Learning SciPy for Numerical and Scientific Computing

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A PERSONAL PERSPECTIVE

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Scientific Computing is concerned with constructing mathematical models and quantitative analysis techniques, and using computers to analyze and solve scientific problems.

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- ► Find a reliable way to communicate through software.

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- Trust the Computer Scientist: Writing low-level code from scratch seldom guarantees best results.
- ► Find a reliable way to communicate through software.
- Big guns: Solving these problems usually require massive amounts of calculations and are often executed on supercomputers or distributed computing platforms.

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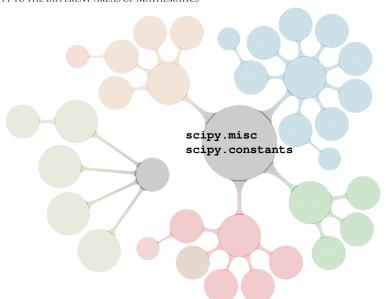
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 - ▶ with PyLab: ipython + NumPy + SciPy + matplotlib

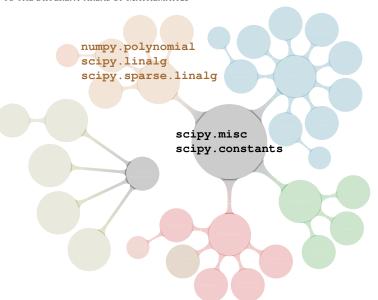
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 - ▶ with PyLab: ipython + NumPy + SciPy + matplotlib
 - ▶ with scikits and Pandas on top of that

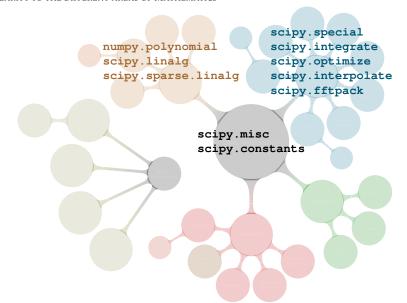
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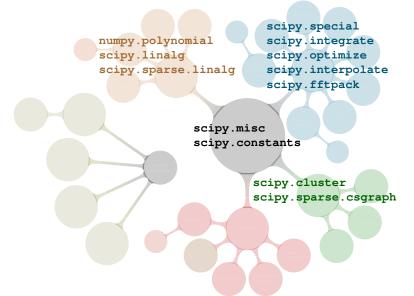
THE STRUCTURE OF SCIPY

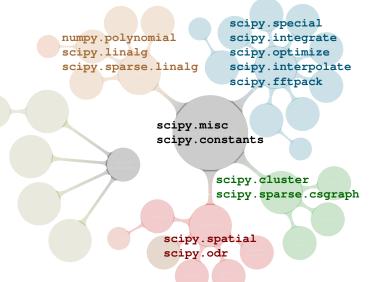












SIMILARITY TO THE DIFFERENT AREAS OF MATHEMATICS

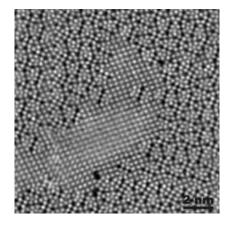
numpy.polynomial scipy.linalg scipy.sparse.linalg scipy.special scipy.integrate scipy.optimize scipy.interpolate scipy.fftpack

scipy.signal scipy.ndimage scipy.stats scipy.stats.mstats scipy.misc scipy.constants

> scipy.cluster scipy.sparse.csgraph

scipy.spatial scipy.odr

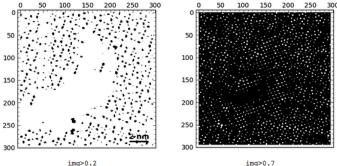
Extract the structural model of a molecule of $\mathit{Nb}_{4}\mathit{W}_{13}\mathit{O}_{47}$



COMPUTATION OF STRUCTURAL MODELS

We take the following (naïve) approach:

► Segmentation of the atoms by thresholding and morphological operations.



COMPUTATION OF STRUCTURAL MODELS

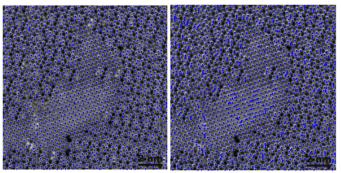
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- Connected component labeling to extract each atom for posterior examination.

