

In [4]:

```
from IPython.display import HTML,Image,SVG,YouTubeVideo
```

Eyes in the nature

In [1]:

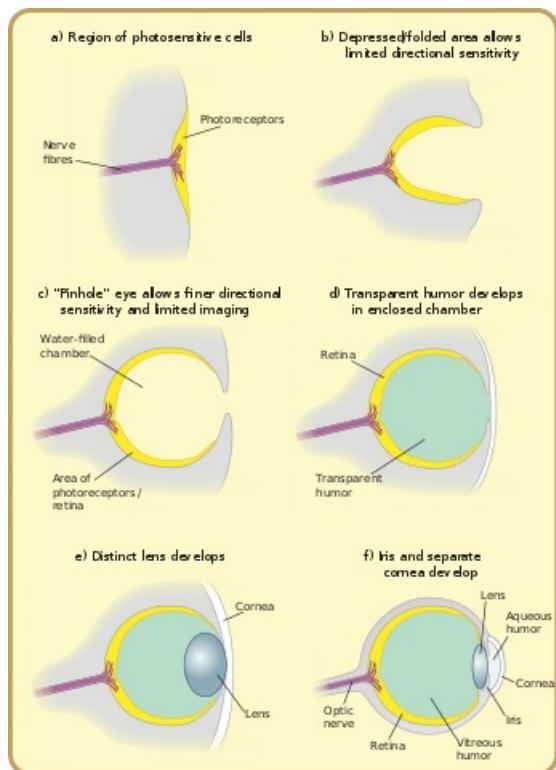
```
from IPython.display import HTML,Image,SVG,YouTubeVideo
```

Evolution has produced a large variety of light sensors among living organisms. From the simplest, where sensitive cells are directly on the skin surface, to more complex ones where the light sensing cells are embedded into a *camera obscura* like organ.

In [2]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/b/b6/Diagram_of_eye_evolution.svg/344px-Diagram_of_eye_evolution.svg.png')
```

Out[2]:



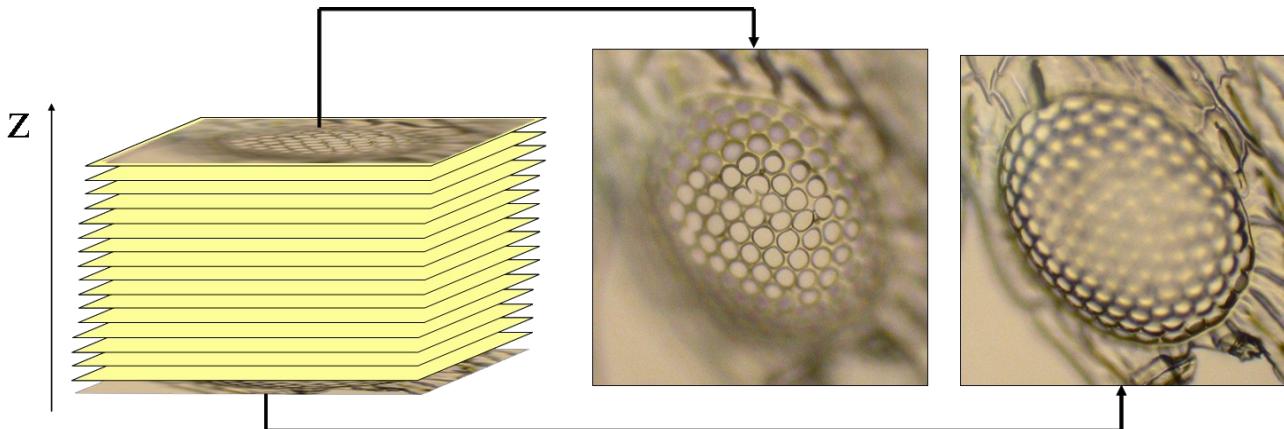
Wikimedia Commons (https://en.wikipedia.org/wiki/Evolution_of_the_eye#/media/File:Diagram_of_eye_evolution.svg)

Eye, can be compounded, like insect vision, or non-compound (simple eyes) when a single lens-system focus light on all the sensible cells.

In [3]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/stack.png')
```

Out[3]:

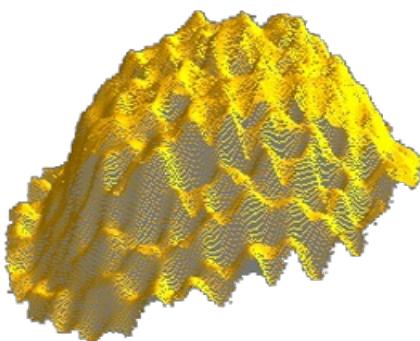


The figure above, illustrate how to reconstruct the 3D shape of an ant eye (a polymer mold is taken from the ant and acquired using a brightfield microscope), several Z focus position are used, by analysing the position of in focus point of the image, a 3D representation can be build (see below).

In [4]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/fourmis_3d.png')
```

Out[4]:



In fact, evolution has followed two different path to add vision capabilities, for example our mammalian eye has a blind spot due to the reversed optical nerve implantation.

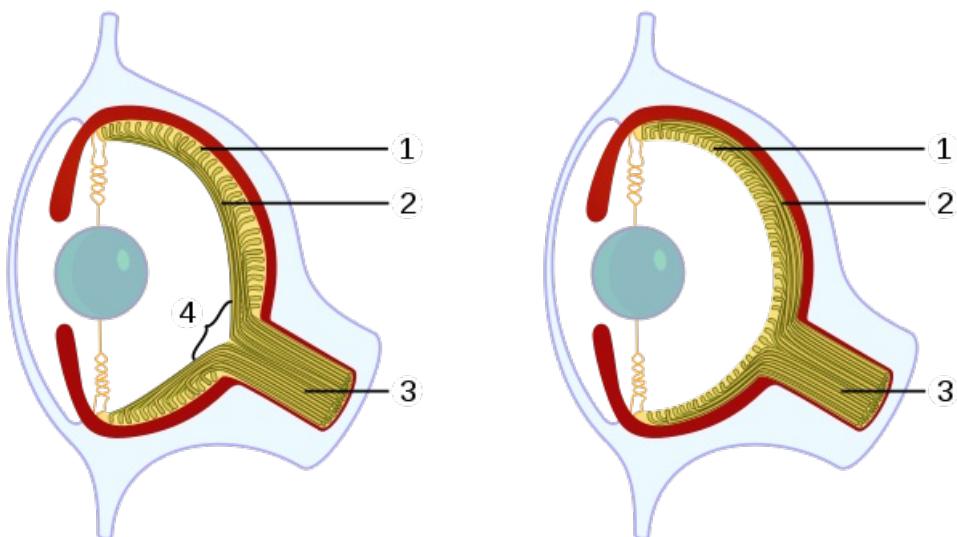
Vertebrates and octopuses developed the camera eye independently.

In the vertebrate version (left figure) the nerve fibers pass in front of the retina, and there is a blind spot where the nerves pass through the retina. In the vertebrate example, **4** represents the blind spot, which is notably absent from the octopus eye (right figure). In vertebrates, **1** represents the retina and **2** is the nerve fibers, including the optic nerve (**3**), whereas in the octopus eye, **1** and **2** represent the nerve fibers and retina respectively.[\[evolution of the eye - wikipedia \(https://en.wikipedia.org/wiki/Evolution_of_the_eye\)\]](https://en.wikipedia.org/wiki/Evolution_of_the_eye)

In [5]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/9/9b/Evolution_eye.svg/640px-Evolution_eye.svg.png')
```

Out[5]:



Wikimedia Commons (https://en.wikipedia.org/wiki/Evolution_of_the_eye#/media/File:Evolution_eye.svg)

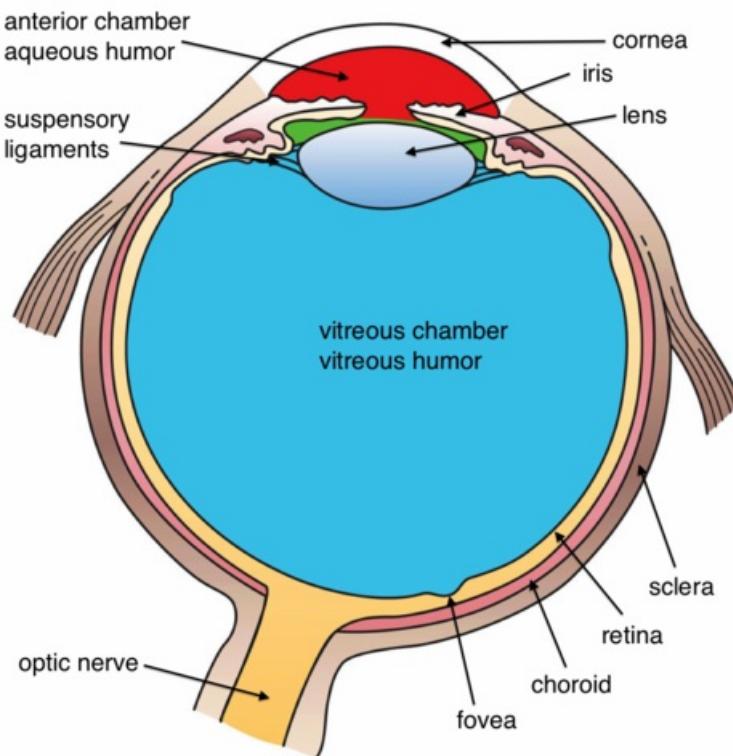
Human vision

Human vision rely on two principal organs: the eyes, and the brain.

In [6]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/8/8a/Three_Internal_chambers_of_the_Eye.png/469px-Three_Internal_chambers_of_the_Eye.png')
```

Out[6]:



wikimedia commons (https://commons.wikimedia.org/wiki/File:Three_Internal_chambers_of_the_Eye.png)

Sensitivity

Eye sensitivity is due to two types of cells: rods and cones. Rods cells are highly sensitive to brightness, their spectrum sensitivity is roughly situated in the green part of the visible spectrum (see below). Rods are spread all over the retina excepted in the fovea and of course in the blind spot.

Cones are cells responsible for the color discrimination, the highest cones concentration is located inside the fovea (see figure).

The ~100 million rods cover the entire retina, while the ~10 million cones cover only the 5° solid angle viewed by the fovea (in fact most of them are inside the 1° angle).

Actually, the eye does not give a full resolution image on the complete field of view at once, to have a complete picture (literally), the brain has to merge image information taken by the most spatially resolved part of the eye, the fovea, while the eye is *scanning* the scene.

In [7]:

Image('https://www.sfu.ca/cmns/courses/2011/325/Resources/Optics/Visual%20System/04%20Rods-and-Cones.jpg')

Out[7]:

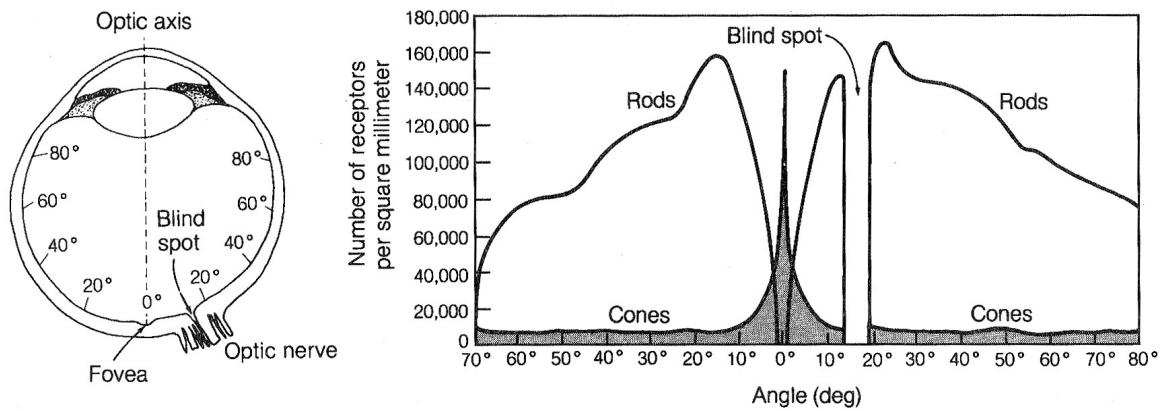


FIGURE 3-7 The distribution of rods and cones in the human retina. The left figure gives the locations on the retina of the “angle” relative to the optic axis on the right figure (based on Lindsay & Norman, 1977).

image source (<https://www.sfu.ca/cmns/courses/2011/325/Resources/Optics/Visual%20System/04%20Rods-and-Cones.jpg>)

While rods cells can be grouped in bundle to the optical nerve, meaning that the spatial localisation of the information is lower than the one given by cones having their own nervous termination.

The eye has also a system for light limitation, the iris. Similarly to the camera, the iris role is to modulate the quantity of light entering the eye, to increase its dynamic range (from very low light to complete sun exposed scene). Iris diameter is modulated from 2 to 8 mm giving a global dynamic up to 10^{10} .

But the retina itself has a limited dynamic range, i.e. the total intensity variation that it is capable of discriminating. The actual retina dynamic range is less than 200, this explains some technological choices that have been made for image storage (see in next chapters).

In [8]:

```
from skimage.data import camera
import matplotlib.pyplot as plt
import matplotlib.cm as cm

g = camera()
g_poster = (g>>3)

plt.figure(figsize=[12,5])
plt.subplot(1,2,1)
plt.imshow(g,cmap=cm.gray)
plt.colorbar()
plt.subplot(1,2,2)
plt.imshow(g_poster,cmap=cm.gray)
plt.colorbar();
```

The illustration above, shows the same image with a dynamic of 256 levels (left) and only 32 levels (right). With only 32 levels, one can see false contours appearing in large continuous regions (e.g. the sky). But for the left image, it is almost impossible to discriminate very close graylevels from each other.

However thanks to the agile displacement of the eye and the reconstruction of the brain, we are able to look at scenes with a very high dynamic range as illustrated below (with much more than 255 levels!).

In [9]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/4/46/HDR_image_%2B_3_source_pictures_%28Cerro_Tronador%2C_Argentina%29.jpg/542px-HDR_image_%2B_3_source_pictures_%28Cerro_Tronador%2C_Argentina%29.jpg')
```

Out[9]:



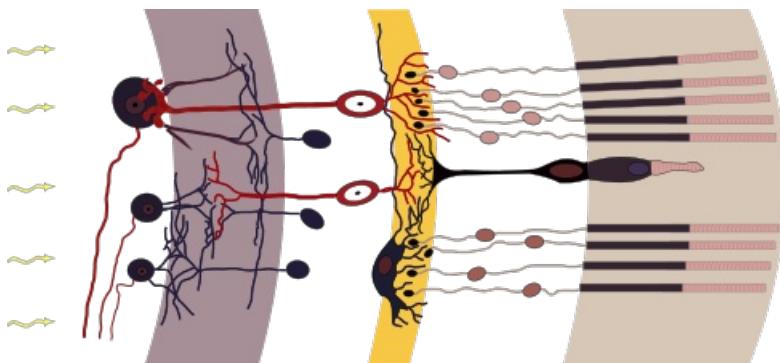
[wikimedia commons \(https://commons.wikimedia.org/wiki/File:HDR_image_%2B_3_source_pictures_\(Cerro_Tronador,_Argentina\).jpg\)](https://commons.wikimedia.org/wiki/File:HDR_image_%2B_3_source_pictures_(Cerro_Tronador,_Argentina).jpg)

The anatomy of the retina is composed of several layers, it is interesting to note that a typical nervous signal speed is ~ 100 m/s, and its commutation time is about 25 Hz. We can therefore be astonished by how fast our vision is able to process some information. In fact, the natural vision *processing power* is hidden in the high parallelism achieved by the complete chain of 100.000.000.000 interconnected neurons!

In [10]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/d/d6/Retina-diagram.svg/500px-Retina-diagram.svg.png')
```

Out[10]:



[image source \(http://www.slideshare.net/schwartzcm/ch-10-senses-part-ii\)](http://www.slideshare.net/schwartzcm/ch-10-senses-part-ii)

The light acquisition is done by the eyes only, but the processing is shared by both organs.

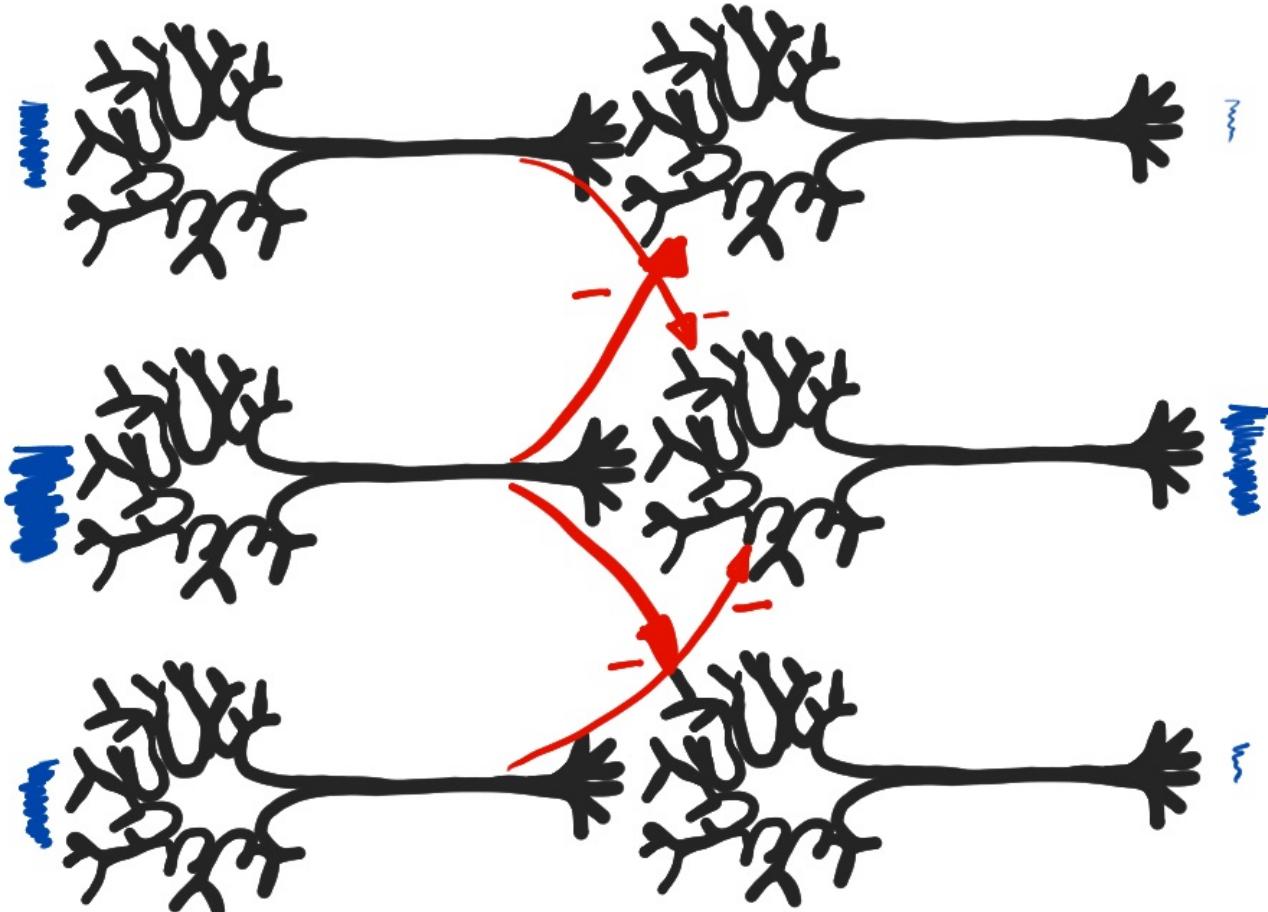
Indeed some low-level image processing is done at the retina level, for example the first stage of direction selectivity is accomplished in the inner layer of the retina see [lee2006 \(<http://www.sciencedirect.com/science/article/pii/S0896627306006283>\)](http://www.sciencedirect.com/science/article/pii/S0896627306006283).

The phenomenon of **lateral inhibition** occurs when a stimulus is applied to neighbour neurons. Each neuron will negatively influence its neighbours (relative to its own signal intensity). The resulting pattern has a high contrast.

In [11]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/lat_inhib3.png')
```

Out[11]:



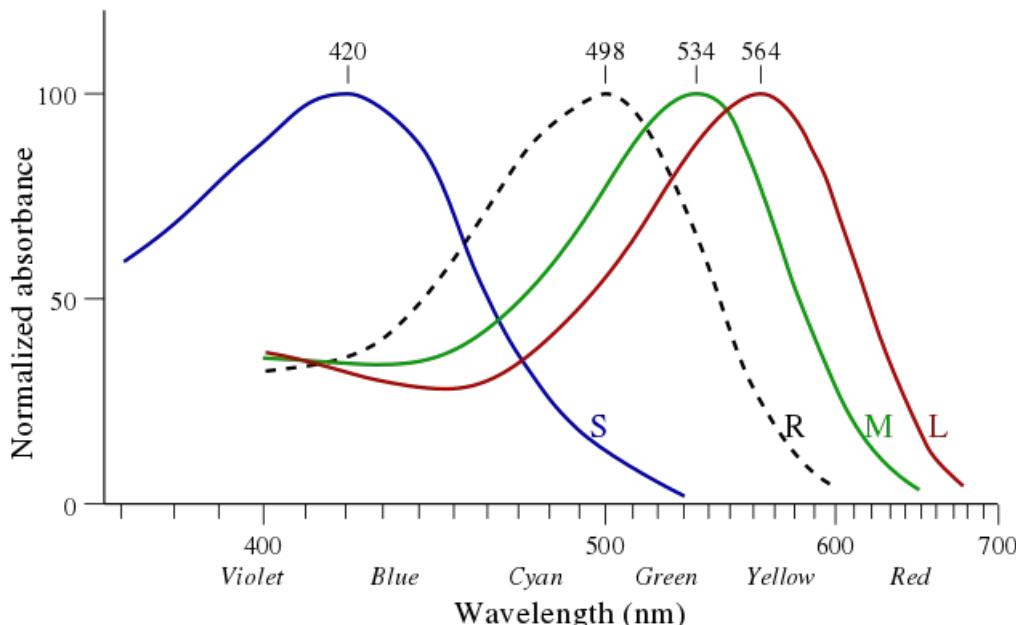
Color perception

Color perception is enabled by the presence of 3 types of cones in the retina. The sensitivity spectrum of these cones is spread over the visible spectrum as illustrated below.

In [12]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/2/26/Cone-response-en.svg/640px-Cone-response-en.svg.png')
```

Out[12]:



wikimedia commons (<https://commons.wikimedia.org/wiki/File:Cone-response.svg>)

Our spectrum of acquisition is limited on one hand at the ultraviolet light, and on the other by the near infrared light. We will see that image analysis is not limited to this range, depending of the image sensor used.

Other species developed a vision in a shifted spectrum (e.g. bees can see some UV).

In [13]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/0/0d/UV_and_Vis_Sunscreen.jpg/640px-UV_and_Vis_Sunscreen.jpg')
```

Out[13]:



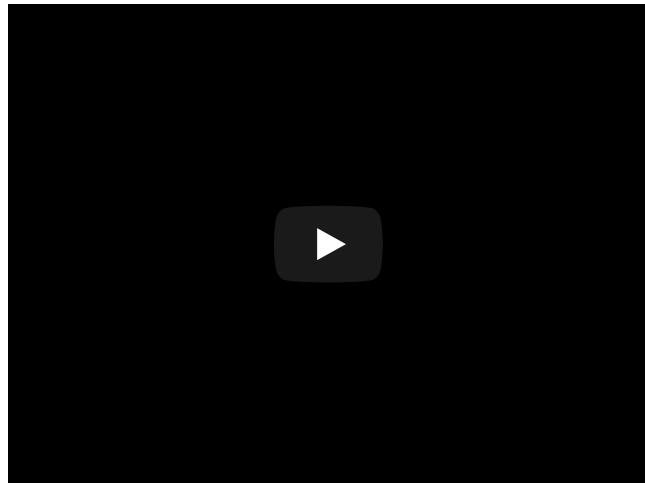
wikimedia commons (https://commons.wikimedia.org/wiki/File:UV_and_Vis_Sunscreen.jpg)

Persistence

In [14]:

```
YouTubeVideo('Md78s0I1-r8')
```

Out[14]:



Because of the, so to speak, slow speed of firing of the chemico-physical processes involved in the vision, a fast change in light intensity may be impossible to be detected. This is why, as illustrated above, a fast fast rotating stripe of leds can give the illusion of being a 2D disk (rem. actually the video present already a remanent object, due to the camera limited acquisition speed and frame integration time...).

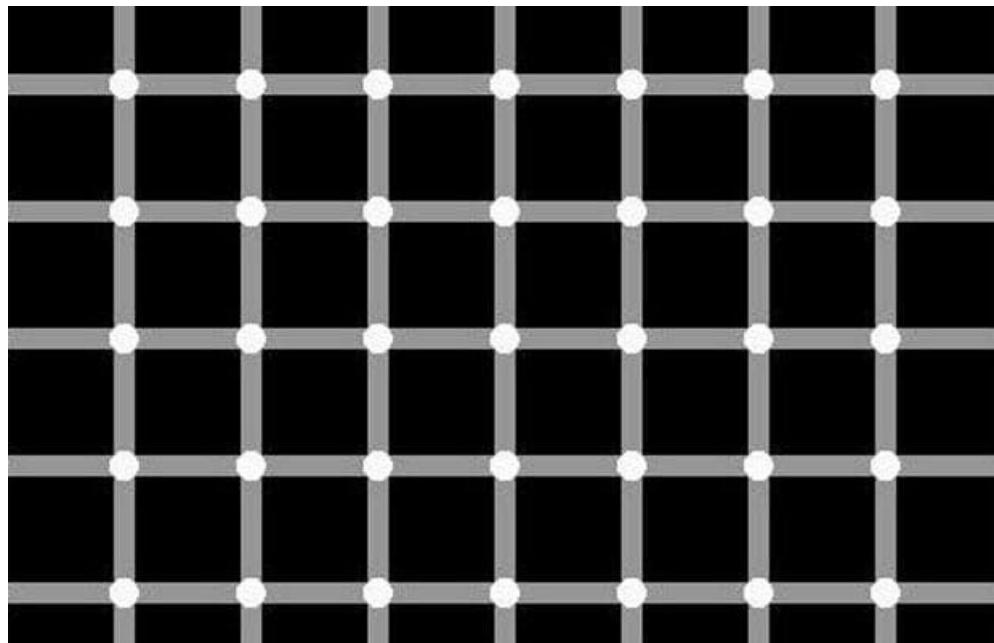
Vision limitations and optical illusions

Here are some classical optical illusions that can trick your eye, maybe machine vision can be used here to avoid some troubles...

In [15]:

```
Image('http://i.telegraph.co.uk/multimedia/archive/01120/blackballs_1120775i.jpg')
```

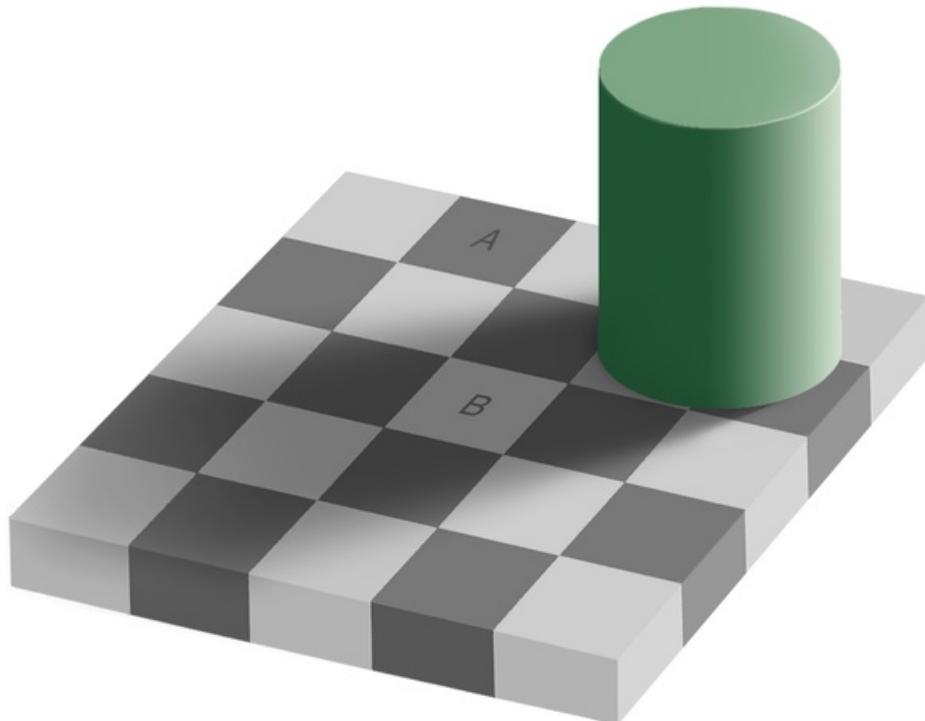
Out[15]:



In [16]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/6/60/Grey_square_optical_illusion.PNG/618px-Grey_square_optical_illusion.PNG')
```

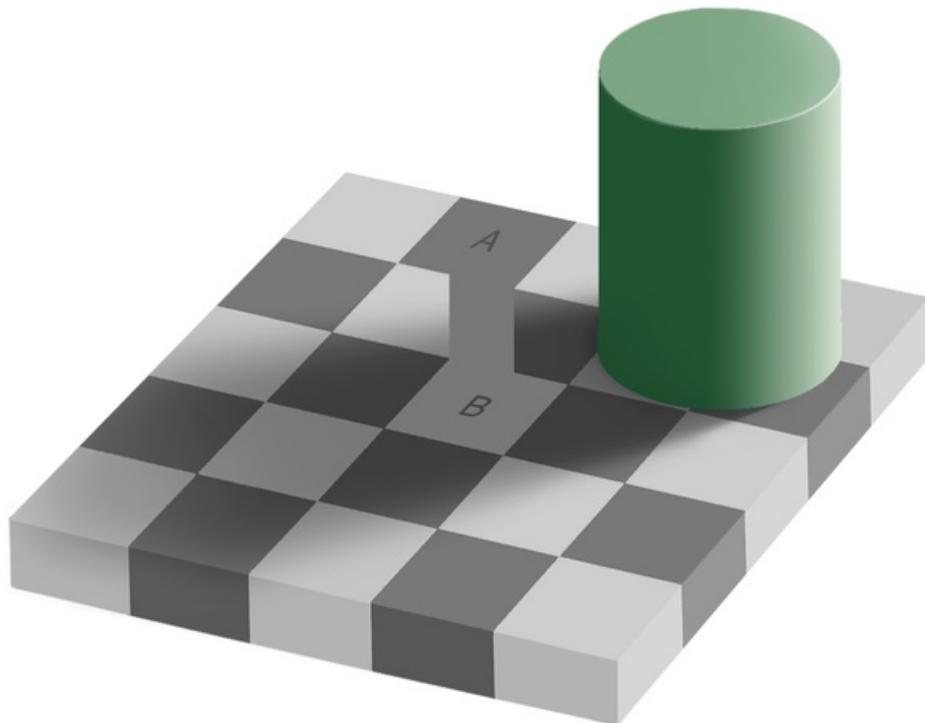
Out[16]:



In [17]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/e/ed/Same_color_illusion_proof2.png/618px-Same_color_illusion_proof2.png')
```

Out[17]:

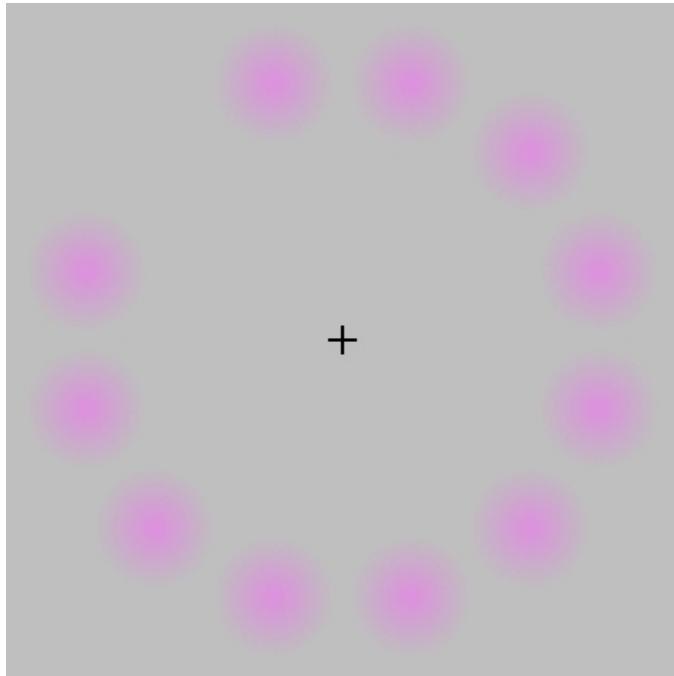


By Original by Edward H. Adelson, this file by Gustavb [Copyrighted free use], via Wikimedia Commons

In [18]:

```
Image(url='https://upload.wikimedia.org/wikipedia/commons/6/6e/Lilac-Chaser.gif')
```

Out[18]:

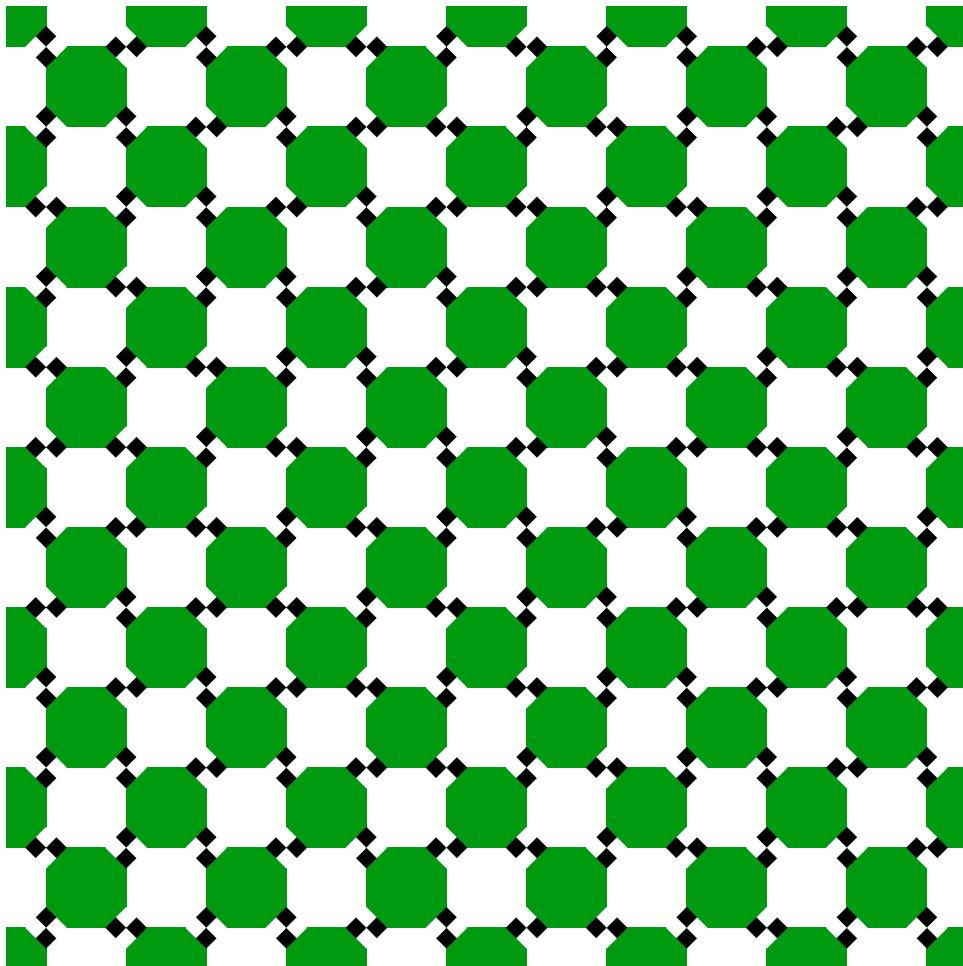


TotoBaggins at the English language Wikipedia [GFDL](http://www.gnu.org/copyleft/fdl.html) (<http://www.gnu.org/copyleft/fdl.html>) or [CC-BY-SA-3.0](http://creativecommons.org/licenses/by-sa/3.0/) (<http://creativecommons.org/licenses/by-sa/3.0/>), via Wikimedia Commons

In [19]:

```
SVG('https://upload.wikimedia.org/wikipedia/commons/b/b8/0ptical-illusion-checkerboard-twisted-cord.svg')
```

Out[19]:



By AnonMoos [Public domain], via Wikimedia Commons

In []:

In [111]:

```
from IPython.display import HTML,Image,SVG,YouTubeVideo
```

Image sources

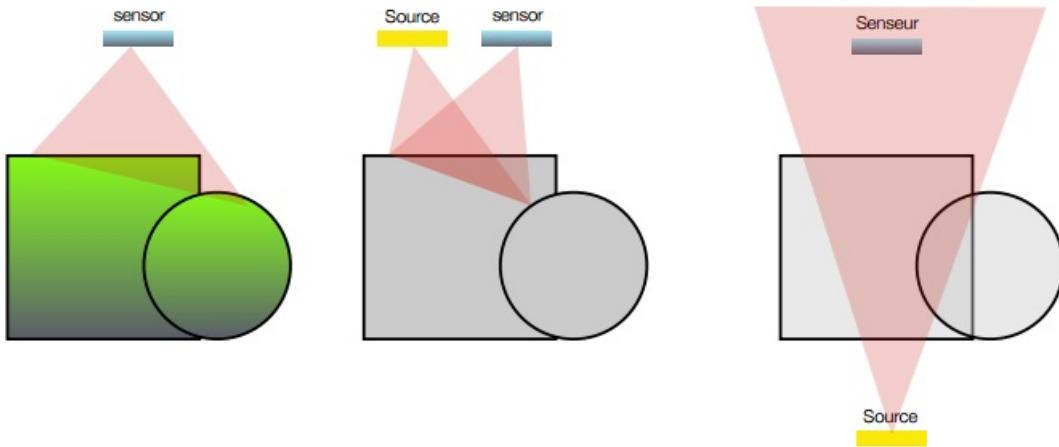
Passive vs active imaging

- the object is the source of photon (SPECT, stars,...)
- the object reflects/react to light given by a external source (flash, fluorescence)
- the object is traversed by the light and diffuses/asborbes it (X-ray)

In [112]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/trans_reflect.png')
```

Out[112]:



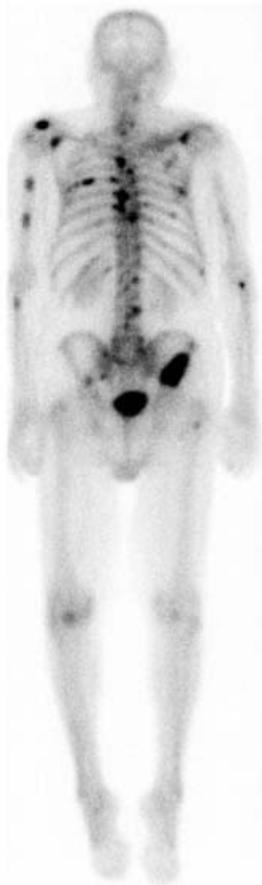
Object as a source

Nuclear imaging is a good example of the first setup, here an injection of radio-tracer will accumulates to some region of interest (due to specific biochemical affinity). The following example shows how the radio-tracer identifies bone metastasis of a prostate cancer using a gamma camera.

In [113]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/a/ae/Prostate-mets-102.jpg')
```

Out[113]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:Prostate-mets-102.jpg\)](https://commons.wikimedia.org/wiki/File:Prostate-mets-102.jpg)

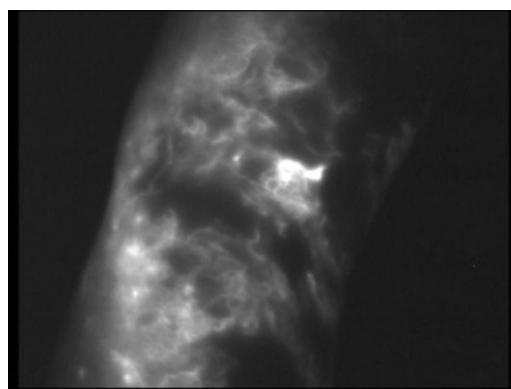
The source can also be the result of an external excitation i.e. an absorption and a re-emission of another photon (fluorescence).

