

Mathematical characterization of galaxies using IFS data: II - adding spatial information

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INAOE, 02/09/2014

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Outline

1 Binning

- Voronoi
- HII explorer
- BaTMAAn

2 Morphological characterization

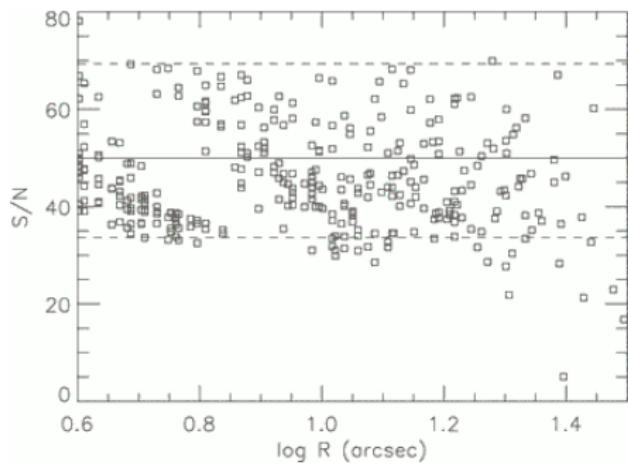
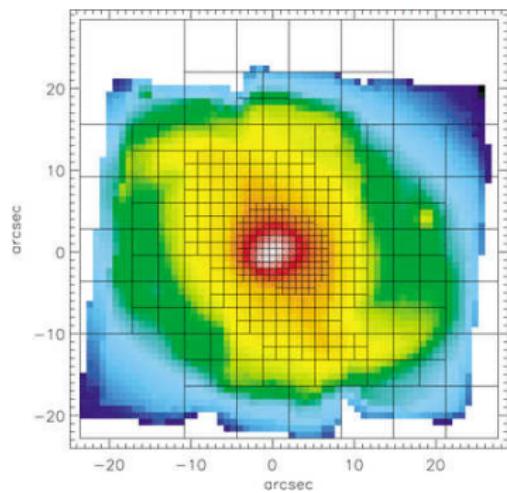
Voronoi binning

(Cappellari & Copin 2003)

Motivation

Constant signal-to-noise

$$(S/N)_b \simeq (S/N)_0$$



Motivation

Constant signal-to-noise

$$(S/N)_b \simeq (S/N)_0$$

Desired properties

- Topological tessellation
- Morphological compactness
- Uniformity

Algorithm

Weighted average

$$(S/N)_{A+B} = \frac{w_A S_A + w_B S_B}{\sqrt{w_A^2 N_A^2 + w_B^2 N_B^2}}$$

“Optimal” weighting

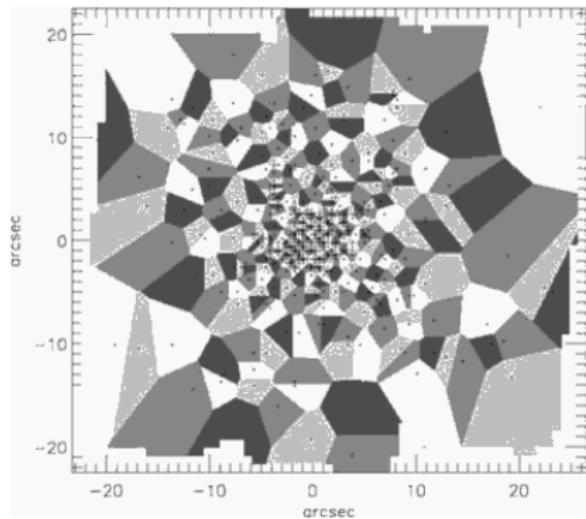
$$w_i \propto S_i / N_i^2$$

$$(S/N)_{A+B} = \sqrt{(S/N)_A^2 + (S/N)_B^2}$$

(S/N) “density”

$$\rho(\vec{x}) \equiv (S/N)^2(\vec{x})$$

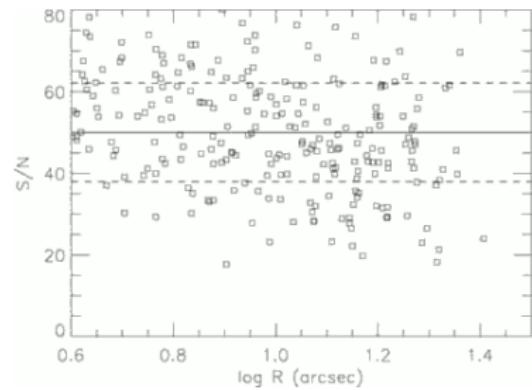
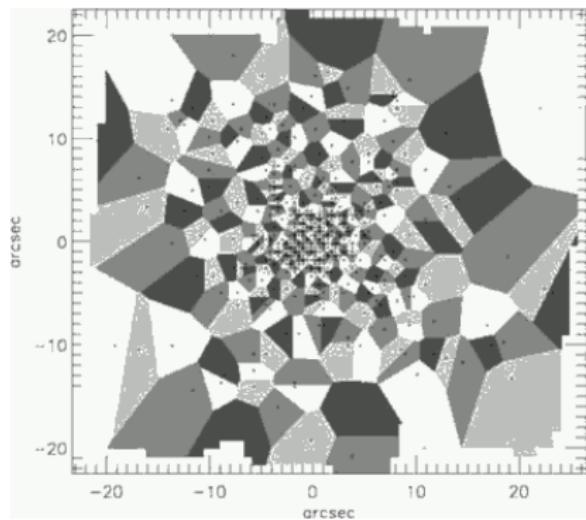
Algorithm



