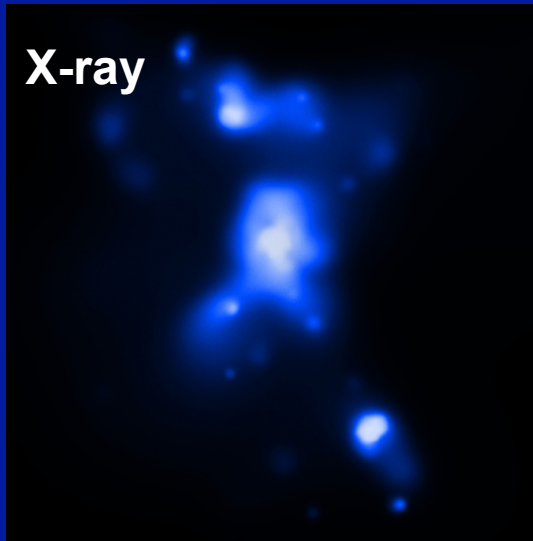


Chandra analysis of 4C+29.30

X-ray



RA=08 40 02.35; DEC=29 49 02.6

$z=0.065$

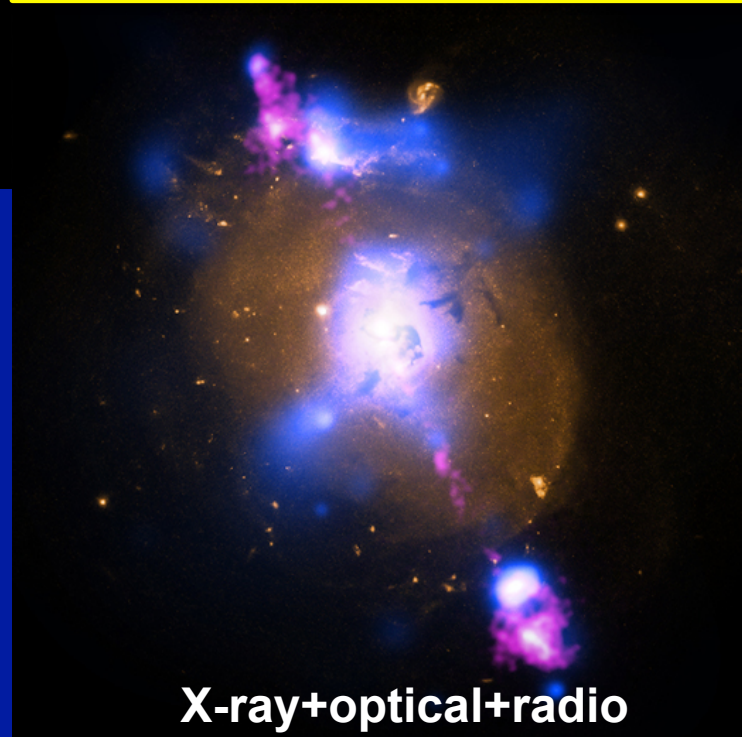
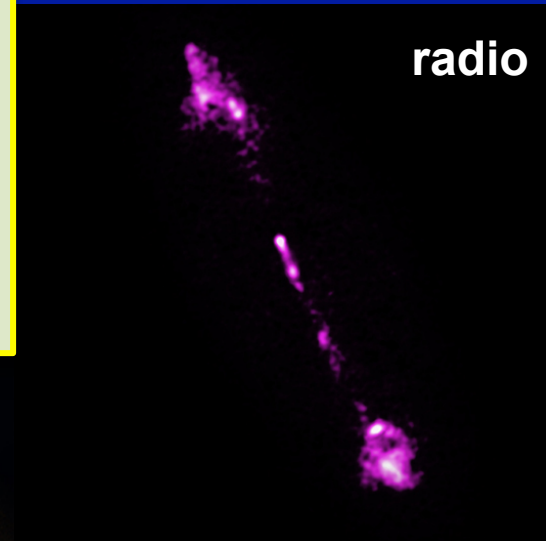
$N_{\text{H,Gal}}=4.23 \times 10^{20} \text{ cm}^{-2}$