Fluorescence lymphography is an example of imaging using an external excitation, here, infrared light is used to excite fluorophore injected in the lymph system. Fluorophore can in turn re-emit infrared (at a longer wavelength). By using adapted filter, one can observe the lymph displacement inside the lymph network (close to the skin surface).

In [114]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/fluoroscopy.png')
```

Out[114]:



J.P.Belgrado

An other example, where fluorescence is used: the fluorescence microscopy.

In [115]:

```
#add example
```

Object reflects / diffuses the light from an external source

This is the more common acquisition setup, external light source flood the scene with visible photons that are reflected by the objects, these photons are then acquired by a sensor.

Object attenuates the source

Source and sensor can be placed on both side of the object being imaged, a good example is the X-Ray imaging, where a X-Ray source project photon trough a patient, these photon interact with the matter in such a way that tissue density and composition (bones vs soft tissues) can give a contrast variation at the sensor level.

image source (<https://www.flickr.com/photos/tracemeek/5327224133>)

Direct image acquisition

CCD - coupled charge device

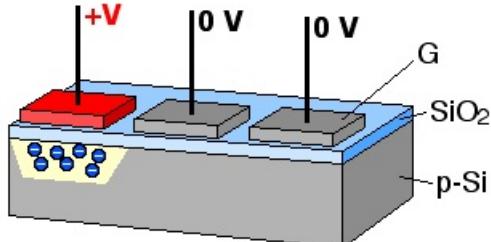
Charges are liberated by light interaction with the semiconductor inside photoactive region, for each pixel of the sensor grid. In order to digitize the amount of charges (proportional to light captured, CCD devices will move the charges along the substrate up to a charge to voltage converter.

Coupled Charge Device uses electrode potentials to move charges inside silicon substrate as illustrated below.

In [116]:

```
Image(url='https://upload.wikimedia.org/wikipedia/commons/6/66/CCD_charge_transfer_animation.gif')
```

Out[116]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:CCD_charge_transfer_animation.gif\)](https://commons.wikimedia.org/wiki/File:CCD_charge_transfer_animation.gif)

Image sensor can have essentially two types of geometry:

- linear: typically used when the sensor is translated (flatbed scanner, but also bank note scanner, satellite, photo finish)
- rectangular: almost every other camera

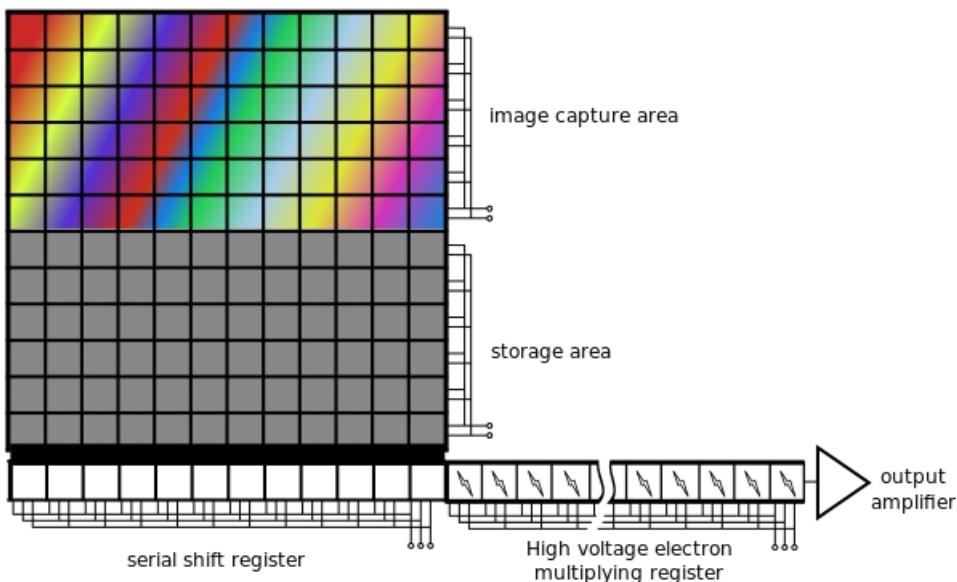
In order to move charges along the dimensions of the CCD sensor, charges are moved along each image line, a perpendicular buffer is used to discharge all these pixels in column into an amplifier that transform each charge into a voltage. The voltage is then converted by an ADC circuit.

Because all the pixels charges are compared using the same circuit, the CCD sensor provide a very constant specification on the complete sensor. The other main advantage of the sensor is the coverage factor of the sensor (the ratio between the sensor surface and the total pixel surface), almost the surface is devoted to light acquisition (no extra circuitry needed).

In [117]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/e/e1/EMCCD2_color_en.svg/640px-EMCCD2_color_en.svg.png')
```

Out[117]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File%3ACcd_schematic.JPG\)](https://commons.wikimedia.org/wiki/File%3ACcd_schematic.JPG)

In [118]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/3/30/CCD_line_sensor.JPG/320px-CCD_line_sensor.JPG')
```

Out[118]:

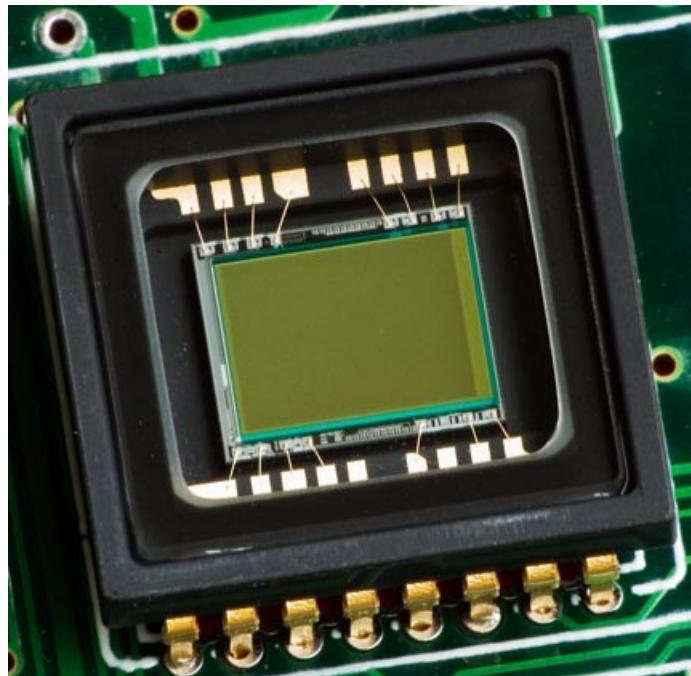


Linear CCD sensor. [wikimedia commons \(https://commons.wikimedia.org/wiki/File:CCD_line_sensor.JPG\)](https://commons.wikimedia.org/wiki/File:CCD_line_sensor.JPG)

In [119]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/1/17/CCD_in_camera.jpg')
```

Out[119]:



CCD line sensor in a ceramic dual in-line package. [wikimedia commons \(https://commons.wikimedia.org/wiki/File:CCD_in_camera.jpg\)](https://commons.wikimedia.org/wiki/File:CCD_in_camera.jpg)

CMOS

The CMOS technology embeds a photo-detector and a charge amplifier for each sensor pixel, the voltage being then transmitted by electrical conductors.

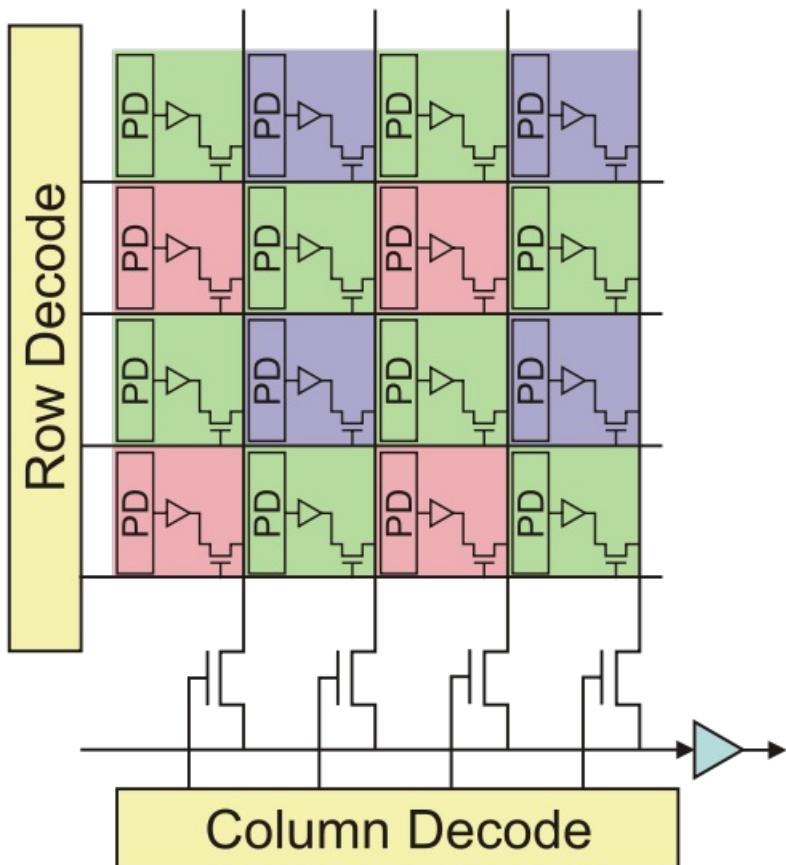
This strategy enables a greater variety of sensor usage, e.g. addressing a part of the sensor (for low resolution and higher speed).

Because the conversion is done separately for each pixels, no charge shifting is needed, but discrepancy between charge amplifier may exist, giving unequal pixel sensitivity and noise.

In [120]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/c/c0/CMOS_Image_Sensor_Mechanism_Illustration.svg/500px-CMOS_Image_Sensor_Mechanism_Illustration.svg.png')
```

Out[120]:



wikimedia commons (https://commons.wikimedia.org/wiki/File:CMOS_Image_Sensor_Mechanism_Illustration.svg)

CMOS vs CCD

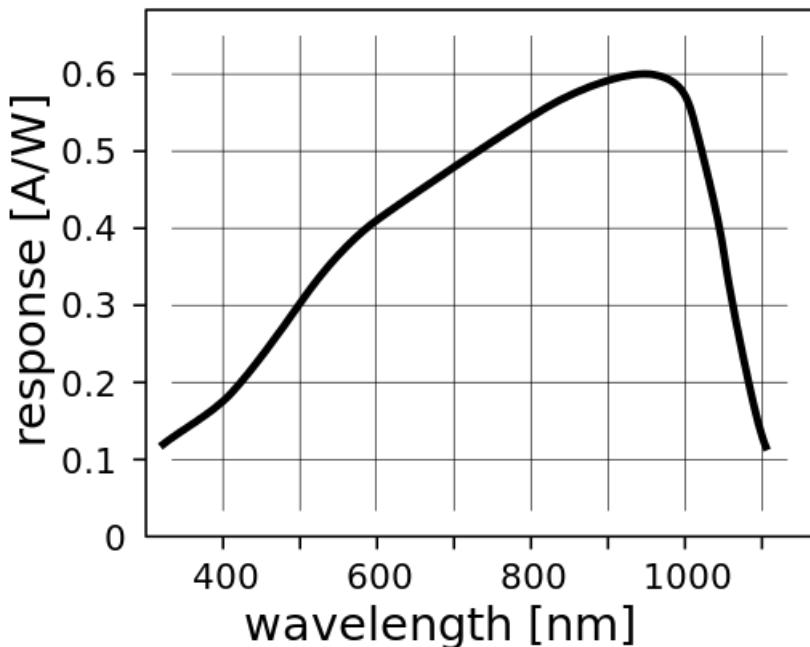
feature	CDD	CMOS
Signal out of pixel	Electron packet	Voltage
Fill factor	high	moderate
Amplifier mismatch	none	moderate
Noise	low	moderate
system complexity	high	low
sensor complexity	low	high
dynamic range	high	moderate
uniformity	high	moderate
speed	moderate	high

CMOS + CDD : high sensitivity to near infrared, therefore, most of the sensors are equipped with a NIR filter.

In [121]:

```
Image(url='https://upload.wikimedia.org/wikipedia/commons/thumb/4/41/Response_silicon_photodiode.svg/544px-Response_silicon_photodiode.svg.png')
```

Out[121]:



[wikimedia commons \(https://upload.wikimedia.org/wikipedia/commons/thumb/4/41/Response_silicon_photodiode.svg/544px-Response_silicon_photodiode.svg.png\)](https://upload.wikimedia.org/wikipedia/commons/thumb/4/41/Response_silicon_photodiode.svg/544px-Response_silicon_photodiode.svg.png)

Multispectral acquisition

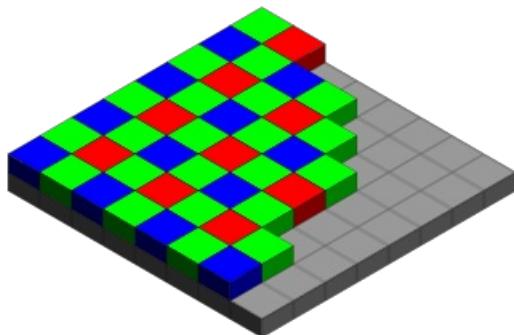
Color acquisition is done by acquiring several images at different wavelength, one common (and cheap) approach is to cover sensors pixels by colored dyes (red, green and blue). The figure above illustrated such filters (bayer), where on each 2x2 pixel square, one pixel is sensitive to the red part of the spectrum, one to the blue part of the spectrum, and finally 2 pixels sensitive to the green part of the spectrum.

The choice of duplicating green is done for symmetry purposes and also because the intensity sensitivity of the eye (see rods) is correlated to the green part of the spectrum.

In [122]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/3/37/Bayer_pattern_on_sensor.svg/320px-Bayer_pattern_on_sensor.svg.png')
```

Out[122]:



[wiki commons \(https://en.wikipedia.org/wiki/Bayer_filter#/media/File:Bayer_pattern_on_sensor.svg\)](https://en.wikipedia.org/wiki/Bayer_filter#/media/File:Bayer_pattern_on_sensor.svg)

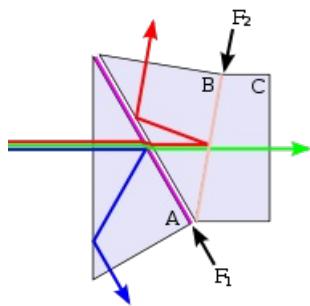
One limitation of the dye approach is the resolution limitation, indeed the image resolution is divided by 4.

The other method used is based on three CCD coupled on the same optical axis and having three different dyes (red, green, blue) as illustrated bellow.

In [123]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/e/ef/Dichroic-prism.svg/200px-Dichroic-prism.svg.png')
```

Out[123]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:Dichroic-prism.svg\)](https://commons.wikimedia.org/wiki/File:Dichroic-prism.svg)

The big advantage of this approach is to keep the sensor native resolution for each color channel.

The number of spectral bands can be higher than three, for example satellite imagery offers many wavelength inside but also next to it (UV and near-IR).

Quick-bird (environmental imagery, pixel = 0.65m)

- Pan: 450-900 nm
- Blue: 450-520 nm
- Green: 520-600 nm
- Red: 630-690 nm
- Near IR: 760-900 nm

IKONOS (commercial earth observation satellite)

resolution

- 0.8 m panchromatic (1-m PAN)
- 4-meter multispectral (4-m MS)

spectrum

- Blue: 0.445–0.516 μm
- Green: 0.506–0.595 μm
- Red: 0.632–0.698 μm
- Near IR: 0.757–0.853 μm

Landsat 8 (American Earth observation satellite)

- Band 1 - Coastal / Aerosol 0.433 - 0.453 μm 30 m
- Band 2 - Blue 0.450 - 0.515 μm 30 m
- Band 3 - Green 0.525 - 0.600 μm 30 m
- Band 4 - Red 0.630 - 0.680 μm 30 m
- Band 5 - Near Infrared 0.845 - 0.885 μm 30 m
- Band 6 - Short Wavelength Infrared 1.560 - 1.660 μm 30 m
- Band 7 - Short Wavelength Infrared 2.100 - 2.300 μm 30 m
- Band 8 - Panchromatic 0.500 - 0.680 μm 15 m
- Band 9 - Cirrus 1.360 - 1.390 μm 30 m

AVIRIS - airborne visible/infrared imaging spectrometer

- four linear spectrometers (614-pixel wide) / 224 adjacent spectral bands.

In [124]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/4/48/HyperspectralCube.jpg')
```

Out[124]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:HyperspectralCube.jpg\)](https://commons.wikimedia.org/wiki/File:HyperspectralCube.jpg)

Depth acquisition

Depth imaging is traditionnaly used in stereo application, such for robot vision. Recently depth sensor became widely available thanks to game applications. The main technologies used are:

- stereovision
- laser triangulation
- structured light projection
- Time-Of-Flight (TOF) imaging

The information provided by these sensors is of two types: a rgb image of the scene, and a depth estimation (usually at a coarser resolution).

example of a high resolution laser triangulation scanner:

In [125]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/scanner.png')
```

Out[125]:

Laser scanner



When high speed is needed, structured light may be a solution.

For example, the first generation of the Kinect sensor uses the principle of structured light projection, a pseudo-random pattern is projected in the near-infrared spectrum(i.e. invisible to human eye) and acquired by a IR sensitive camera. The depth image is produced with a video framerate compatible with gaming.

In [126]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/6/67/Xbox-360-Kinect-Standalone.png/320px-Xbox-360-Kinect-Standalone.png')
```

Out[126]:



wikimedia commons (<https://commons.wikimedia.org/wiki/File:Xbox-360-Kinect-Standalone.png>)

In [127]:

```
Image('http://www.mattcutts.com/images/ir-projection.jpg')
```

Out[127]:

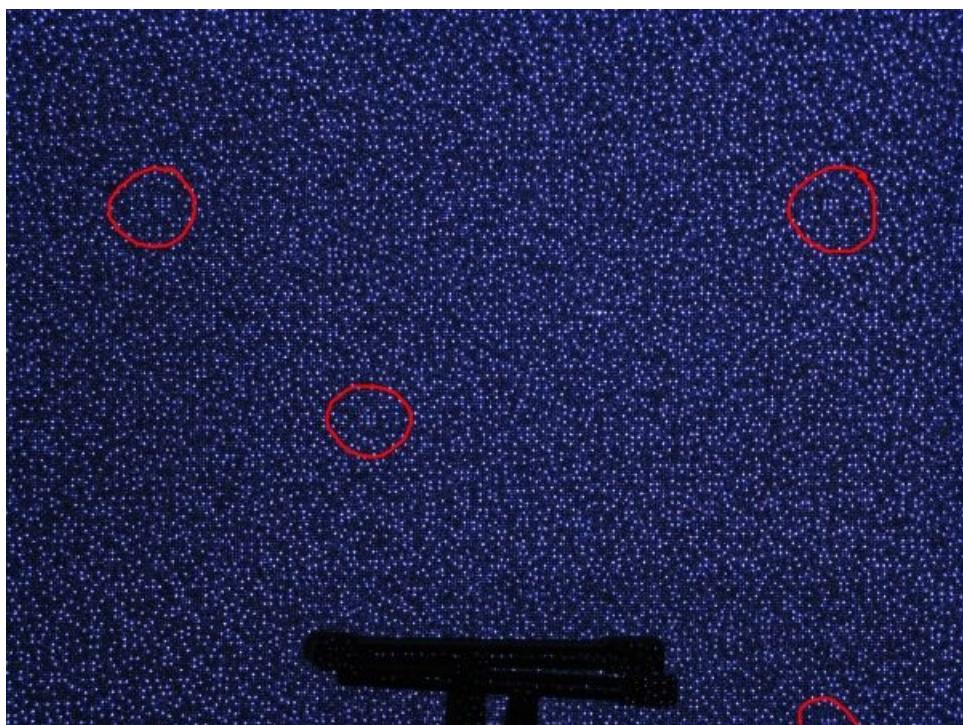


[image source \(https://www.mattcutts.com/blog/open-kinect-contest/\)](https://www.mattcutts.com/blog/open-kinect-contest/)

In [128]:

```
Image('http://3.bp.blogspot.com/_PsITwyT0c4Y/T0UvQTQE7WI/AAAAAAAAPg/cGHQtXou2yw/s1600/Kinect+Pattern_IMG_0073.jpg')
```

Out[128]:



[image source \(http://image-sensors-world.blogspot.be/2010_11_01_archive.html\)](http://image-sensors-world.blogspot.be/2010_11_01_archive.html)

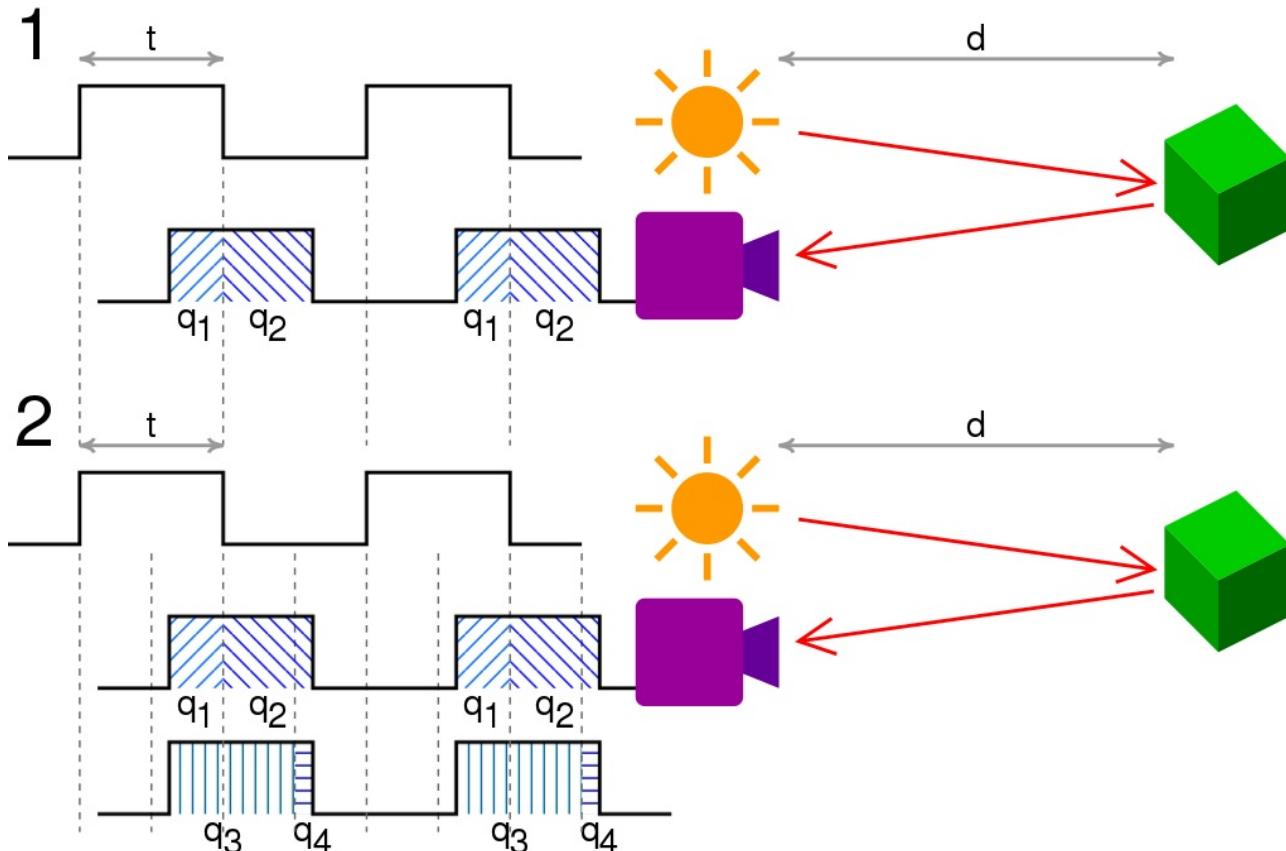
The depth is computed by triangulation thanks to the identification of specific pattern in the image.

The second generation of sensors is based on a completely different technology, the Time-Of-Flight (TOF). To estimate the distance between the sensor and the scene, a light wave is send and received by the sensor. The phase difference between a modulated light pattern sended by the source and the signal received by the camera gives a measure of the scene depth.

In [129]:

Image(url='https://upload.wikimedia.org/wikipedia/commons/thumb/7/79/Time_of_flight_camera_principle.svg/1024px-Time_of_flight_camera_principle.svg.png')

Out[129]:



How to measure distance with light ?

Continuous wave demodulation

- retrieve phase shift by demodulation of the received signal
- demodulation by cross-correlation of the received signal with the emitted signal
- emitted signal is

$$g(t) = \cos(\omega t)$$

with ω the modulation frequency

- received signal after the return trip to the scene surface:

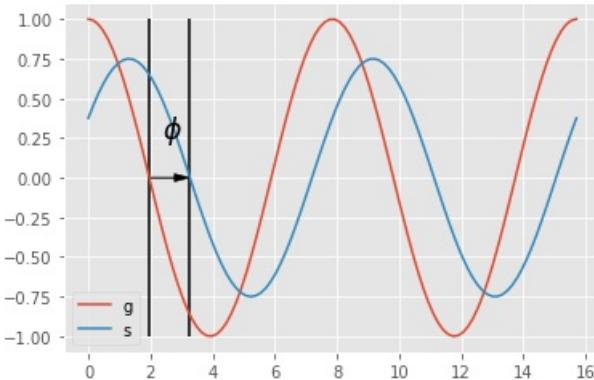
$$s(t) = b + a \cos(\omega t + \phi)$$

where a is an unknown attenuation, ϕ the phase shift **i.e. a value proportional to the scene distance** and b an unknown acquisition noise (neglected here).

In [130]:

```
%matplotlib inline
import numpy as np
import matplotlib.pyplot as plt
from matplotlib import cm
from skimage.data import camera
plt.style.use('ggplot')

plt.figure()
omega = .8
t = np.linspace(0,5*np.pi,100)
g = np.cos(omega*t)
a = .75
phi = -np.pi/3
s = a * np.cos(omega*t+phi)
plt.plot(t,g,label='g')
plt.plot(t,s,label='s')
plt.gca().arrow(np.pi*.5/omega, 0, -phi/omega-.5, 0, head_width=0.05, head_length=0.5, fc='k', ec='k')
plt.vlines([.5*np.pi/omega,(.5*np.pi-phi)/omega],-1,1)
plt.text(.6*np.pi/omega,.25,'$\phi$',size='xx-large')
plt.legend();
```



- cross correlation of both emitted and received signal becomes:

$$d(\tau) = s * g = \int_{-\infty}^{+\infty} s(t) \cdot g(t + \tau) dt$$

with τ an internal offset

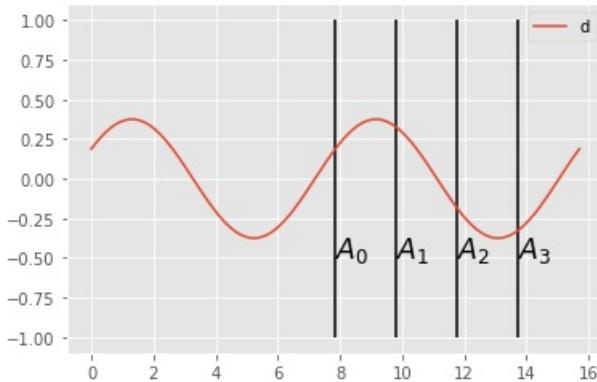
$$d(\tau) = \frac{a}{2} \cos(\omega t + \phi) + b$$

- sample $d(\tau)$ at 4 distinct moments (phase offsets):

$$A_i = d(i \cdot \frac{\pi}{2}) \text{ with } i = 0, \dots, 3$$

In [131]:

```
plt.figure()
d = a/2 * np.cos(omega*t+phi)
plt.plot(t,d,label='d')
samples = 1/omega * (np.pi*2+np.arange(0,2*np.pi,np.pi/2))
plt.vlines(samples,-1,1)
for i,pos in enumerate(samples):
    plt.text(pos,-.5,'$A_{%d}$'%i, size='xx-large')
plt.legend();
```



- phase and attenuation are then:

$$\phi = \arctan\left(\frac{A_3 - A_1}{A_0 - A_2}\right)$$

and

$$a = \frac{1}{2} \sqrt{(A_3 - A_1)^2 + (A_0 - A_2)^2}$$

- scene distance is then:

$$dist = \frac{c}{4\pi\omega} \phi$$

where c is the speed of light.

What a depth image looks like ?

In [132]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/1/19/TOF_Kamera_3D_Gesicht.jpg')
```

Out[132]:



Indirect image acquisition

Image can also be the result of a mathematical reconstruction based on an indirect acquisition, the sensor do not acquire an image directly.

For example, computed tomography, uses a series of 1D density profile acquisition enable a 2D reconstruction of the slice.

In [133]:

```
Image('http://130.237.83.53/medicaldevices/album/Ch%207%20Medical%20images/slides/F%207-10%20Computer%20tomography.jpg')
```

Out[133]:

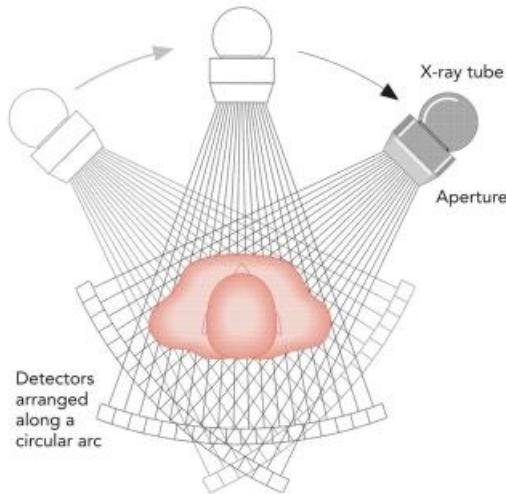


Figure 7-10 Computer tomography

[image source \('http://130.237.83.53/medicaldevices/album/Ch%207%20Medical%20images/slides/F%207-10%20Computer%20tomography.jpg'\)](http://130.237.83.53/medicaldevices/album/Ch%207%20Medical%20images/slides/F%207-10%20Computer%20tomography.jpg)

Echography is another example of indirect imaging, where mechanical wave propagation are transformed in a 2D image showing the presence of interfaces between tissue of different acoustic impedance.

In [134]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/c/c7/CRL_Crown_rump_length_12_weeks_ecografia_Dr._Wolfgang_Moroder.jpg')
```

Out[134]:



image source ('https://commons.wikimedia.org/wiki/File:CRL_Crown_rump_length_12_weeks_ecografia_Dr._Wolfgang_Moroder.jpg')

other example: MRI image reconstruction

Synthetic images

Image can also be the result of the grouping of a huge number of localized data, for example, one can imagine a network of temperature sensors spread over a complete country, then the temperature measurements can be grouped on a 2D map (and interpolated to have a complete coverage).

Visualized data can be from various nature, the common aspect is that these data are placed in a geometric space (usually 2D or 3D).

In [68]:

```
from IPython.display import HTML,Image,SVG,YouTubeVideo
```

Image representation

A digital image is discrete. It means that there is somewhere in the acquisition process, most of the time, sampling that occurs.

A digital image is basically a multidimensional array of numbers. Each picture element store a numerical value, with 2D images we speak about **pixels** (from Picture Elements) and for 3D images we use **voxels** (from VOlume EElements).

The lattice of pixels are usually rectangular (or square if pixels are squares) or hexagonal, hexagonal lattice has a unique distance between a pixel and its neighbors.

The image dimensionality will depend on:

- the spatial dimensionality: 1D or profile, 2D image, 3D volume
- the temporal dimension: add one dimension for the time when dealing with **sequences**
- the spectral dimension: the number of spectral values associated with one image element

The spatial dimension is given, typically, by the grid step of the sensor. A 640x480 CMOS sensor will produce a 640x480 pixels grid.

For a flatbed scanner, one dimension will be given by the number of sensor along the acquisition line, while the second dimension will be given by the spatial repetition of a line acquisition.

One remark concerning these images: the shape of the pixels is not always a square, depending on the sensor geometry and/or the sampling speed (i.e. for the scanners). One have to pay attention to that, in particular when we will extract measures from images (e.g. distance or surface).

When acquisition is a sequence one add one dimension for the time.

Similarly, image can be composed of different spectral band.