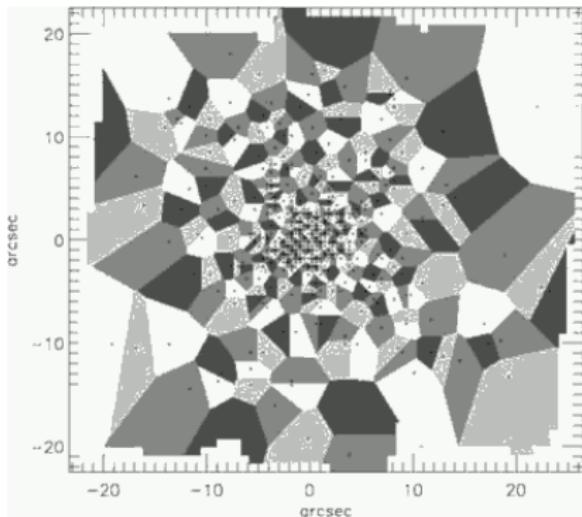
Modified k-means

- ① Select random seed
- ② Tessellate
- ③ $\vec{x}_b = \frac{\int_b \vec{x} \rho^2(\vec{x}) d\vec{x}}{\int_b \rho^2(\vec{x}) d\vec{x}}$
- ④ Iterate to convergence

Algorithm



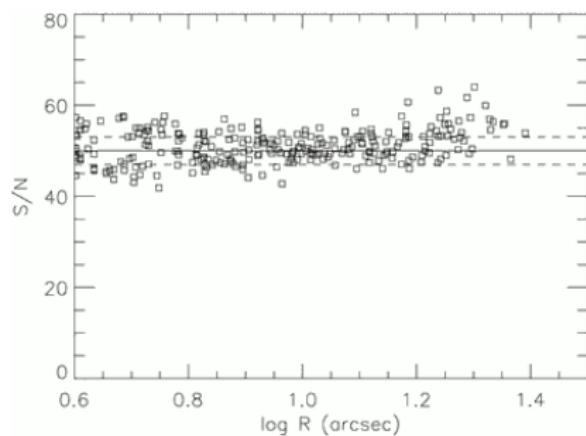
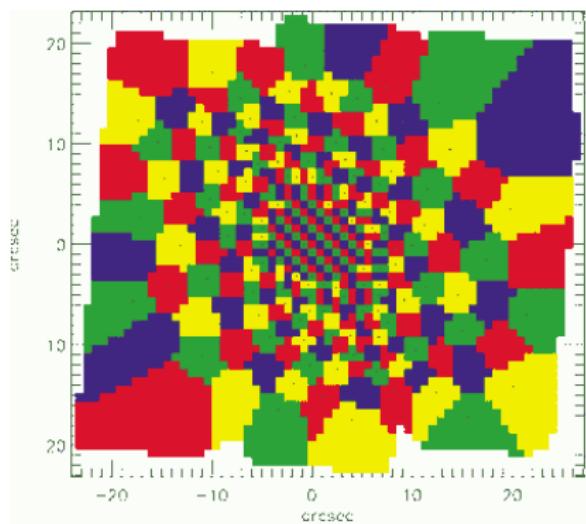
Algorithm



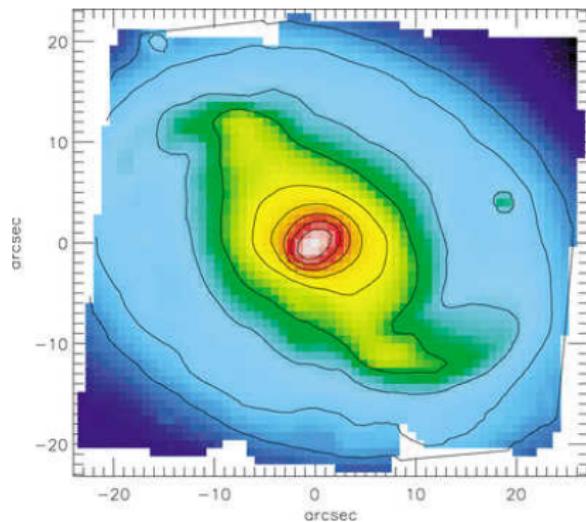
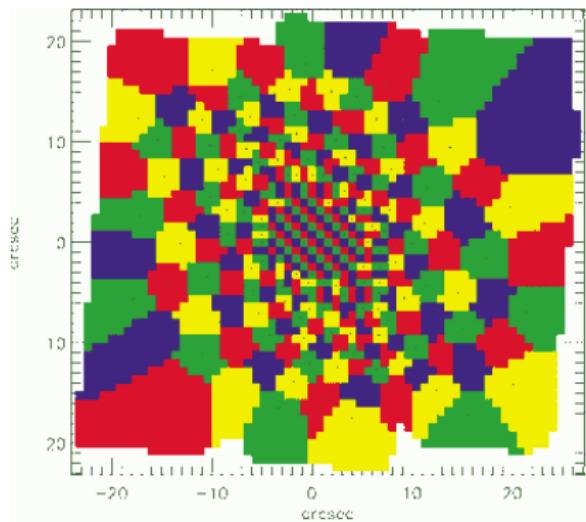
Initial seeds