4 long exposures *Chandra*, 284.5ks

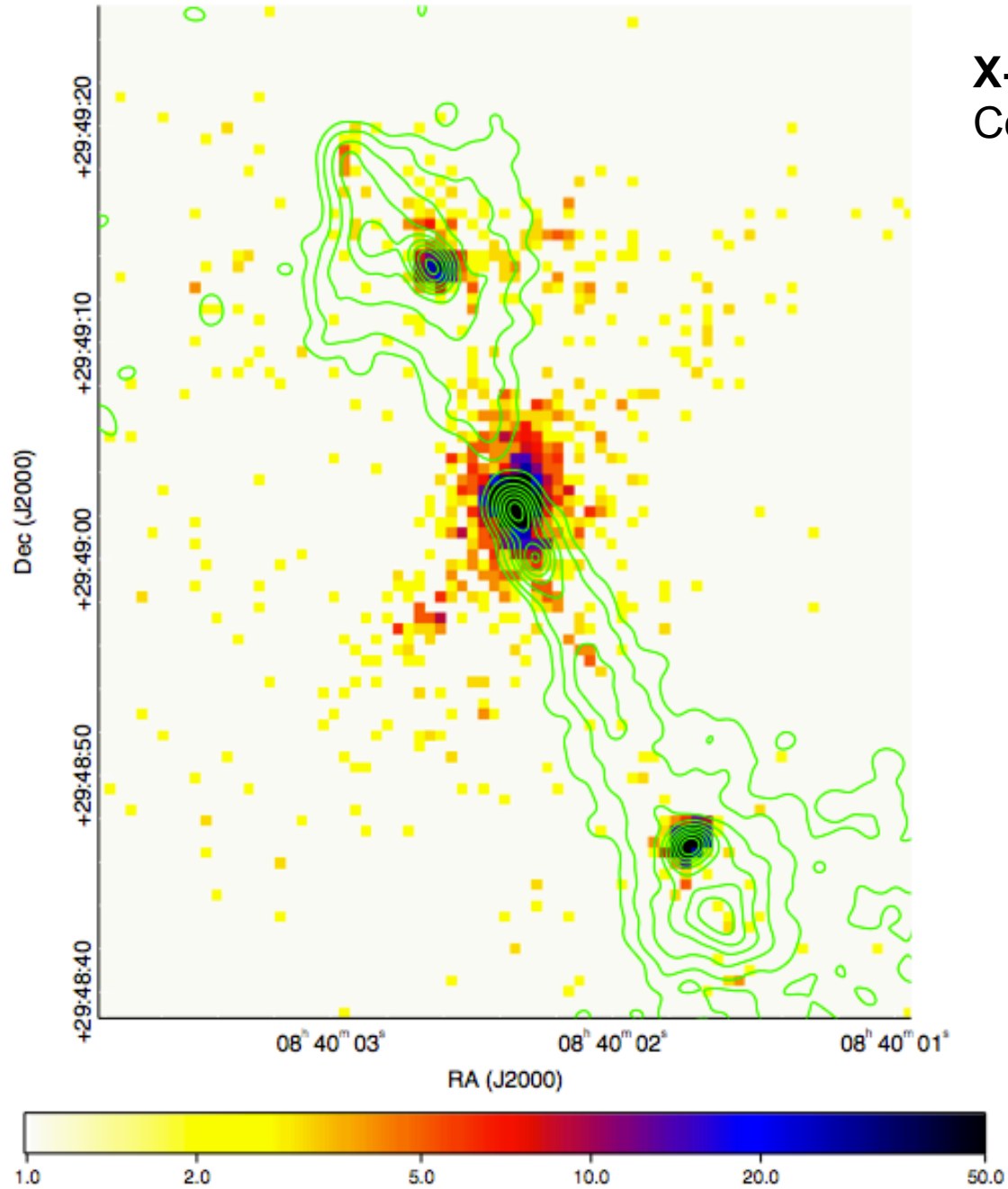
Transitional radio morphology FRI-

FRII ($L_{\text{R}} \approx 10^{42} \text{ erg/s}$)