For example, a time-lapse acquisition of fluorescent confocal microscopy can have 5-D:

- three X,Y and Z spatial dimensions (voxels)
- one spectral dimension , i.e. several fluorophore channels (e.g. dapi, alexa, GFP)
- one time dimension (time-lapse)

Image sampling

In [89]:

```
from scipy.ndimage.filters import gaussian_filter
from skimage.data import camera
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np

def profile(im,p0,p1,num):
    n = np.linspace(p0[0],p1[0],num)
    m = np.linspace(p0[1],p1[1],num)
    return [n,m,ima[m.astype(int),n.astype(int)]]]

im = camera()[-1::-2,:,:2]

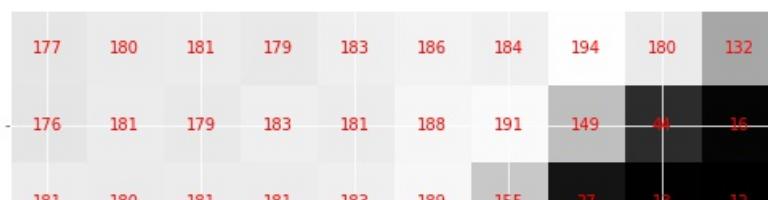
#detail
p0 = (60, 190)
p1 = (70, 180)
sub = im[p1[1]:p0[1],p0[0]:p1[0]]

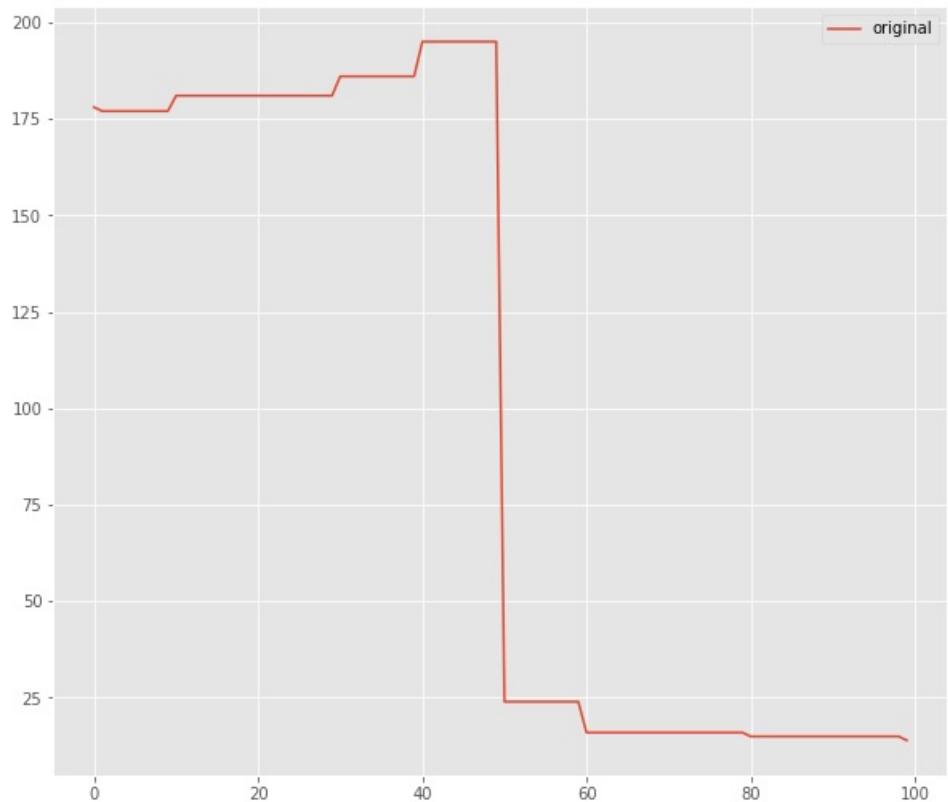
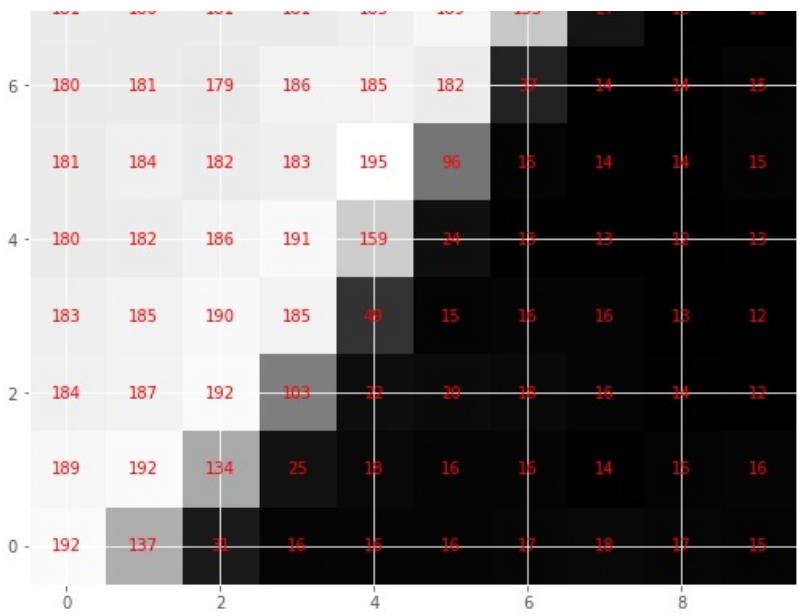
[x,y,p] = profile(im, p0, p1,100)

plt.figure(figsize=[10,30])
plt.subplot(3,1,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)

plt.subplot(3,1,2)
plt.imshow(sub,interpolation='nearest',cmap=cm.gray,origin='lower')
mm,nn = sub.shape
for m in range(mm):
    for n in range(nn):
        plt.text(n,m,'%d'%sub[m,n],color='r',horizontalalignment='center',verticalalignment='center')

plt.subplot(3,1,3)
plt.plot(p,label='original')
plt.legend();
```





In [70]:

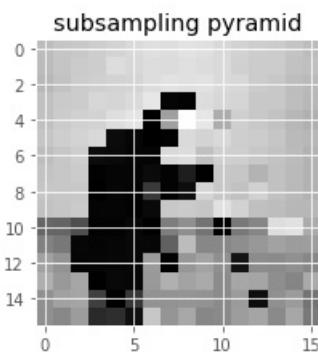
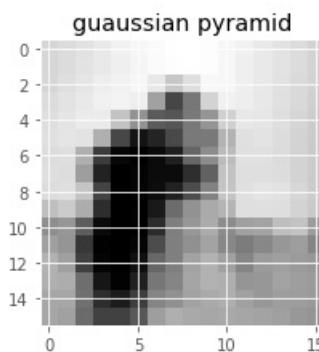
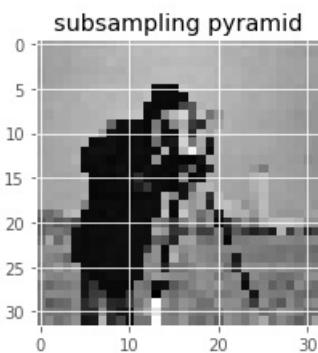
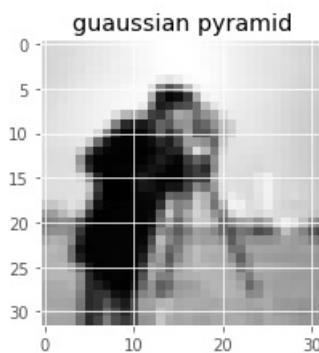
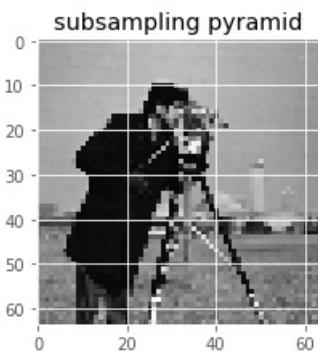
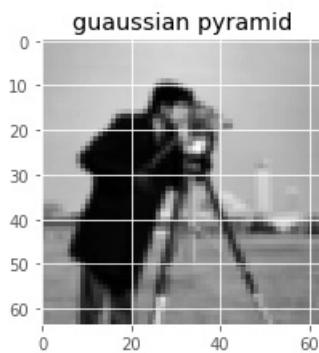
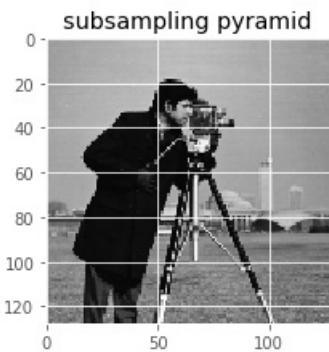
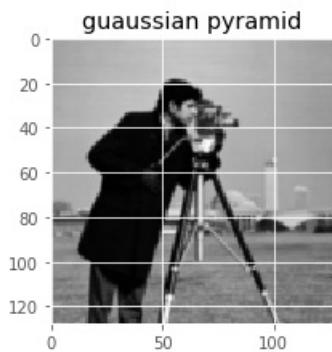
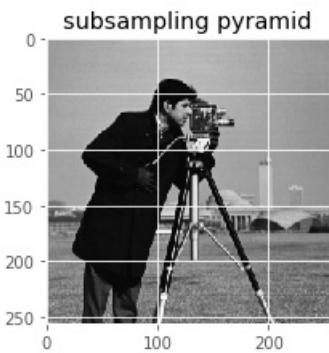
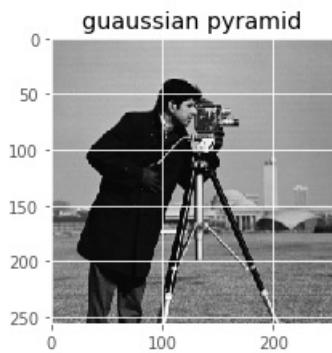
```
def build_gaussian_pyramid(ima,levelmax):
    """return a list of subsampled images (using gaussian pre-filter
    """
    r = [ima]
    current = ima
    for level in range(levelmax):
        lp = gaussian_filter(current,1.0)
        sub = lp[::2,::2]
        current = sub
        r.append(current)
    return r

def build_pyramid(ima,levelmax):
    """return a list of subsampled images (using gaussian pre-filter
    """
    r = [ima]
    current = ima
    for level in range(levelmax):
        sub = current[::2,::2]
        current = sub
        r.append(current)
    return r

im = camera()[::2,::2]

#build filtered and non-filtered pyramids
N = 4
fpyramid = build_gaussian_pyramid(im,N)
nfpymad = build_pyramid(im,N)

for f,nf in zip(fpyramid,nfpymad):
    plt.figure(figsize=[7,7])
    plt.subplot(1,2,1)
    plt.imshow(f,cmap=plt.cm.gray,interpolation='nearest')
    plt.title('gaussian pyramid')
    plt.subplot(1,2,2)
    plt.imshow(nf,cmap=plt.cm.gray,interpolation='nearest')
    plt.title('subsampling pyramid');
```



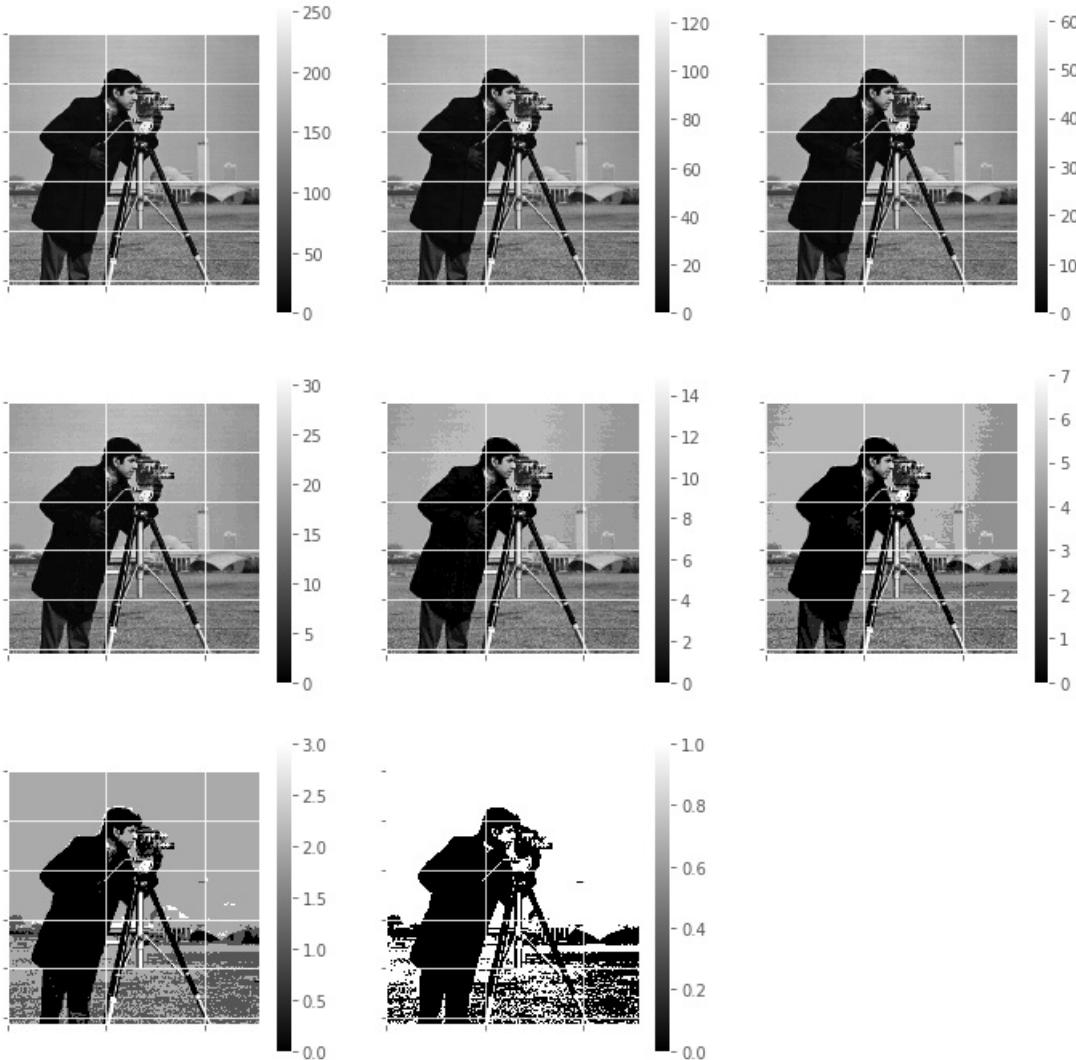
Level sampling

In [71]:

```
g = camera()
plt.figure(figsize=[12,12])

for s in range(0,8):
    g_poster = (g>>s)

    plt.subplot(3,3,s+1)
    plt.imshow(g_poster,cmap=cm.gray)
    plt.colorbar()
    plt.gca().set_xticklabels([])
    plt.gca().set_yticklabels([])
```



Information

entropy of a signal is given by:

$$h = - \sum_i p_i \log p_i$$

where p_i is the probability of occurrence of a symbol i .

For a gray scale image, one can consider the gray level distribution as the 'probability' of occurrence of a gray level. The following example illustrate how image entropy vary with respect to the graylevel distribution.

If the logarithm base used is 2, the entropy corresponds to the number of bits required to encode the signal.

see also:

- image entropy [IPAMV \(./00-Preface/06-References.ipynb#\[IPAMV\]\)](#) p19

In [72]:

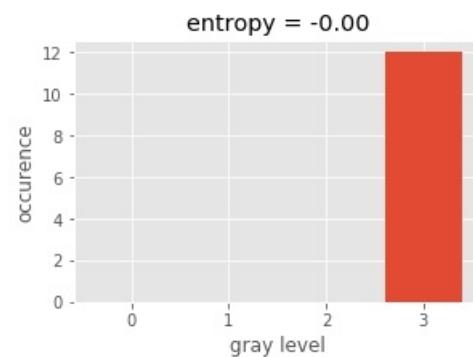
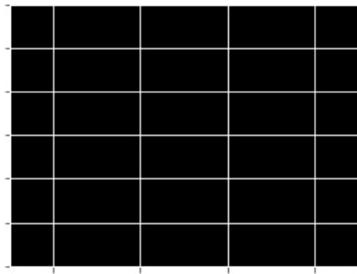
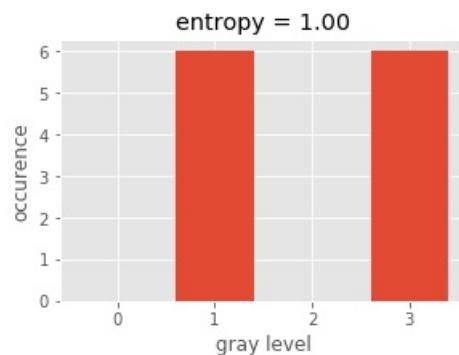
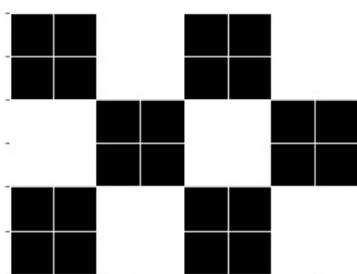
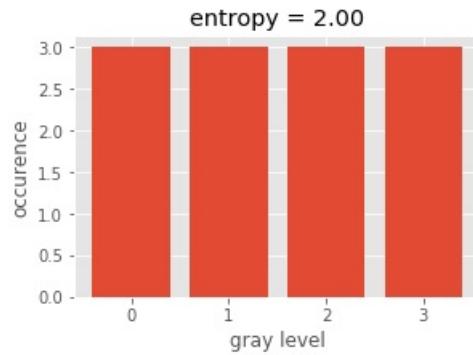
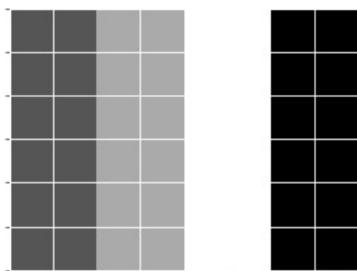
```
import numpy as np
import matplotlib.pyplot as plt
from matplotlib import cm
from skimage.data import camera
plt.style.use('ggplot')

def display_hist(ima,nbin):
    hist,bins = np.histogram(ima.flatten(),bins=range(0,nbin+1))
    norm_hist = 1.*hist/np.sum(hist) # normalized histogram
    # entropy
    idx = norm_hist>0
    h = -np.sum(norm_hist[idx]*np.log2(norm_hist[idx]))

    # display the results
    plt.figure(figsize=[10,3])
    ax = plt.subplot(1,2,1)
    plt.imshow(ima,cmap=cm.gray,interpolation='nearest')
    ax.set_xticklabels([])
    ax.set_yticklabels([])
    ax = plt.subplot(1,2,2)
    plt.bar(bins[:-1],hist,.8)
    if len(bins)<10:
        ax.set_xticks(bins[:-1])
        ax.set_xticklabels( bins[:-1] )
    plt.xlabel('gray level')
    plt.ylabel('occurrence');
    plt.title('entropy = %.2f'%h)

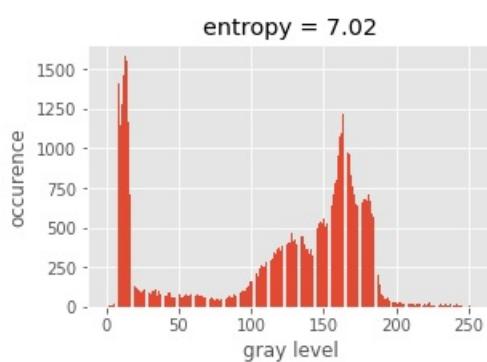
i1 = np.array([[1,2,3,0],[1,2,3,0],[1,2,3,0]])
i2 = np.array([[1,3,1,3],[3,1,3,1],[1,3,1,3]])
i3 = np.array([[3,3,3,3],[3,3,3,3],[3,3,3,3]])

display_hist(i1,4)
display_hist(i2,4)
display_hist(i3,4)
```



In [73]:

```
from skimage.data import camera
ima = camera()[:,::2]
display_hist(ima,255)
```



The entropy of the above image is 7.05, it means that we need a bit more than 7 bits to encode graylevel for this image. Which is consistent with the chosen data storage (8 bit per pixel).

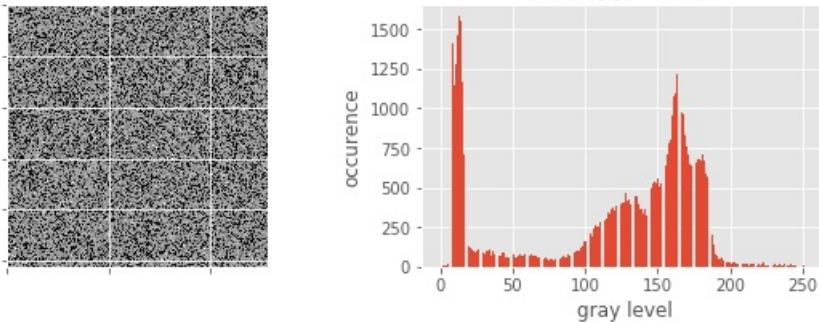
Entropy is often known as an information measure.

But what would be the entropy of an image where all the pixels are randomly permuted ?

In [74]:

```
# randomly permutes all the image pixels
d = ima.flatten()
shuffled_ima = np.random.permutation(d.flatten())
shuffled_ima = shuffled_ima.reshape(ima.shape)

display_hist(shuffled_ima,255)
```



Since the entropy is computed on the graylevel **distribution**, nothing changes. So, from the entropy point of view, information carried by the above image is the same as the one of the cameraman, something about the way we define the information is wrong here...

In fact, when we shuffle the pixels of the image, we lose the localisation information, pixels that were located close to each other are splitted.

In order to take this collocality between pixels, one can use the coocurrence matrix.

In [75]:

```
#coocurrence matrix
```

```
def cooc(im,dx,dy):
    rim = np.roll(im,dy, axis=0)
    rim = np.roll(rim,dx, axis=1)
    G1 = im.flatten()
    G2 = rim.flatten()

    histo2D = np.zeros((256,256))

    for g1,g2 in zip(G1,G2):
        histo2D[g1,g2] = histo2D[g1,g2]+1

    return histo2D, rim

dx = -10
dy = -5
histo2D,rim = cooc(ima,dx=dx,dy=dy)
plt.figure(figsize=[7,7])
plt.imshow(rim,cmap=cm.gray,interpolation='nearest');
plt.arrow(200,50,-dx,-dy,color='b',head_width=5);
```



For an image I one define a coocurance matrix by:

$$C_{i,j}^{\Delta x, \Delta y} = \sum_{p=1}^n \sum_{q=1}^m \begin{cases} 1, & \text{if } I(p, q) = i \text{ and } I(p + \Delta x, q + \Delta y) = j \\ 0, & \text{otherwise} \end{cases}$$

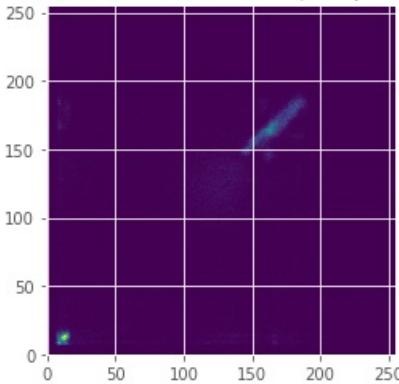
in other words, this matrix count the number of pixels having a gray level i and its neighbor, defined by the $(\Delta x, \Delta y)$ translation, has a gray level j .

Usually close pixels share a similar gray level, for small $(\Delta x, \Delta y)$ the matrix is close to the diagonal.

In [76]:

```
plt.imshow(histo2D)
plt.ylim([0,255])
plt.title('Coocurence matrix $\Delta = (%d,%d)$' %(dx,dy));
```

Coocurence matrix $\Delta = (-10, -5)$



Exercice:

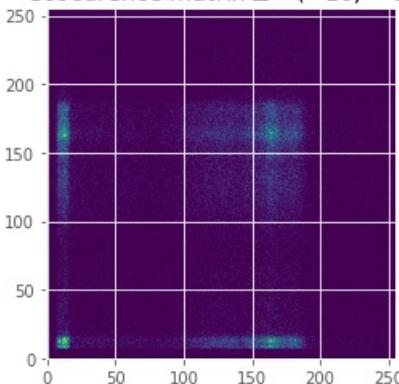
1. what contains the diagonal of a coocurence matrix with $(\Delta x, \Delta y) = (0, 0)$?

What about the impact of our previous shuffling on the coocurence matrix ?

In [77]:

```
histo2D,_ = cooc(shuffled_ima,dx=dx,dy=dy)
plt.imshow(histo2D)
plt.ylim([0,255])
plt.title('Coocurence matrix $\Delta = (%d,%d)$' %(dx,dy));
```

Coocurence matrix $\Delta = (-10, -5)$



Exercice:

1. evaluate the entropy of the coocurrence matrices of the cameraman image
2. do the same with its shuffled version.

see also:

- cooccurrence matrix [IPAMV \(../00-Preface/06-References.ipynb#\[IPAMV\]\)](#) p45

Compression

The aim of the compression can be to limit the size of the image in memory, for the storage, for the transmission. Depending on its use, compression can be lossless, i.e. the data remains untouched after decompression, or, the compression can be lossy if one tolerate some data degradation to have better compression. Image compression used redundancy present in the image to limit the actual number of bits to be used, for example continuity in an image, or in a sequence (video compression).

Lossless compression

We give here two examples of compression that can be used to diminish the amount of bits to be used to store/transmit a data without changing its content.

Huffman encoding

This method compresses a sequence of symbols (i.e. one can think of pixel graylevels) such that one takes advantage of the unequal distribution of the occurrence of symbols. Indeed if a symbol is very common, let's a gray level often used, it could be interesting to store its value with a very short (in terms of bits) symbol. On the contrary, a very rare symbol could use more bits without penalizing the average total message length.

In the given example hereunder, four symbols have a variable frequency of occurrence, from a1 the most frequent to a4 the least one.

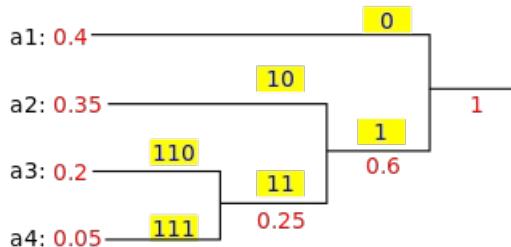
With the Huffman method we first associate the shortest code '0' to the most frequent symbol 'a1', all the other symbols will be coded by a word beginning by '1'.

The same process is done recursively for the remaining symbols. Finally the rarest symbol will be coded by 3 bits, whereas normally only 2 bits are sufficient for coding 1 symbol out of 4. But statistically, because a very short word is used for the most frequent symbol, the length of the total message will be shorter.

In [78]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/7/74/Huffman_coding_example.svg/320px-Huffman_coding_example.svg.png')
```

Out[78]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:Huffman_coding_example.svg\)](https://commons.wikimedia.org/wiki/File:Huffman_coding_example.svg)

Exercice:

1. apply the Huffman compression to a grayscale 8-bits image.

see also:

- Huffman encoding [DIPM \(../00-Preface/06-References.ipynb#\[DIPM\]\)](#) pp261-269

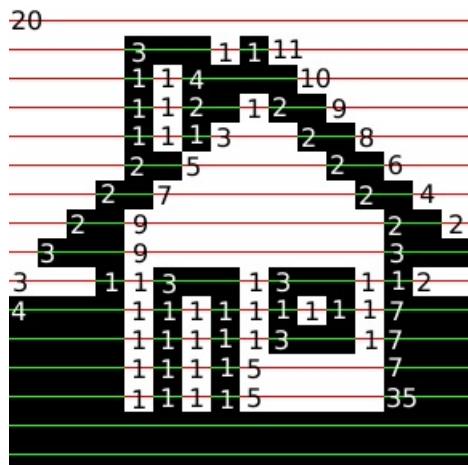
Run length encoding

When image is binary, one can group consecutive similar pixels (0 or 1) and encode them by saving the length of the sequence only. Of course the encoded image is stored in integer, big enough to store the longest segment (35 in the given example). If the image has a lot of continuous parts, this compression can be very efficient.

In [79]:

```
Image('http://pippin.gimp.org/image_processing/images/rle.png')
```

Out[79]:



[image source \(http://pippin.gimp.org/image_processing/images/rle.png\)](http://pippin.gimp.org/image_processing/images/rle.png)

Exercice:

1. propose a method for labelled image

see also:

- Run-length encoding [IPH \(./00-Preface/06-References.ipynb#\[IPH\]\)](#) p396

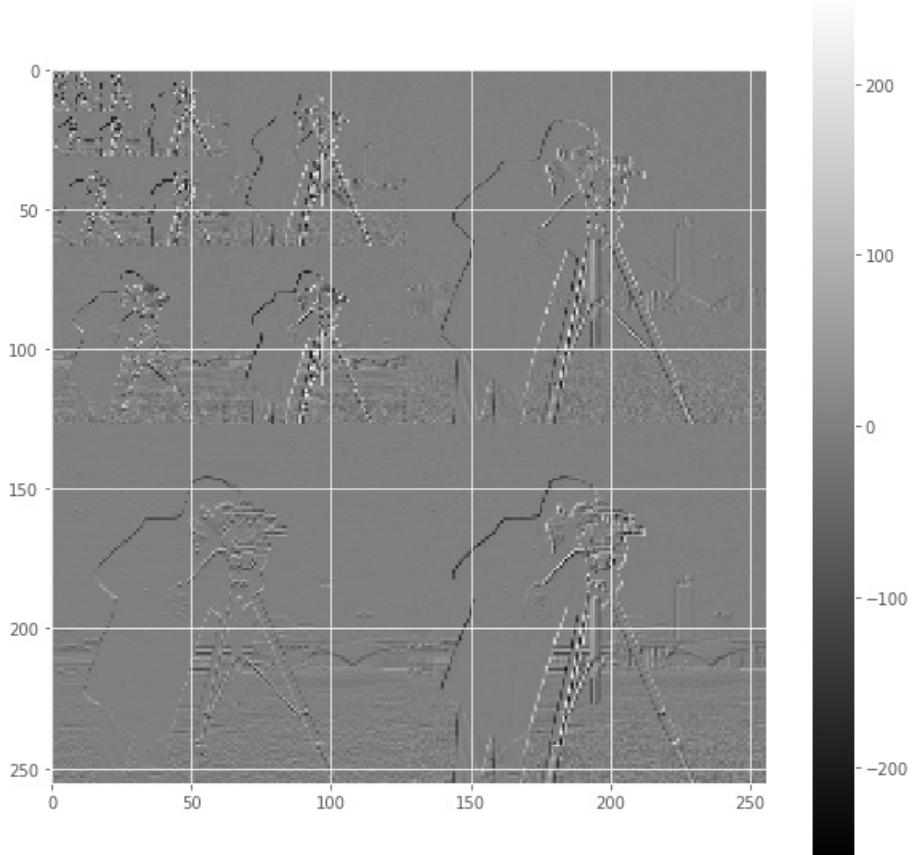
Image pyramid

In [80]:

```
def split(im):
    if im.shape[0] > 1:
        a = im[0:-1:2,0:-1:2]
        b = im[0:-1:2,1::2]-a
        c = im[1::2,0:-1:2]-a
        d = im[1::2,1::2]-a
        R = np.vstack((np.hstack((split(a),b)),np.hstack((c,d))))
    else:
        R = im
    return R

im = camera()[:,::2,::2].astype(np.int)
s = split(im)

fig = plt.figure(figsize=[10,10])
plt.imshow(s,interpolation='nearest',cmap=cm.gray,vmin=-255, vmax=255)
plt.colorbar();
```



Exercices :

1. rebuild the original image from its compressed pyramid.
2. compute image entropy of the compressed pyramid

Lossy image compression

For human vision, some details are of less importance(e.g. the precise color of a small object in a large scene), therefore lossy compression has been developed that enable very high compression ratio with relatively few perceptual loss. For example the JPEG compression can achieve 1:10 to 1:20 compression ratio while keeping a good image quality. This compression ratio can be much higher if we can tolerate some minor image degradation.

One must pay attention to the fact that, while these compressions are done for visual perception, the result of the compression can be disastrous for machine vision.

see also:

- image pyramid (compression) [DIPM \(../00-Preface/06-References.ipynb#\[DIPM\]\)](#) p304
- lossy image compression (jpeg/dct) [IPH \(../00-Preface/06-References.ipynb#\[IPH\]\)](#) p126

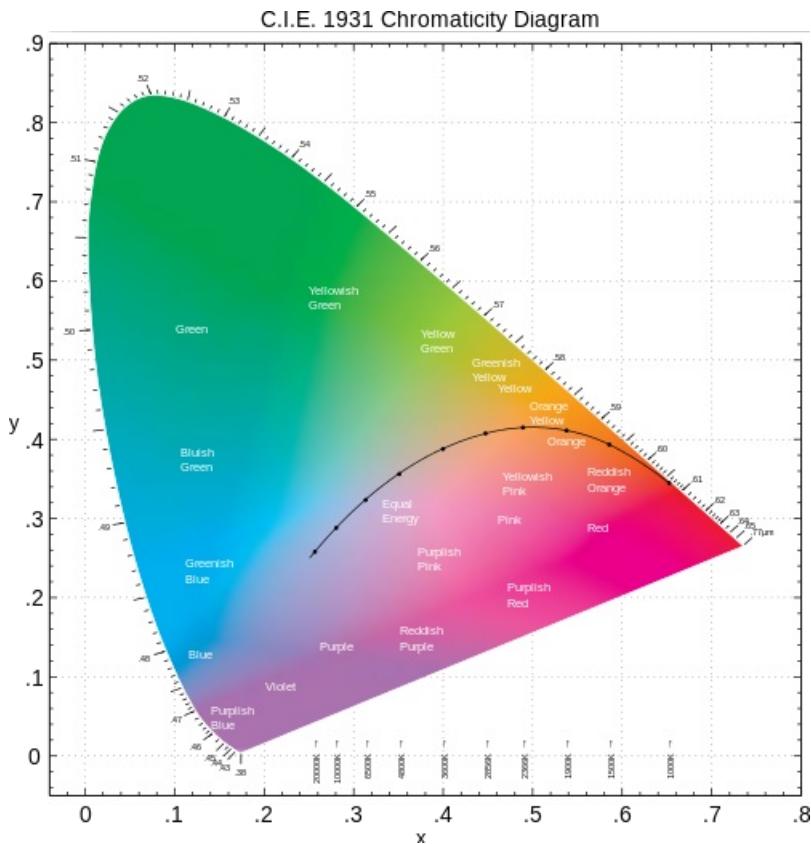
Color representation

Color vision is due to the ability of the eye to discriminate three part of the spectrum thanks to specialized light sensitive cells, the cones. Color image are based on the same principle, the image is composed of the contributions of the light in several spectral bands. For usual picture these bands are red, green and blue. For the human vision, every color is a combination of these (additive) colors.

In [81]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/5/5f/CIE-1931_diagram_in_LAB_space.svg/500px-CIE-1931_diagram_in_LAB_space.svg.png')
```

Out[81]:



wikimedia commons (https://commons.wikimedia.org/wiki/File:CIE-1931_diagram_in_LAB_space.svg)

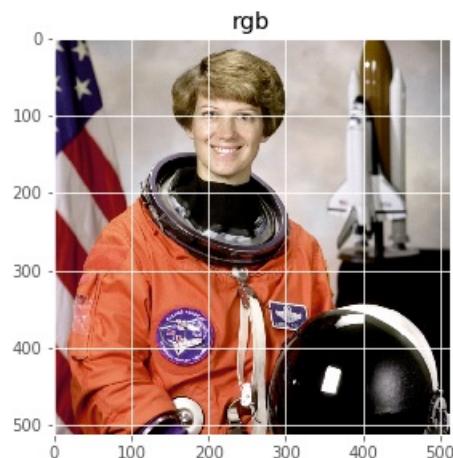
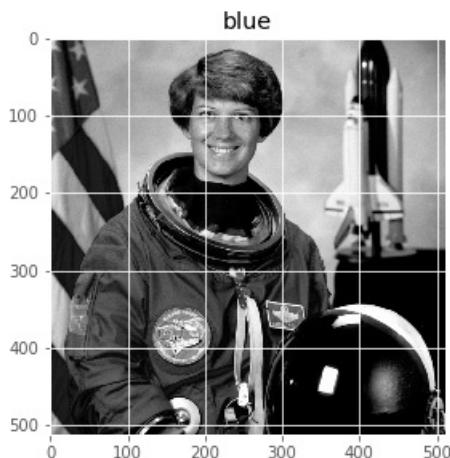
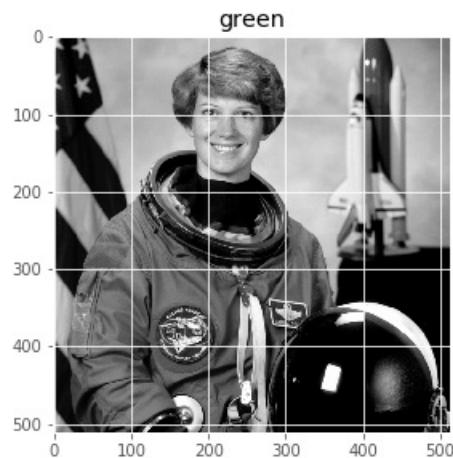
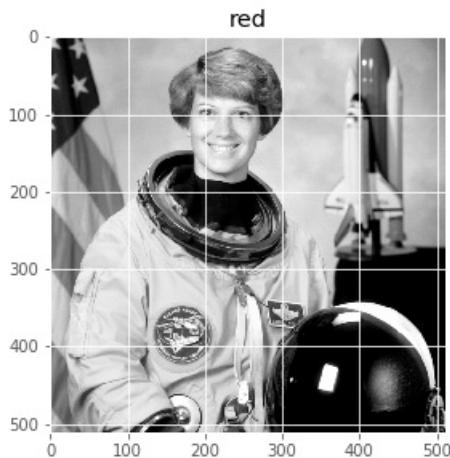
color systems

