

# Exorcising the Ghost in the Machine

Synthetic Spectral Data Cubes for Assessing Big Data Algorithms

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No free lunch!

- data consuming, or...
- highly dependent of prior knowledge
- verifiable (labelled) real data is scarce
- more advanced  $\sim$  more complex
- more flexible  $\sim$  more parameters
- data analysis science (study)
- parameter tuning is a nightmare

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We need synthetic data!

# Astronomical SYnthetic Data Observations (ASYDO)

- **Synthetic** ALMA-like data generator
- **Simple/mock** astrophysical models
- **Arbitrary** data cubes observations (FITS)
- Opportunity for Machine Learning
  - ▶ Labelled and reliable data
  - ▶ Unbounded number of samples
  - ▶ Data-driven sensitivity analysis
- Opportunity for information technologies in general
  - ▶ Assessment of bulk-data transfer, compression, etc.
  - ▶ Assessment of image analysis techniques
  - ▶ Assessment of storage systems

# Data Cube Characterization

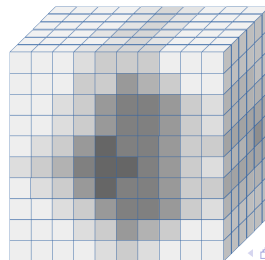
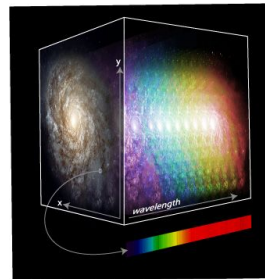
A spectroscopic data cube with calibrated temperatures, with two spatial axes and a frequency axis.

Basic model:

$$C(x, y, f) = \hat{C}(x, y, f) + \mathcal{N}(0, \sigma) \quad (1)$$

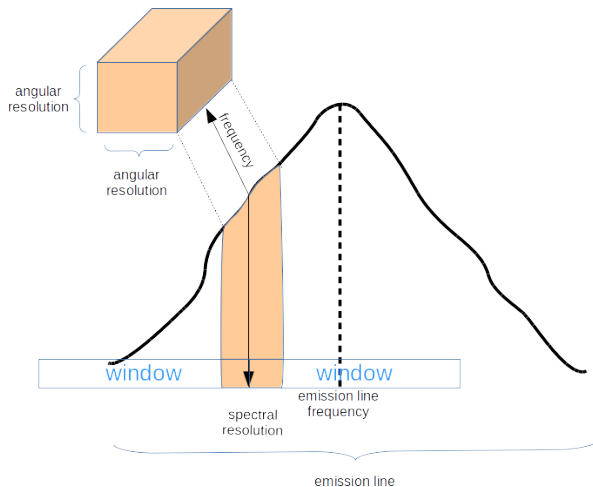
What to simulate in  $\hat{C}(x, y, f)$ ?

- Emission lines frequency, energy, etc (DB)
- Local radial velocity gradients
- Red-shift correction
- Broadening functions (frequency distribution model)
- Reasonable spatial distribution models





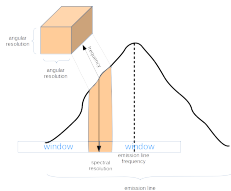
# Data Cell Characterization



The temperature of a cell is given by the following model:

$$C(x, y, f) = \sum_{l \in \mathcal{L}(x, y, f, l)} \int_{\nu_f - \Delta\nu/2}^{\nu_f + \Delta\nu/2} Br(\nu, l) d\nu + \epsilon \quad (2)$$

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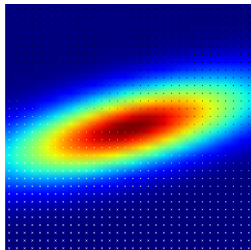
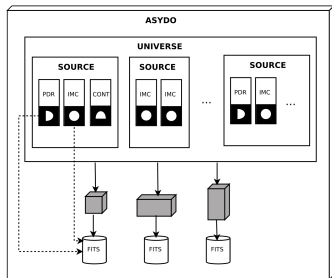
## Example

