical depths.
- **model** (*callable*) – Model to fit to the continuum.
- **p0** – Starting values for the model to be fit to the continuum.
- **wy** (*array*) – Weights for the y values. (Optional)
- **mask** (*array*) – Mask to apply to the x and y values.

**Returns** y/model - 1.

**Return type** array

`crllpy.crlls.tryint(str)`

Returns True if *str* is an integer.

**Parameters** **str** (*string*) – String to check.

**Returns** True is str can be cast to an int.

**Return type** bool

`crrlpy.crpls.vel2freq(f0, vel)`

Convert a velocity axis to a frequency axis given a central frequency. Uses the radio definition,  $\nu = f_0(1 - v/c)$ .

**Parameters**

- **f0** (*float*) – Rest frequency in Hz.
- **vel** (*float or array*) – Velocity to convert in m/s.

**Returns** The frequency which is equivalent to vel.

**Return type** float or array

`crrlpy.crpls.voigt(x, sigma, gamma, center, amplitude)`

The Voigt line shape in terms of its physical parameters.

**Parameters**

- **x** – independent variable
- **sigma** – HWHM of the Gaussian
- **gamma** – HWHM of the Lorentzian
- **center** – the line center
- **amplitude** – the line area

`crrlpy.crpls.voigt_(x, y)`

`crrlpy.crpls.voigt_area(amp, fwhm, gamma, sigma)`

Returns the area under a Voigt profile. This approximation has an error of less than 0.5%

`crrlpy.crpls.voigt_area_err(area, amp, damp, fwhm, dfwhm, gamma, sigma)`

Returns the error of the area under a Voigt profile. Assumes that the parameter c has an error of 0.5%.

`crrlpy.crpls.voigt_fwhm(dD, dL)`

Computes the FWHM of a Voigt profile. [http://en.wikipedia.org/wiki/Voigt\\_profile#The\\_width\\_of\\_the\\_Voigt\\_profile](http://en.wikipedia.org/wiki/Voigt_profile#The_width_of_the_Voigt_profile)

$$FWHM_V = 0.5346dL + \sqrt{0.2166dL^2 + dD^2}$$

**Parameters**

- **dD** (*array*) – FWHM of the Gaussian core.
- **dL** (*array*) – FWHM of the Lorentz wings.

**Returns** The FWHM of a Voigt profile.

**Return type** array

`crrlpy.crpls.voigt_fwhm_err(dD, dL, ddD, ddL)`

Computes the error in the FWHM of a Voigt profile. [http://en.wikipedia.org/wiki/Voigt\\_profile#The\\_width\\_of\\_the\\_Voigt\\_profile](http://en.wikipedia.org/wiki/Voigt_profile#The_width_of_the_Voigt_profile)

**Parameters**

- **dD** (*array*) – FWHM of the Gaussian core.
- **dL** (*array*) – FWHM of the Lorentz wings.
- **ddD** (*array*) – Error on the FWHM of the Gaussian.
- **ddL** (*array*) – Error on the FWHM of the Lorentzian.

**Returns** The FWHM of a Voigt profile.

**Return type** array

`crrlpy.crpls.voigt_peak(A, alphaD, alphaL)`

Gives the peak of a Voigt profile given its Area and the Half Width at Half Maximum of the Gaussian and Lorentz profiles.

**Parameters**

- **A** (array) – Area of the Voigt profile.
- **alphaD** (array) – HWHM of the Gaussian core.
- **alphaL** (array) – HWHM of the Lorentz wings.

**Returns** The peak of the Voigt profile.

**Return type** array

`crrlpy.crpls.voigt_peak2area(peak, alphaD, alphaL)`

Converts the peak of a Voigt profile into the area under the profile given the Half Width at Half Maximum of the profile components.

**Parameters**

- **peak** (array) – Peak of the Voigt profile.
- **alphaD** (array) – HWHM of the Gaussian core.
- **alphaL** (array) – HWHM of the Lorentz wings.