- ➊ Start with highest S/N
- ➋ Accrete nearest pixel while
 - adjacent
 - “roundness”
 $R = \frac{r_{max}}{r_{eff}} - 1 < 0.3$
 - $(S/N)_b$ approaches $(S/N)_0$
- ➌ Mark as unsuccessful if $(S/N)_b < 0.8(S/N)_0$
- ➍ Start new bin with nearest pixel

Results



Results



HII explorer

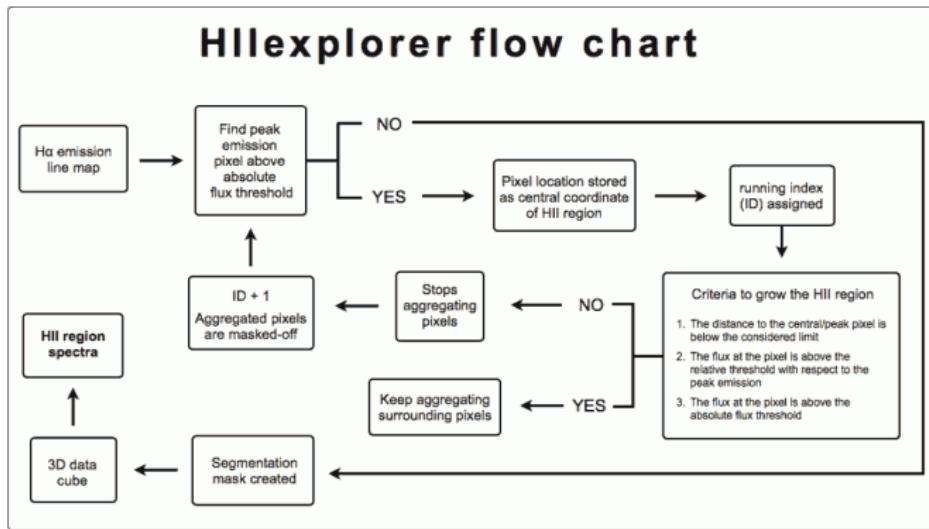
(Sánchez et al. 2012)

Algorithm

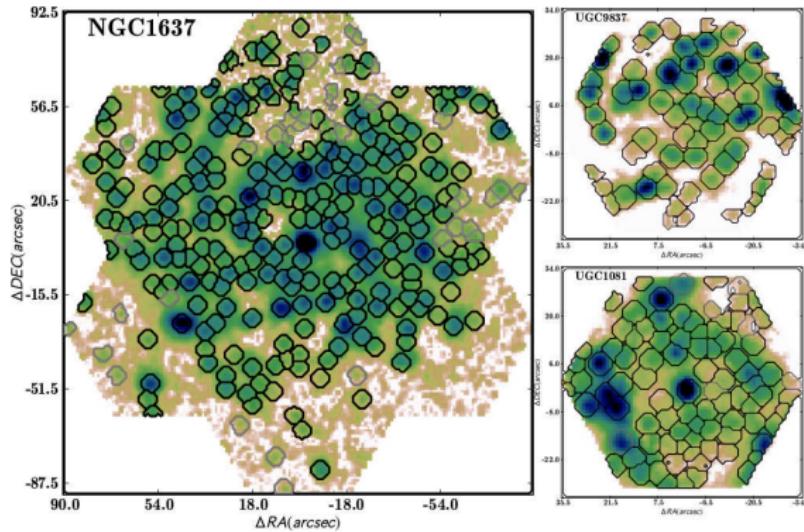
Input

- Emission-line map ($H\alpha$)
- Minimum peak intensity
- Intensity contrast
- Maximum radius
- Minimum flux

Algorithm



Results



Bayesian Technique for Multi-image Analysis

(Casado et al., in prep.)

A long-standing battle

Frequentist approach

- Likelihood
- Confidence intervals

Bayesian approach

- Likelihood
- Prior prob.
- Model evidence
- Posterior prob.