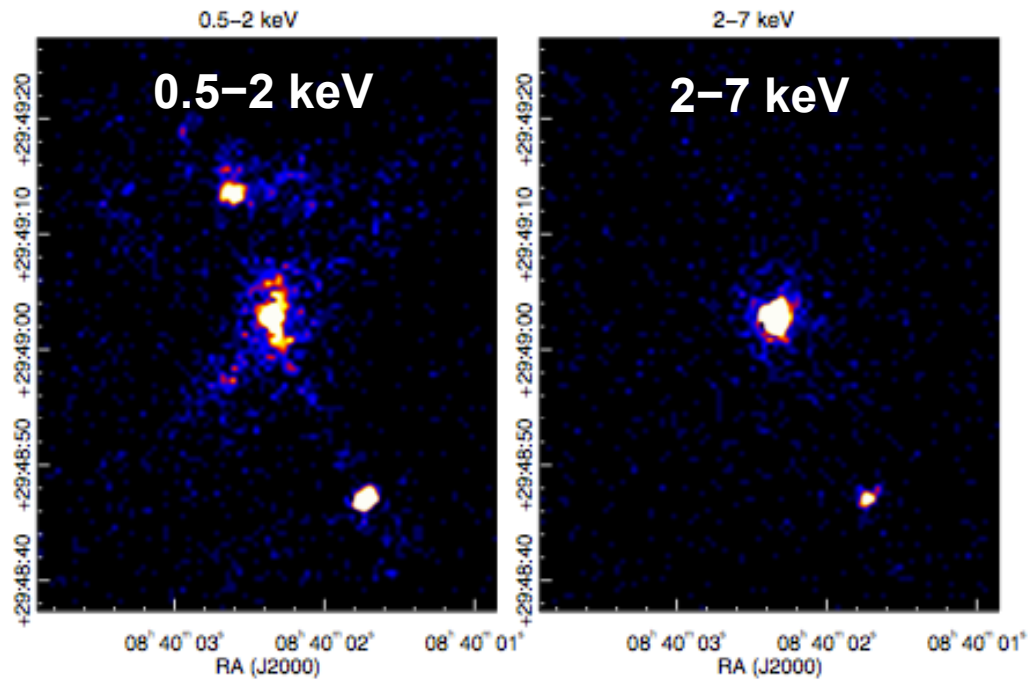
radio



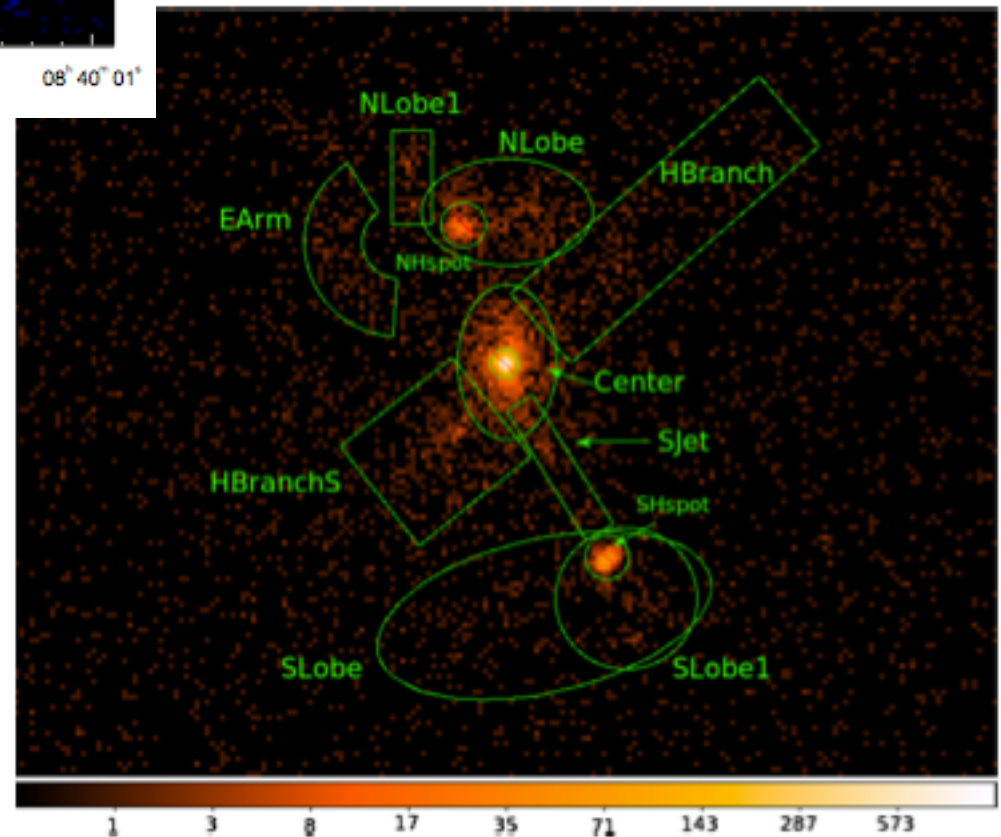
X-ray+optical+radio



X-ray: 0.5-7 keV combined image
Contours from **VLA** 1.45 GHz map



Significant structures in the soft band
 Most of the hard emission from the
 nucleus