**Returns** The area under the Voigt profile.

**Return type** array

`crrlpy.crpls.voigt_peak_err(peak, A, dA, alphaD, dalphaD)`

Gives the error on the peak of the Voigt profile. It assumes no correlation between the parameters and that they are normally distributed.

**Parameters**

- **peak** (array) – Peak of the Voigt profile.
- **A** – Area under the Voigt profile.
- **dA** (array) – Error on the area A.
- **alphaD** (array) – HWHM of the Gaussian core.



## CRRLPY.MODELS.RRLMOD MODULE

`crrlpy.models.rrlmod.I_Bnu(specie, Z, n, Inu_func, *args)`

Calculates the product  $B_{n+\Delta n, n} I_\nu$  to compute the line broadening due to a radiation field  $I_\nu$ .

### Parameters

- **specie** (*string*) – Atomic specie to calculate for.
- **n** (*int or list*) – Principal quantum number at which to evaluate  $\frac{2}{\pi} \sum_{\Delta n} B_{n+\Delta n, n} I_{n+\Delta n, n}(\nu)$ .
- **Inu\_func** (*function*) – Function to call and evaluate  $I_{n+\Delta n, n}(\nu)$ . It's first argument must be the frequency.
- **args** – Arguments to *Inu\_func*. The frequency must be left out. The frequency will be passed internally in units of MHz. Use the same unit when required. *Inu\_func* must take the frequency as first parameter.

**Returns** (Hz)

**Return type** `array`

### Example

```
>>> I_Bnu('CI', 1., 500, I_broken_plaw, 800, 26*u.MHz.to('Hz'), -1., -2.6)
array([ 6.65540582])
```

`crrlpy.models.rrlmod.I_broken_plaw(nu, Tr, nu0, alpha1, alpha2)`

Returns the blackbody function evaluated at nu. As temperature a broken power law is used. The power law shape has parameters: Tr, nu0, alpha1 and alpha2.

### Parameters

- **nu** ((Hz) or `astropy.units.Quantity`) – Frequency. (Hz) or `astropy.units.Quantity`
- **Tr** – Temperature at nu0. (K) or `astropy.units.Quantity`
- **nu0** – Frequency at which the spectral index changes. (Hz) or `astropy.units.Quantity`
- **alpha1** – spectral index for  $\nu < \nu_0$
- **alpha2** – spectral index for  $\nu \geq \nu_0$

**Returns** Specific intensity in  $\text{erg cm}^{-2} \text{Hz}^{-1} \text{s}^{-1} \text{sr}^{-1}$ .  
`tripy.analytic_functions.blackbody.blackbody_nu`

See `as-`

**Return type** `astropy.units.Quantity`

`crrlpy.models.rrlmod.I_cont(nu, Te, tau, I0, unitless=False)`

Computes the specific intensity due to a blackbody at temperature  $T_e$  and optical depth  $\tau$ . It considers that there is background radiation with  $I_0$ .

**Parameters**

- **nu** ((Hz) or [astropy.units.Quantity](#)) – Frequency.
- **Te** – Temperature of the source function. (K) or [astropy.units.Quantity](#)
- **tau** – Optical depth of the medium.
- **I0** – Specific intensity of the background radiation. Must have units of erg / (cm<sup>2</sup> Hz s sr) or see *unitless*.
- **unitless** – If True the return

**Returns** The specific intensity of a ray of light after traveling in an LTE medium with source function  $B_\nu(T_e)$  after crossing an optical depth  $\tau_\nu$ . The units are erg / (cm<sup>2</sup> Hz s sr). See [astropy.analytic\\_functions.blackbody.blackbody\\_nu](#)

```
crrlpy.models.rrlmod.I_external (nu, Tbkg, Tff, tau_ff, Tr, nu0=<Quantity 100000000.0 MHz>,
                                   alpha=-2.6)
```