Likelihood

$\mathcal{L}_m(\vec{x}|\vec{\theta}_m) d\vec{x}$: probability of observing the datum \vec{x} , given model m with parameters $\vec{\theta}_m$

Prior probability

$\pi_m(\vec{\theta}_m) d\vec{\theta}_m$: expected probability distribution for $\vec{\theta}_m$

Model evidence

$E_m(\vec{x}) = \int \pi(\vec{\theta}_m) \mathcal{L}_m(\vec{x}|\vec{\theta}_m) d\vec{\theta}_m$: probability of \vec{x} , given m

Posterior probability

$p(\vec{\theta}_m|\vec{x}) = \frac{\pi(\vec{\theta}_m) \mathcal{L}_m(\vec{x}|\vec{\theta}_m)}{E_m(\vec{x})}$: probability of $\vec{\theta}_m$, given m and \vec{x}

Dear Sir or Madam?

Data

The subject gave birth to twins

Likelihood

Probability of a woman having twins $\sim 10^{-5}$

Probability of an amoeba having twins ~ 1

Bayesian approach

Frequentist approach

The hypothesis that the subject is a woman can be ruled out at the 99.999% level

- Prior: $\pi_m = \pi_w = 0.5$
- $E = \pi_m \mathcal{L}_m + \pi_w \mathcal{L}_w = 5 \times 10^{-6}$
- Posterior: $p_m = 0; p_w = 1$

Model comparison

Data

N measurements with errors: $\{x_i, \sigma_i\}_{i=1,N}$

Model 1

- One object with actual value x
- $\pi(x) dx = \frac{dx}{L}$ if $0 < x < L$
- $\mathcal{L}_1(\{x_i\}|x) = \prod_{i=1}^N \frac{e^{-\frac{(x_i-x)^2}{2\sigma_i^2}}}{\sqrt{2\pi\sigma_i^2}}$

Model 2

- Two objects, with x_A and x_B
- $\pi(x_A) = \pi(x_B) = \pi(x)$
- “labels” $\{l_i\}_{i=1,N} = A|B$
- $\mathcal{L}_2(\{x_i\}|x) = \prod_{i=1}^N \frac{e^{-\frac{(x_i-x_{l_i})^2}{2\sigma_i^2}}}{\sqrt{2\pi\sigma_i^2}}$

$$E_1(\{x_i\}) = \int \prod_{i=1}^N \frac{e^{-\frac{(x_i-x)^2}{2\sigma_i^2}}}{\sqrt{2\pi\sigma_i^2}} \frac{dx}{L}$$

$$E_2(\{x_i\}) = \int \int \prod_{i=1}^N \frac{e^{-\frac{(x_i-x_{l_i})^2}{2\sigma_i^2}}}{\sqrt{2\pi\sigma_i^2}} \frac{dx_A}{L} \frac{dx_B}{L}$$

BaTMAAn

Formulation of the problem

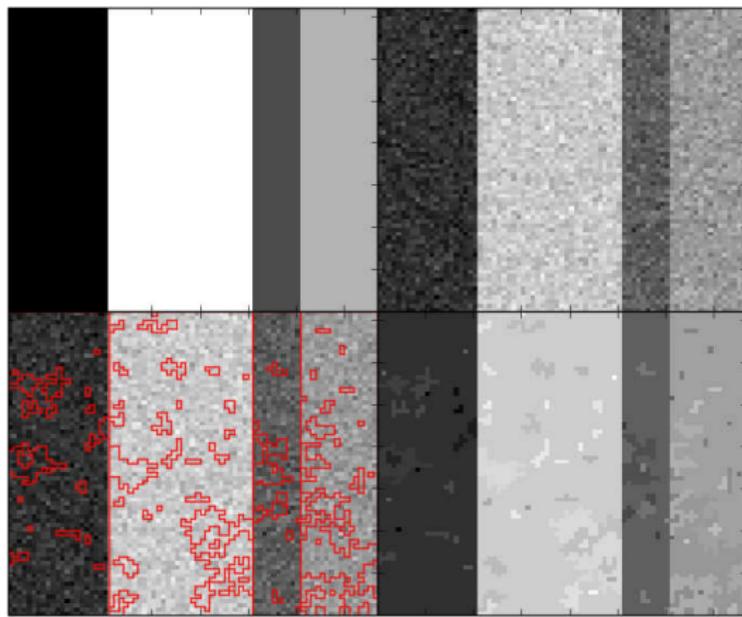
- Data: N measurements (pixels) with
 - intensity $\{x_i\}_{i=1,N}$
 - errors $\{\sigma_i\}_{i=1,N}$
- Model: B regions (bins) where $x = x_b$
 - $\pi(x_b) dx_b \sim \frac{dx_b}{\max(x_i) - \min(x_i)}$
 - “labels” $\{b_i\}_{i=1,N}$
 - $$\mathcal{L}(\{x_i\}_{i=1,N} | \{x_b\}_{b=1,B}) = \prod_{i=1}^N \frac{e^{-\frac{(x_i - x_{b_i})^2}{2\sigma_i^2}}}{\sqrt{2\pi\sigma_i^2}}$$