RGB

In [82]:

```
from mpl_toolkits.mplot3d import Axes3D
from skimage import color
from skimage.data import astronaut,immunohistochemistry

rgb = astronaut()
plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(rgb[:, :, 0], cmap=cm.gray)
plt.title('red')
plt.subplot(2,2,2)
plt.imshow(rgb[:, :, 1], cmap=cm.gray)
plt.title('green')
plt.subplot(2,2,3)
plt.imshow(rgb[:, :, 2], cmap=cm.gray)
plt.title('blue')
plt.subplot(2,2,4)
plt.imshow(rgb, cmap=cm.gray)
plt.title('rgb');
```



In [83]:

```
rgb = immunohistochemistry()

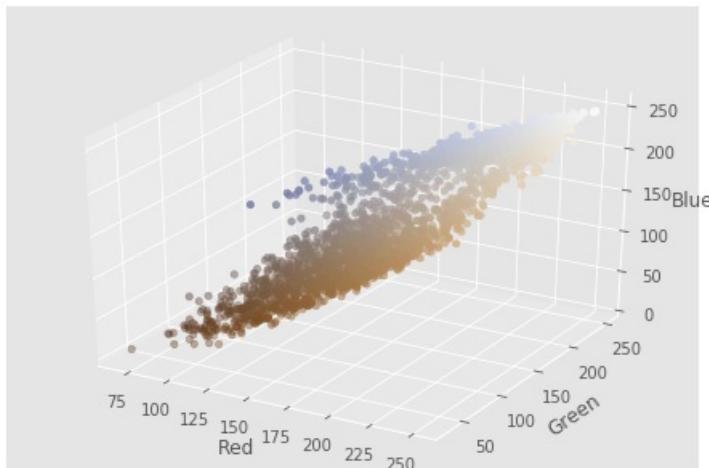
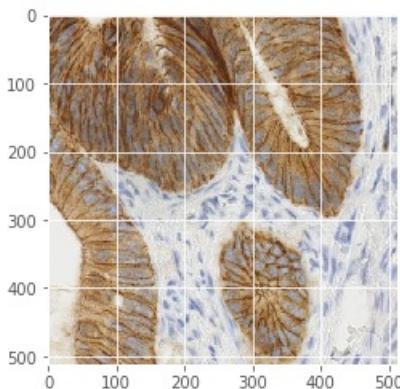
fig = plt.figure(0)
plt.imshow(rgb,origin='lower')
plt.gca().invert_yaxis()

#subsample
print(rgb.shape)
rgb = rgb[::10,::10,:]

r = rgb[:, :, 0].flatten()
g = rgb[:, :, 1].flatten()
b = rgb[:, :, 2].flatten()

fig = plt.figure()
ax = Axes3D(fig)
col = np.vstack((r,g,b)).T/255.0
ax.scatter(r,g,b,c=col)
ax.set_xlabel('Red')
ax.set_ylabel('Green')
ax.set_zlabel('Blue');
```

(512, 512, 3)

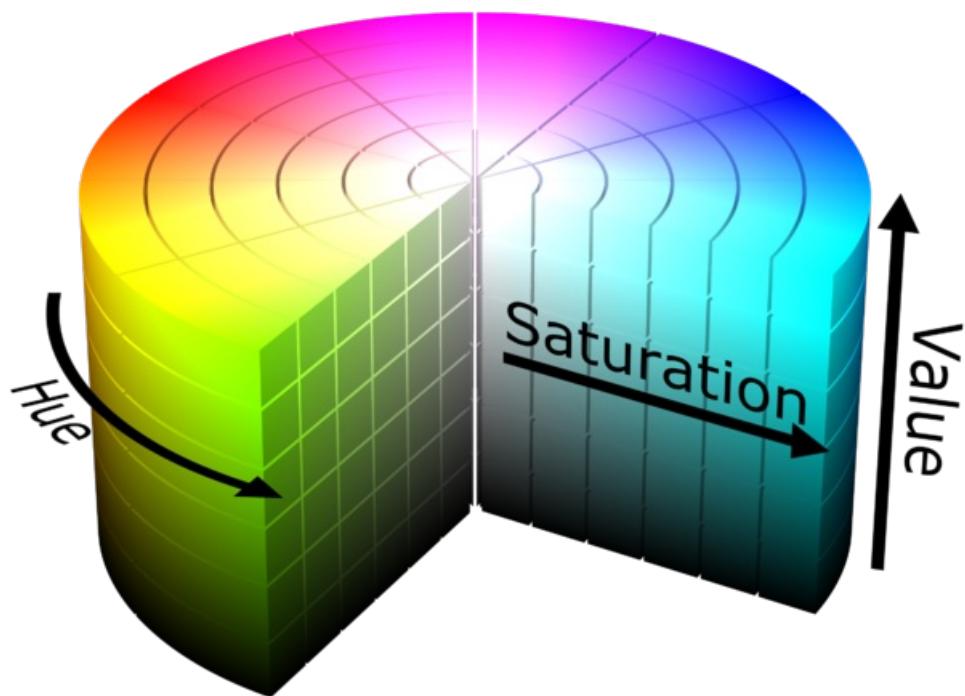


HSV

In [84]:

```
Image('https://upload.wikimedia.org/wikipedia/commons/thumb/4/4e/HSV_color_solid_cylinder.png/640px-HSV_color_solid_cylinder.png')
```

Out[84]:



[wikimedia commons \(https://commons.wikimedia.org/wiki/File:HSV_color_solid_cylinder_alpha_lowgamma.png\)](https://commons.wikimedia.org/wiki/File:HSV_color_solid_cylinder_alpha_lowgamma.png)

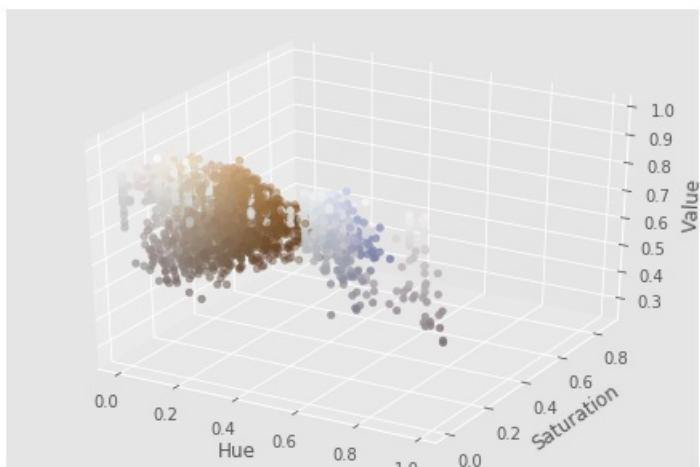
In [85]:

```
hsv = color.rgb2hsv(rgb)

fig = plt.figure()
ax = Axes3D(fig)
h = hsv[:, :, 0].flatten()
s = hsv[:, :, 1].flatten()
v = hsv[:, :, 2].flatten()
ax.scatter(h, s, v, c=col)
ax.set_xlabel('Hue')
ax.set_ylabel('Saturation')
ax.set_zlabel('Value'),
```

Out[85]:

```
(Text(0.5, 0, 'Value')),)
```

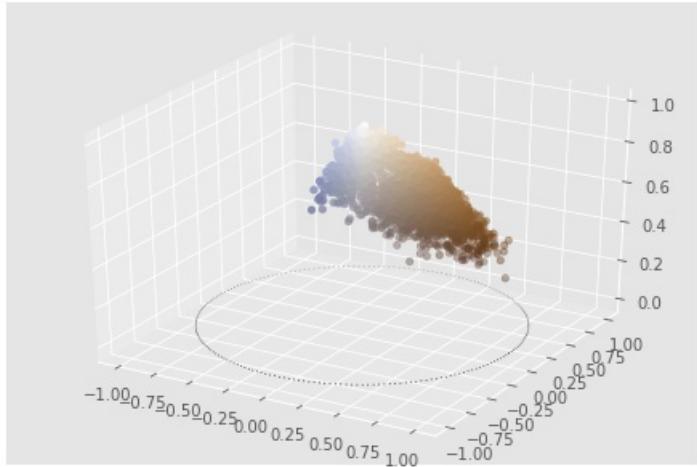


XYZ

In [86]:

```
xyz = color.rgb2xyz(rgb)
fig = plt.figure()
ax = Axes3D(fig)
x = s*np.cos(h*2*np.pi)
y = s*np.sin(h*2*np.pi)
z = v
ax.scatter(x,y,z,c=col)

th = np.linspace(0,6.28,200)
ax.scatter(np.cos(th),np.sin(th),0,c='k',s=.1);
```



color unmixing

to be continued...

In [1]:

```
%matplotlib inline
from IPython.display import HTML,Image,SVG,YouTubeVideo
```

Histogram

In [2]:

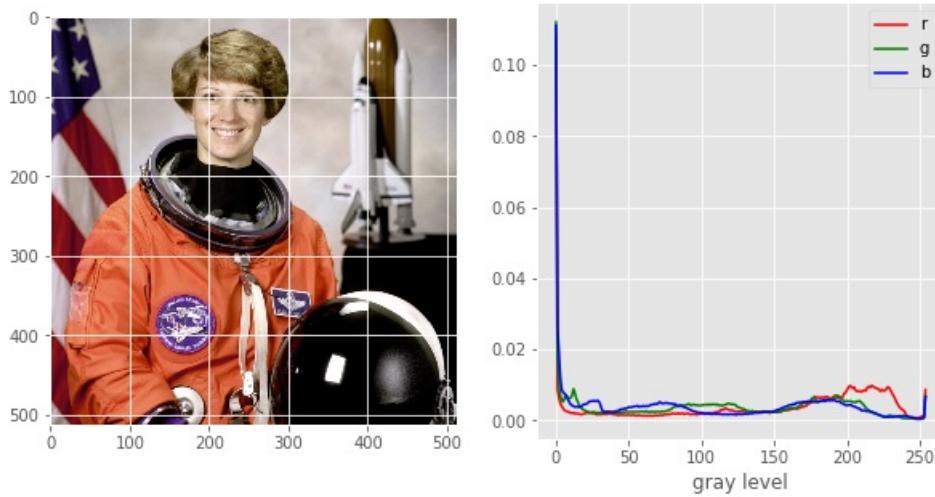
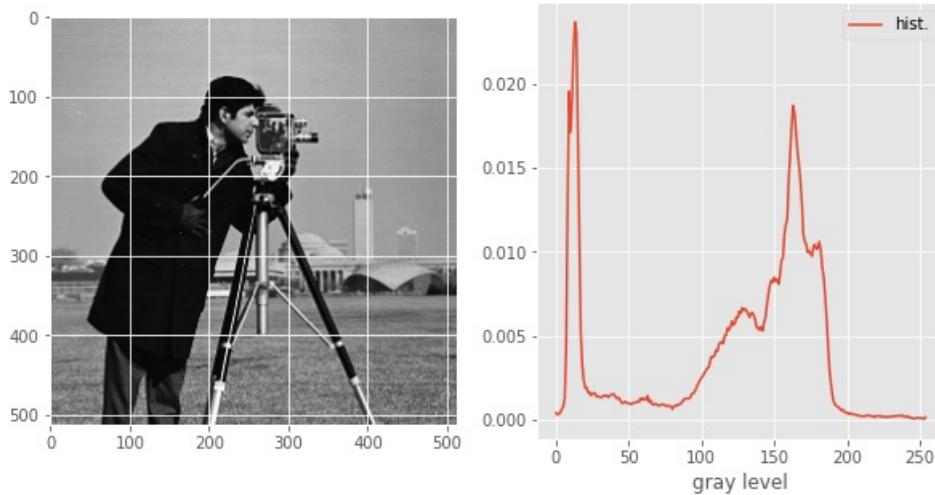
```
import numpy as np
import matplotlib.pyplot as plt
from matplotlib import cm
from skimage.data import camera,astronaut
plt.style.use('ggplot')
```

In [3]:

```
def norm_hist(ima):
    hist,bins = np.histogram(ima.flatten(),range(256)) # histogram is computed on a 1D distribution --> flatten()
    return 1.*hist/np.sum(hist) # normalized histogram

def display_hist(ima,vmin=None,vmax=None):
    plt.figure(figsize=[10,5])
    if ima.ndim == 2:
        nh = norm_hist(ima)
    else:
        nh_r = norm_hist(ima[:, :, 0])
        nh_g = norm_hist(ima[:, :, 1])
        nh_b = norm_hist(ima[:, :, 2])
    # display the results
    plt.subplot(1,2,1)
    plt.imshow(ima,cmap=cm.gray,vmin=vmin,vmax=vmax)
    plt.subplot(1,2,2)
    if ima.ndim == 2:
        plt.plot(nh,label='hist.')
    else:
        plt.plot(nh_r,color='r',label='r')
        plt.plot(nh_g,color='g',label='g')
        plt.plot(nh_b,color='b',label='b')
    plt.legend()
    plt.xlabel('gray level');

display_hist(camera())
display_hist(astronaut())
```

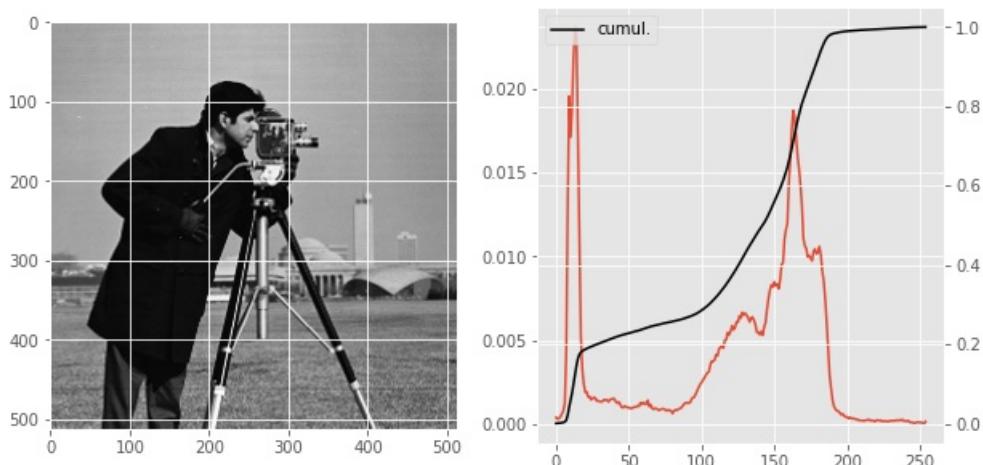


In [4]:

```
def display_hist2(ima):
    nh = norm_hist(ima)
    cumul_hist = np.cumsum(nh)

    plt.figure(figsize=[10,5])
    plt.subplot(1,2,1)
    plt.imshow(ima,cmap=cm.gray)
    ax1 = plt.subplot(1,2,2)
    plt.plot(nh)
    ax2 = ax1.twinx()
    plt.plot(cumul_hist,label='cumul.',color='k')
    plt.legend()

display_hist2(camera())
```



Look-Up-Table

Example are given for 8-bits images but can of course be generalized for any kind of integer image, however, due to memory limitation, LUT method will be used only with bit-depth limited images.

In [5]:

```
def apply_lut(ima,lut,vmin=None,vmax=None):
    nh = norm_hist(ima)
    lima = lut[ima]
    nh_lima = norm_hist(lima)

    plt.figure(figsize=[10,5])
    plt.subplot(1,2,1)
    plt.imshow(lima,cmap=cm.gray,vmin=vmin,vmax=vmax)
    ax1 = plt.subplot(1,2,2)
    plt.plot(nh,label='ima')
    plt.plot(nh_lima,label='lut[ima]')
    plt.legend(loc='upper left')
    ax2 = ax1.twinx()
    plt.plot(lut,label='lut',color='k')
    plt.legend()
```

Negative

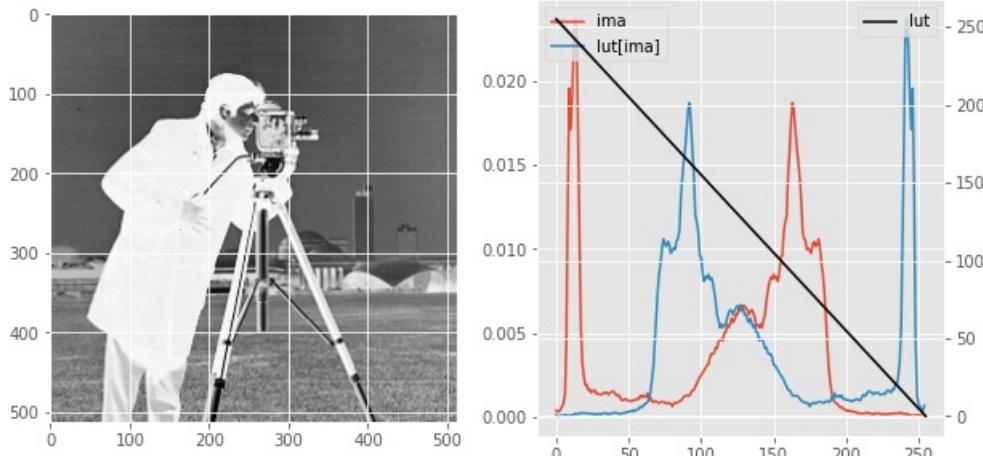
$$g_{out} = 255 - g_{in}$$

To apply this inversion tp the complete image, one use the Look Up Table (LUT) method which consist in pre-computng the transformed levels for all the 255 possible gray level into one vector. Image transformation is then a simple vector addressing.

```
lut = np.arange(255,-1,-1)
g_out = lut[g_in]
```

In [6]:

```
# LUT inversion
ima = camera()
lut = np.arange(255,-1,-1)
apply_lut(ima,lut)
```



Threshold

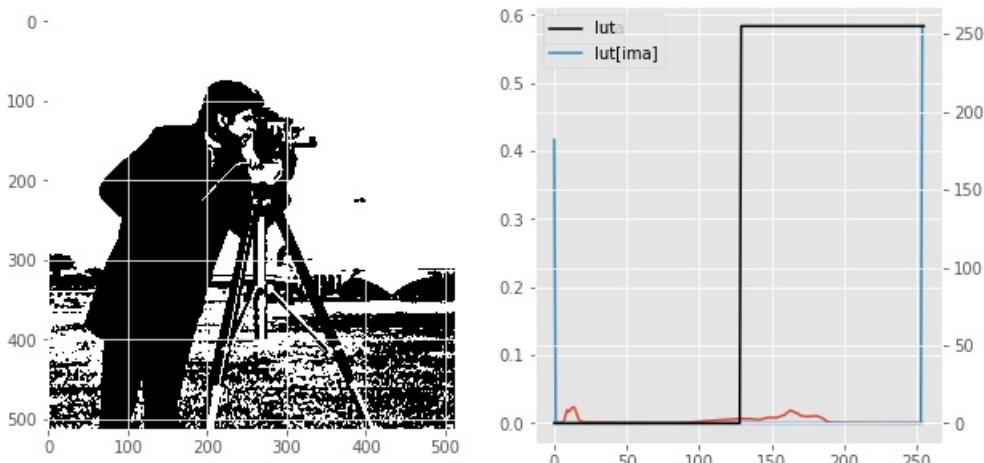
Look up table for a simple threshold is:

$$g_{out} = \begin{cases} 255, & \text{if } g_{in} > th \\ 0, & \text{otherwise} \end{cases}$$

In [7]:

```
def lut_threshold(th):
    lut = np.arange(0,256)
    lut = 255 * (lut > th)
    return lut

apply_lut(im1,lut_threshold(128))
```



Semi-threshold

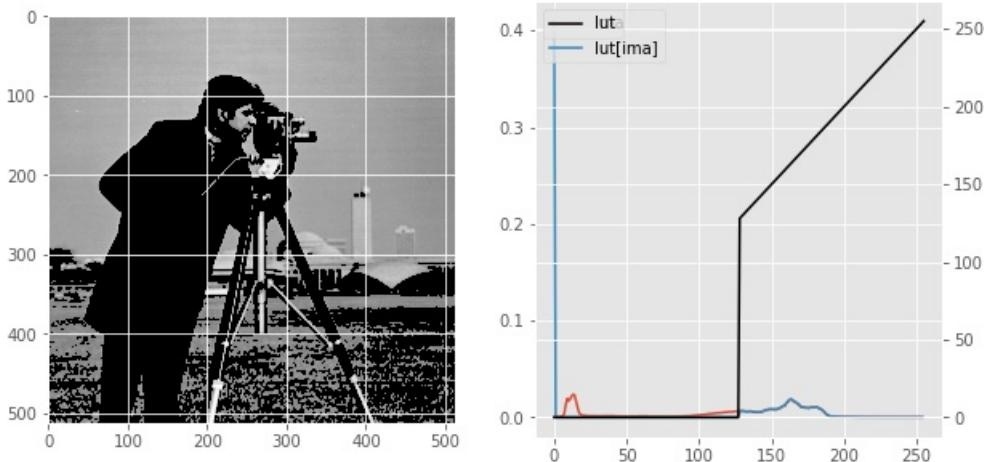
Look up table for a simple threshold is:

$$g_{out} = \begin{cases} g_{in}, & \text{if } g_{in} > th \\ 0, & \text{otherwise} \end{cases}$$

In [8]:

```
def lut_semi_threshold(th):
    lut = np.arange(0,256)
    lut[lut < th] = 0
    return lut

apply_lut(im1,lut_semi_threshold(128))
```



Gamma correction

Gamma transform is used to reinforce contrast of the image, level trasform is given by:

$$g_{out} = A g_{in}^\gamma$$

where

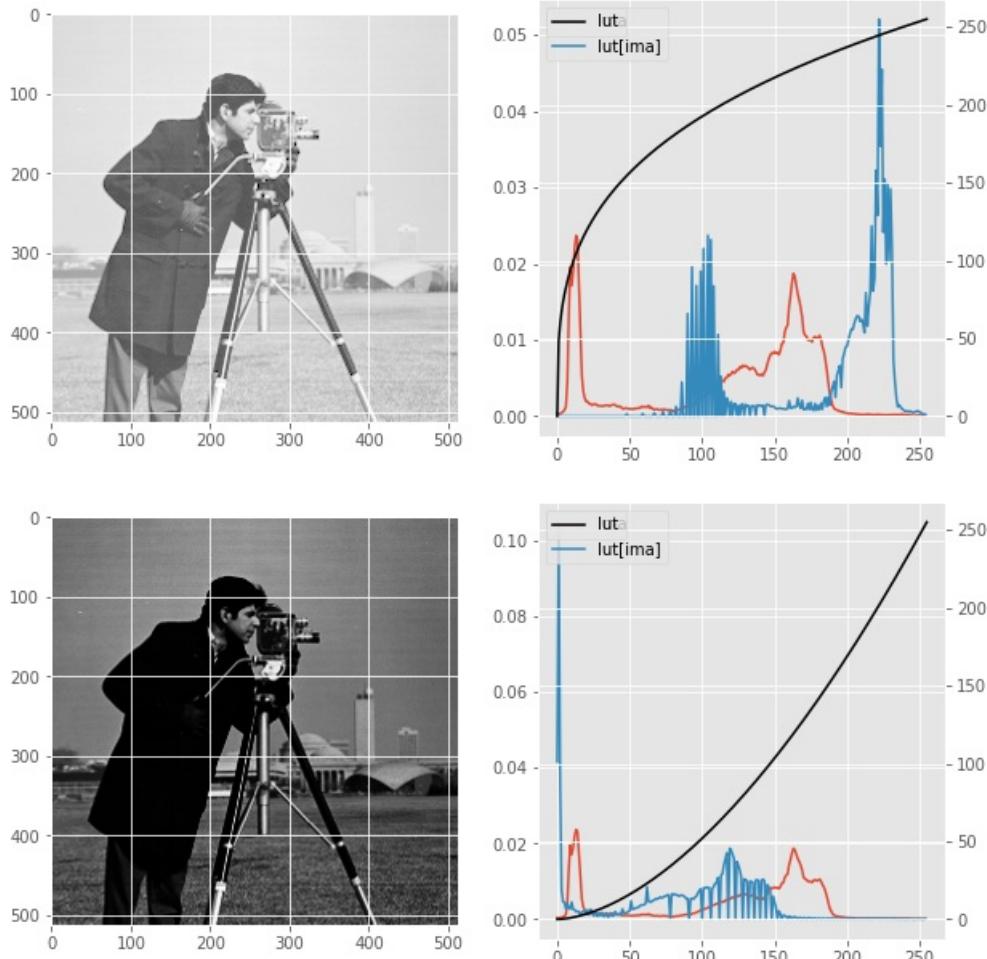
$$A = 255^{(1-\gamma)}$$

if $\gamma < 1$ the low-level are contrasted, reversely if $\gamma > 1$ bright part of the image gains in contrast.

In [9]:

```
def lut_gamma(gamma):
    lut = np.power(np.arange(0,256),gamma) * np.power(255,1-gamma)
    return lut

apply_lut(ima,lut_gamma(.3))
apply_lut(ima,lut_gamma(1.7))
```



Auto-level

Auto-level map the complete image dynamic to the full dynamic scale:

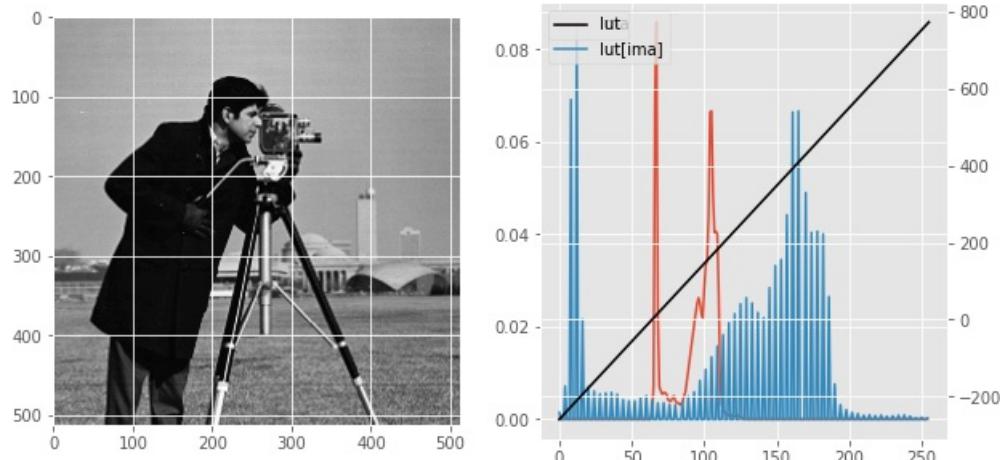
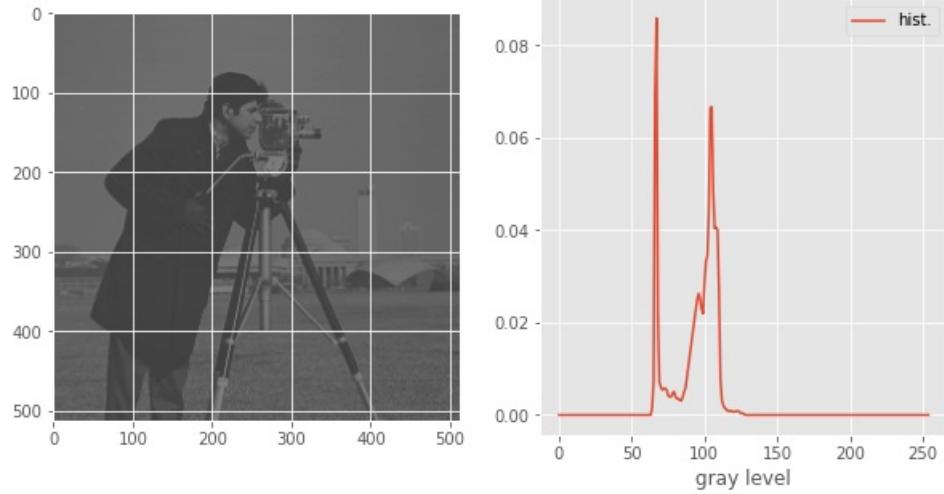
$$g_{out} = 255 \frac{g_{in} - g_{min}}{g_{max} - g_{min}}$$

where g_{min} and g_{max} are respectively minimimal and maximal value present in the image.

In [10]:

```
def lut_autolevel(ima):
    g_min = np.min(ima)
    g_max = np.max(ima)
    lut = 255*(np.arange(0,256)-g_min)/(1.*g_max-g_min)
    return lut

ima=camera()
t_ima = (ima/4+64).astype(np.uint8)
display_hist(t_ima,vmin=0,vmax=255)
apply_lut(t_ima,lut_autolevel(t_ima))
```



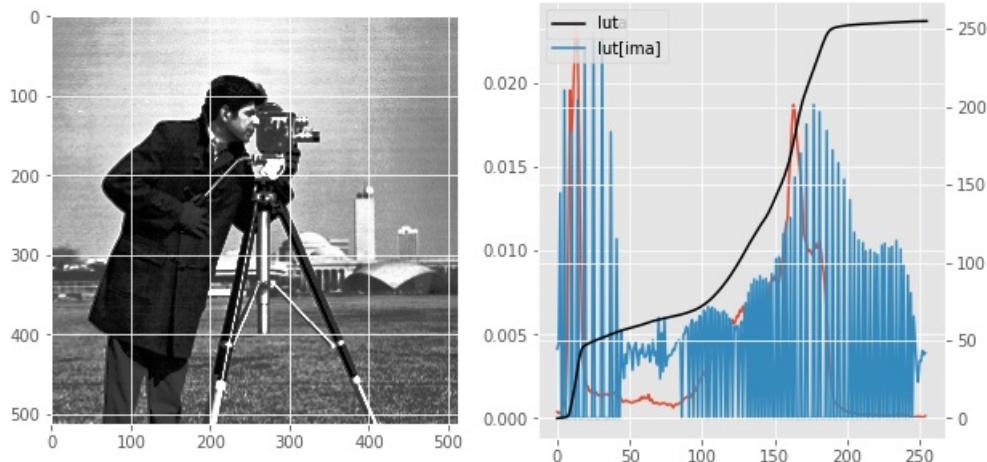
Equalization

One may be interested in using as many gray levels possible for frequent levels, and in grouping rare levels. This is called histogram equalization since, after the operation, the histogram distribution is more equal.

The next figure illustrate equalization done of the cameraman picture.

In [11]:

```
def lut_equalization(ima):
    nh = norm_hist(ima)
    ch = np.append(np.array(0), np.cumsum(nh))
    lut = 255*ch
    return lut
apply_lut(ima,lut_equalization(ima))
```



We can see that levels frequently observed (in the sky) are now more spread (more contrast is visible), the same inside the cameraman where details are now visible (hand). The histogram is not perfectly equal, this is due to the technique used (the look up table), indeed pixel having an equal gray level are transformed similarly, they cannot be separated.

If we look to the code used to achieve the equalization, we see that we simply used, as look up table, the summed histogram !

Here is the justification of that:

- the gray level(arbitrarily set in $[0, 1]$) probability is given by:

$$p_I(r) = \frac{n_r}{n} \quad 0 \leq r \leq 1$$

where n_r is the number of pixels having the value r and n the total number of image pixels.

- let's consider a transform T that maps graylevels r to graylevel s , $T(r)$ is considered as monotonically increasing on $0 \leq r \leq 1$.

$$\begin{aligned} s &= T(r) \\ 0 &\leq T(r) \leq 1 \\ r &= T^{-1}(s) \end{aligned}$$

We also assume that $T^{-1}(s)$ is monotonically increasing on $0 \leq s \leq 1$ and bounded to $[0, 1]$.

- from probability theory, the probability density function of the transformed gray level is:

$$p_s(s) = \left[p_I(r) \frac{dr}{ds} \right]_{r=T^{-1}(s)}$$

- if we consider the following transform function:

$$s = T(r) = \int_0^r p_I(w) dw \quad 0 \leq r \leq 1$$

- then

$$\frac{ds}{dr} = p_I(r)$$

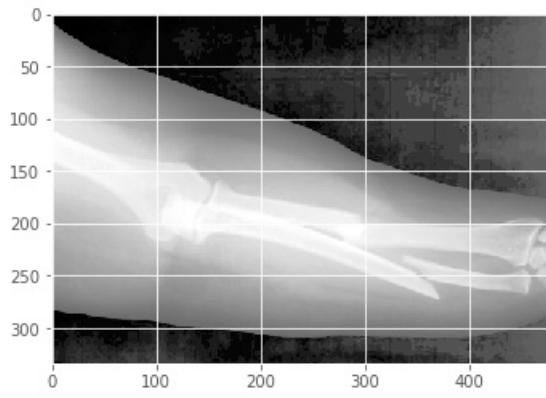
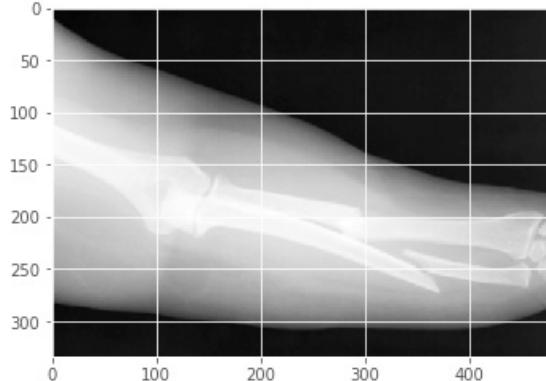
- we can substitute this fraction in the previous equation:

$$\begin{aligned} p_s(s) &= \left[p_I(r) \frac{1}{p_I(r)} \right]_{r=T^{-1}(s)} \\ &= [1]_{r=T^{-1}(s)} \\ &= 1 \quad 0 \leq s \leq 1 \end{aligned}$$

which is uniform on the interval.

In [12]:

```
#other example
from skimage.io import imread
ima = imread('http://homepages.ulb.ac.be/~odebeir/data/bones.png')
lut = lut_equalization(ima)
plt.figure()
plt.imshow(ima,cmap=cm.gray)
plt.figure()
plt.imshow(lut[ima],cmap=cm.gray);
```



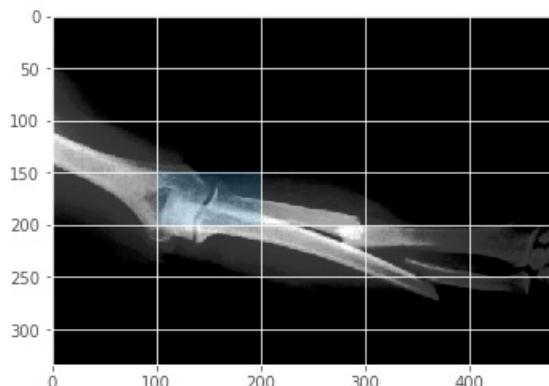
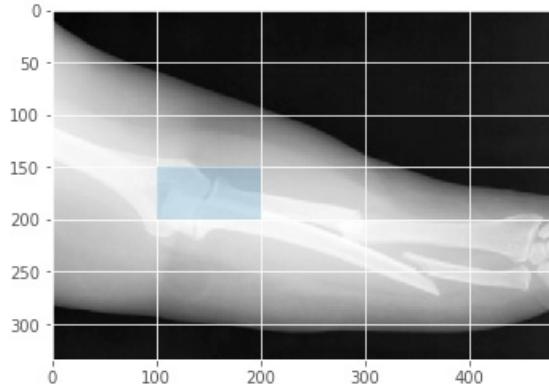
if we need to increase the contrast in a certain part of the image, equalization LUT may be restricted to a certain area:

In [13]:

```
roi = [(100,150),100,50]
sample = ima[roi[0][1]:roi[0][1]+roi[2],roi[0][0]:roi[0][0]+roi[1]]
lut = lut_equalization(sample)

plt.figure()
plt.imshow(ima,cmap=cm.gray)
rect = plt.Rectangle(*roi, facecolor=None, alpha=.25)
plt.gca().add_patch(rect)

plt.figure()
plt.imshow(lut[ima],cmap=cm.gray);
rect = plt.Rectangle(*roi, facecolor=None, alpha=.25)
plt.gca().add_patch(rect);
```



see also:

- histogram based methods [IPAMV \(../00-Preface/06-References.ipynb#\[IPAMV\]\)](#) pp58-61

In []:

In [49]:

```
%matplotlib inline
import sys
sys.path.insert(0,'..')
from IPython.display import HTML,Image,SVG,YouTubeVideo
from helpers import header,compare
```

Linear filtering

In the previous chapter, the processed value of a pixel was only a function of its original value. Here the processed value will also take into account the surrounding pixels as well. We speak about pixel *neighborhood*.