This method is equivalent to the IDL routine

**Parameters** **nu** – Frequency. (Hz) or [astropy.units.Quantity](#)

```
crrlpy.models.rrlmod.I_total (nu, Te, tau, I0, eta)
```

```
crrlpy.models.rrlmod.Mdn (dn)
```

Gives the  $M(\Delta n)$  factor for a given  $\Delta n$ . ref. Menzel (1968)

**Parameters** **dn** –  $\Delta n$ .

**Returns**  $M(\Delta n)$

**Return type** [float](#)

**Example**

```
>>> Mdn(1)
0.1908
>>> Mdn(5)
0.001812
```

```
crrlpy.models.rrlmod.broken_plaw (nu, nu0, T0, alpha1, alpha2)
```

Defines a broken power law.

$$T(\nu) = T_0 \left( \frac{\nu}{\nu_0} \right)^{\alpha_1} \quad \text{if } \nu < \nu_0$$
$$T(\nu) = T_0 \left( \frac{\nu}{\nu_0} \right)^{\alpha_2} \quad \text{if } \nu \geq \nu_0$$

**Parameters**

- **nu** – Frequency.
- **nu0** – Frequency at which the power law breaks.
- **T0** – Value of the power law at nu0.
- **alpha1** – Index of the power law for nu<nu0.
- **alpha2** – Index of the power law for nu>=nu0.

**Returns** Broken power law evaluated at nu.

```
crrlpy.models.rrlmod.eta (freq, Te, ne, nion, Z, Tr, trans, n_max=1500)
```

Returns the correction factor for the Planck function.

```
crllpy.models.rllmod.itau(temp, dens, line, n_min=5, n_max=1000, other='', verbose=False,
                           value='itau')
```

Gives the integrated optical depth for a given temperature and density. The emission measure is unity. The output units are Hz.

#### Parameters

- **temp** (*string*) – Electron temperature. Must be a string of the form ‘8d1’.
- **dens** (*float*) – Electron density.
- **line** (*string*) – Line to load models for.
- **n\_min** (*int*) – Minimum n value to include in the output. Default 1
- **n\_max** (*int*) – Maximum n value to include in the output. Default 1500, Maximum allowed value 9900
- **other** (*string*) – String to search for different radiation fields and others.
- **verbose** (*bool*) – Verbose output?
- **value** (*string*) – ['itau','bbnMdn','None'] Value to output. itau will output the integrated optical depth. bbnMdn will output the  $\beta_{n,n'} b_n$  times the oscillator strenght  $M(\Delta n)$ .

**Returns** The principal quantum number and its asociated value.

```
crllpy.models.rllmod.itau_h(temp, dens, trans, n_max=1000, other='', verbose=False,
                             value='itau')
```

Gives the integrated optical depth for a given temperature and density. The emission measure is unity. The output units are Hz.

```
crllpy.models.rllmod.itau_norad(n, te, b, dn, mdn)
```

Returns the optical depth with only the approximate solution to the radiative transfer problem.

```
crllpy.models.rllmod.j_line_lte(n, ne, nion, Te, Z, trans)
```

```
crllpy.models.rllmod.kappa_cont(freq, Te, ne, nion, Z)
```

Computes the absorption coefficient for the free-free process.

```
crllpy.models.rllmod.kappa_cont_base(nu, Te, ne, nion, Z)
```

```
crllpy.models.rllmod.kappa_line(Te, ne, nion, Z, Tr, trans, n_max=1500)
```

Computes the line absorption coefficient for CRRLs between levels  $n_i$  and  $n_f$ ,  $n_i > n_f$ . This can only go up to  $n_{\max}$  1500 because of the tables used for the Einstein Anm coefficients.

#### Parameters

- **Te** (*float*) – Electron temperature of the gas. (K)
- **ne** (*float*) – Electron density. ( $\text{cm}^{-3}$ )
- **nion** (*float*) – Ion density. ( $\text{cm}^{-3}$ )
- **Z** (*int*) – Electric charge of the atoms being considered.
- **Tr** (*float*) – Temperature of the radiation field felt by the gas. This specifies the temperature of the field at 100 MHz. (K)
- **trans** (*string*) – Transition for which to compute the absorption coefficient.
- **n\_max** (*int*<1500) – Maximum principal quantum number to include in the output.