BaTMAAn

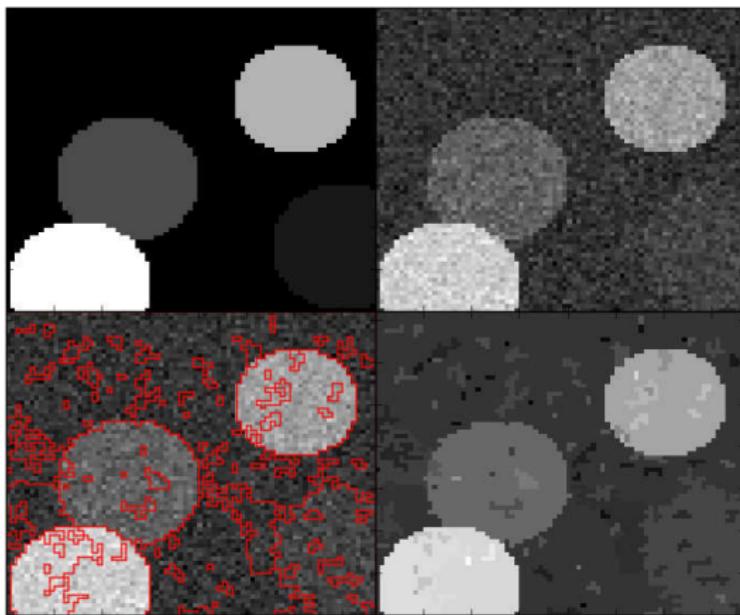
Algorithm

- ① Start with $B = N$
- ② While $E_{B-1} > E_B$:
 - merge the two adjacent regions that yield the highest E_{B-1}