PLAN

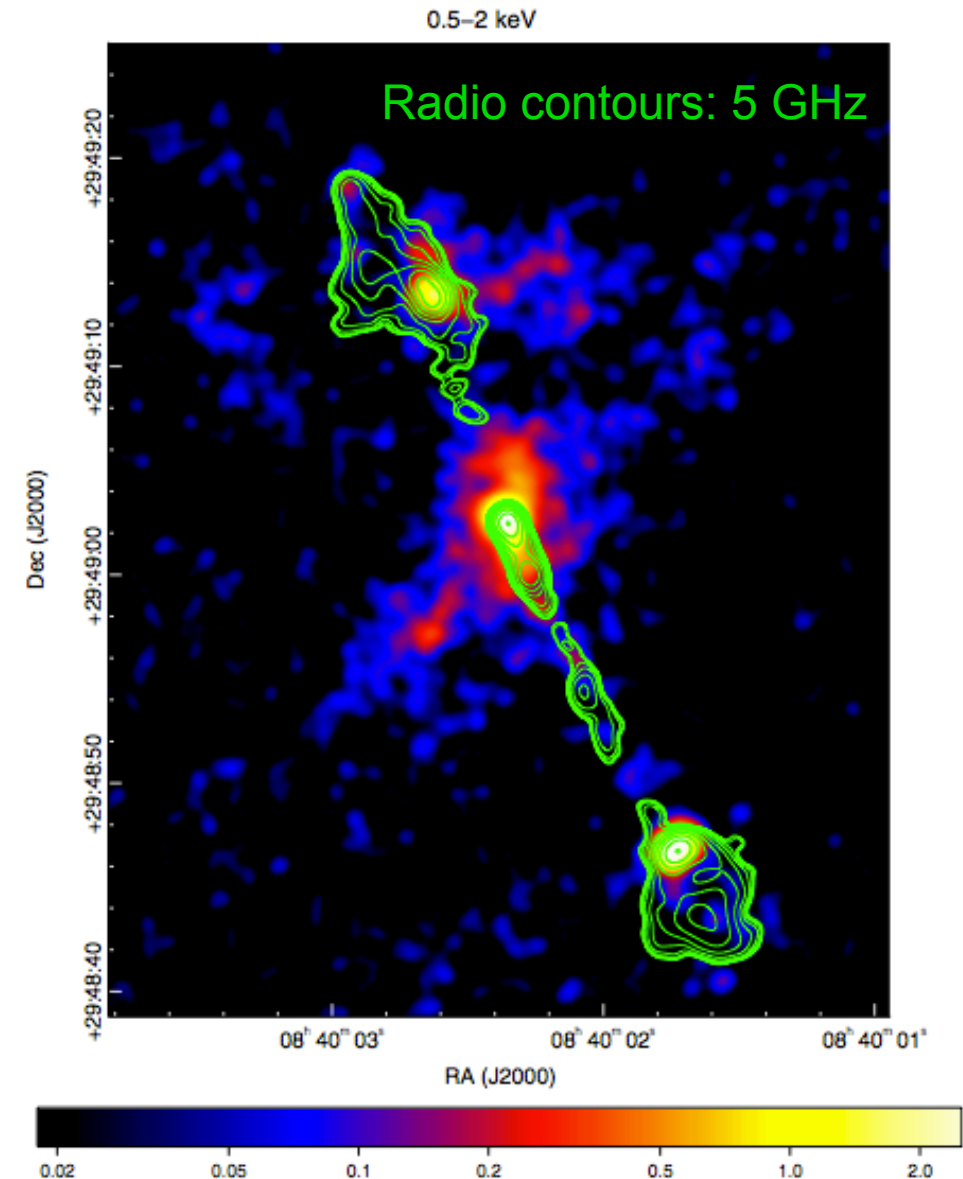
MAIN

1. Compare the X-ray emitting regions to the radio components from 1.5 and 5GHz maps
2. Extraction and analysis of *Chandra* spectra from the mosaic image (core, jets, lobes, extended features, ...)

Proper (but longer) way to proceed: spectral extraction from the four individual pointings, then merge the X-ray spectra

OPTIONAL

1. Extraction of spectra using XMM data (nucleus, extended emission, etc.), spectral analysis and comparison to *Chandra* results
2. Simultaneous *Chandra* (or XMM) and *Swift*/BAT (70-month data) spectral analysis of the nucleus



How to proceed (I)

Proper way: all spectral products should be extracted from the individual pointings and then combined (**MATHPHA**, **ADDARF**, **ADDRMF** – FTOOLS – and **COMBINE_SPECTRA** within CIAO). The spectral extraction is typically done using the CIAO tool **SPEXTRACT**.

A way to overcome this lengthy procedure consists of

- i. extracting the spectra (source and background) from the mosaic using **DMEXTRACT**;
- ii. producing ARF and RMF from one of the 4 pointings (choosing the longest one, 11688, T=123 ks) using the **SPEXTRACT** tool (see the *Chandra* tutorial for details); this would create the source and background spectra (which you will not use – you use those from the mosaic) and the ARF+RMF (which you will use);
- iii. associating the source spectrum (from the mosaic) to the background file (from the mosaic) and the ARF and RMF matrices (from the individual pointing) using the ftool **GRPPHA**, and group the data to a minimum number of counts to apply χ^2 statistics;