#### Returns

**Return type** `array`

`crrlpy.models.rrlmod.kappa_line_lte` (*nu*, *Te*, *ne*, *nion*, *Z*, *Tr*, *line*, *n\_min=1*, *n\_max=1500*)

Returns the line absorption coefficient under LTE conditions.

#### Parameters

- **nu** (*array*) – Frequency. (Hz)
- **Te** (*float*) – Electron temperature of the gas. (K)
- **ne** (*float*) – Electron density. ( $\text{cm}^{-3}$ )
- **nion** (*float*) – Ion density. ( $\text{cm}^{-3}$ )
- **Z** (*int*) – Electric charge of the atoms being considered.
- **Tr** (*float*) – Temperature of the radiation field felt by the gas. This specifies the temperature of the field at 100 MHz. (K)
- **trans** (*string*) – Transition for which to compute the absorption coefficient.
- **n\_max** (*int*<1500) – Maximum principal quantum number to include in the output.

#### Returns

**Return type** `array`

`crrlpy.models.rrlmod.level_pop_lte` (*n*, *ne*, *nion*, *Te*, *Z*)

Returns the level population of level *n*. The return has units of  $\text{cm}^{-3}$ .

`crrlpy.models.rrlmod.load_betabn` (*temp*, *dens*, *other=''*, *trans='CIalpha'*, *verbose=False*)

Loads a model for the CRRL emission.

`crrlpy.models.rrlmod.load_betabn_h` (*temp*, *dens*, *other=''*, *trans='alpha'*, *verbose=False*)

Loads a model for the HRRL emission.

`crrlpy.models.rrlmod.load_bn` (*temp*, *dens*, *other=''*)

Loads the bn values from the CRRL models.

`crrlpy.models.rrlmod.load_bn2` (*temp*, *dens*, *other=''*)

Loads the bn values from the CRRL models.

`crrlpy.models.rrlmod.load_itaui_all` (*trans='CIalpha'*, *n\_min=5*, *n\_max=1000*, *verbose=False*,  
*value='itau'*)

Loads all the available models for Carbon.

`crrlpy.models.rrlmod.load_itaui_all_hydrogen` (*trans='alpha'*, *n\_max=1000*, *ver-*  
*bose=False*, *value='itau'*)

Loads all the available models for Hydrogen.

`crrlpy.models.rrlmod.load_itaui_all_match` (*trans\_out='alpha'*, *trans\_tin='beta'*,  
*n\_max=1000*, *verbose=False*, *value='itau'*)

Loads all *trans\_out* models that can be found in *trans\_tin*. This is useful when analyzing line ratios.

`crrlpy.models.rrlmod.load_itaui_all_norad` (*trans='alpha'*, *n\_max=1000*)

Loads all the available models.

`crrlpy.models.rrlmod.load_itaui_dict` (*dict*, *trans*, *n\_min=5*, *n\_max=1000*, *verbose=False*,  
*value='itau'*)

Loads the models defined by *dict*.

`crrlpy.models.rrlmod.load_itaui_nelim` (*temp*, *dens*, *trad*, *trans*, *n\_max=1000*, *verbose=False*,  
*value='itau'*)

Loads models given a temperature, radiation field and an upper limit for the electron density.

`crrlpy.models.rrlmod.load_models` (*models*, *trans*, *n\_max=1000*, *verbose=False*, *value='itau'*)

Loads the models in backwards compatible mode. It will sort the models by *Te*, *ne* and *Tr*.



```

crrlpy.models.rrlmod.make_betabn (line, temp, dens, n_min=5, n_max=1000, other='')
crrlpy.models.rrlmod.make_betabn2 (line, temp, dens, n_min=5, n_max=1000, other='')
crrlpy.models.rrlmod.plaw (x, x0, y0, alpha)
    Returns a power law.