By assuming a Gaussian line profile,

$$C(x, y, f) = \sum_{l \in \mathcal{L}(x, y, f, l)} \int_{\nu_f - \Delta\nu/2}^{\nu_f + \Delta\nu/2} \frac{T_l(x, y)}{S_l(x, y) \sqrt{2\pi}} \exp\left(-\frac{(\nu - \nu_l(1 + Z_l(x, y)))^2}{2S_l(x, y)^2}\right) d\nu + \epsilon \quad (3)$$

For each line  $l$  we need to generate  $T_l$  (temperature),  $Z_l$  (redshift) and broadening parameters  $\Phi_l$  (i.e.  $S$  in example)

# ASYDO Elements



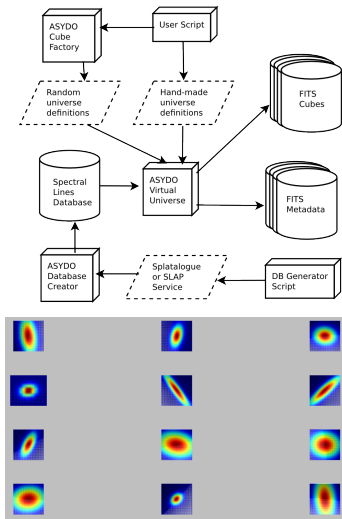
## Elements:

- Persistent object called Virtual Universe (VU)
- Sources that belongs to VU ( $\alpha_S, \delta_S, z_S$ ).
- Sources have several components (structures)
- Components use a models
  - ▶ Molecular Clouds, PDR, blackbody, continuum
- A model generates each  $T_l$ ,  $Z_l$  and  $\phi_l$
- Arbitrary observations
  - ▶ Angular position ( $\alpha, \delta$ )
  - ▶ Angular resolution  $\Delta\theta$  and the Field of View (FOV)
  - ▶ Central frequency  $\nu$ , spectral resolution ( $\delta\nu$ ) and spectral bandwidth (BW)

## Modules:

- `asydopy.vu` virtual universe (asydo core)
- `asydopy.db` line database manipulation (SLAP service)
- `asydopy.factory` generate randomized cubes

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# Tools for the Models

- Spatial structures

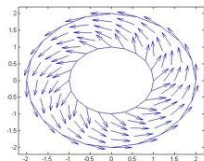
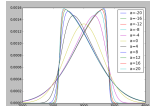
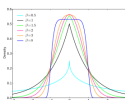
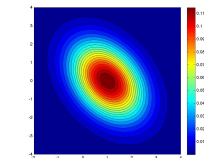
- ▶ Gaussian 2D
- ▶ Generalized Gaussian 2D (saturated, Lorentzian, etc)
- ▶ Exponential
- ▶ Soft-edge rings (TODO)
- ▶ Random Clouds (TODO)

- Spectral form

- ▶ Skew-Normal Distribution (1D)

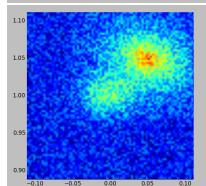
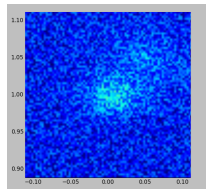
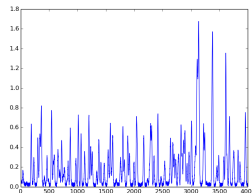
- Local shift functions

- ▶ Linear
- ▶ Exponential



# What we save in the FITS?

- A 3D image (cube)
- For each component (and subcomponent) we have
  - ▶ 2D images with the original matrices
    - ★ Temperature ( $T_m$ )
    - ★ Red-shift ( $Z_m$ )
    - ★ Broadening ( $\Phi_m$ )
  - ▶ A FITS table with each line of the component
  - ▶ This include:
    - ★ unique line code
    - ★ molecule name
    - ★ chemical name
    - ★ rest frequency
    - ★ observed frequency
    - ★ base red-shift
    - ★ temperature