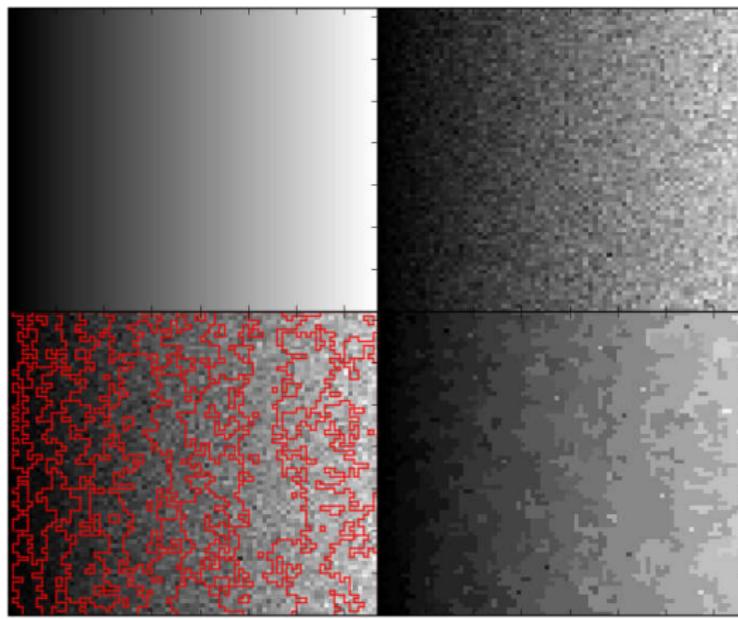
Results



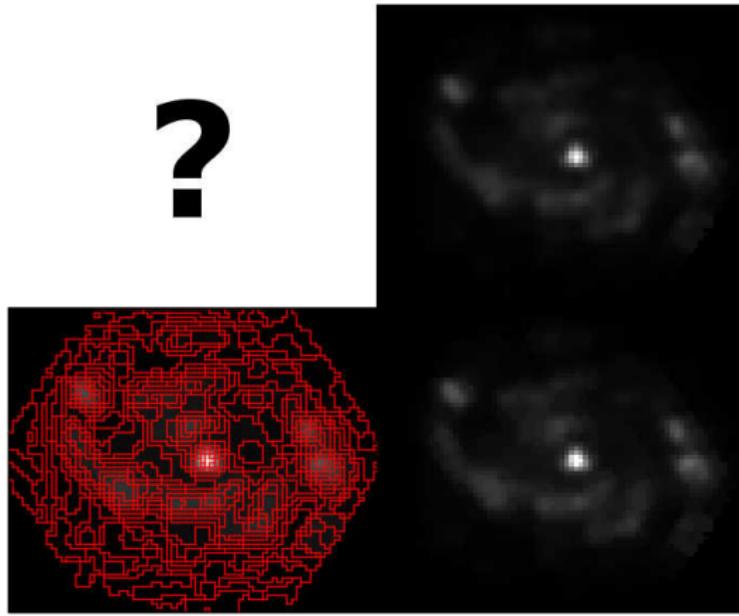
Results



Results

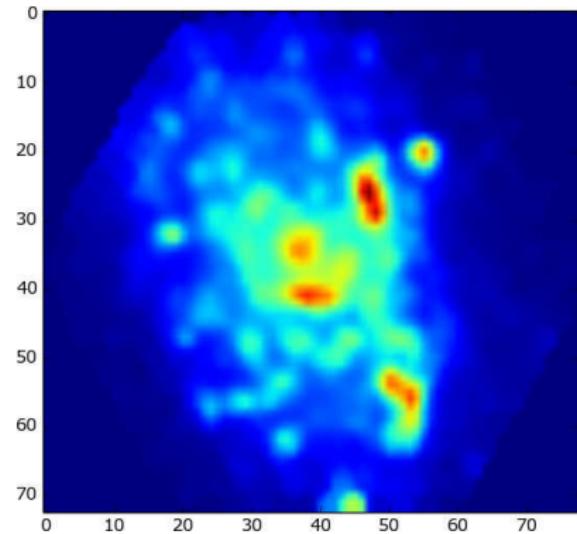
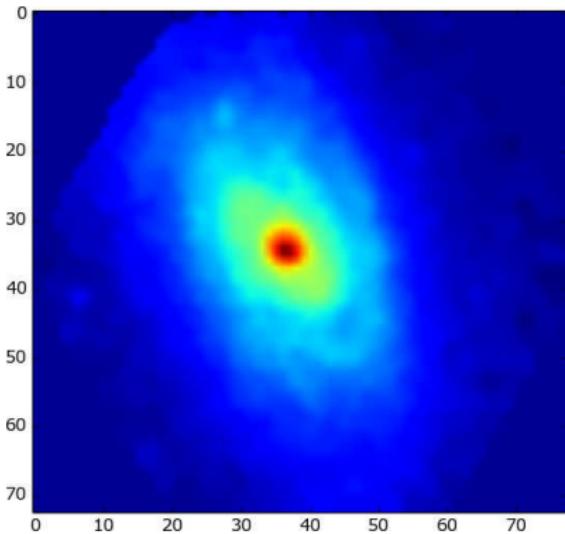


Results



Morphological characterization

Multi-wavelength morphology



Multi-wavelength morphology

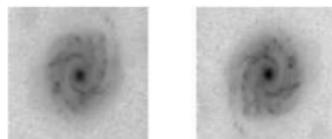
What for?

- ① Classification and comparison
- ② Statistics of the galaxy population
- ③ Input for chemical evolution models
- ④ Low signal-to-noise kinematics

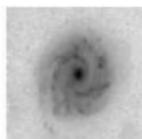
How?

- ① Statistical descriptors
- ② Parametric models
- ③ Non-parametric models

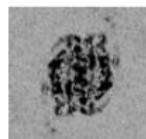
Statistical descriptors



I

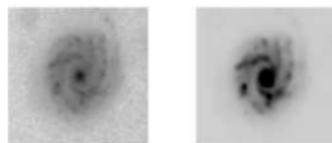


R



abs(I-R)

$$A = \frac{\text{abs}(I-R)}{I}$$



I

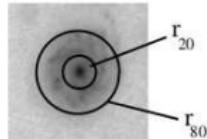


B



I-B

$$S = \frac{I-B}{I}$$



$$C = 5 \log\left(\frac{r_{80}}{r_{20}}\right)$$

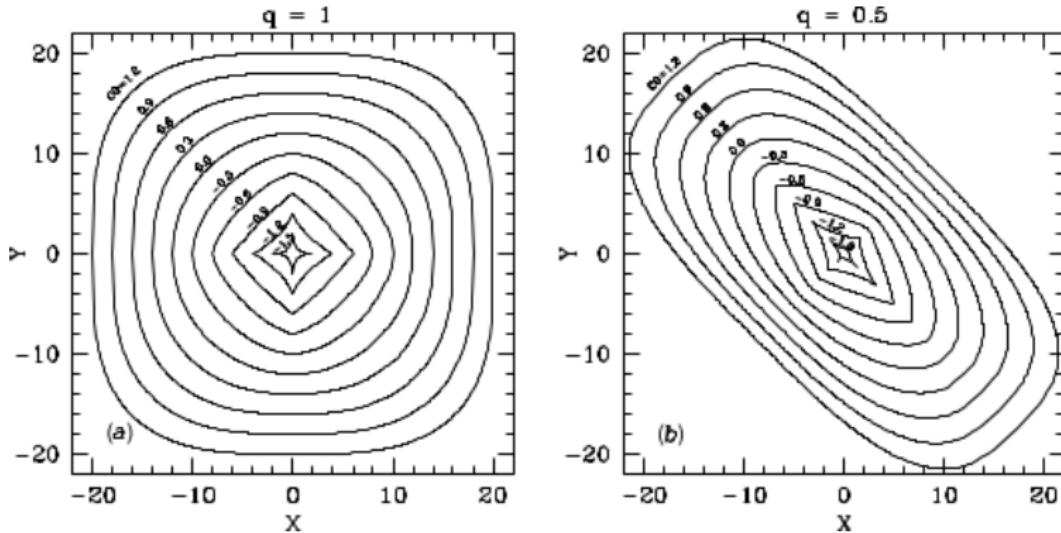
[Conselice \(2003\)](#)