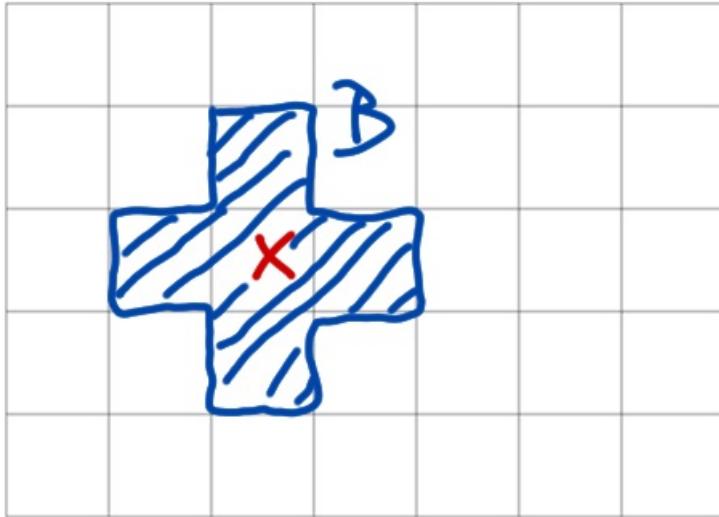
Neighborhood

Pixel neighborhood is defined by a binary structuring element B having an origin.

In [50]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b.png')
```

Out[50]:

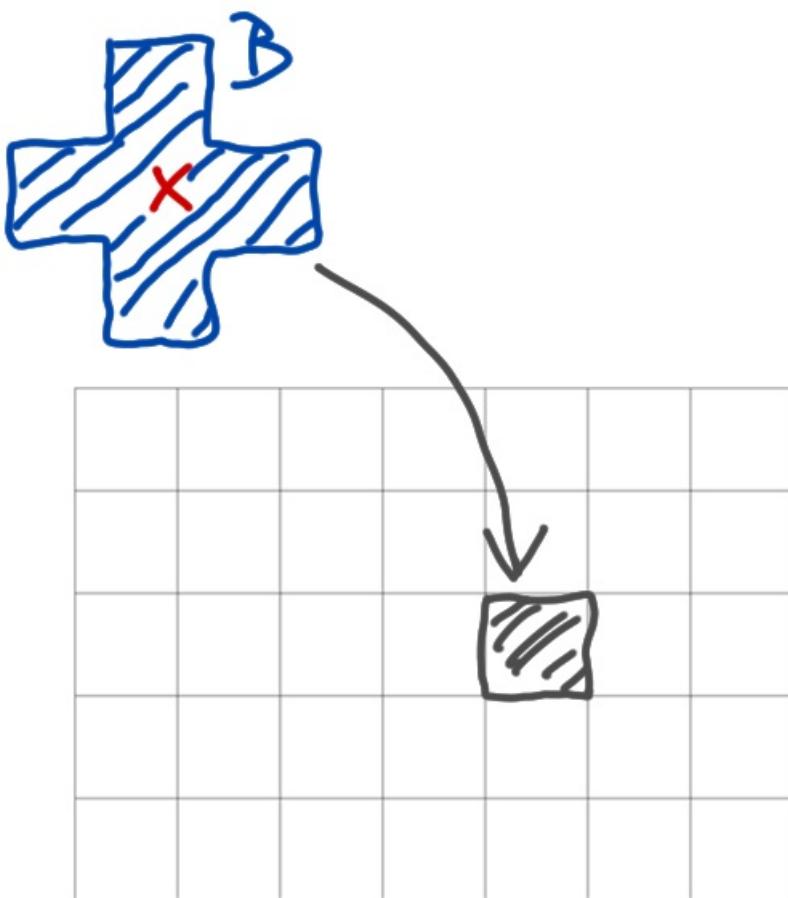


For a specific pixel (the gray one below), the neighborhood defined by B ...

In [51]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b1.png')
```

Out[51]:

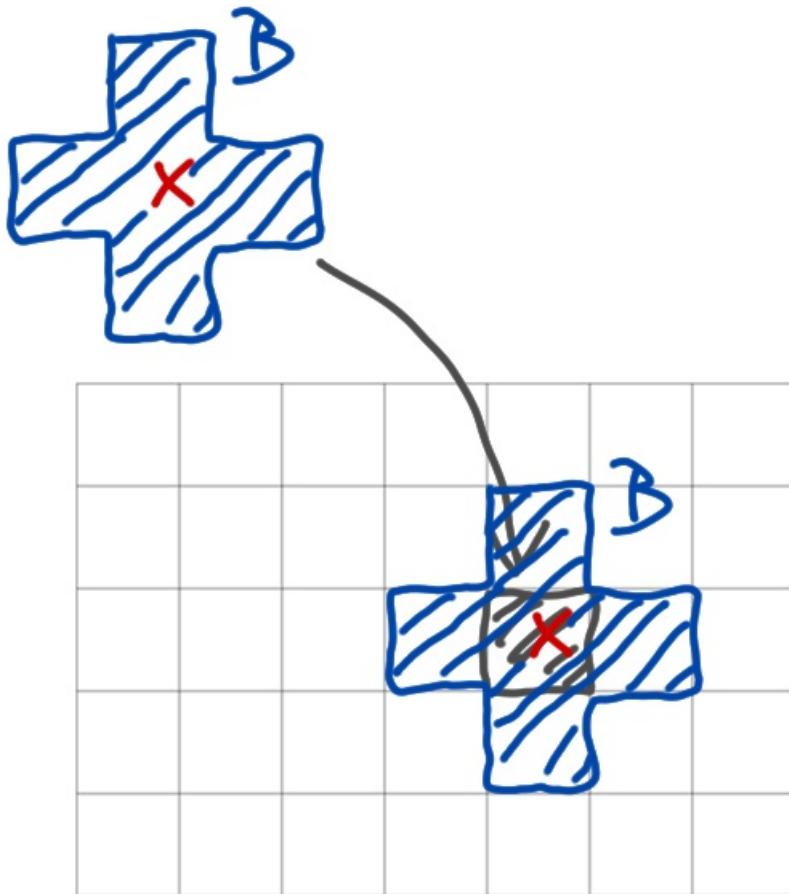


is the ensemble of pixels belonging to B when its origin is moved on the pixel of interest.

In [52]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b2.png')
```

Out[52]:



Structuring elements can have any size and shape. Often the size of the structuring element will directly impact the process and also the processing time.

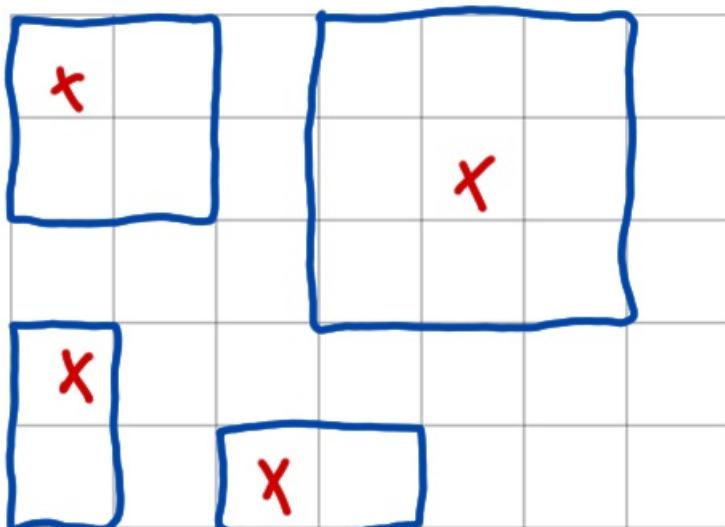
The previous element is known as the 4-neighbors.

Common structuring element are:

In [53]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b3.png')
```

Out[53]:



the upper right is known as the 8-neighbors, we understand why...

One also use bigger element, such as disk with a bigger radius such as:

In [54]:

```
from skimage.morphology import disk
import numpy as np
import matplotlib.pyplot as plt
from matplotlib import cm
from skimage.data import camera
plt.style.use('ggplot')
```

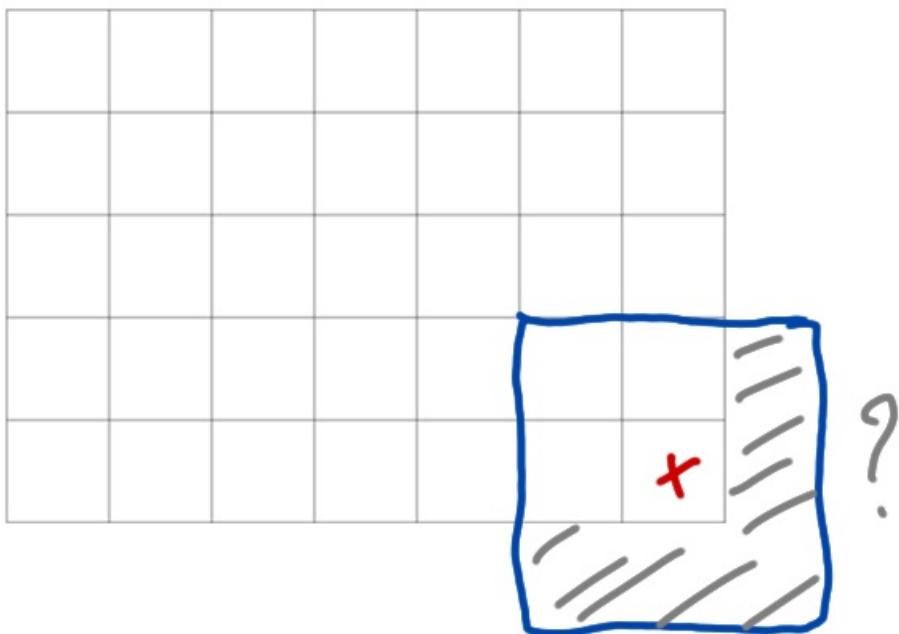
Question:

- what happens close to the borders?

In [55]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b5.png')
```

Out[55]:

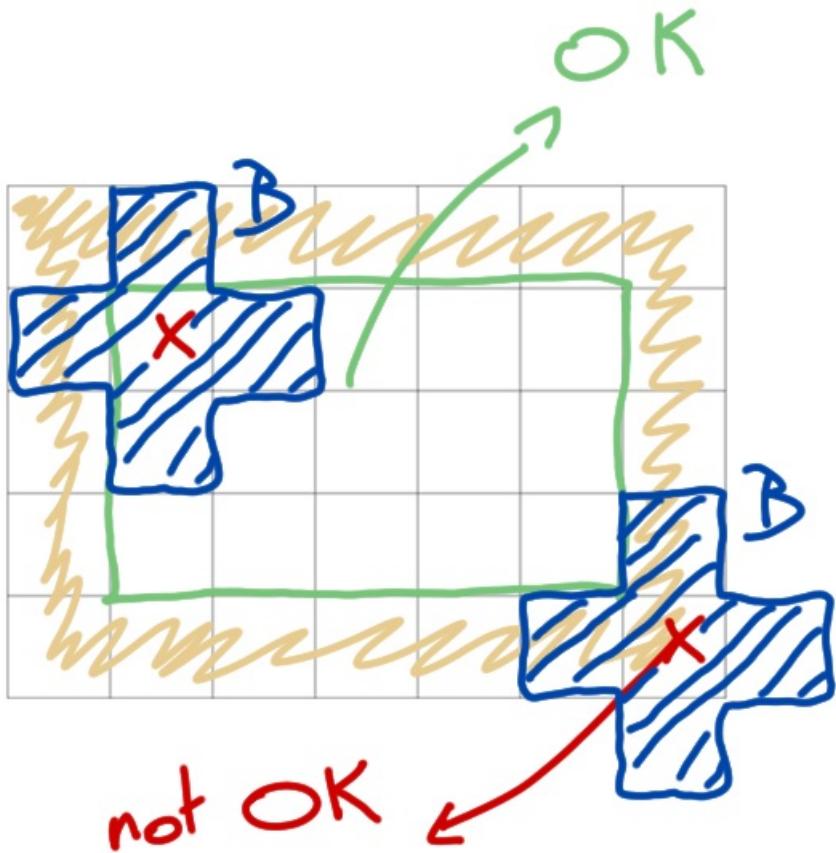


Because the neighborhood is not defined when one pixel of B is outside the image, only the inner part of the image can be processed by a structuring element.

In [56]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/b6.png')
```

Out[56]:



Local processing

A first local process we can do using a structuring element is a weighted sum of the pixels value inside the neighborhood.

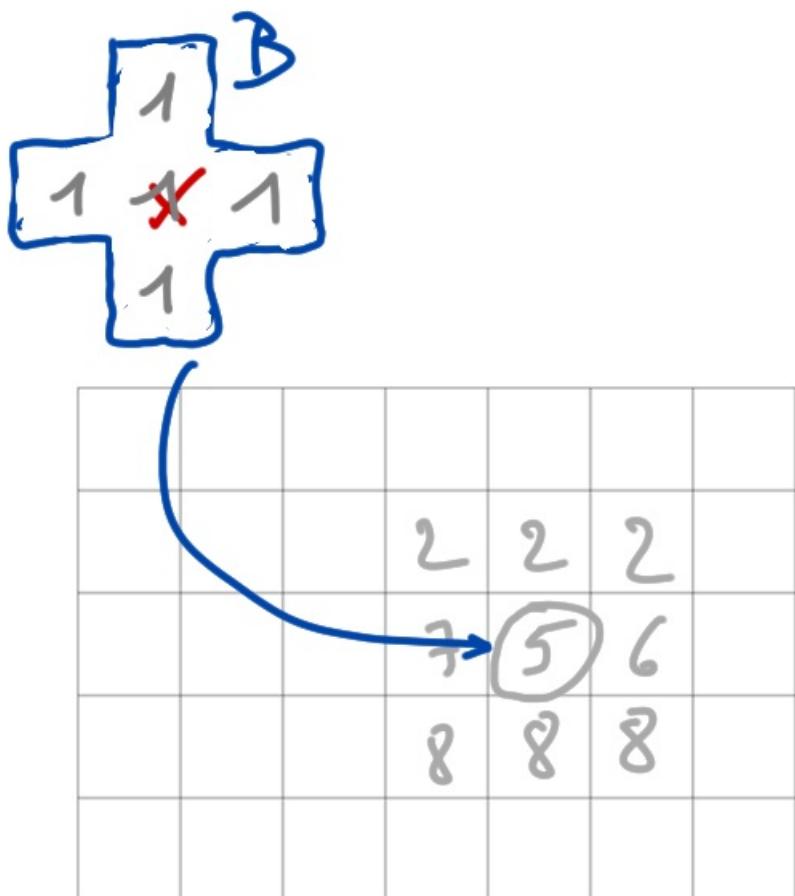
We associate, with each element of the structuring element, a weight.

This is also known as the convolution of the image with the structuring element.

In [57]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/conv1.png')
```

Out[57]:



$$\begin{aligned} & 1 \times 2 + 1 \times 7 + 1 \times 5 + 1 \times 5 \\ & + 1 \times 8 = 28 \end{aligned}$$

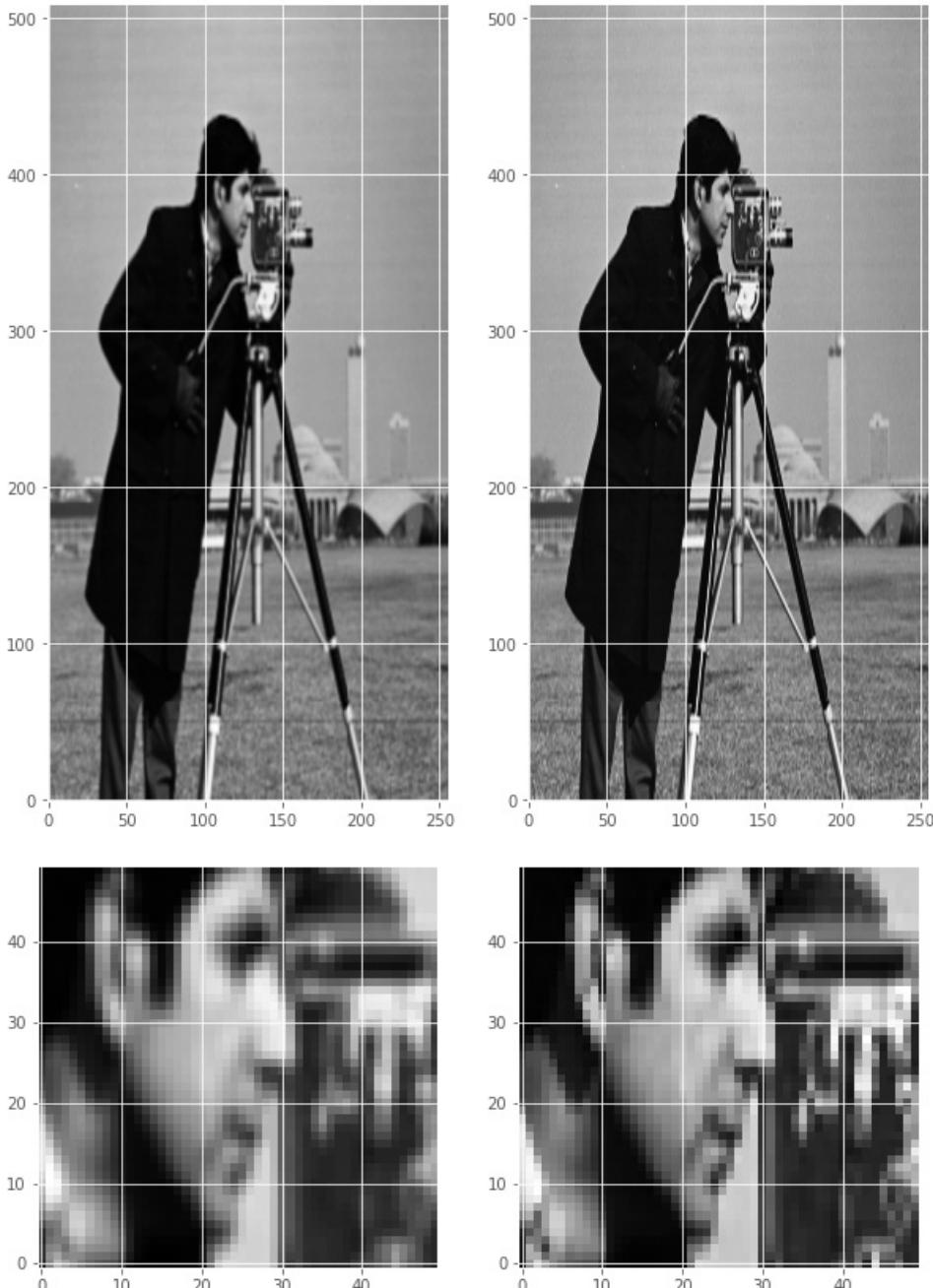
Let's have a look on a real example ... the cameraman, we apply the mean filter which is basically the sum of the pixels inside the structuring element, divided by the number of pixels of the structuring element.

In [58]:

```
from skimage.morphology import disk
from skimage.data import camera
from skimage.filters.rank import mean

g = camera()[-1:2:-1,:,:2]
f = mean(g,disk(1))

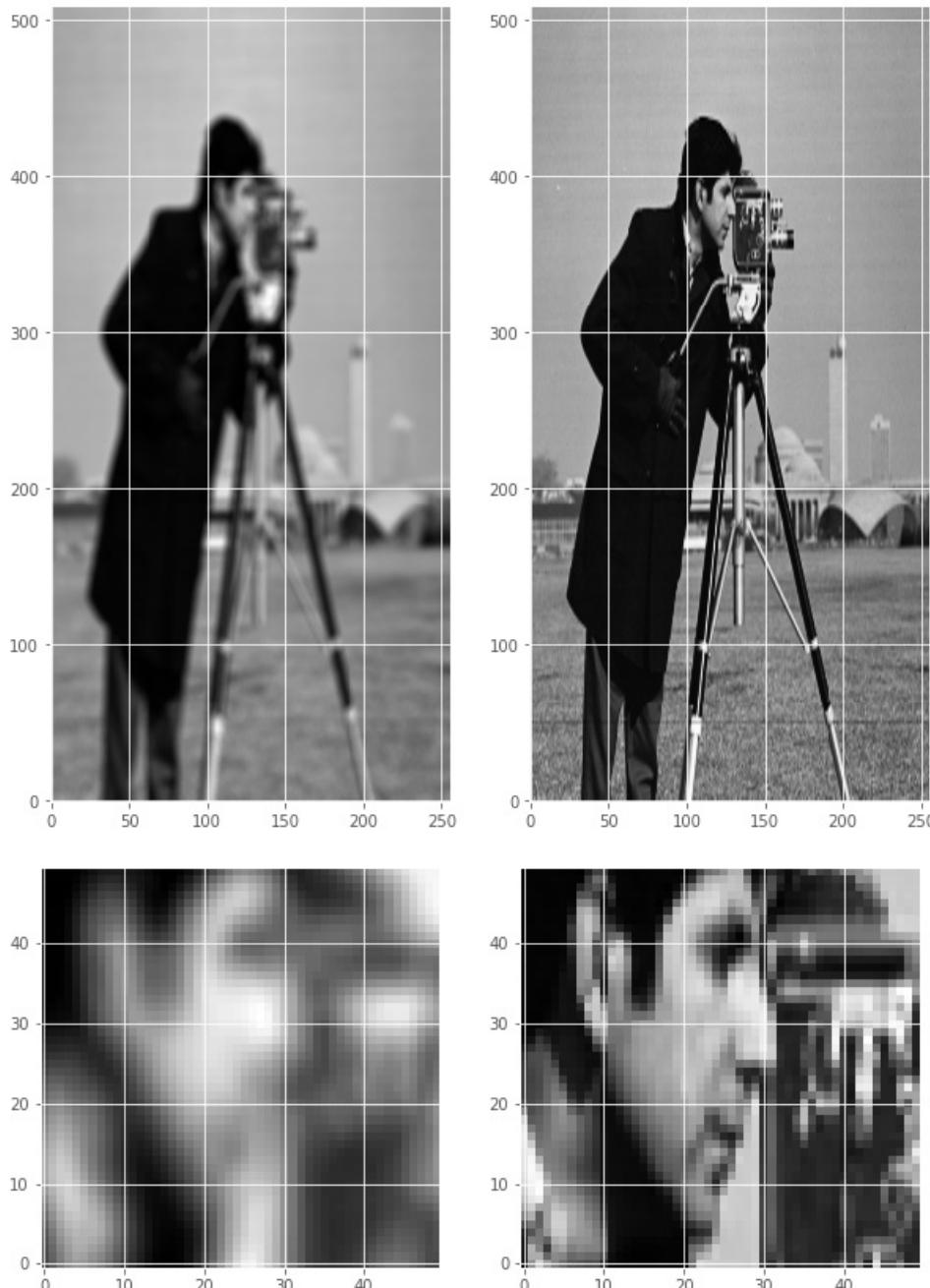
compare(f,g,roi=[350,400,100,150])
```



We observe that noise is smoothed and that small details can disappear. If we increase the radius of the structuring element we see how bigger borders are blurred.

In [59]:

```
f = mean(g,disk(4))
compare(f,g,roi=[350,400,100,150])
```



One can implement the convolution algorithm using embedded loops for a structuring element 3x3:

```
I = [source image]
F = [destination image]
n = 9
for x in 1:sizex-1:
    for y in 1:sizey-1:
        sum = 0
        for i in -1:+1:
            for j in -1:+1:
                sum += I[n+i,m+j]
        F[n,m] = sum/n
```

as the structuring element increases in size, the number of loops is increasing fast...

Questions:

- How the previous example returns a value for pixels close to the border ?
- what should be the destination image type ?

Fourier transform

recall

Fourier analysis allows to play with a signal in the frequency domain.

In 1D signal, the continuous Fourier transform of a function $f(x)$ is given by:

$$F(u) = \int_{-\infty}^{\infty} f(x) e^{-i2\pi ux} dx,$$
$$e^{-i2\pi ux} = \cos(2\pi ux) - i\sin(2\pi ux)$$

the u variable is in the frequency domain,

in general, the Fourier transform of a real signal is complex,

$$F(u) = |F(u)| e^{i\phi(u)}$$

with an amplitude:

$$|F(u)| = [R(u) + I(u)]^{1/2}$$

and a phase:

$$\phi(u) = \tan^{-1}\left(\frac{I(u)}{R(u)}\right)$$

the inverse Fourier transform is given by:

$$f(x) = F^{-1}(F(u)) = \int_{-\infty}^{\infty} F(u) e^{i2\pi ux} du,$$

the magnitude function $|F(u)|^2$ is called the Fourier spectrum or power spectrum of $f(x)$.

Power spectrum for several 1D step.

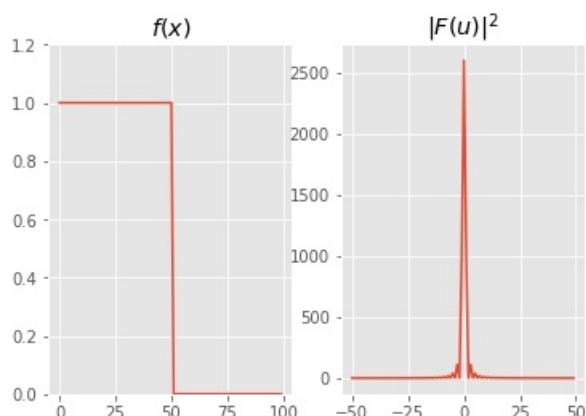
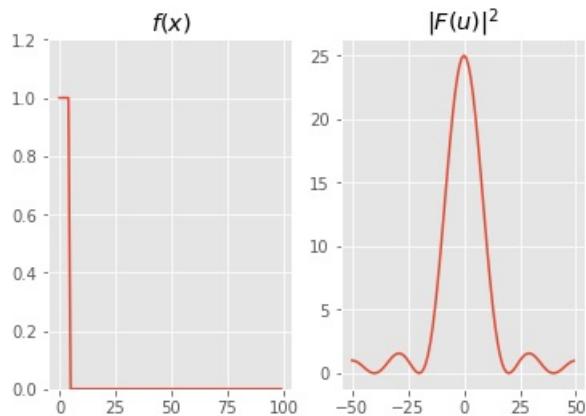
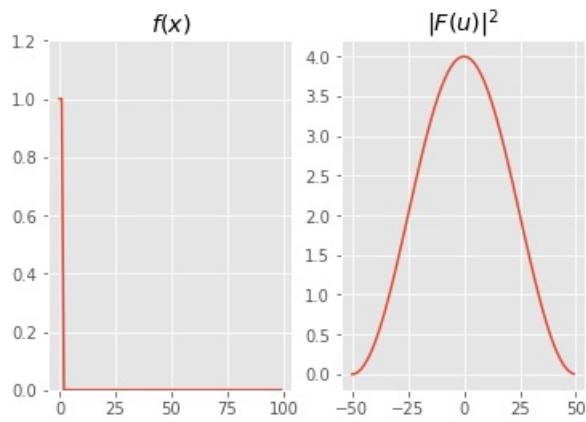
In [60]:

```
import numpy as np
from numpy.fft import fft,fftshift,ifftshift
import matplotlib.pyplot as plt
import matplotlib.cm as cm

def step_power(length):
    nsample = 100
    t = np.arange(nsample)
    x = t<=length
    fx = fftshift(fft(x))
    power = np.abs(fx)**2
    angle = np.angle(fx)

    plt.figure()
    plt.subplot(1,2,1)
    plt.gca().set_yscale([0,1.2])
    plt.title('$f(x)$')
    plt.plot(x)
    plt.subplot(1,2,2)
    plt.plot(np.arange(-nsample/2,nsample/2),power);
    plt.title('$|F(u)|^2$')

step_power(1)
step_power(4)
step_power(50)
```



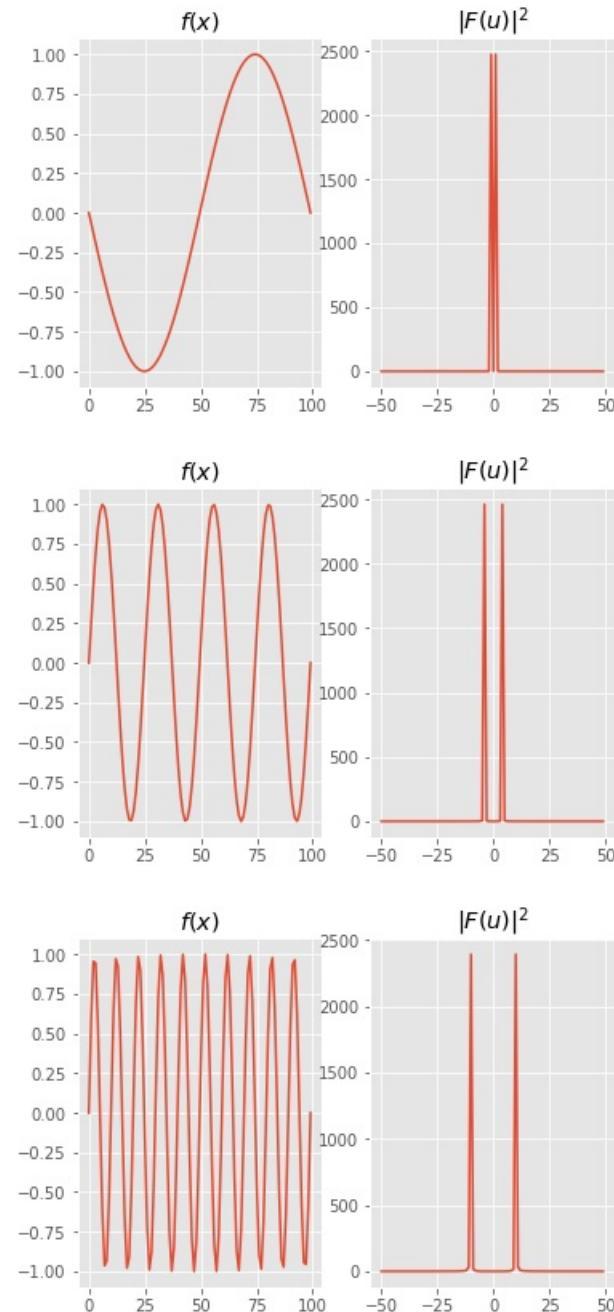
Power spectrum of a sine waves for various wave length, the peak position in the Fourier power spectrum corresponds to the sine frequency.

In [61]:

```
def sine_power(freq):
    nsample = 100
    t = np.linspace(-np.pi,np.pi,nsample)
    x = np.sin(t*freq)
    fx = fftshift(fft(x))
    power = np.abs(fx)**2
    angle = np.angle(fx)

    plt.figure()
    plt.subplot(1,2,1)
    plt.gca().set_yscale([-1.1,1.1])
    plt.title('$f(x)$')
    plt.plot(x)
    plt.subplot(1,2,2)
    plt.plot(np.arange(-nsample/2,nsample/2),power);
    plt.title('$|F(u)|^2$')

sine_power(1)
sine_power(4)
sine_power(10)
```



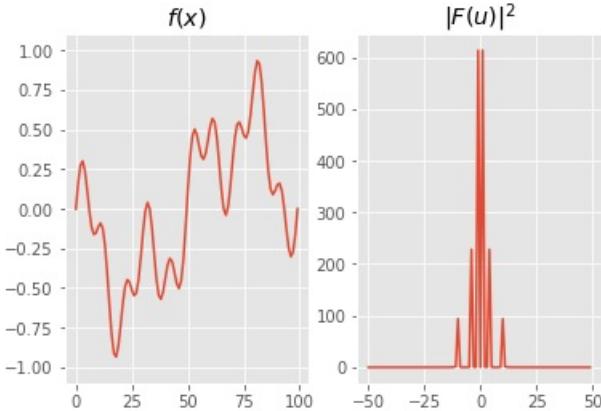
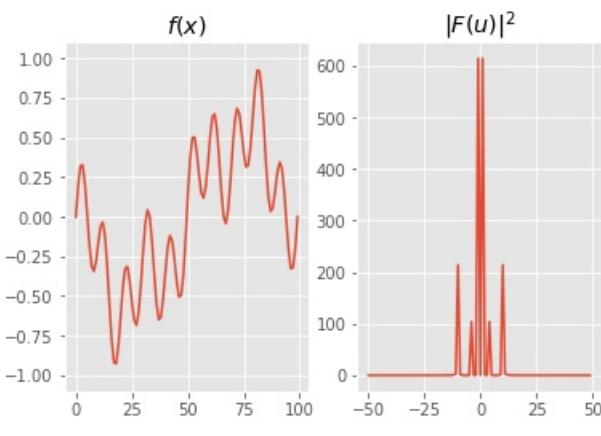
Power spectrum of two differnt mixtures of three sine waves:

In [62]:

```
def mix_sine_power(freq):
    nsample = 100
    t = np.linspace(-np.pi,np.pi,nsample)
    x = np.zeros_like(t)
    for f,a in freq:
        x += np.sin(t*f)*a
    fx = fftshift(fft(x))
    power = np.abs(fx)**2
    angle = np.angle(fx)

    plt.figure()
    plt.subplot(1,2,1)
    plt.gca().set_yscale([-1.1,1.1])
    plt.plot(x)
    plt.title('f(x)')
    plt.subplot(1,2,2)
    plt.plot(np.arange(-nsample/2,nsample/2),power);
    plt.title('|F(u)|^2')

mix_sine_power([(1,.5),(4,.2),(10,.3)])
mix_sine_power([(1,.5),(4,.3),(10,.2)])
```



2D Fourier transform

Similarly, 2D Fourier transform is defined as:

$$F(u, v) = F(f(x, y)) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(x, y) e^{-i2\pi(ux+vy)} dx dy$$

the equivalent step transform is given below:

In [63]:

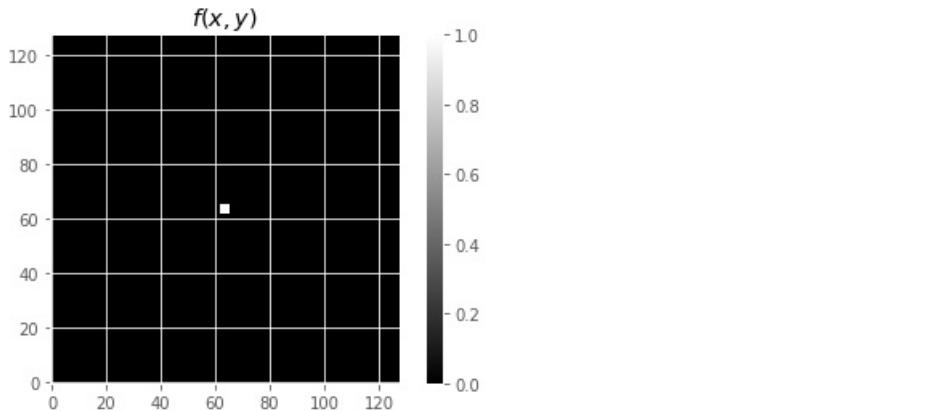
```
import matplotlib.pyplot as plt
import matplotlib.pylab as pylab
import matplotlib.cm as cm
import scipy as scp
import numpy as np
from numpy.fft import fft2, ifft2, fftshift, ifftshift
from scipy import ndimage
from mpl_toolkits.mplot3d import axes3d

w = 2
n = 128
r = np.zeros((n,n), dtype = np.complexfloating)
r[int(n/2)-w:int(n/2)+w, int(n/2)-w:int(n/2)+w] = 1.0 + 0.0j