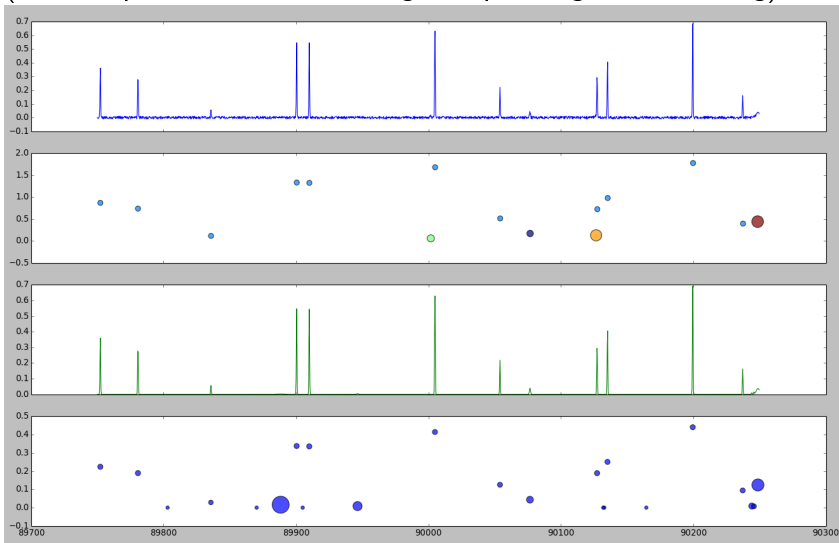
# Examples

## Supervised Learning Example:

- Pick a 2GHz frequency window ( $\sim 300$  GHz)
- Select randomly ( $p=0.3$ ) if a cube have a molecule
- Force Phosphatidylylidyne existence and absence (Binary Class)
- 30000 cubes,  $25 \times 25 \times 1000$  size each
- Naive approach: Use a SVM to train and test
- Result: 62% (something)
- The ML approach is insanely simple

# Examples

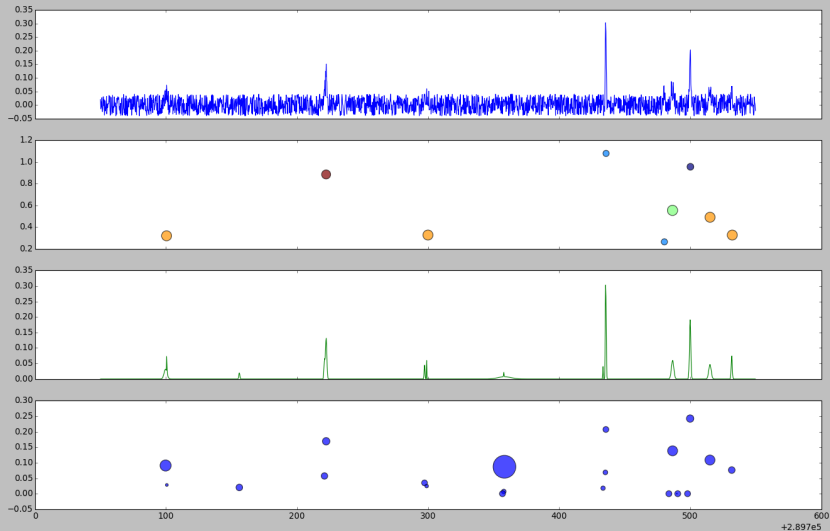
## Line Identification Example (wavelet-peak-detect, levenberg-marquardt gaussian fitting)