Model-based decomposition



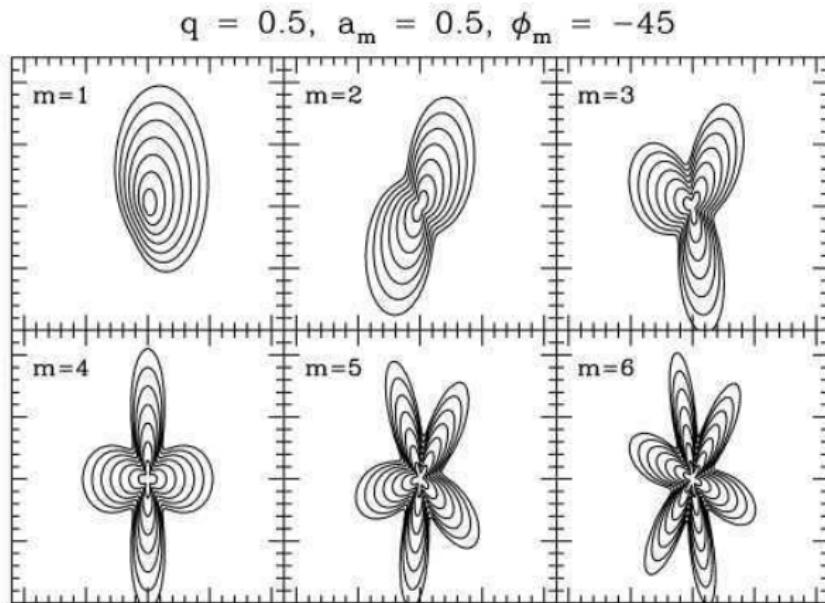
[de Souza et al. \(2004\)](#)

Generalized ellipsoids



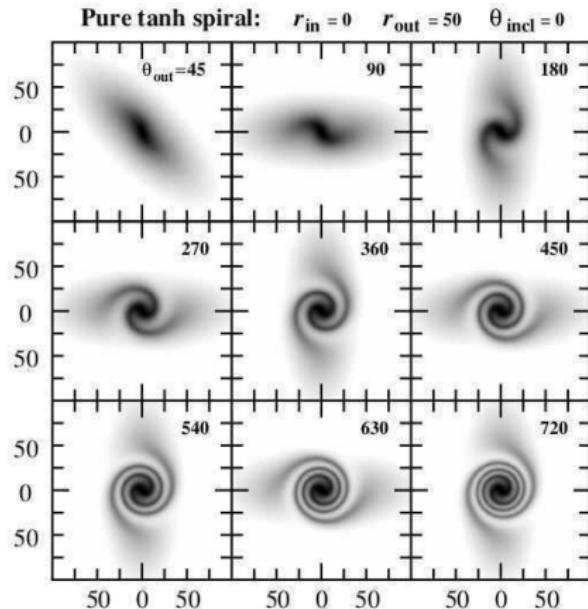
[Peng et al. \(2010\)](#)

Fourier modes



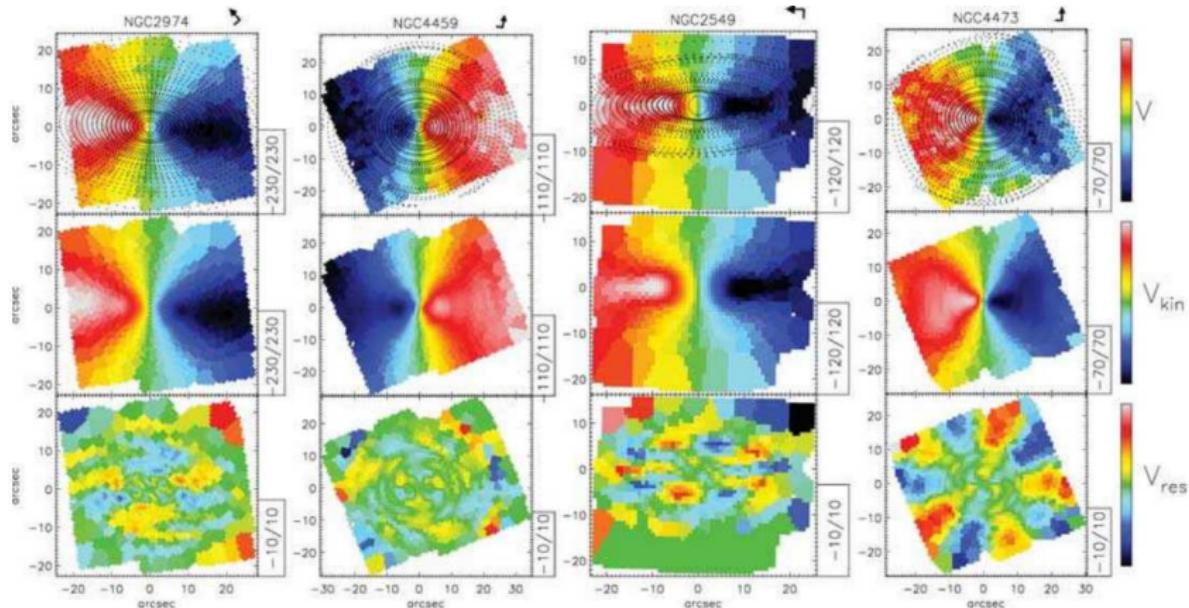
Peng et al. (2010)