- iv. ARFs should be produced for every “source of interest” in the field (e.g., the core, the lobes, the hot spots, etc.), again from a single pointing.

How to proceed (II)

SOURCE

```
punlearn dmextract
pset dmextract infile="../4c29.30_2015_merged_evt.fits.gz[sky=region(nucleus.reg)]
[bin pi]"
pset dmextract outfile=4c29_30_r1p5_nucleus.pi
pset dmextract verbose=2
dmextract clobber+
```

define the region of interest



BACKGROUND

```
punlearn dmextract
pset dmextract infile="../4c29.30_2015_merged_evt.fits.gz[sky=region(back.reg)][bin
pi]"
pset dmextract outfile=4c29_30_r1p5_nucleus_bgd.pi
pset dmextract verbose=2
dmextract clobber+
```

GRPPHA

```
grppha 4c29_30_r1p5_nucleus.pi 4c29_30_r1p5_nucleus_c20.pi comm="group min
20 & chkey BACKFILE 4c29_30_r1p5_nucleus_bgd.pi & chkey RESPFILE
4c29_30_r1p5_nucleus.rmf & chkey ANCRFILE 4c29_30_r1p5_nucleus.corr.arf &
exit"
```

How to proceed (III)

CHECKS on the BACKSCAL (ratio of back vs. source areas)

```
dmkeypar 4c29_30_r1p5_nucleus.pi BACKSCAL echo+  
--> 4.3548145383623e-07
```

```
dmkeypar 4c29_30_r1p5_nucleus_bgd.pi BACKSCAL echo+  
--> 1.0533010170724e-05
```

$BACKSCAL = 15^2 / 3.05^2 = 24.2$
vs. $1.05e-5 / 4.35e-7 = 24.2$

(where 15 and 3.05 are the background and source extraction radius, in pixels, respectively)

OK!

Main publications

- ① Siemiginowska et al. 2012, ApJ, 750, 124
Chandra data
- ② Sobolewska et al. 2012, ApJ, 758, 90
XMM-Newton + Swift/BAT data