## **HAZEL2: "inversion" mode**

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## **Basic Functions of HAZEL2**

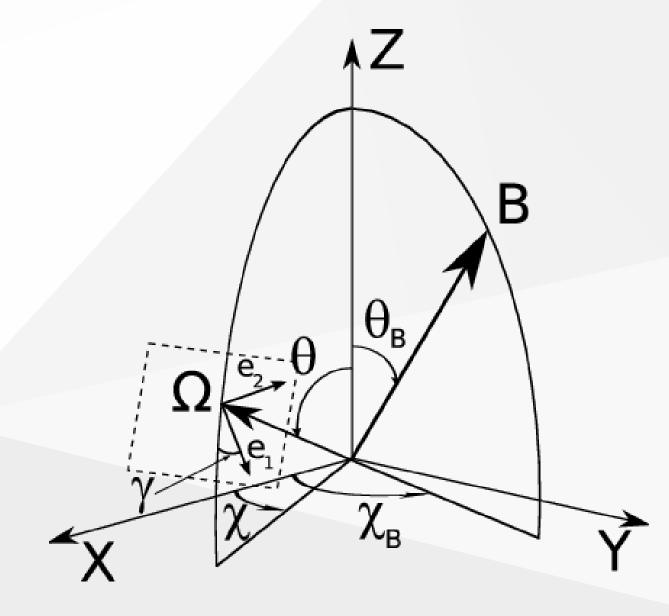
- Synthesis (Core function)
- **Inversion** (Call synthesis repeatedly, and return the values with best match.)

# Input Dangersion

- Stokes profiles (I, Q, U, V)
- Observer angle  $(\theta, \chi, \gamma)$

Hazel2 documents > 5.3.1.

The geometry for the scattering event



## Inversion Dutput

- ullet Magnetic field vector ( $B_x, B_y, B_z$ )
- Thermal velocity affecting the width of the line
- Bulk velocity of the plasma leading to a redshift / blueshift
- Optical depth of the line
- Line damping parameter

## Steps to invert

- Data file
- 2 Initial guess file
- 3 Configuration file
- 4 Invert
- 5 Read results

## Data file

```
noise = 2<mark>e-4</mark>
f = open('10830aStokes.1d', 'wb')
f.write(b'# LOS theta_LOS, phi_LOS, gamma_LOS\n')
f.write(b'0 0 90\n')
f.write(b'\n')
f.write(b'# Boundary condition I/Ic(mu=1), Q/Ic(mu=1), U/Ic(mu=1), V/Ic(mu=1)\n')
f.write(b'1 0 0 0\n')
f.write(b'\n')
f.write(b'# SI SQ SU SV sigmaI sigmaQ sigmaU sigmaV\n')
tmp = np.vstack([stokes, noise*np.ones(stokes.shape)])
np.savetxt(f, tmp.T)
f.close()
```

Noise can be different for all Stokes, all wavelength.

## Data file

```
-n 10830aStokes.1d
%cat
    1 # LOS theta_LOS, phi_LOS, gamma_LOS
       # Boundary condition I/Ic(mu=1), Q/Ic(mu=1), U/Ic(mu=1), V/Ic(mu=1)
       # SI SQ SU SV sigmal sigmaQ sigmaU sigmaV
```

# 2 Initial guess file

hazel2/examples/chromospheres/model\_chromosphere.1d

```
%cat -n model_chromosphere.1d

1 Bx By Bz tau v deltav beta a ff
2 0 0 10.0 1.0 0.0 8.0 1.0 0.5 1
```

- Initial values should be within ranges.
- A very good guess (e.g. do a second round using the results of first round as initial values) cost less computation time and may lead to slightly better fitting results.

# **3** Configuration file

hazel2/examples/configurations

#### 3.1 Working mode

```
%cat -n conf.ini
1  # Hazel configuration File
3  [Working mode]
4  Output file = output.h5
5  Number of cycles = 2
```

Number of cycles to carry out during inversion. Cycle 1 is to invert Stokes I only. Cycle 2 is to invert Stokes Q, U, and V.

#### 3.2 Spectral regions

```
14 [Spectral regions]
15         [[Region 1]]
16         Name = spec1
17         Wavelength = 10828, 10831, 50
18         Topology = ch1
19         Observations file = '10830aStokes.1d'
20         Weights Stokes I = 1, 0
21         Weights Stokes Q = 0, 1
22         Weights Stokes U = 0, 1
23         Weights Stokes V = 0, 1
```

Weights are float numbers. E.g. Use 0.5 for noisy Stokes V.

#### 3.3 Atmospheres

Relative path refers to the current folder. HAZEL goes to the folder that contains the configuration file to find files such as 'model\_chromosphere.1d'.

#### Ranges

Narrower ranges may be faster, with more accurate results.

If some values in your results are very close to range limits: Make sure initial values are in range. Expand ranges and redo inversion.

#### **Nodes**

In the first cycle, use Stokes I to invert tau, v, deltav, a. In the last cycle, use Stokes Q, U, and V to invert B. beta and ff are not changed. O or 1 for chromosphere.

# 4 Invert

```
modi = hazel.Model('conf.ini', working_mode='inversion')
modi.read_observation()
modi.open_output()
modi.invert()
modi.write_output()
modi.close_output()
```

```
# OSError: Unable to create file (unable to truncate a file which is already open)
```

To fix this error, in 5, explicitly close .h5 files when no longer in use.

## 5 Read results

Print model results.

```
res = h5py.File('output.h5', 'r') # explicitly close when no longer in use.
sto = ['Bx','By','Bz','tau','v','deltav','beta','a','ff']
stp = ''
for i in sto:
    sti = res['ch1'][i][0,0,0]
    stp += ', '+i+':'+f'{sti:.2f}'
chi2 = res['spec1']['chi2'][0,0,0]
print(stp[2:]+', chi2:'+f'{chi2:.2f}')
# Bx:-199.58, By:53.06, Bz:100.03, tau:1.02, v:-3.01,
# deltav:6.13, beta:1.00, a:0.47, ff:1.00, chi2:0.61
```

## 5 Read results

Plot fitting lines.

```
for i in range(4):
    plt.subplot(221+i)
    plt.plot(ms.wavelength_axis, res['spec1']['stokes'][0,0,i])
    plt.xlabel('Wavelength [$\AA$]')
    plt.ylabel(iq[i]+'/Ic')
res.close()
```

Explicitly close .h5 files when no longer in use.

This series of presentations focus on **basic functions** of HAZEL2. **Everything required** for HAZEL2 synthesis and inversion has been shown.

You will find things that are not covered in the presentations (E.g. weights file, extra lines in configuration file). They are **optional**, and more information can be found in <u>Hazel2 documents</u>.

Now, let's piece all the code snippets together.

<u>hazel2simple.ipynb</u>