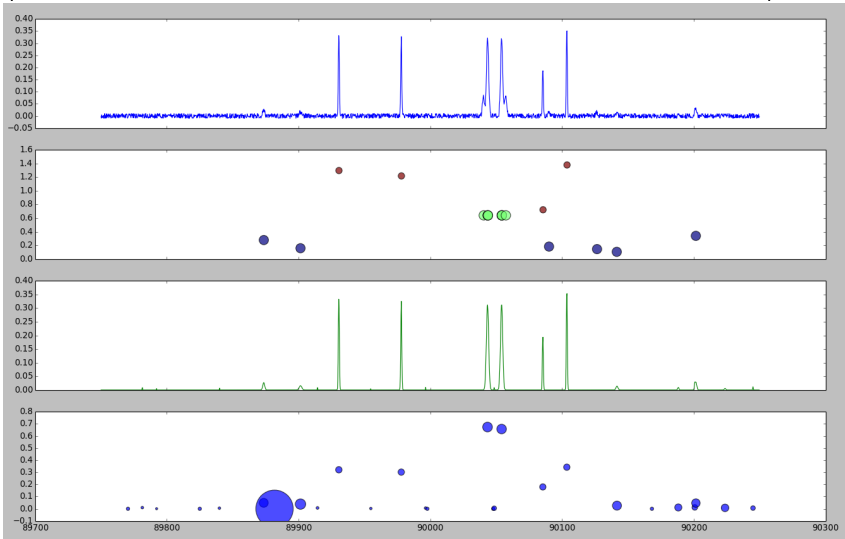
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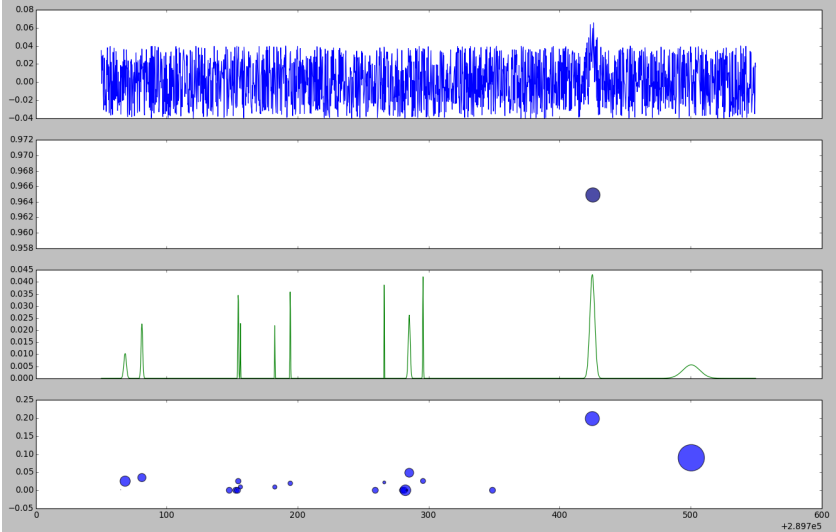
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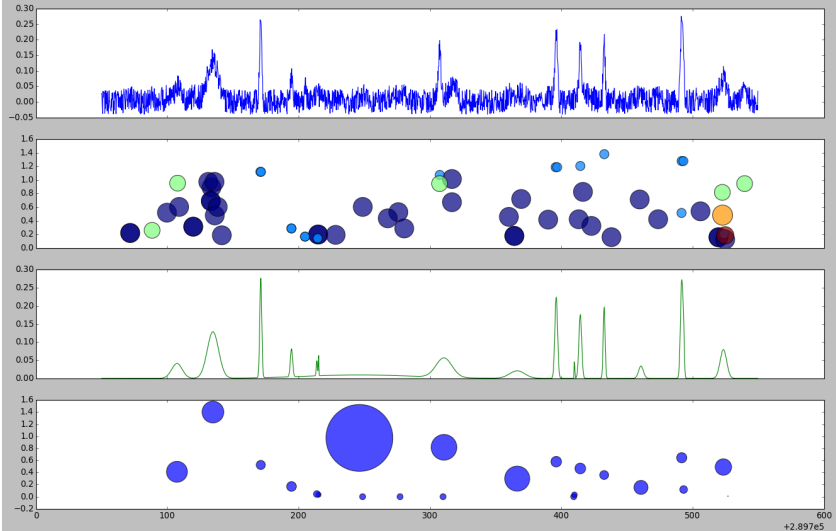
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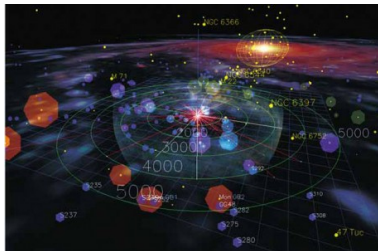
# Examples

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# Future Work

- Virtual Universe Service!