F = fft2(r) #2D FFT of the image
rr = ifft2(F) #2D inverse FFT of F

real_part = F.real
imag_part = F.imag

fig = plt.figure(1)
plt.imshow(r.real, interpolation='nearest', origin='lower', cmap=cm.gray)
plt.colorbar()
plt.title('$f(x,y)$');
```



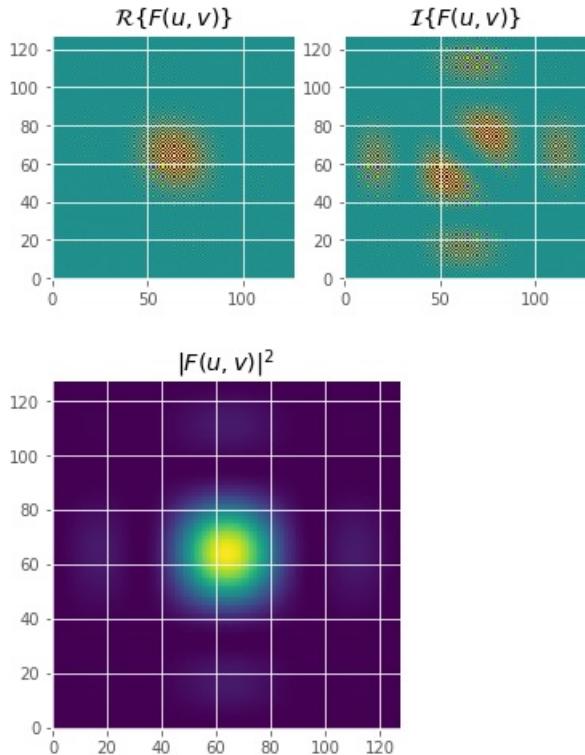
The original image consist of a 128x128 image containing zeros everywhere except in its center (a 2x2 square).

In [64]:

```
fig = plt.figure(2)
plt.subplot(1,2,1)
plt.imshow(fftshift(real_part),interpolation='nearest',origin='lower')
plt.title('$\mathcal{R}\{F(u,v)\}$')

plt.subplot(1,2,2)
plt.imshow(fftshift(imag_part),interpolation='nearest',origin='lower')
plt.title('$\mathcal{I}\{F(u,v)\}$')

fig = plt.figure(3)
plt.imshow(np.abs(fftshift(F))**2,interpolation='nearest',origin='lower')
plt.title('$|F(u,v)|^2$');
```

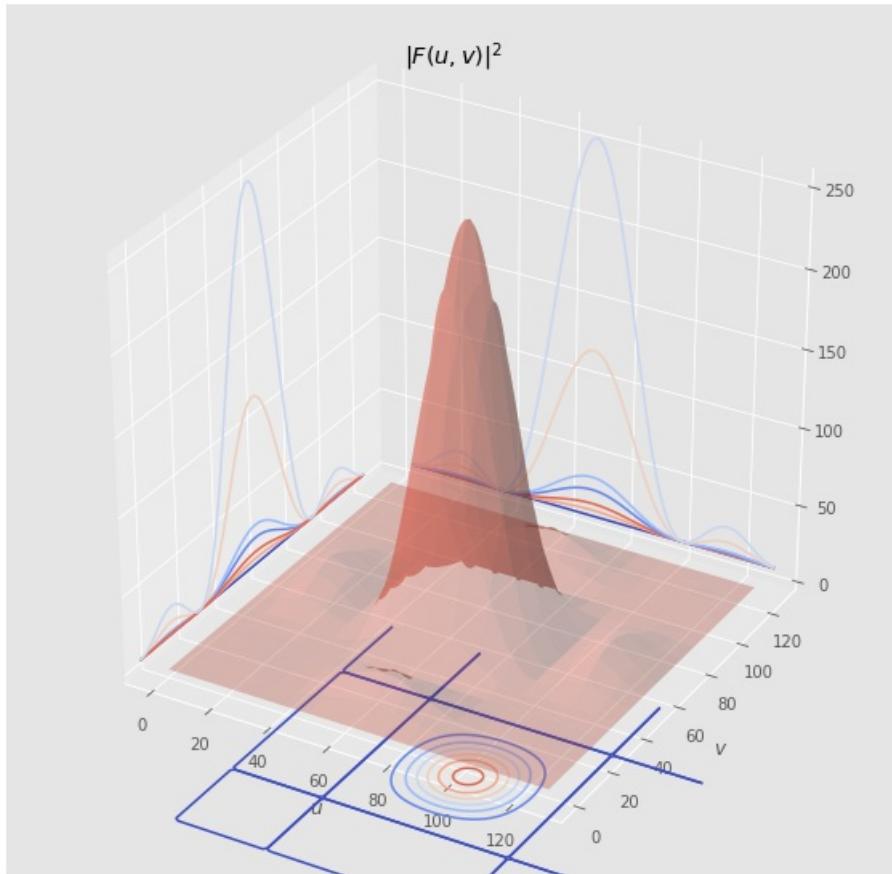


In the Fourier spectrum, the zero spatial frequency is in the center. We see that most of the energy is in the central part of the spectrum (a great part of the image has no transition), however, because of the sharp border of the pixel we have also higher frequencies available, the figure below illustrates the $\sin(x)/x$ aspect of the power spectrum of the 2D step function.

In [65]:

```
fig = plt.figure(figsize=[10,10])
ax = fig.gca(projection='3d')
X,Y = np.meshgrid(range(n),range(n))
Z = np.abs(fftshift(F))**2
ax.plot_surface(X, Y, Z, rstride=8, cstride=8, alpha=0.3)
cset = ax.contour(X, Y, Z, zdir='z', offset=-100, cmap=cm.coolwarm)
cset = ax.contour(X, Y, Z, zdir='x', offset=-10, cmap=cm.coolwarm)
cset = ax.contour(X, Y, Z, zdir='y', offset=140, cmap=cm.coolwarm)

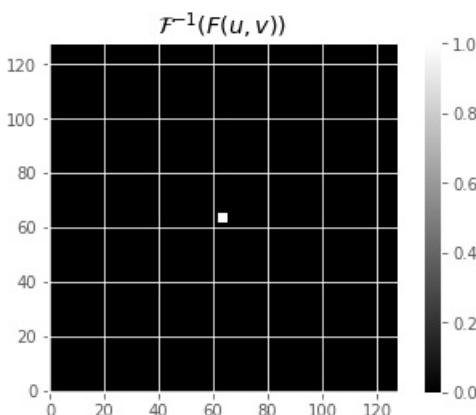
plt.xlabel('$u$')
plt.ylabel('$v$')
plt.title('$|F(u,v)|^2$');
```



The inverse transform is illustrated below, this is the result of the FFT inverse transform of the previous transformed image. We get the original image back.

In [66]:

```
fig = plt.figure()
plt.imshow(rr.real, interpolation='nearest', origin='lower', cmap=cm.gray)
plt.colorbar()
plt.title('$\mathcal{F}^{-1}(F(u,v))$');
```



Phase and amplitude

What appends if we add noise to amplitude or to phase ?

In [67]:

```
ima = camera()[-1::-1,:]
F = fft2(ima)
amplitude = np.abs(F)
angle = np.angle(F)

noise = np.random.random(ima.shape)
n_amplitude = amplitude * (1+10*noise)

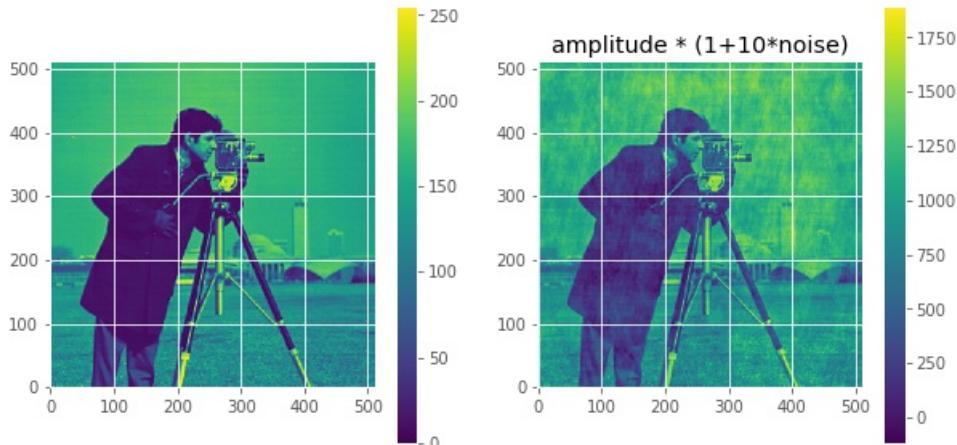
real = n_amplitude * np.cos(angle)
imag = n_amplitude * np.sin(angle)

n_F = real + 1j *imag

n_ima = np.real(ifft2(n_F))

fig = plt.figure(figsize=[10,5])
plt.subplot(1,2,1)
plt.imshow(ima,interpolation='nearest',origin='lower')
plt.colorbar()

plt.subplot(1,2,2)
plt.imshow(n_ima,interpolation='nearest',origin='lower')
plt.colorbar();
plt.title('amplitude * (1+10*noise)');
```



In [68]:

```
ima = camera()[-1::-1,:]
F = fft2(ima)
amplitude = np.abs(F)
angle = np.angle(F)

noise = np.random.random(ima.shape)
n_angle = angle * (1+.5*noise)

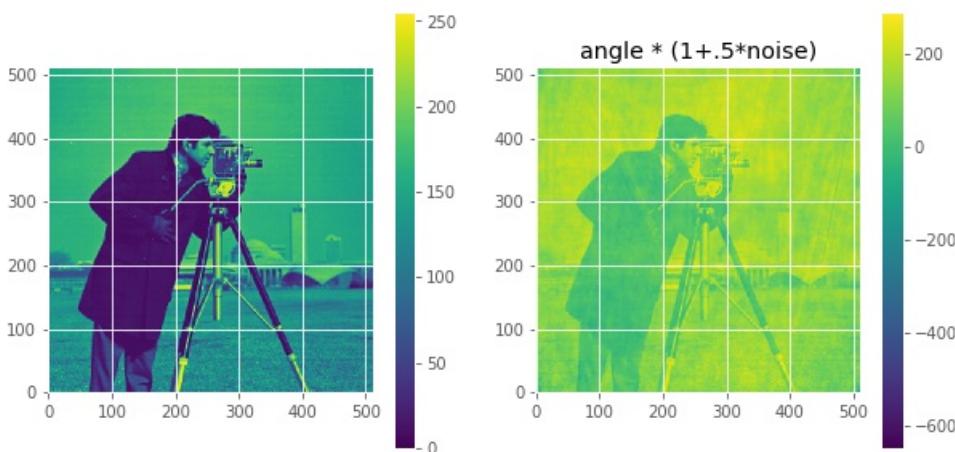
real = amplitude * np.cos(n_angle)
imag = amplitude * np.sin(n_angle)

n_F = real + 1j *imag

n_ima = np.real(ifft2(n_F))

fig = plt.figure(figsize=[10,5])
plt.subplot(1,2,1)
plt.imshow(ima,interpolation='nearest',origin='lower')
plt.colorbar()

plt.subplot(1,2,2)
plt.imshow(n_ima,interpolation='nearest',origin='lower')
plt.colorbar();
plt.title('angle * (1+.5*noise)');
```



Discrete Fourier transform

This is the Fourier transform for discrete signals, if $f(x)$ is a discrete function with N samples $x = 0, \dots, N-1$.

The discrete Fourier transform is given by:

$$F(u) = \frac{1}{N} \sum_{x=0}^{N-1} f(x) e^{-i2\pi ux/N}$$

and the corresponding inverse transform is:

$$f(x) = \sum_{u=0}^{N-1} F(u) e^{i2\pi ux/N}$$

the 2D extension is :

$$\begin{aligned} F(u, v) &= \frac{1}{MN} \sum_{x=0}^{M-1} \sum_{y=0}^{N-1} f(x, y) e^{-i2\pi(ux/M + vy/N)} \\ f(x, y) &= \sum_{u=0}^{M-1} \sum_{v=0}^{N-1} F(u, v) e^{i2\pi(ux/M + vy/N)} \end{aligned}$$

The Fast Fourier Transform algorithm

The FFT algorithm enable to drastically diminish the number of operation for the discrete Fourier transform of a function. Without going into details (see [DIP] p 120), the complexity of the original direct Fourier transform algorithm goes from N^2 to $N \log_2 N$ for FFT method, which is a major improvement especially when image size N is big.

Because image are discrete functions, and because we are interested in doing convolution with them, if the size of the structuring element is big, one may use the FFT in place of naïve convolution techniques as described above.

The FFT algorithm changes the order of the coefficients, in order to have the 0 frequency back in the center of the image we use the `fftshift()` function.

Convolution

Local processing as described before is in fact a convolution, thanks to the nice properties of the Fourier domain, convolution can be computed in an other way.

Convolution of two function is defined as:

$$f(x) * g(x) = \int_{-\infty}^{\infty} f(u)g(x-u)du$$

the corresponding operation in the frequency domain is:

$$f(x) * g(x) \Rightarrow F(u)G(u)$$

in other word, the convolution in the signal space is equivalent to a term by term product in the frequency domain.

Similarly for 2D images we have:

$$f(x, y) * g(x, y) \Rightarrow F(u, v)G(u, v)$$

so the convolution of an image f by a structuring element g consist of two discrete Fourier transform, a simple term-by-term matrix product and an inverse discrete Fourier transform.

The following example illustrate a low pass filter (only frequencies near the center of the image are kept).

In [69]:

```
from skimage.data import camera
from skimage.filters import rank as skr
from skimage.morphology import disk

im = camera()[-1::-1,:,:]
w,h = im.shape
n = 32

#square filter
s = np.zeros(im.shape,dtype = np.complexfloating)
s[int(w/2)-n:int(w/2)+n,int(h/2)-n:int(h/2)+n] = 1.0 + 0.0j

#circular filter
c = np.zeros(im.shape,dtype = np.complexfloating)
for i in range(w):
    for j in range(h):
        if ((i-w/2)**2 + (j-h/2)**2)<(n*n):
            c[i,j] = 1.0 + 0.0j

#smooth filter borders
c = skr.mean(np.real(c*255).astype('uint8'),disk(10))
c = c.astype(np.complexfloating)/(255.0+0j)

F1 = fft2(im.astype(np.complexfloating))
F3 = F1*ifftshift(s)
F4 = F1*ifftshift(c)

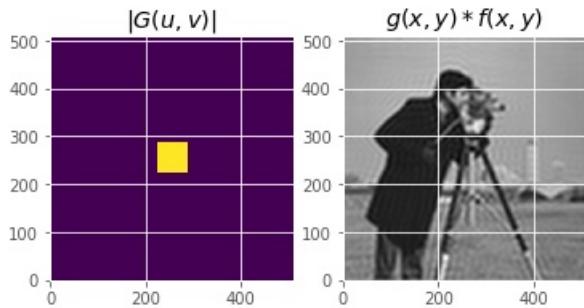
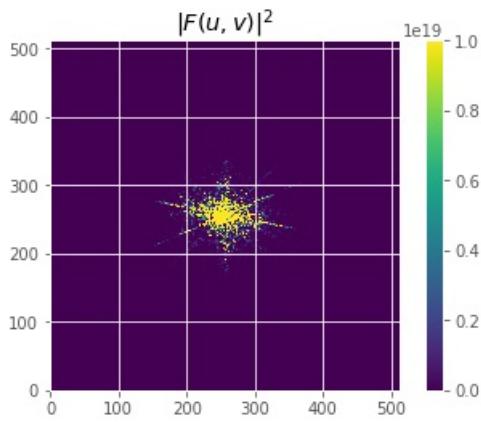
#high pass using the complement of c
F5 = F1*ifftshift((1.0 + 0j)-c)

psF1 = (F1**2).real

low_pass_rec = ifft2(F3)
low_pass_circ = ifft2(F4)
high_pass_circ = ifft2(F5)

fig = plt.figure(1)
plt.imshow(np.abs(fftshift(psF1))**2,interpolation='nearest',origin='lower',vmax = 10e18)
plt.title('$|F(u,v)|^2$')
plt.colorbar()

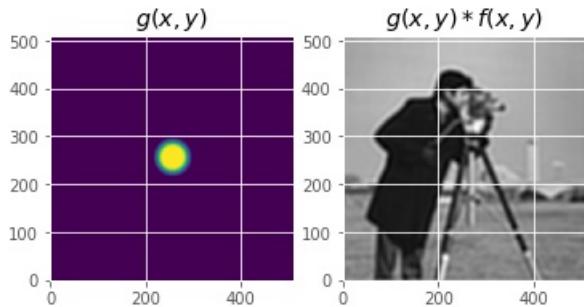
fig = plt.figure(2)
plt.subplot(1,2,1)
plt.imshow(s.real,interpolation='nearest',origin='lower')
plt.title('$|G(u,v)|$')
plt.subplot(1,2,2)
plt.imshow(low_pass_rec.real,interpolation='nearest',origin='lower',cmap=cm.gray)
plt.title('$g(x,y)*f(x,y)$');
```



One remark the presence of oscillation in tje filtered image, this is due to the sharpness of the filter used, bellow the same low-pass filter but with smoother border.

In [70]:

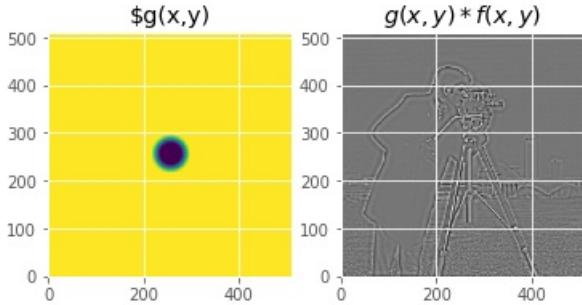
```
fig = plt.figure(3)
plt.subplot(1,2,1)
plt.imshow(c.real, interpolation='nearest', origin='lower')
plt.title('$g(x,y)$')
plt.subplot(1,2,2)
plt.imshow(low_pass_circ.real, interpolation='nearest', origin='lower', cmap=cm.gray)
plt.title('$g(x,y)*f(x,y)$');
```



If we suppress the central part of the spectrum we obtain a high pass filter, i.e. only border are kept.

In [71]:

```
fig = plt.figure(4)
plt.subplot(1,2,1)
plt.imshow(1.0-c.real,interpolation='nearest',origin='lower')
plt.title('$g(x,y)$')
plt.subplot(1,2,2)
plt.imshow(high_pass_circ.real,interpolation='nearest',origin='lower',cmap=cm.gray)
plt.title('$g(x,y)*f(x,y)$');
```



Correlation

Similarly, correlation can benefit from the Fourier domain properties.

Correlation is defined as:

$$f(x) \circ g(x) = \int_{-\infty}^{\infty} f^*(u)g(x+u)du$$

where f^* is the complex conjugate of f (i.e. f for real image).

the corresponding operation in the frequency domain is:

$$f(x) \circ g(x) \Leftrightarrow F^*(u)G(u)$$

In [72]:

```
from skimage.data import camera

im = camera().astype(np.complexfloating)
target_center = (314,375)
w = 16
crop = im[target_center[1]-w:target_center[1]+w,target_center[0]-w:target_center[0]+w]
m,n = im.shape

g = np.zeros((m,n),dtype = np.complexfloating)
g[int(n/2)-w:int(n/2)+w,int(m/2)-w:int(m/2)+w] = crop - np.mean(im) + .0j

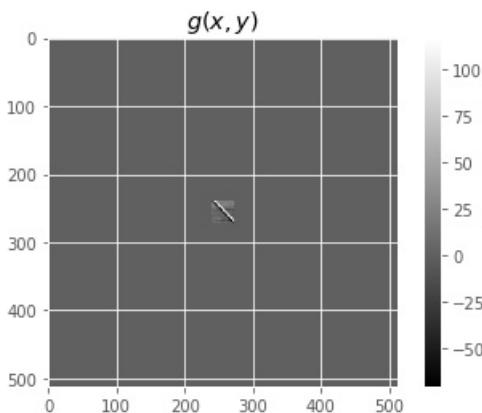
#normalize
f = im-np.mean(im)

plt.figure(2)
plt.imshow(g.real,interpolation='nearest',cmap=cm.gray)
plt.colorbar()
plt.title('g(x,y)$');

F = fft2(f)
G = fft2(np.conjugate(g))

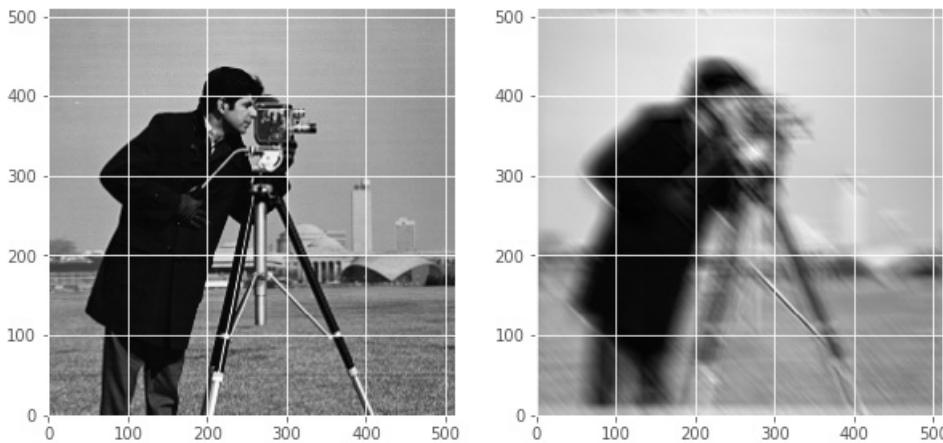
R = F*G
r = fftshift(np.real(ifft2(R)))

corr = 255*((r-np.min(r))/(np.max(r)-np.min(r)))
#corr = 255*(255*((r-np.min(r))/(np.max(r)-np.min(r)))>250)
```



In [73]:

```
compare(np.real(im[-1::-1,:]),np.real(corr[-1::-1,:]))
```



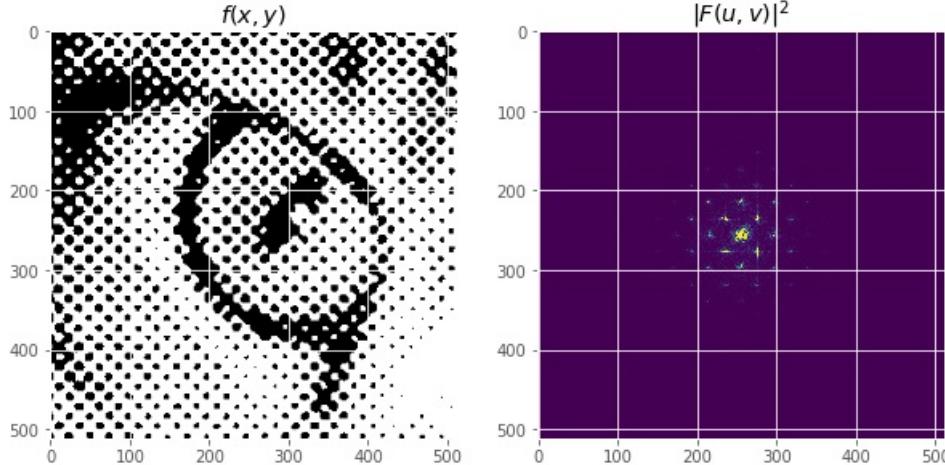
Adapted filter

In [74]:

```
from skimage.io import imread
im = imread('http://homepages.ulb.ac.be/~odebeir/data/moire1.png').astype(np.float)
f = fft2(im)

power = fftshift(np.abs(f)**2)
pmax = np.max(power)

plt.figure(figsize=[10,5])
plt.subplot(1,2,1)
plt.imshow(im,cmap = cm.gray)
plt.title('$f(x,y)$');
plt.subplot(1,2,2)
plt.imshow(power,vmin=0,vmax=pmax/5000.);
plt.title('$|F(u,v)|^2$');
```



Question:

- How to eliminate the repetitive raster pattern ?

In []:

In [2]:

```
%matplotlib inline
import sys
sys.path.insert(0,'..')
from IPython.display import HTML,Image,SVG,YouTubeVideo
from helpers import compare
```

Rank based filters

Rank vs. weighted sum

If the local processing consists in something different from a weighted sum of neighbor pixels, one can speak of non-linear filters.

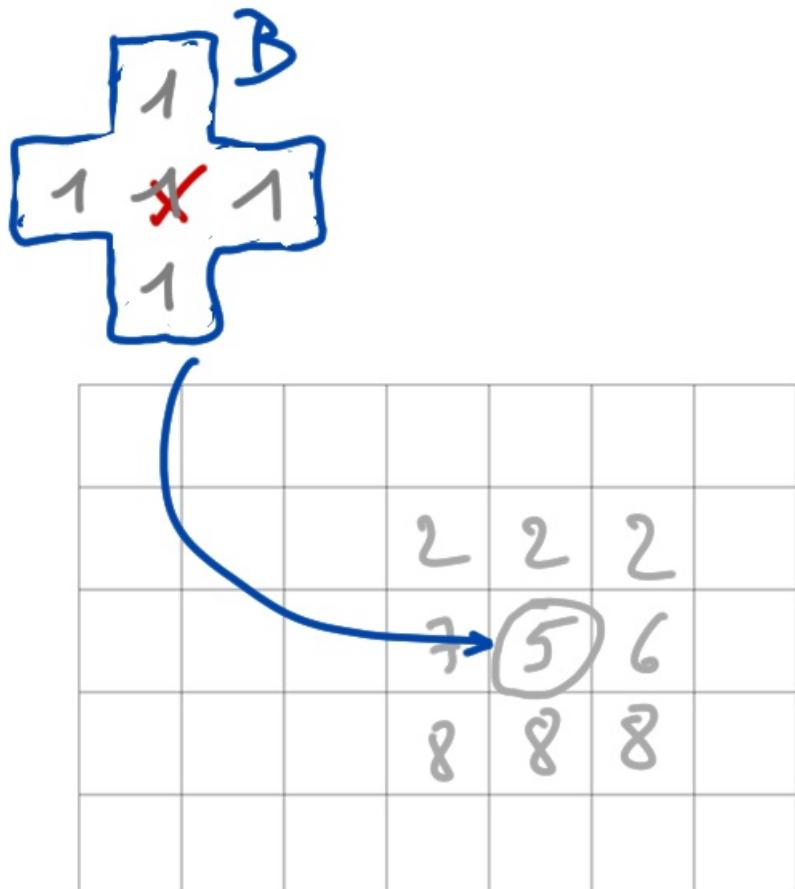
One important category of non-linear filters are the *rank filters*.

These filters use the ranked levels from the neighborhood.

In [3]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/median.png')
```

Out[3]:



2, 2, 2, 5, 6, 7, 8, 8, 8
↑
median

In [4]:

```
from skimage.data import camera
import matplotlib.pyplot as plt
import numpy as np

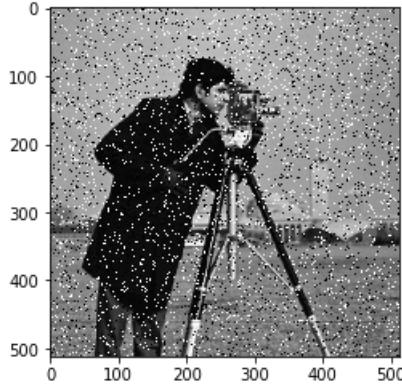
# Salt and pepper noise filtering
ima = camera()
plt.imshow(ima,cmap=plt.cm.gray)

n = np.random.random(ima.shape)

noised_ima = ima.copy()

noised_ima[n<.05] = 0
noised_ima[n>.95] = 255

plt.imshow(noised_ima,cmap=plt.cm.gray);
```

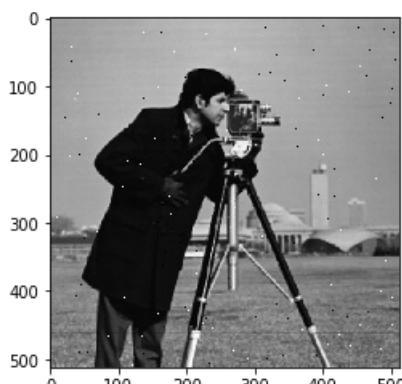


In [5]:

```
from skimage.filters import rank as skr
from skimage.morphology import disk

filtered_im = skr.median(noised_ima,disk(1))

plt.imshow(filtered_im,cmap=plt.cm.gray);
```



In [6]:

```
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np
from scipy import ndimage

from skimage.data import camera
import skimage.filters.rank as skr
from skimage.morphology import disk

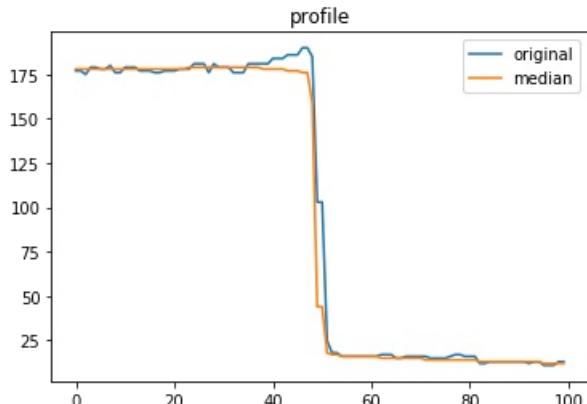
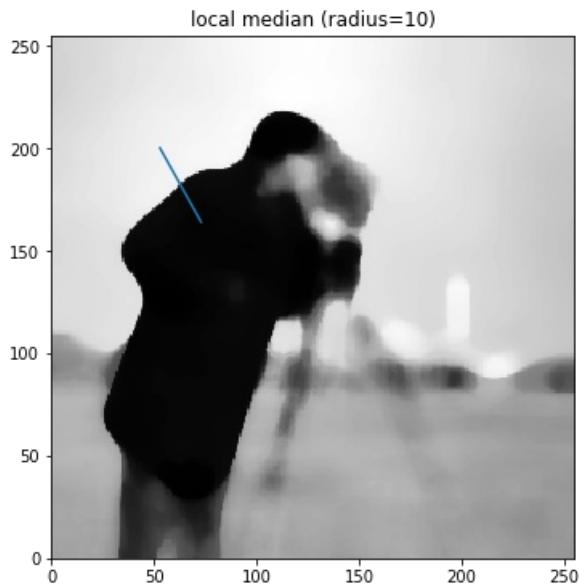
def profile(ima,p0,p1,num):
    n = np.linspace(p0[0],p1[0],num)
    m = np.linspace(p0[1],p1[1],num)
    return [n,m,ndimage.map_coordinates(ima, [m,n], order=0)]

im = camera()[-1::-2,:,:2]

#filtered version
rank = skr.median(im,disk(10))
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank] = profile(rank,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[6,6])
plt.imshow(rank,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('local median (radius=10)')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank,label='median')
plt.title('profile')
plt.legend();
```

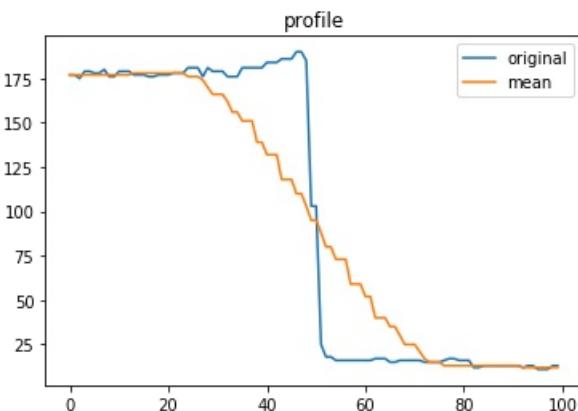
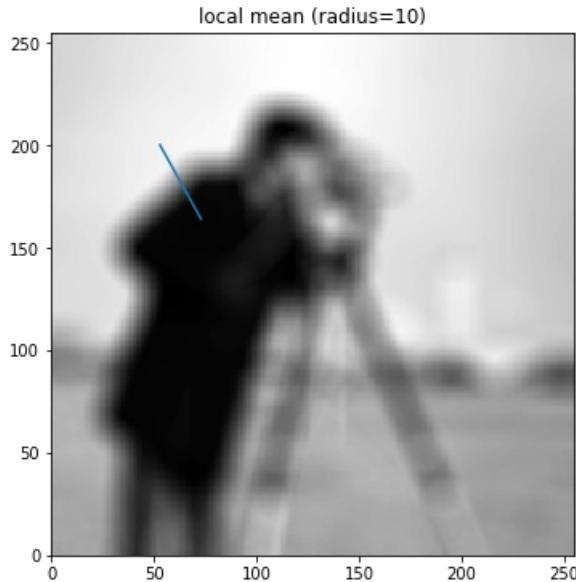


In [7]:

```
#filtered version
mean = skr.mean(im,disk(10))
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,pmean] = profile(mean,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[6,6])
plt.imshow(mean,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('local mean (radius=10)')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(pmean,label='mean')
plt.title('profile')
plt.legend();
```



Question:

- what can we conclude about the border sharpness of the filtered image between the linear smoothing and the median filter?