Spiral modes



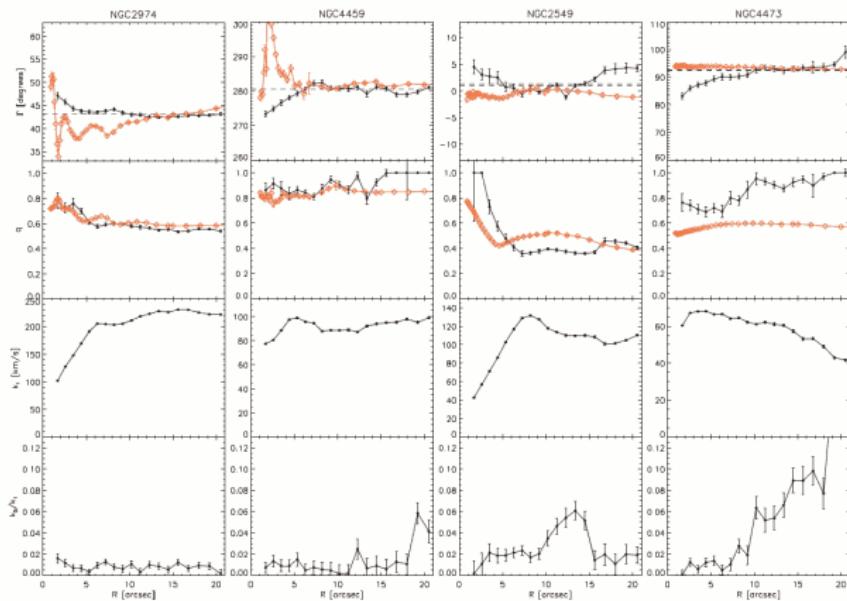
[Peng et al. \(2010\)](#)

Kinometry



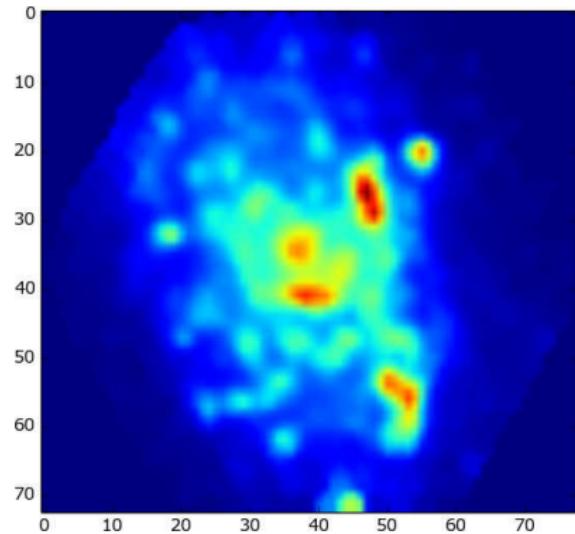
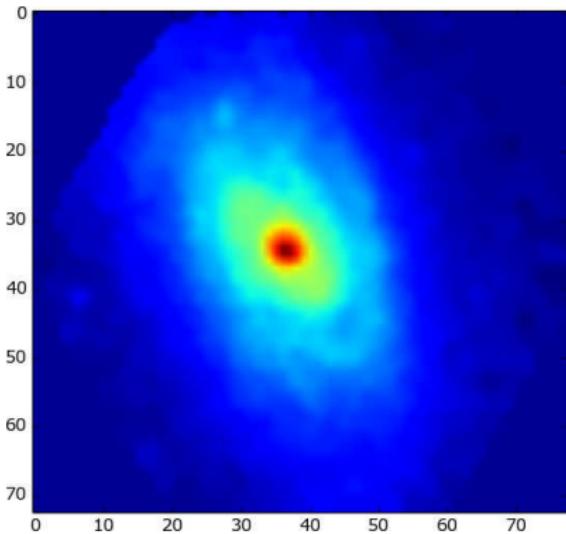
[Krajnović et al. \(2006\)](#)

Kinometry



[Krajnović et al. \(2006\)](#)

What next?



Summary

Binning

Bayes' theorem

Morphological characterization

There is no need to re-invent the wheel...
...but there is plenty of information to recover!