- IVOA-like synthetic data generation standard (web)
- Include more models and tools
- Integration with *astropy* and/or *CASA*
- Train using **synthetic data**, test using **real data**

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# Skew-normal Distribution

- The pdf of the skew normal (SN) distribution is:

$$f(x) = \frac{2}{\omega} \phi\left(\frac{x - \xi}{\omega}\right) \Phi\left(\alpha \left(\frac{x - \xi}{\omega}\right)\right) \quad (4)$$

where  $\phi(x) = \frac{1}{\sqrt{2\pi}} e^{-\frac{x^2}{2}}$  is the standard normal pdf and  $\Phi(x) = \frac{1}{2} \left[1 + \operatorname{erf}\left(\frac{x}{\sqrt{2}}\right)\right]$  the standard normal cdf.

- The parameters of  $SN(\xi, \omega, \alpha)$  are  $\xi$ =location,  $\omega$ =scale and  $\alpha$ =shape
- We propose first to reparametrize as follows

$$\mu = E[X] = \xi + \omega \delta \sqrt{\frac{2}{\pi}} \quad (5)$$

$$\sigma^2 = E[(X - \mu)^2] = \omega^2 \left(1 - \frac{2\delta^2}{\pi}\right) \quad (6)$$

$$\delta = \frac{\alpha}{\sqrt{1 + \alpha^2}} \quad (7)$$

which gives the following form

$$SN'(\mu, \sigma, a) = SN\left(\mu - \frac{\sigma \delta}{\sqrt{1 - \frac{2\delta^2}{\pi}}} \cdot \sqrt{\frac{2}{\pi}}, \frac{\sigma}{\sqrt{1 - \frac{2\delta^2}{\pi}}}, \alpha\right) \quad (8)$$

# Example: simple model

Defining  $T_l$ ,  $Z_l$  y  $\Phi_l$  matrices for each line is tedious. Group by molecules:

- molecule intensity maps  $T_m$ ,
- local molecule redshift maps  $Z_m$ ,
- and molecule broadening maps  $S_m$ .

The molecular model need to define (simple example):

$$T_l = f(T_m, l) = T_m \exp\left(-\frac{|e_l - t|}{t}\right)$$

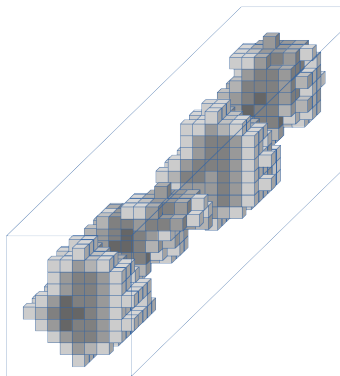
$$Z_l = g(Z_m, l) = Z_m$$

$$\phi_l = h(\phi_m, l) = \frac{S_m \nu_l}{2\sqrt{2 \ln 2} c}$$

$$T_m = 2D\text{Gauss}(\sigma_x, \sigma_y, \theta)$$

$$Z_m = \text{Linear}(\alpha, \beta, \theta)$$

$$S_m = s_m$$





# Spectral Line Database

$\mathcal{L}$  is the set of all lines, and  $\nu_l$  the central frequency of  $l \in \mathcal{L}$ . If the frequency range is constrained to  $\mathcal{R} = [\nu_{min}, \nu_{max}]$ , the set of potentials peaks in  $\mathcal{R}$  is:

$$\mathcal{L}_{\mathcal{R}} = \{l \in \mathcal{L} | \nu_l \in \mathcal{R}\}$$

A line  $l$  has other associated values in the DB such as the transition temperature  $e_l$  or the molecule species  $m_l$  (formula). For example, the set of species in an arbitrary region  $\mathcal{R}$  is defined as

$$\mathcal{M}_{\mathcal{R}} = \{m \in \mathcal{M} | \exists l \in \mathcal{L}_{\mathcal{R}}, m = m_l\},$$

where  $\mathcal{M}$  is the set of all the molecules in the DB.



Species	Chemical Name	Central Freq. (GHz)	Transition	Energy Level (MHz)	Line ID	Line ID	Line ID
1	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
2	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
3	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
4	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
5	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
6	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
7	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
8	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
9	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
10	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
11	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
12	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
13	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0
14	CH <sub>3</sub> OH	77.0000	10-100-1-1-0-0-0-0-0	0.0000	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0	10-100-1-1-0-0-0-0

## Assumptions

- **S1:** the DB contains all the observable lines
- **S2:** each transition has an associated molecule