Local histogram method

In [8]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/histo1.png')
```

Out[8]:

0 0
1 1
2 2
3 1
4 1
5 1
6 1
7 0
8 0

4	5	3	1	2	3	5
2	x	1	4	6	6	6
6	5	4	4	6	7	8
3	4	5	5	6	5	4
2	2	1	1	1	3	2

In [9]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/histo2.png')
```

Out[9]:

#

0	000
1	112
2	211
3	111
4	213
5	111
6	100
7	000
8	000

- ↓ + ↓

4	5	3	1	2	3	5
2	2	X	4	6	6	6
6	5	4	4	6	7	8
3	4	5	5	6	5	4
2	2	1	1	1	3	2

In [10]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/histo3.png')
```

Out[10]:

4	5	3	1	2	3	5	-
2	2	1	4	6	6	6	
6	5	4	4	6	X	8	
3	4	5	5	6	5	4	+
2	2	1	1	1	3	2	

Local maximum, local minimum

Local minimum value and local maximum value are special case of the ranked gray levels.

The figure below illustrates the effect of respectively replacing the pixel:

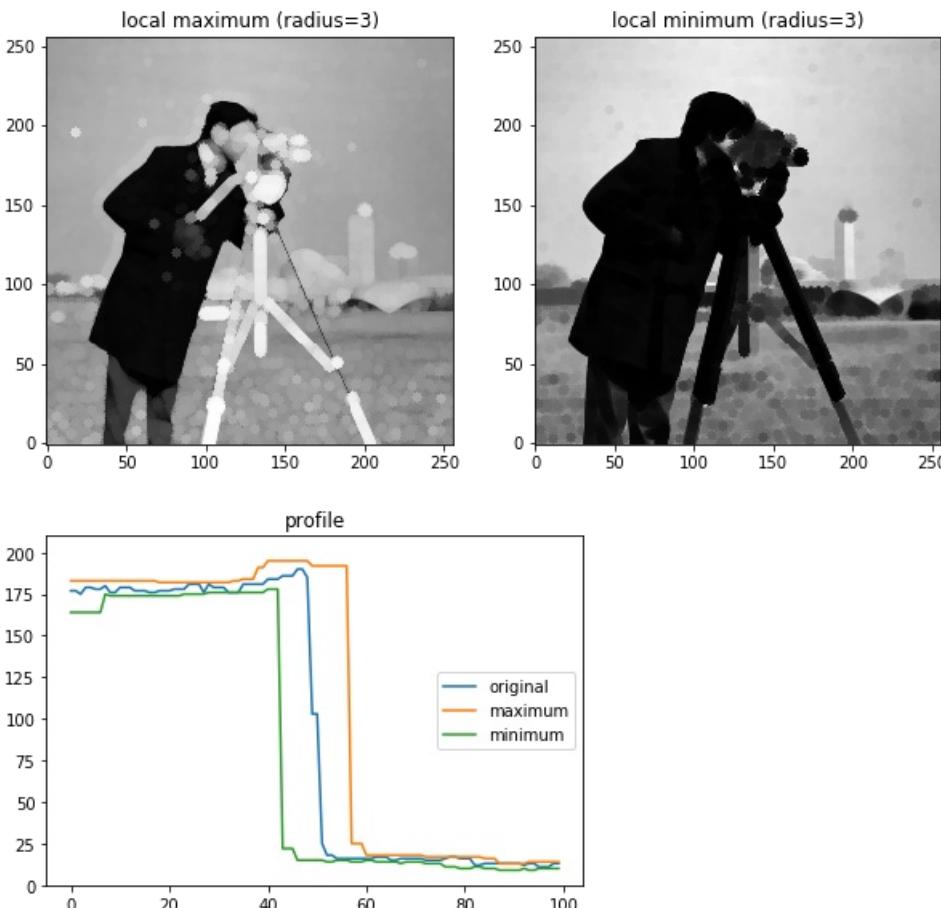
- by the highest value of its neighborhood, the local maximum,
- by the lowest value of its neighborhood, the local minimum.

In [11]:

```
#filtered version
radius = 3
selem = disk(radius)
rank1 = skr.maximum(im,selem)
rank2 = skr.minimum(im,selem)
rank3 = skr.gradient(im,selem)
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank1] = profile(rank1,(53,200),(73,164),100)
[x,y,prank2] = profile(rank2,(53,200),(73,164),100)
[x,y,prank3] = profile(rank3,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[10,10])
plt.subplot(1,2,1)
plt.imshow(rank1,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('local maximum (radius=%d)'%radius)
plt.subplot(1,2,2)
plt.imshow(rank2,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('local minimum (radius=%d)'%radius)

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank1,label='maximum')
plt.plot(prank2,label='minimum')
plt.title('profile')
plt.gca().set_ylim([0,210])
plt.legend(loc=5);
```



Question:

- How to compute borders using local maximum and local minimum ?

Local contrast enhancement

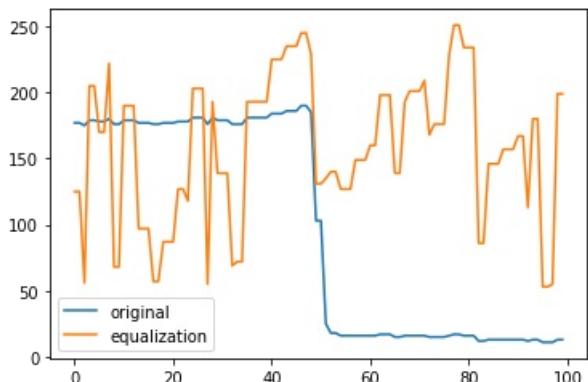
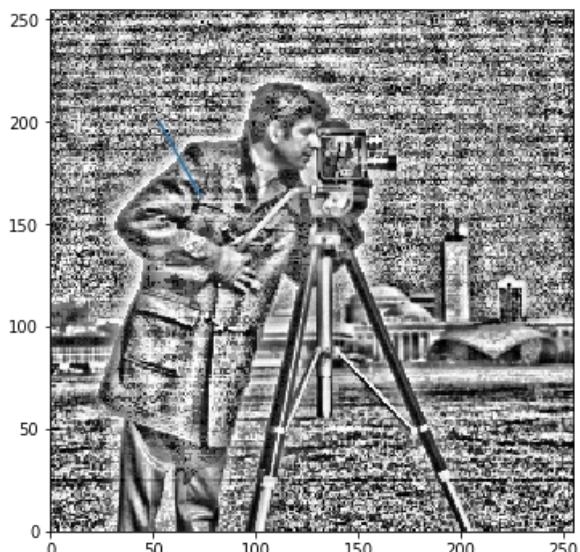
Local contrast equalization

In [12]:

```
#local equalization
rank = skr.equalize(im,disk(10))
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank] = profile(rank,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[6,6])
plt.imshow(rank,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank,label='equalization')
plt.legend();
```



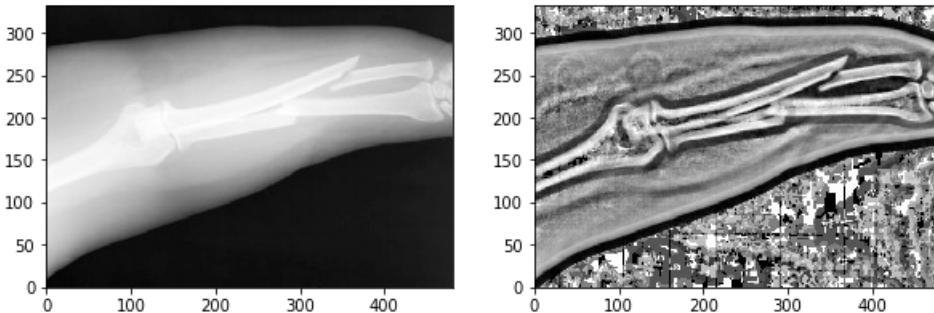
Question:

- How to compute an histogram equalization?

In [13]:

```
#local auto-level
from skimage.io import imread
ima = imread('http://homepages.ulb.ac.be/~odebeir/data/bones.png')
rank = skr.autolevel(ima,disk(10))

compare(ima,rank)
```



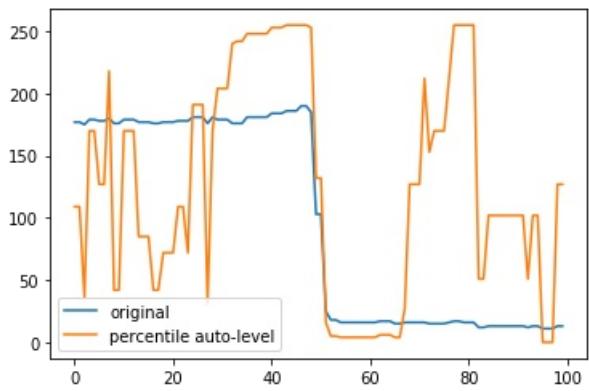
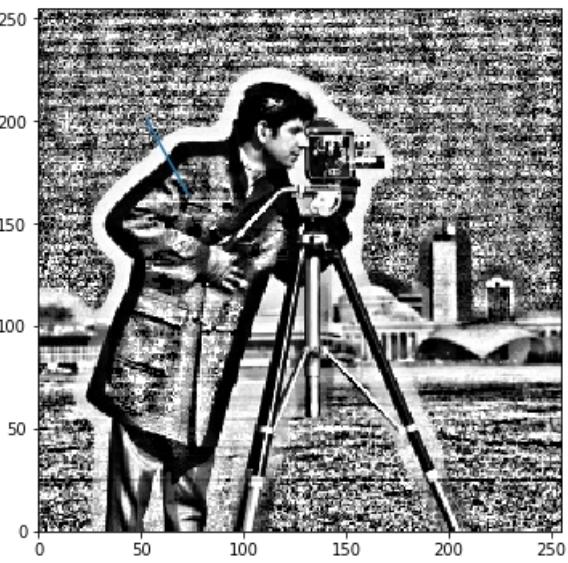
Local autolevel (with percentile)

In [14]:

```
#local soft autolevel
rank = skr.autolevel_percentile(im,disk(10),p0=.1,p1=.9)
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,p_rank] = profile(rank,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[6,6])
plt.imshow(rank,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(p_rank,label='percentile auto-level')
plt.legend();
```



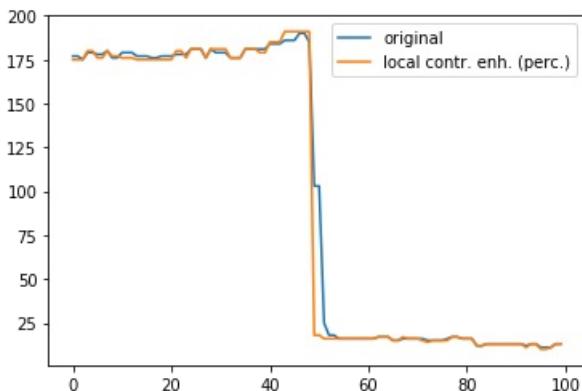
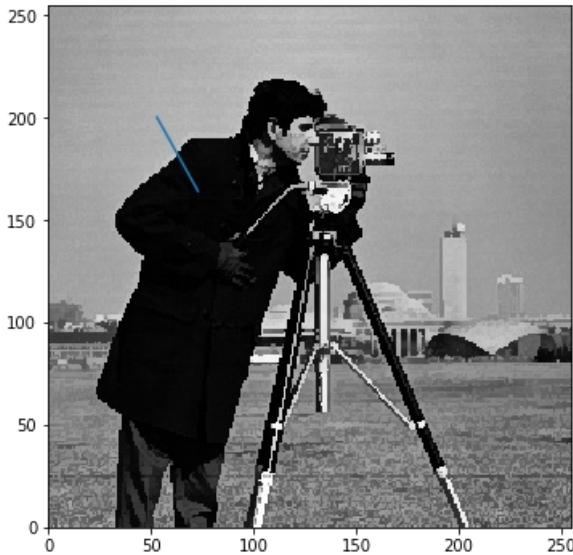
Local morphological contrast enhancement

In [15]:

```
#local soft autolevel
rank = skr.enhance_contrast_percentile(im,disk(2),p0=.1,p1=.9)
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank] = profile(rank,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[6,6])
plt.imshow(rank,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank,label='local contr. enh. (perc.)')
plt.legend();
```



Local threshold

In [16]:

```
p0 = (53,200)
p1 =(73,164)

[x,y,p] = profile(im, p0, p1,100)
[x,y,prank] = profile(rank, p0, p1,100)

fig = plt.figure(0)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(1)
plt.imshow(rank,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

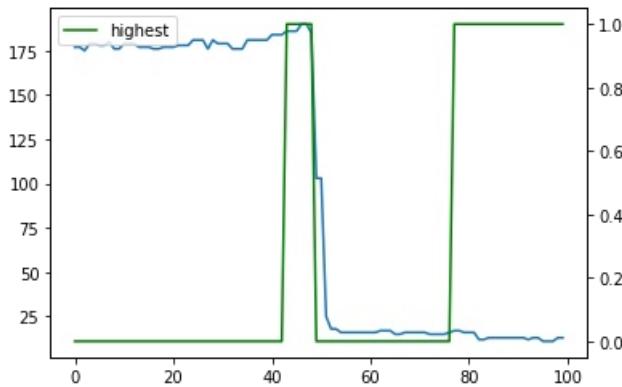
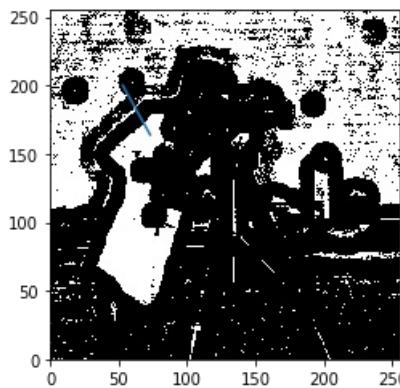
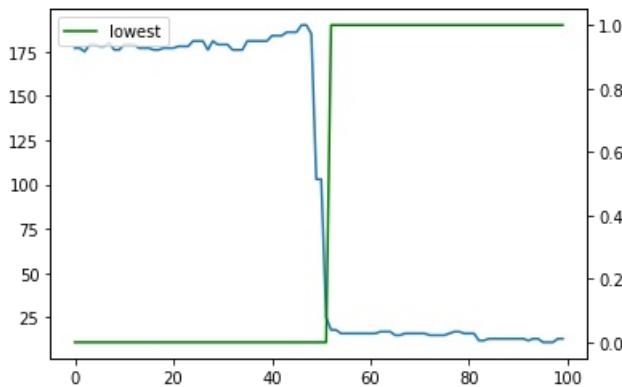
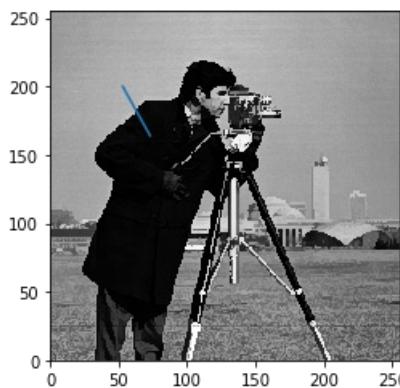
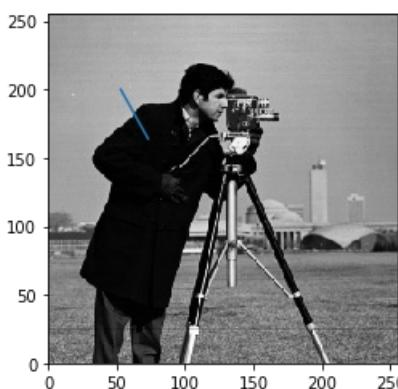
low = im<=(skr.minimum(im,disk(10))+10)
[x,y,p] = profile(im, p0, p1,100)
[x,y,prank] = profile(low, p0, p1,100)

fig = plt.figure(2)
ax1 = plt.subplot(1, 1, 1)
ax2 = ax1.twinx()
ax1.plot(p,label='original')
ax2.plot(prank,'g',label='lowest')
ax2.legend()

high = im>=(skr.maximum(im,disk(10))-10)
[x,y,p] = profile(im, p0, p1,100)
[x,y,prank] = profile(high, p0, p1,100)

fig = plt.figure(3)
plt.imshow(high,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim((0,255))
plt.gca().set_ylim((0,255))

fig = plt.figure(4)
ax1 = plt.subplot(1, 1, 1)
ax2 = ax1.twinx()
ax1.plot(p,label='original')
ax2.plot(prank,'g',label='highest')
ax2.legend();
```



others non linear-filters

Bi-lateral filtering

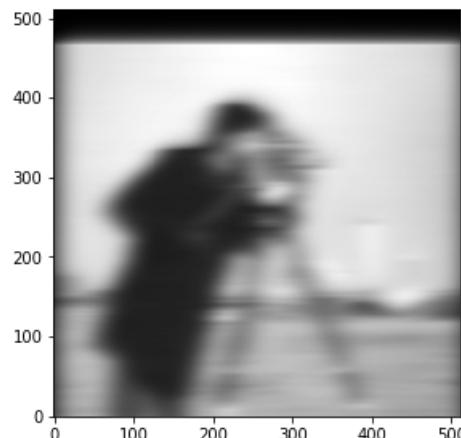
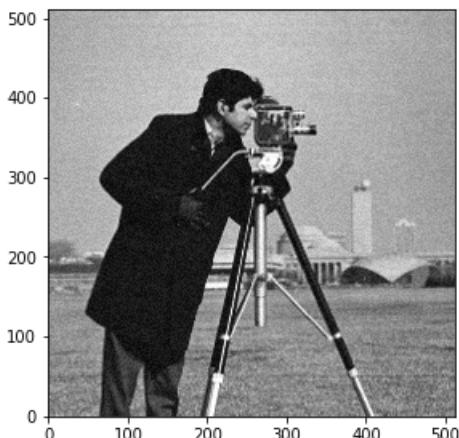
We define the neighbourhood of a pixel as:

- a local spatial neighbourhood and
- a spectral (gray-level) neighbourhood

In [17]:

```
from skimage.restoration import denoise_bilateral
from skimage import img_as_float
ima = img_as_float(camera()[-1:-1])
#add noise
noisy = np.clip(ima+1*np.random.random(ima.shape),0,1)
bilat = denoise_bilateral(noisy, sigma_spatial=15,multichannel=False)

compare(255*noisy,255*bilat)
```



see also:

- bi-lateral filtering [Paris08 \(../00-Preface/06-References.ipynb#\[Paris08\]\)](#)

Anisotropic filter

Nagao

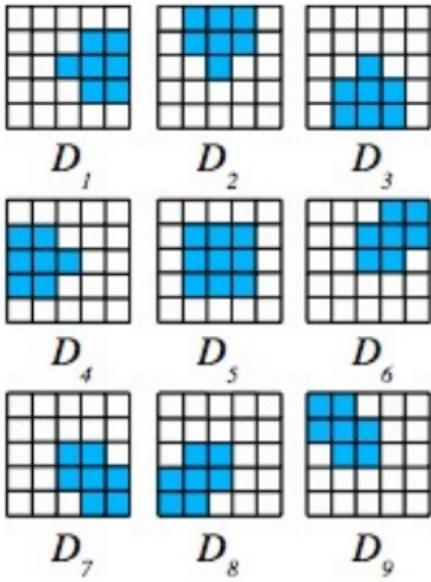
This is an edge preserving smoothing.

- one define 5 anisotropic 5x5 complementary filters (<>directions)
- mean and variance are computed on filter
- filtered value is the mean of the lowest variance filter

In [18]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/nagao.png')
```

Out[18]:



see also:

- Edge preserving smoothing. Makoto Nagao and Takashi Matsuyama. Computer Graphics and Image Processing. Volume 9, Issue 4, April 1979, Pages 394-407
- Rotating filter [IPAMV \(..//00-Preface/06-References.ipynb#\[IPAMV\]\)](#) pp72

Diffusion filter

- smoothing is formulated as a diffusive process
- smoothing is performed at intra regions
- and suppressed at region boundaries

the iteration is given by:

$$I_t = \operatorname{div}(D(|\nabla I|)\nabla u)$$

where I_t is the time derivative of I , the image, D is a diffusion function and ∇ denotes the gradient.

Diffusion function can be defined such as:

$$D(\nabla I) = e^{-(|\nabla I|/k)^2}$$

or

$$D(\nabla I) = \frac{1}{1 + (\frac{|\nabla I|}{k})^2}$$

Solution is found using an iterative approach on the discretized image grid (e.g. for 3D volume):

$$I_{x,y,z}^{t+1} = I_{x,y,z}^t + \lambda \sum_{R=1}^6 [D(\nabla_R I) \nabla_R I]$$

and the gradient evaluated via finite differences:

$$\nabla_1 I_{x,y,z} = I_{x-1,y,z} - I_{x,y,z}, \quad \nabla_2 I_{x,y,z} = I_{x+1,y,z} - I_{x,y,z}, \quad \nabla_3 I_{x,y,z} = I_{x,y-1,z} - I_{x,y,z}, \quad \nabla_4 I_{x,y,z} = I_{x,y+1,z} - I_{x,y,z}, \quad \nabla_5 I_{x,y,z} = I_{x,y,z-1} - I_{x,y,z}, \quad \nabla_6 I_{x,y,z} = I_{x,y,z+1} - I_{x,y,z}$$

The following figure illustrates the application of the diffusion function:

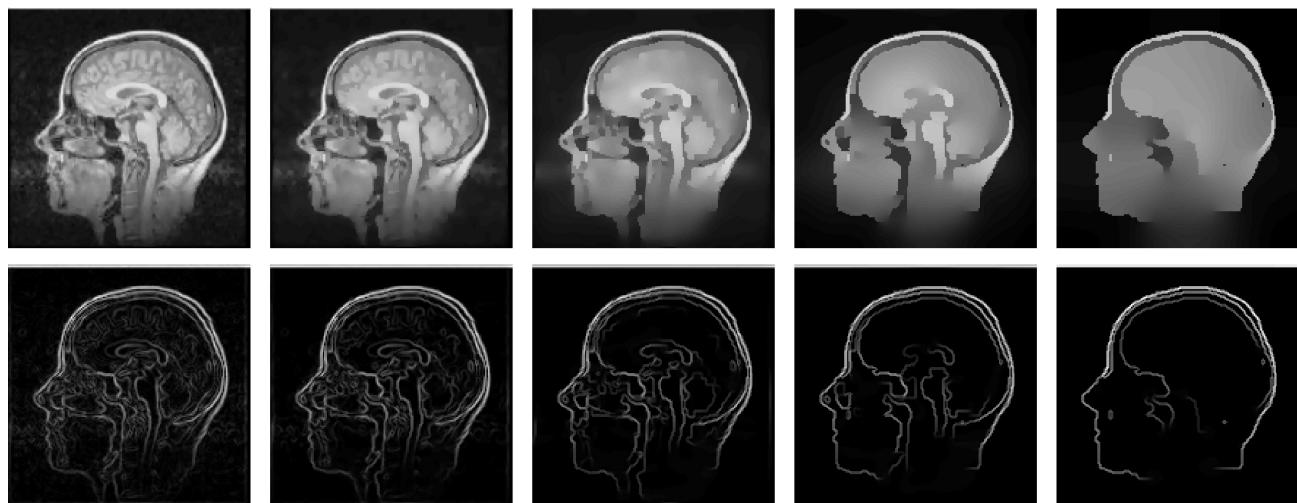
$$D(\nabla I) = e^{-(|\nabla I|/k)^2}$$

after a few iterations, the noise is removed while the heavy borders are conserved.

In [19]:

```
Image('http://www.cs.utah.edu/~jfishbau/advimproc/project2/images/image_k_7.659825.png')
```

Out[19]:



[image source \(http://www.cs.utah.edu/~jfishbau/advimproc/project2/\)](http://www.cs.utah.edu/~jfishbau/advimproc/project2/)

Question:

- what is the equivalent iteration for a 2D image?

see also:

- Ovidiu Ghita, Kevin Robinson, Michael Lynch, Paul F. Whelan . MRI diffusion-based filtering: a note on performance characterisation. Computerized Medical Imaging and Graphics 29 (2005) 267-277
- diffusion based edge detection [IVP \(../00-Preface/06-References.ipynb#\[IVP\]\)](#) p433, [HCVA \(../00-Preface/06-References.ipynb#\[HCVA\]\)](#) vol2 p425
- Pietro Perona and Jitendra Malik (July 1990). "Scale-space and edge detection using anisotropic diffusion". IEEE Transactions on Pattern Analysis and Machine Intelligence, 12 (7): 629-639

Non-local image denoising

- filtered image is a weighted mean of pixels belonging to the “neighbourhood”
- similarity between pixel i and j is function of their local gray level distribution, the pixels with a similar grey level neighbourhood have larger weights in the average

$$f_{new}(p) = \frac{1}{W(p)} \sum_{q \in N(p)} w(p, q) f(q)$$

with

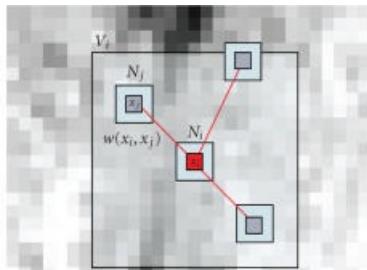
$$W(p) = \sum_{q \in N(p)} w(p, q)$$

- generalized distance defined as the weighted Euclidian distance between two i, j surrounding pixels

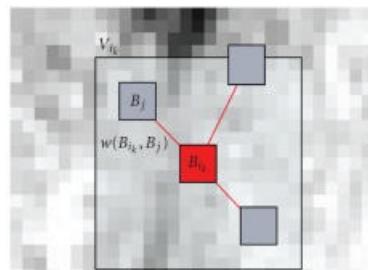
In [20]:

```
Image('http://openi.nlm.nih.gov/imgs/512/290/2292807/2292807_IJBI2008-590183.001.png')
```

Out[20]:



(a)



(b)

[image source \(http://openi.nlm.nih.gov/imgs/512/290/2292807/2292807_IJBI2008-590183.001.png\)](http://openi.nlm.nih.gov/imgs/512/290/2292807/2292807_IJBI2008-590183.001.png)

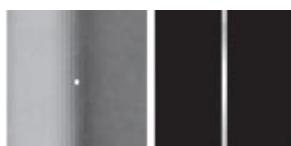
In [21]:

```
#Image('http://deliveryimages.acm.org/10.1145/1950000/1941513/figs/f4.jpg')
Image('http://deliveryimages.acm.org/10.1145/1950000/1941513/figs/f4.jpg', embed=False)
```

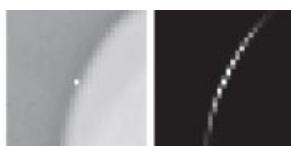
Out[21]:



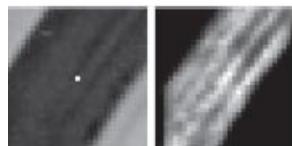
(a)



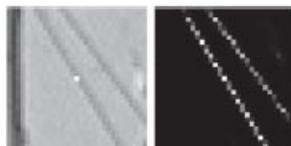
(b)



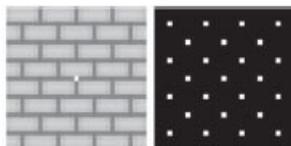
(c)



(d)



(e)



(f)

[image source \(http://deliveryimages.acm.org/10.1145/1950000/1941513/figs/f4.jpg\)](http://deliveryimages.acm.org/10.1145/1950000/1941513/figs/f4.jpg)

see also:

- A non-local algorithm for image denoising [Buades05 \(../00-Preface/06-References.ipynb#\[Buades05\]\)](#)

In [2]:

```
%matplotlib inline
import sys
sys.path.insert(0, '..')
from IPython.display import HTML, Image, SVG, YouTubeVideo
```

Image restoration

Image acquisition is rarely perfect, physics or external condition may deform the image being acquired, here are some example of typical problems:

Deformation model

The original image is $f(x, y)$ undergoes a deformation, given by H , and an additive noise $\eta(x, y)$ the acquired image is $g(x, y)$.

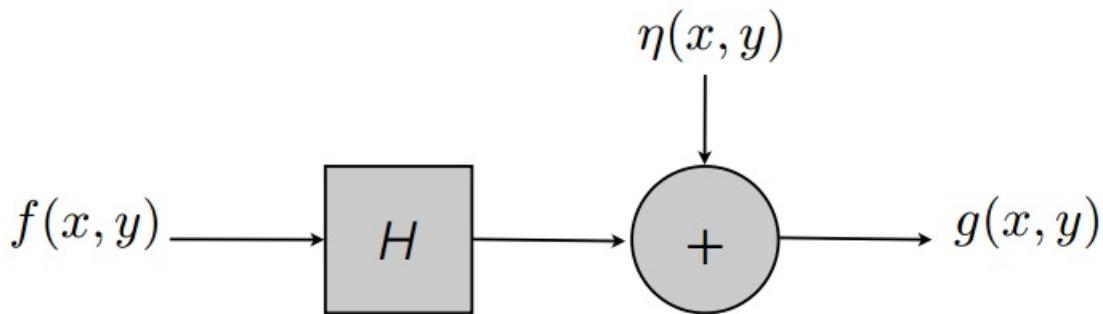
The restoration problem can be stated as follow:

how to recover a good approximation of $f(x, y)$ from $g(x, y)$?

In [3]:

```
Image('http://homepages.ulb.ac.be/~odebeir/data/restauration.png')
```

Out[3]:



Some examples:

$$g(x, y) = H[f(x, y)] + \eta(x, y)$$

H properties:

- linear

$$H[k_1 f_1(x, y) + k_2 f_2(x, y)] = k_1 H[f_1(x, y)] + k_2 H[f_2(x, y)]$$

- additive

$$H[f_1(x, y) + f_2(x, y)] = H[f_1(x, y)] + H[f_2(x, y)]$$

- homogeneous

$$H[k_1 f_1(x, y)] = k_1 H[f_1(x, y)]$$

- spatially invariant

$$H[f(x - \alpha, y - \beta)] = g(x - \alpha, y - \beta)$$

Point Spread Function (PSF)

We can rewrite $f(x, y)$ as a sum of Dirac function.

$$f(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) \delta(x - \alpha, y - \beta) d\alpha d\beta$$

if there is no additive noise:

$$g(x, y) = H[f(x, y)] = H \left[\int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) \delta(x - \alpha, y - \beta) d\alpha d\beta \right]$$

by linearity property:

$$g(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} H[f(\alpha, \beta) \delta(x - \alpha, y - \beta)] d\alpha d\beta$$

$f(\alpha, \beta)$ is independent of x and y :

$$g(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) H[\delta(x - \alpha, y - \beta)] d\alpha d\beta$$

the impulse response of H , also known as point spread function (PSF), is:

$$h(x, \alpha, \beta) = H[\delta(x - \alpha, y - \beta)]$$

after substitution we have:

$$g(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) h(x, \alpha, y, \beta) d\alpha d\beta$$

this expression means that, if the response H of an impulse is known, the response of any input $f(\alpha, \beta)$ is known.

if h is spatially invariant (see above):

$$g(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) h(x - \alpha, y - \beta) d\alpha d\beta$$

by adding the noise:

$$g(x, y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\alpha, \beta) h(x - \alpha, y - \beta) d\alpha d\beta + \eta(x, y)$$

see also:

- [DIP \(..00-Preface/06-References.ipynb#\[DIP\]\)](#) p254

Restoration

inverse filtering

If noise is negligible and PSF is known (in the Fourier domain):

$$\hat{F}(u, v) = G(u, v)H(u, v)\hat{f}(x, y) = F^{-1}[\hat{F}(u, v)] = F^{-1}[G(u, v)H(u, v)]$$

else:

$$\hat{F}(u, v) = F(u, v) + N(u, v)H(u, v)$$

and noise is increasing when H is low, restoration is limited where H is big.

if define a restoration transform $M(u, v)$ as:

$$\hat{F}(u, v) = (G(u, v) + N(u, v))M(u, v)$$

one can use:

$$M(u, v) = \begin{cases} 1/H(u, v), & u^2 + v^2 \leq w_0^2, \\ 0, & u^2 + v^2 > w_0^2 \end{cases}$$

with w_0 being a distance to the origin in the Fourier space.

see also:

- [DPP \(..00-Preface/06-References.ipynb#\[DPP\]\)](#) p276

Wiener filtering

To avoid arbitrary setting of a parameter for the inverse transform such as w_0 , one can use the Wiener approach which consists in minimizing by least square some error function.

Without going into the details, the restoration transform is:

$$M(u, v) = H^*(u, v) |H(u, v)|^2 + S_{vv}(u, v)S_{gg}(u, v)$$

where $H^*(u, v)$ is the complex conjugate of $H(u, v)$,

$S_{vv}(u, v)$ is the spectral density of the noise and

$S_{gg}(u, v)$ is the spectral density of the degraded image.

see also:

- [IPAMV \(../00-Preface/06-References.ipynb#\[IPAMV\]\)](#) p107

Blind deconvolution

If PSF is unknown, it has to be estimated. This is called blind deconvolution.

In [4]:

```
Image('http://bigwww.epfl.ch/algorithms/deconvolutionlab/meta/splash.png')
```

Out[4]:

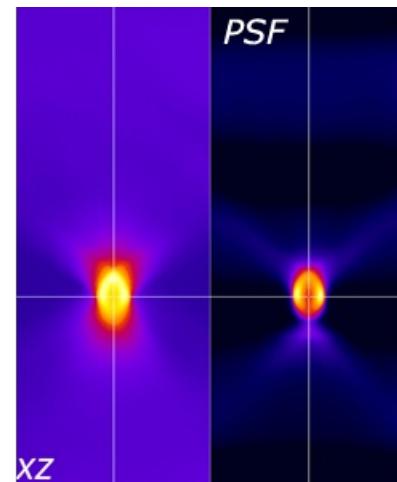
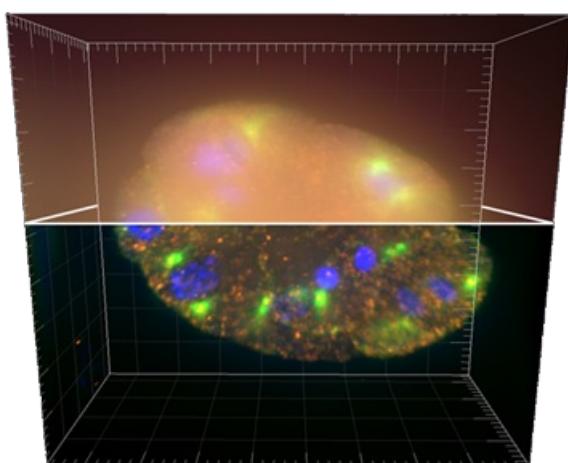


image source (<http://bigwww.epfl.ch/algorithms/deconvolutionlab/>)

In []:

In [1]:

```
%matplotlib inline
import sys
sys.path.insert(0, '..')
from IPython.display import HTML, Image, SVG, YouTubeVideo
```

Edge detection

Edges are important features in an image, this is one of the most saillant feature that our eye catches.

Edges are also highly correlated with object borders, this is why a lot of different thechniques have been developped.