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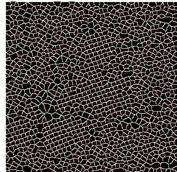
- Segmentation of the atoms by thresholding and morphological operations.
- Connected component labeling to extract each atom for posterior examination.
- ► Computation of the centers of mass of each label identified as an atom.



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We take the following (naïve) approach:

- Segmentation of the atoms by thresholding and morphological operations.
- Connected component labeling to extract each atom for posterior examination.
- Computation of the centers of mass of each label identified as an atom.
- Computation of the Voronoi diagram of the lattice formed by the previous points.



```
# Preamble
   import numpy
   import scipy
   from scipy.ndimage import binary_opening, label, center_of_mass,
        distance transform edt
5
6
8
9
10
11
   # Segmentation of each atom
12
13
14
   # Computation of centers of mass of each atom
15
   coords = center of mass(img, segmentation, range(1, segments+1))
16
   17
   vcoords = arrav([x[1] for x in coords])
18
19
20
```

```
import numpy
    import scipy
    from scipy.ndimage import binary_opening, label, center_of_mass,
5
    # Load the image
    img = scipv.misc.imread('NbW-STEM.png')
9
10
11
    # Segmentation of each atom
12
13
14
    # Computation of centers of mass of each atom
15
    coords = center of mass(img, segmentation, range(1, segments+1))
16
    x = array([x[0] for x in coords])
17
    vcoords = arrav([x[1] for x in coords])
18
19
20
```

```
import numpy
    import scipy
    from scipy.ndimage import binary_opening, label, center_of_mass,
5
6
    # Apply a threshold to segment atoms
    BWatoms = (imq > 0.62)
9
10
11
    # Segmentation of each atom
12
13
14
    # Computation of centers of mass of each atom
15
    coords = center of mass(img, segmentation, range(1, segments+1))
16
    x = array([x[0] for x in coords])
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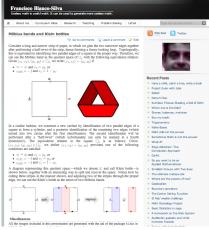
```
import numpy
    import scipy
    from scipy.ndimage import binary_opening, label, center_of_mass,
5
6
8
    # Perform a binary operation to eliminate outliers
9
10
    BWatoms = binary opening(BWatoms, structure=numpy.ones((2,2)))
11
    # Segmentation of each atom
12
13
14
    # Computation of centers of mass of each atom
15
    coords = center of mass(img, segmentation, range(1, segments+1))
16
    x = array([x[0] for x in coords])
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    vcoords = arrav([x[1] for x in coords])
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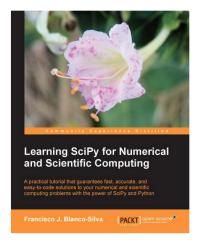
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    import scipy
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    # Segmentation of each atom
12
    structuring_element = [[0,1,0],[1,1,1],[0,1,0]]
13
    segmentation, segments = label (BWatoms, structuring element)
14
    # Computation of centers of mass of each atom
15
    coords = center of mass(img, segmentation, range(1, segments+1))
16
    x = array([x[0] for x in coords])
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    vcoords = arrav([x[1] for x in coords])
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    coords = center of mass(img, segmentation, range(1, segments+1))
16
    x = array([x[0] for x in coords])
17
    ycoords = array([x[1] for x in coords])
18
    # Compute the Voronoi diagram of the lattice
19
20
```

```
import numpy
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   # Segmentation of each atom
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   # Computation of centers of mass of each atom
15
   coords = center of mass(img, segmentation, range(1, segments+1))
16
   17
   vcoords = arrav([x[1] for x in coords])
    # Compute the Voronoi diagram of the lattice
18
   L1, L2 = distance_transform_edt(segmentation==0, return distances=False,
19
        return indices=True)
20
   Voronoi = segmentation[L1,L2]
```

FOR MORE INFORMATION, EXAMPLES, IDEAS, ...





blancosilva.wordpress.com