```

$$y(x) = y_0 \left( \frac{x}{x_0} \right)^\alpha$$

#### Parameters

- **x** (*float or array like*) – x values for which to compute the power law.
- **x0** (*float*) – x value for which the power law has amplitude y0.
- **y0** (*float*) – Amplitude of the power law at x0.
- **alpha** (*float*) – Index of the power law.

**Returns** A power law of index *alpha* evaluated at *x*, with amplitude *y0* at *x0*.

**Return type** float or array

```

crrlpy.models.rrlmod.str2val (str)
    Converts a string representing a number to a float. The string must follow the IDL convention for floats.

```

**Parameters** **str** (*string*) – String to convert.

**Returns** The equivalent number.

**Return type** float

#### Example

```

>>> str2val('2d2')
200

```

```

crrlpy.models.rrlmod.val2str (val)
    Converts a float to the string format required for loading the CRRL models.

```

**Parameters** **val** (*float*) – Value to convert to a string.

**Returns** The value of *val* represented as a string in IDL double format.

**Return type** string

#### Example

```

>>> val2str(200)
'2d2'

```

```

crrlpy.models.rrlmod.valid_ne (trans)
    Checks all the available models and lists the available ne values.

crrlpy.models.rrlmod.xi (n, Te, Z)

```



## CRRLPY.CRRLS.FREC\_CALC MODULE

`crrlpy.frec_calc.line_freq(Z, R_X, n, dn)`

Uses the Rydberg formula to get the frequency of a transition to quantum number  $n$  for a given atom.

### Parameters

- **Z** (*int*) – Charge of the atom.
- **R\_X** (*float*) –
- **n** (*int*) – Principal quantum number of the transition.  $n + \Delta n \rightarrow n$ .
- **dn** (*int*) – Difference between the principal quantum number of the initial state and the final state.  $\Delta n = n_f - n_i$ .

**Returns** The frequency of the transition in MHz.

**Return type** `float`

`crrlpy.frec_calc.main()`

Main body of the program. Useful for calling as a script.

`crrlpy.frec_calc.make_line_list(line, n_min=1, n_max=1500, unitless=True)`

Creates a list of frequencies for the corresponding line. The frequencies are in MHz.

### Parameters

- **line** (*string*) – Line to compute the frequencies for.
- **n\_min** (*int*) – Minimum  $n$  number to include in the list.
- **n\_max** (*int*) – Maximum  $n$  number to include in the list.
- **unitless** (*bool*) – If True the list will have no units. If not the list will be of `astropy.units.Quantity` objects.

**Returns** 3 lists with the line name, principal quantum number and frequency of the transitions.

**Return type** `list`

`crrlpy.frec_calc.set_dn(name)`

Sets the value of Delta  $n$  depending on the transition name.

**Parameters** **name** (*string*) – Name of the transition.

**Returns**  $\Delta n$  for the given transition.

**Return type** `int`

**Example**

```
>>> set_dn('CIalpha')
1
>>> set_dn('CIDelta')
4
```

`crrlpy.frec_calc.set_specie(specie)`

Sets atomic constants based on the atomic specie.

**Parameters** `specie` (*string*) – Atomic specie.

**Returns** Array with the atomic mass in a.m.u., ionization potential, abundance relative to HI,  $V_X - V_H$  and the electric charge.

**Example**

```
>>> set_specie('CI')
[12.0, 11.4, 0.0003, 149.5, 1.0]
```

`crrlpy.frec_calc.set_trans(dn)`

Sets a name depending on the difference between atomic levels.

**Parameters** `dn` (*int*) – Separation between  $n_i$  and  $n_f$ ,  $\Delta n = n_i - n_f$ .

**Returns** alpha, beta, gamma, delta or epsilon depending on  $\Delta n$ .

**Return type** *string*

**Example**

```
>>> set_trans(5)
'epsilon'
```

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