Finite differences

Taylor's theorem:

$$\begin{aligned}f(x+h) &= f(x) + \frac{f'(x)}{1!}h + \frac{f''(x)}{2!}h^2 + \cdots + \frac{f^{(n)}(x)}{n!}h^n + R_n(x) \\f(x+h) &= f(x) + f'(x)h + R_1^+(x) \\f(x-h) &= f(x) - f'(x)h + R_1^-(x)\end{aligned}$$

We neglect R_1 and we substract the two last equations:

$$\begin{aligned}f(x+h) - f(x-h) &\approx 2f'(x)h \\f'(x) &\approx \frac{f(x+h) - f(x-h)}{2h}\end{aligned}$$

Similarly for $f''(x)$

$$f''(x) \approx \frac{f(x+h) - 2f(x) + f(x-h)}{h^2}$$

Finite difference for 2 variables

$$\begin{aligned}f_x(x, y) &\approx \frac{f(x+h, y) - f(x-h, y)}{2h} \\f_y(x, y) &\approx \frac{f(x, y+k) - f(x, y-k)}{2k} \\f_{xx}(x, y) &\approx \frac{f(x+h, y) - 2f(x, y) + f(x-h, y)}{h^2} \\f_{yy}(x, y) &\approx \frac{f(x, y+k) - 2f(x, y) + f(x, y-k)}{k^2} \\f_{xy}(x, y) &\approx \frac{f(x+h, y+k) - f(x+h, y-k) - f(x-h, y+k) + f(x-h, y-k)}{4hk}\end{aligned}$$

Laplacian operator

$$\begin{aligned}\Delta f &= \nabla^2 f = \nabla \cdot \nabla f \\&= \sum_{i=1}^n \frac{\partial^2 f}{\partial x_i^2} \\&= \frac{\partial^2 f}{\partial x^2} + \frac{\partial^2 f}{\partial y^2}\end{aligned}$$

For images, these operators are identical to convolutions with specific structuring element:

example 1D second-derivative is obtained using:

$$\begin{bmatrix} 1 & -2 & 1 \end{bmatrix}$$

2D Laplacian:

$$\begin{bmatrix} 0 & -1 & 0 \\ -1 & +4 & -1 \\ 0 & -1 & 0 \end{bmatrix}$$

2D including diagonals:

$$\begin{bmatrix} -1 & -1 & -1 \\ -1 & +8 & -1 \\ -1 & -1 & -1 \end{bmatrix}$$

3D Laplacian (structuring element is a 3x3x3 cube):

$$\begin{bmatrix} 0 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 0 & 1 & 0 \\ 1 & -6 & 1 \\ 0 & 1 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 0 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

Gradient operator

$$\nabla f = \left(\frac{\partial f}{\partial x_1}, \dots, \frac{\partial f}{\partial x_n} \right)$$

2D

$$\nabla f(x, y) = \left(\frac{\partial f}{\partial x}, \frac{\partial f}{\partial y} \right)$$

3D

$$\nabla f(x, y, z) = \left(\frac{\partial f}{\partial x}, \frac{\partial f}{\partial y}, \frac{\partial f}{\partial z} \right)$$

Amplitude and angle:

$$amplitude = \sqrt{\left(\frac{\partial f}{\partial x}\right)^2 + \left(\frac{\partial f}{\partial y}\right)^2}$$

$$angle = \tan^{-1}\left(\frac{\frac{\partial f}{\partial x}}{\frac{\partial f}{\partial y}}\right)$$

In [2]:

```
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np
from scipy import ndimage,interpolate
from scipy.ndimage.filters import convolve1d
from mpl_toolkits.mplot3d import Axes3D

r = np.arange(-2, 2, 0.2)
x,y = np.meshgrid(r,r)

z = x*np.exp(-x**2-y**2)

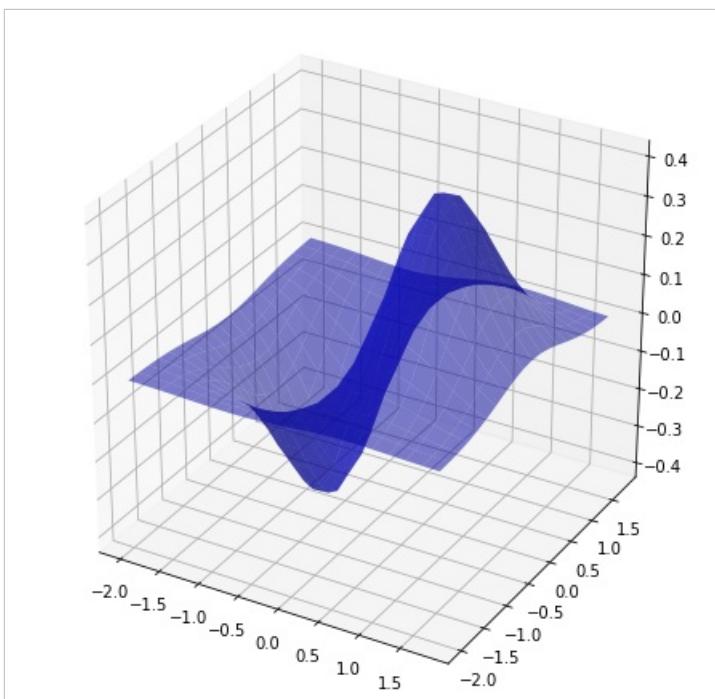
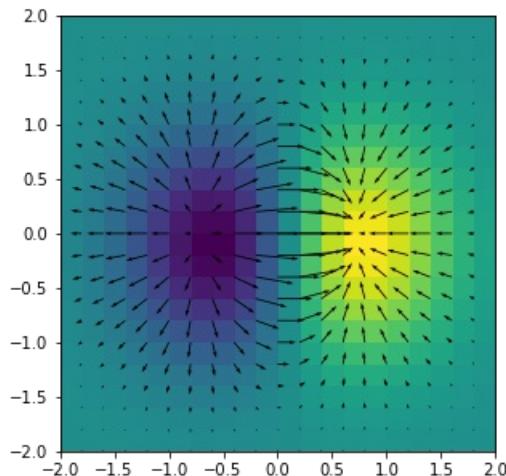
w = np.array([-1,0,+1])

dx = convolve1d(z,-w,axis=1)
dy = convolve1d(z,-w,axis=0)

plt.figure(figsize=[5,5])

plt.imshow(z,extent=[-2,2,-2,2])
plt.quiver(x,y,dx,dy)

fig3d = plt.figure(figsize=[8,8])
ax = fig3d.add_subplot(1, 1, 1, projection='3d')
surf = ax.plot_surface(x, y, z, rstride=1, cstride=1, linewidth=0.2,color=[0.,0.,.8,.5])
```

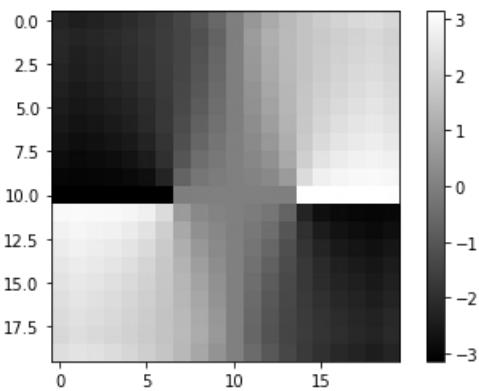


In [3]:

```
plt.imshow(np.arctan2(dy,dx),cmap=plt.cm.gray)
plt.colorbar()
```

Out[3]:

```
<matplotlib.colorbar.Colorbar at 0x7f279f8ca278>
```



In [4]:

```
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as npy
from scipy import ndimage,interpolate
from scipy.ndimage.filters import convolve,convolve1d,gaussian_filter
from mpl_toolkits.mplot3d import Axes3D
from skimage.data import camera

def Cx(ima):
    """x' derivative of image"""
    c = convolve1d(ima,npy.array([-1,0,1]),axis=1,cval=1)
    return c/2.0

def Cy(ima):
    """y' derivative of image"""
    c = convolve1d(ima,npy.array([-1,0,1]),axis=0,cval=1)
    return c/2.0

def grad(ima):
    """gradient of an image"""
    k = npy.array([[0,1,0],[1,0,-1],[0,-1,0]])
    s = convolve(ima,k)
    return s

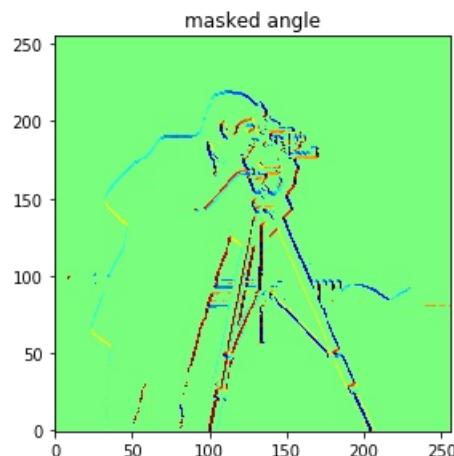
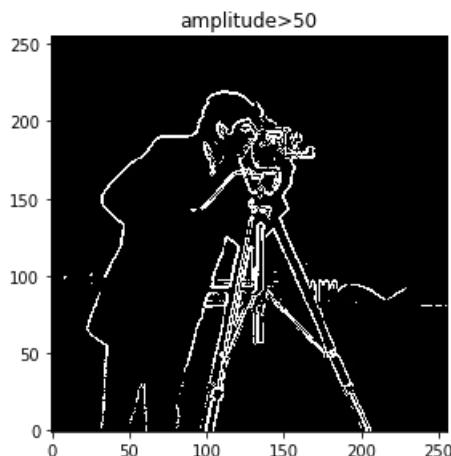
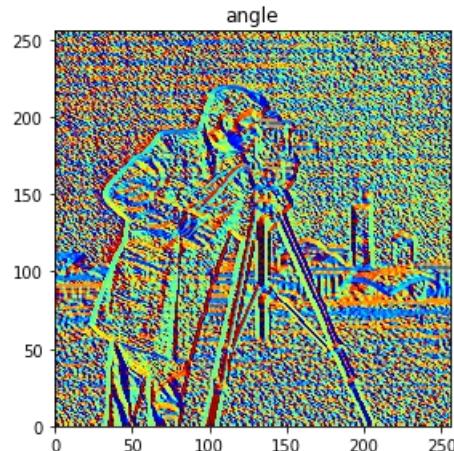
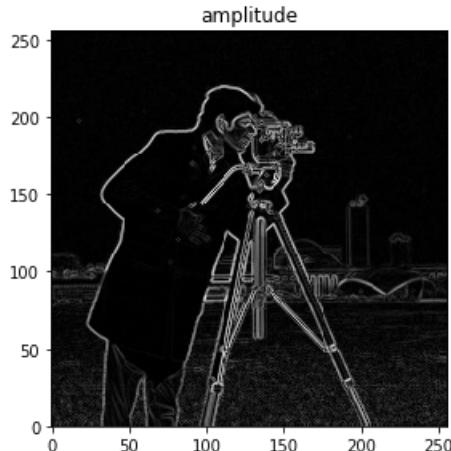
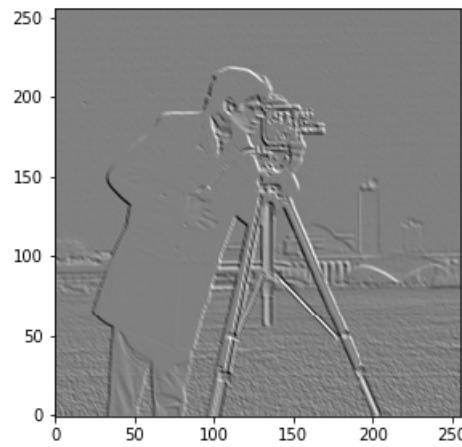
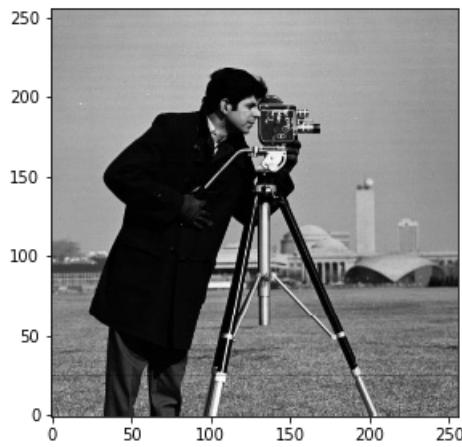
im = camera().astype(np.float)[-1::-2,:,:2]
s = grad(im)

plt.figure(figsize=[10,10])
plt.subplot(1,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(1,2,2)
plt.imshow(s,interpolation='nearest',cmap=cm.gray,origin='lower')

gx = Cx(im)
gy = Cy(im)

magnitude = np.sqrt(gx**2+gy**2)
angle = np.arctan2(gy,gx)
masked_angle = angle.copy()
masked_angle[magnitude<50]=0

plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(magnitude,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('amplitude')
plt.subplot(2,2,2)
plt.imshow(angle,interpolation='nearest',cmap=cm.jet,origin='lower')
plt.title('angle')
plt.subplot(2,2,3)
plt.imshow(magnitude>50,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('amplitude>50')
plt.subplot(2,2,4)
plt.imshow(masked_angle,interpolation='nearest',cmap=cm.jet,origin='lower')
plt.title('masked angle');
```



Question:

- what are the gradient fields for the following images?

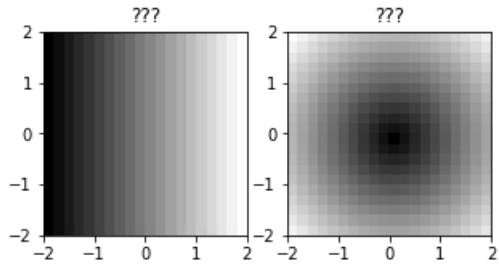
In [5]:

```
plt.figure(figsize=[5,5])

plt.subplot(1,2,1)
z = x
plt.imshow(z,extent=[-2,2,-2,2],cmap=cm.gray)
plt.title('???')

plt.subplot(1,2,2)
z = np.sqrt(x**2+y**2)
plt.imshow(z,extent=[-2,2,-2,2],cmap=cm.gray)
plt.title('???');


```



In []:

Gradient amplitude

$$\vec{\nabla}f = \begin{bmatrix} G_x \\ G_y \end{bmatrix} = \begin{bmatrix} \frac{\partial f}{\partial x} \\ \frac{\partial f}{\partial y} \end{bmatrix}$$

amplitude is given by:

$$\nabla f = ||\vec{\nabla}f|| = [G_x^2 + G_y^2]^{1/2}$$

which can be approximated by (increase processing speed):

$$\nabla f \approx |G_x| + |G_y|$$

Different versions of the gradient amplitude extraction from an image have been proposed, as presented below.

Robert's operator

Robert defines the local image gradient amplitude by:

$$||\vec{\nabla}f|| = |f(x, y) - f(x+1, y+1)| + |f(x+1, y) - f(x, y+1)|$$

which corresponds to the convolution with the two following structuring elements:

$$\begin{bmatrix} 1 & 0 \\ 0 & -1 \end{bmatrix}$$

and

$$\begin{bmatrix} 0 & 1 \\ -1 & 0 \end{bmatrix}$$

Prewitt

Prewitt's operator detect horizontal and vertical borders using:

$$\begin{bmatrix} -1 & -1 & -1 \\ 0 & 0 & 0 \\ 1 & 1 & 1 \end{bmatrix}$$

and

$$\begin{bmatrix} -1 & 0 & 1 \\ -1 & 0 & 1 \\ -1 & 0 & 1 \end{bmatrix}$$

Sobel

Similarly to Prewitt's operator, Sobel border detector is using two orthogonal filters,

$$\begin{bmatrix} 1 & 0 & -1 \\ 2 & 0 & -2 \\ 1 & 0 & -1 \end{bmatrix}$$

$$\begin{bmatrix} 1 & 2 & 1 \\ 0 & 0 & 0 \\ -1 & -2 & -1 \end{bmatrix}$$

In [6]:

```
def sobel(ima):
    """Sobel of image"""
    kx = np.array([[-1,0,1],[-2,0,2],[-1,0,1]])
    ky = np.array([[-1,-2,-1],[0,0,0],[1,2,1]])
    sx = convolve(ima,kx)
    sy = convolve(ima,ky)
    s = np.sqrt(sx**2+sy**2)
    return (sx,sy,s)

def prewitt(ima):
    """Sobel of image"""
    kx = np.array([[-1,0,1],[-1,0,1],[-1,0,1]])
    ky = np.array([[-1,-1,-1],[0,0,0],[1,1,1]])
    sx = convolve(ima,kx)
    sy = convolve(ima,ky)
    s = np.sqrt(sx**2+sy**2)
    return (sx,sy,s)

def roberts(ima):
    """Sobel of image"""
    kx = np.array([[1,0],[0,-1]])
    ky = np.array([[0,1],[-1,0]])
    sx = convolve(ima,kx)
    sy = convolve(ima,ky)
    s = np.sqrt(sx**2+sy**2)
    return (sx,sy,s)

sx,sy,s = sobel(im)

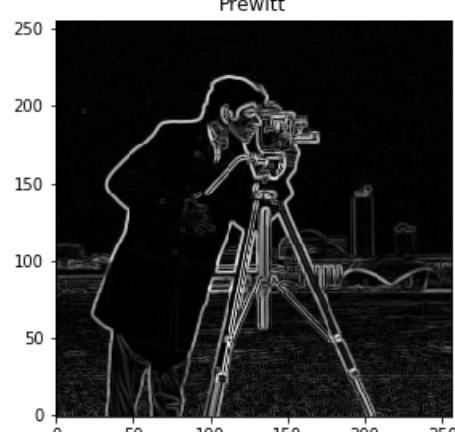
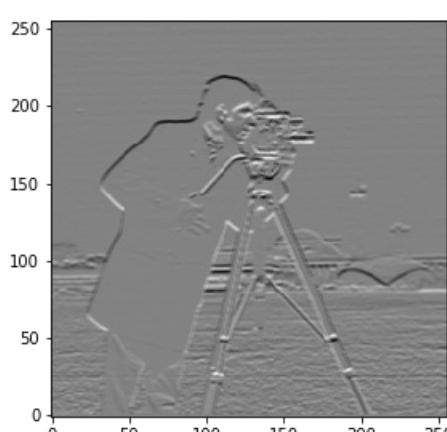
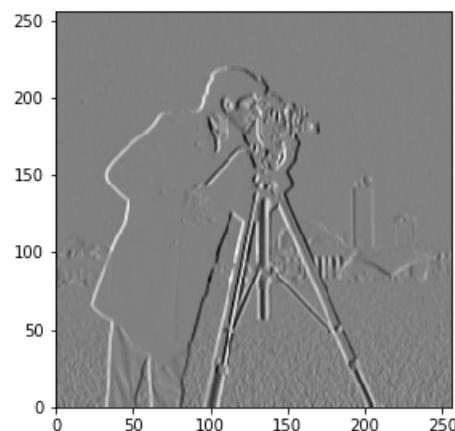
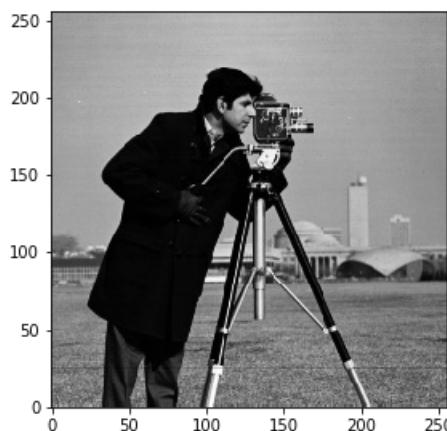
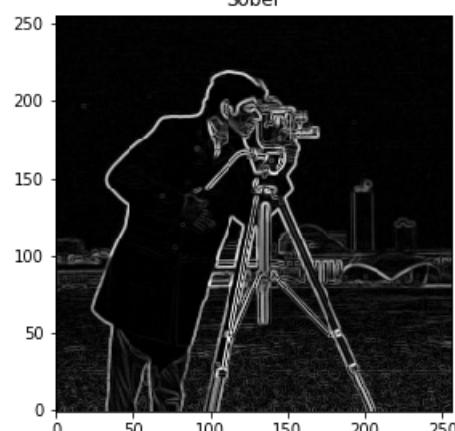
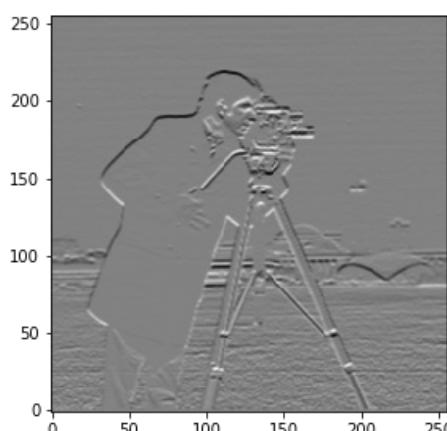
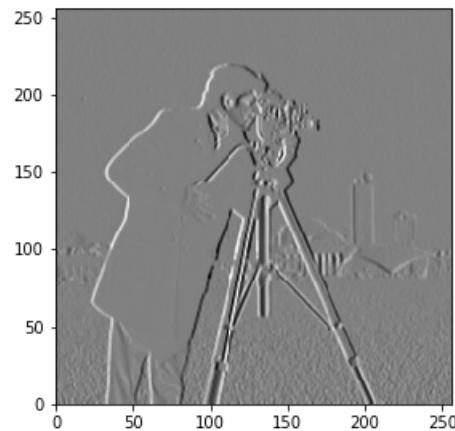
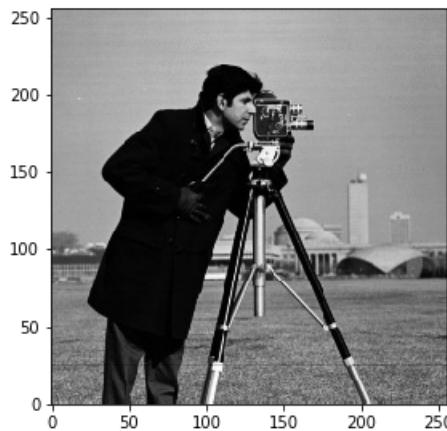
plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,2)
plt.imshow(sx,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,3)
plt.imshow(sy,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,4)
plt.imshow(s,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('Sobel')

sx,sy,s = prewitt(im)

plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,2)
plt.imshow(sx,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,3)
plt.imshow(sy,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,4)
plt.imshow(s,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('Prewitt')

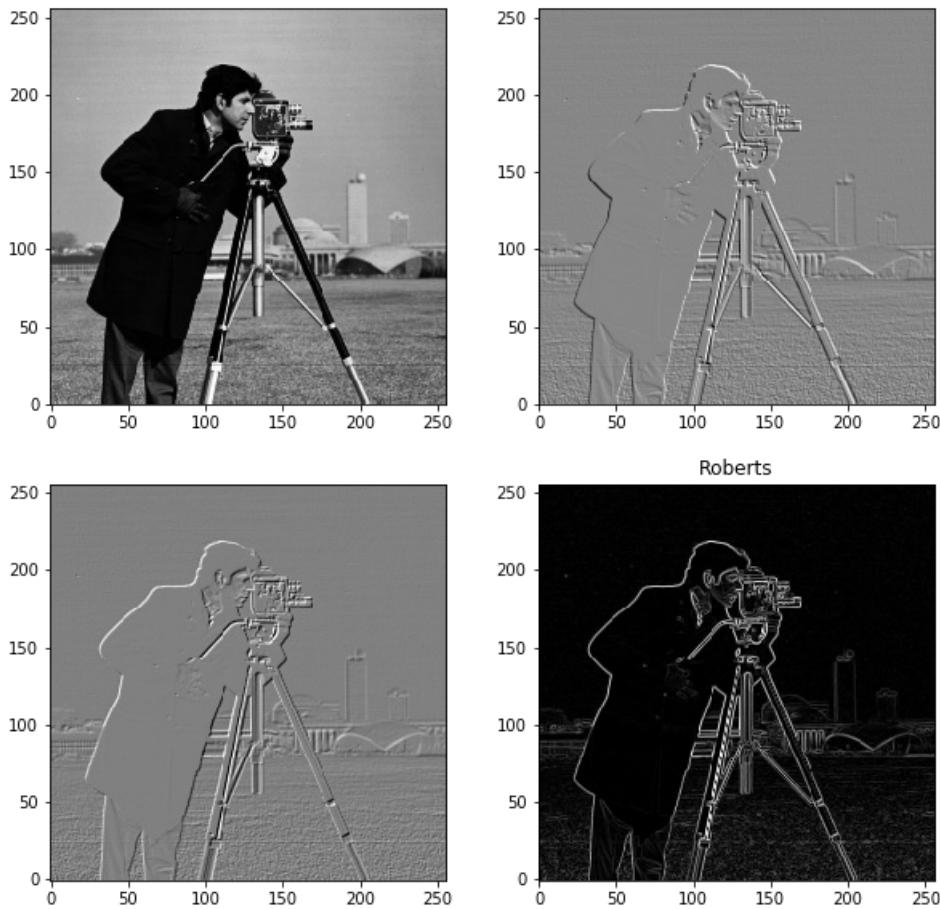
sx,sy,s = roberts(im)

plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,2)
plt.imshow(sx,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,3)
plt.imshow(sy,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.subplot(2,2,4)
plt.imshow(s,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('Roberts');
```



Sobel

Prewitt



Morphological gradient

A gradient can be easily found by subtracting local minimum from local maximum, this is called *morphological gradient* by reference with the morphological operators (see further).

In [7]:

```
from skimage.morphology import disk
import skimage.filters.rank as skr
from scipy import ndimage

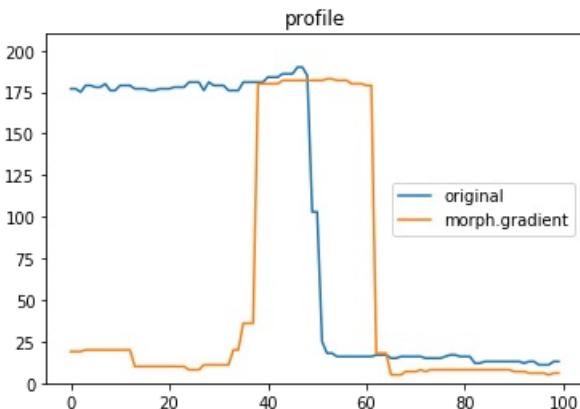
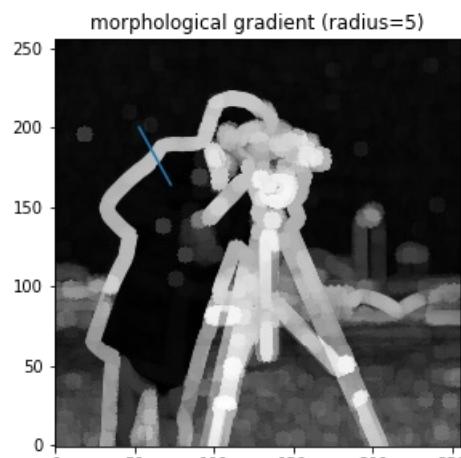
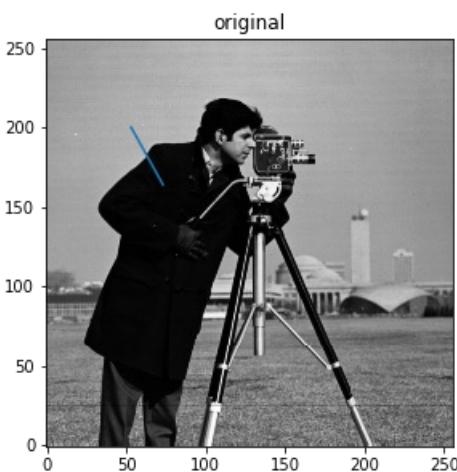
def profile(im,p0,p1,num):
    n = np.linspace(p0[0],p1[0],num)
    m = np.linspace(p0[1],p1[1],num)
    return [n,m,ndimage.map_coordinates(im, [m,n], order=0)]

im = camera()[-1::-2,:,:2]

#filtered version
radius = 5
selem = disk(radius)
rank1 = skr.maximum(im,selem)
rank2 = skr.minimum(im,selem)
rank3 = skr.gradient(im,selem)
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank1] = profile(rank1,(53,200),(73,164),100)
[x,y,prank2] = profile(rank2,(53,200),(73,164),100)
[x,y,prank3] = profile(rank3,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[10,10])
plt.subplot(1,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('original')
plt.plot(x,y)
plt.subplot(1,2,2)
plt.imshow(rank3,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('morphological gradient (radius=%d)'%radius)
plt.plot(x,y)

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank3,label='morph.gradient')
plt.title('profile')
plt.gca().set_ylim([0,210])
plt.legend(loc=5);
```



Two other related morphological gradient are:

- top-hat which is the local maximum - the original image
- bottom-hat which is the original image - the local minimum

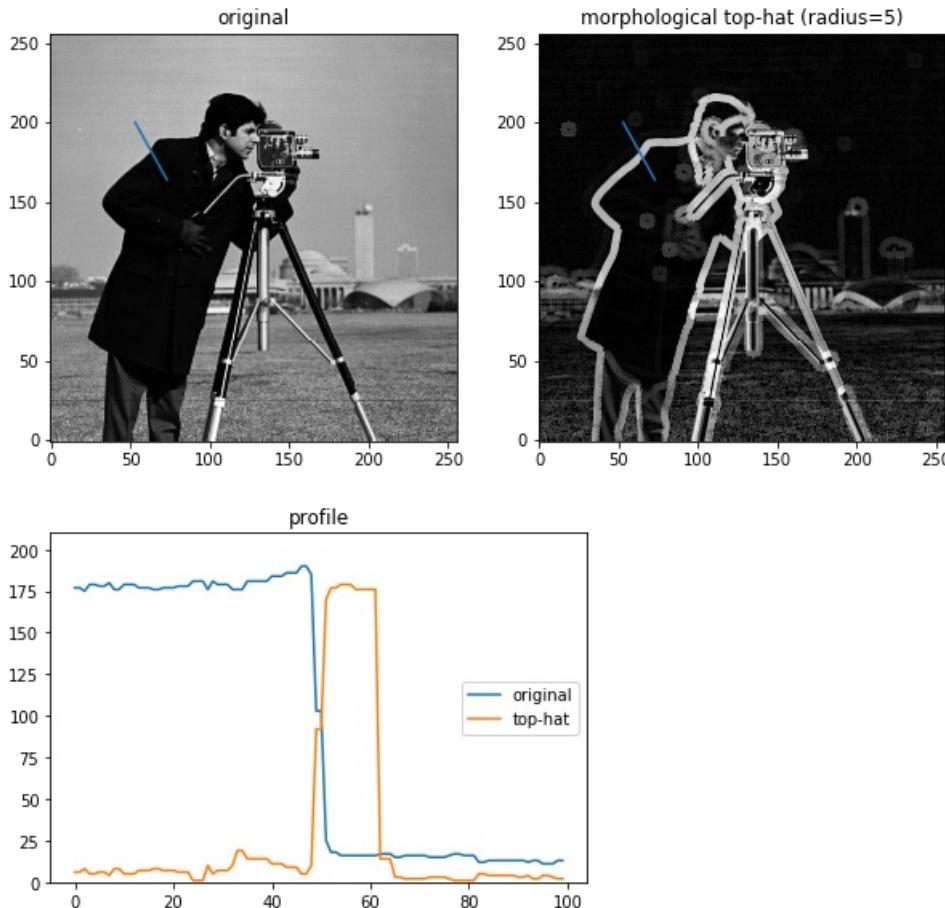
These two filter give thinner borders, but the border are not centered.

In [8]:

```
top_hat = rank1 - im
bottom_hat = im - rank2
[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,prank1] = profile(im,(53,200),(73,164),100)
[x,y,prank2] = profile(top_hat,(53,200),(73,164),100)
[x,y,prank3] = profile(bottom_hat,(53,200),(73,164),100)

fig = plt.figure(1,figsize=[10,10])
plt.subplot(1,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('original')
plt.plot(x,y)
plt.subplot(1,2,2)
plt.imshow(top_hat,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('morphological top-hat (radius=%d)'%radius)
plt.plot(x,y)

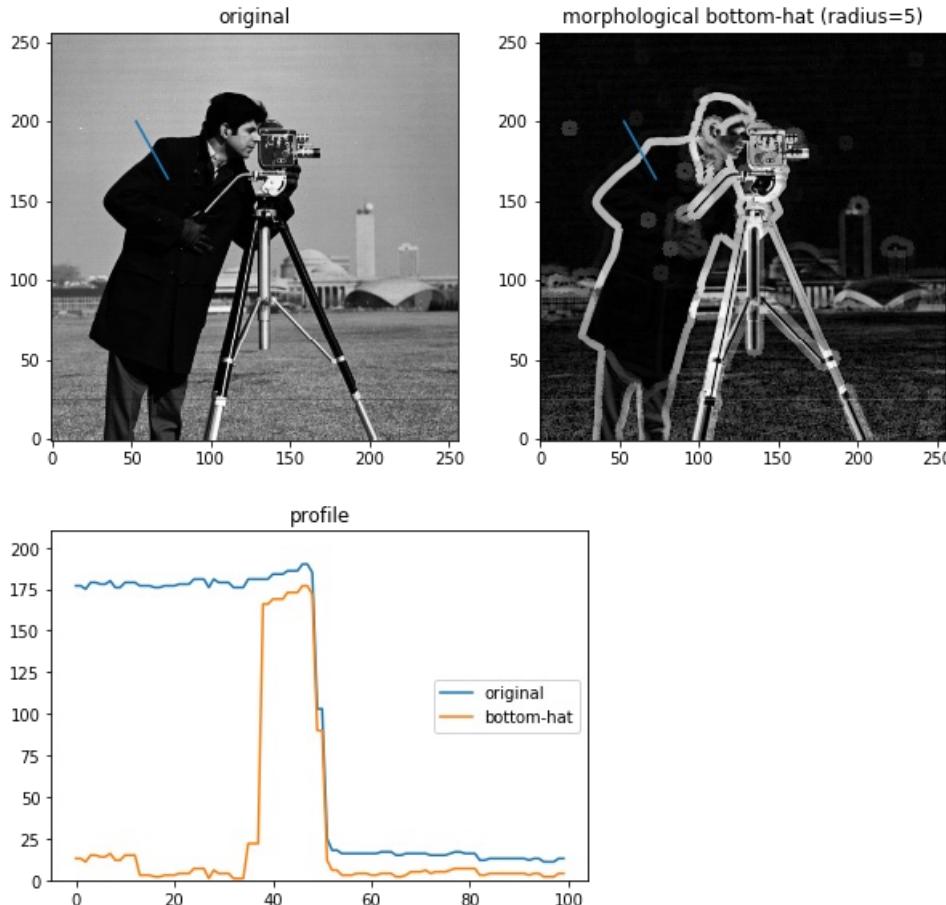
fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank2,label='top-hat')
plt.title('profile')
plt.gca().set_ylim([0,210])
plt.legend(loc=5);
```



In [9]:

```
fig = plt.figure(1,figsize=[10,10])
plt.subplot(1,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('original')
plt.plot(x,y)
plt.subplot(1,2,2)
plt.imshow(top_hat,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('morphological bottom-hat (radius=%d)'%radius)
plt.plot(x,y)

fig = plt.figure(2)
plt.plot(p,label='original')
plt.plot(prank3,label='bottom-hat')
plt.title('profile')
plt.gca().set_ylimits([0,210])
plt.legend(loc=5);
```



Attention must be paid to border detection method used, the size of the detected objects may be influenced, for example, the top-hat transform is over-estimating the size of bright objects and under-estimating the size of dark objects. On the contrary, the bottom-hat is shifting borders in the reverse direction.

Laplacian of gaussian

Laplacian of gaussian is a combination of a high-pass laplacian filter applied on a gaussian low-pass filtered image.

2D gaussian kernel is defined as:

$$G(x, y; \sigma) = \frac{1}{2\pi\sigma^2} e^{-\left(\frac{x^2+y^2}{2\sigma^2}\right)}$$

The Laplacian of Gaussian kernel is then:

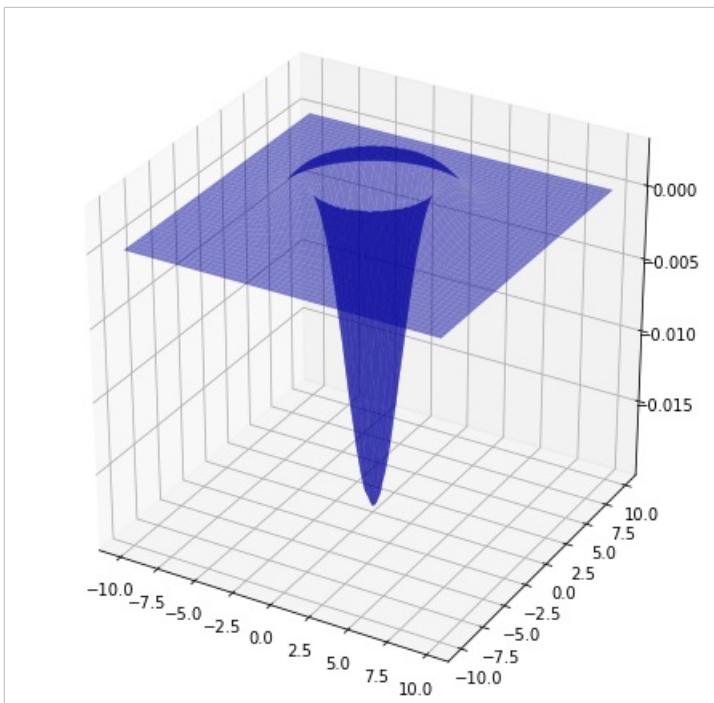
$$\begin{aligned}\Delta f &= \sum_{i=1}^n \frac{\partial^2 f}{\partial x_i^2} \\ LoG(x, y; \sigma) &= \Delta \frac{1}{2\pi\sigma^2} e^{-\left(\frac{x^2+y^2}{2\sigma^2}\right)} \\ &= -\frac{1}{\pi\sigma^4} \left[1 - \frac{x^2 + y^2}{2\sigma^2} \right] e^{-\left(\frac{x^2+y^2}{2\sigma^2}\right)}\end{aligned}$$

In [10]:

```
sigma = 2.
X,Y = np.meshgrid(np.arange(-10.,10.,.1),np.arange(-10.,10.,.1))
e = (X**2+Y**2)/(2*sigma**2)

Z = - 1./(np.pi * sigma**4)*(1-e)*np.exp(-e)

fig3d = plt.figure(figsize=[8,8])
ax = fig3d.add_subplot(1, 1, 1, projection='3d')
surf = ax.plot_surface(X, Y, Z, linewidth=.2,color=[0.,0.,.8,.5])
```



Difference of Gaussian (D.O.G) operator

Gaussian 2D kernel:

$$g(x, y; \sigma) = \frac{1}{2\pi\sigma^2} e^{-\left(\frac{x^2+y^2}{2\sigma^2}\right)}$$

image convolution with a gaussian kernel:

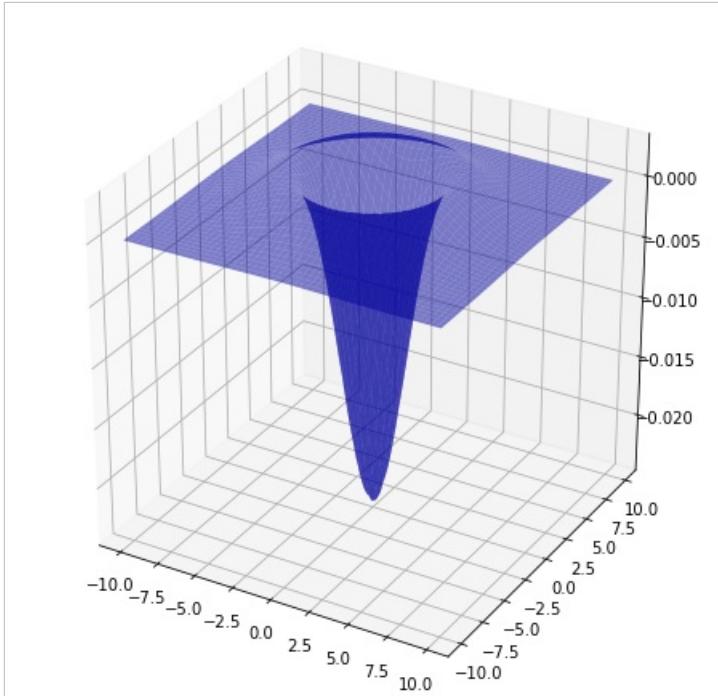
$$L(\cdot, \cdot; \sigma) = g(\cdot, \cdot; \sigma) * f(\cdot, \cdot)$$

In [11]:

```
sigma1 = 2
sigma2 = sigma1*1.6

X,Y = np.meshgrid(np.arange(-10.,10,.1),np.arange(-10.,10,.1))
Z1 = 1./(2*np.pi * sigma1**2)*np.exp(-(X**2+Y**2)/(2*sigma1**2))
Z2 = 1./(2*np.pi * sigma2**2)*np.exp(-(X**2+Y**2)/(2*sigma2**2))

fig3d = plt.figure(figsize=[8,8])
ax = fig3d.add_subplot(1, 1, 1, projection='3d')
surf = ax.plot_surface(X, Y, Z2-Z1, linewidth=.2,color=[0.,0.,.8,.5])
```



In [12]:

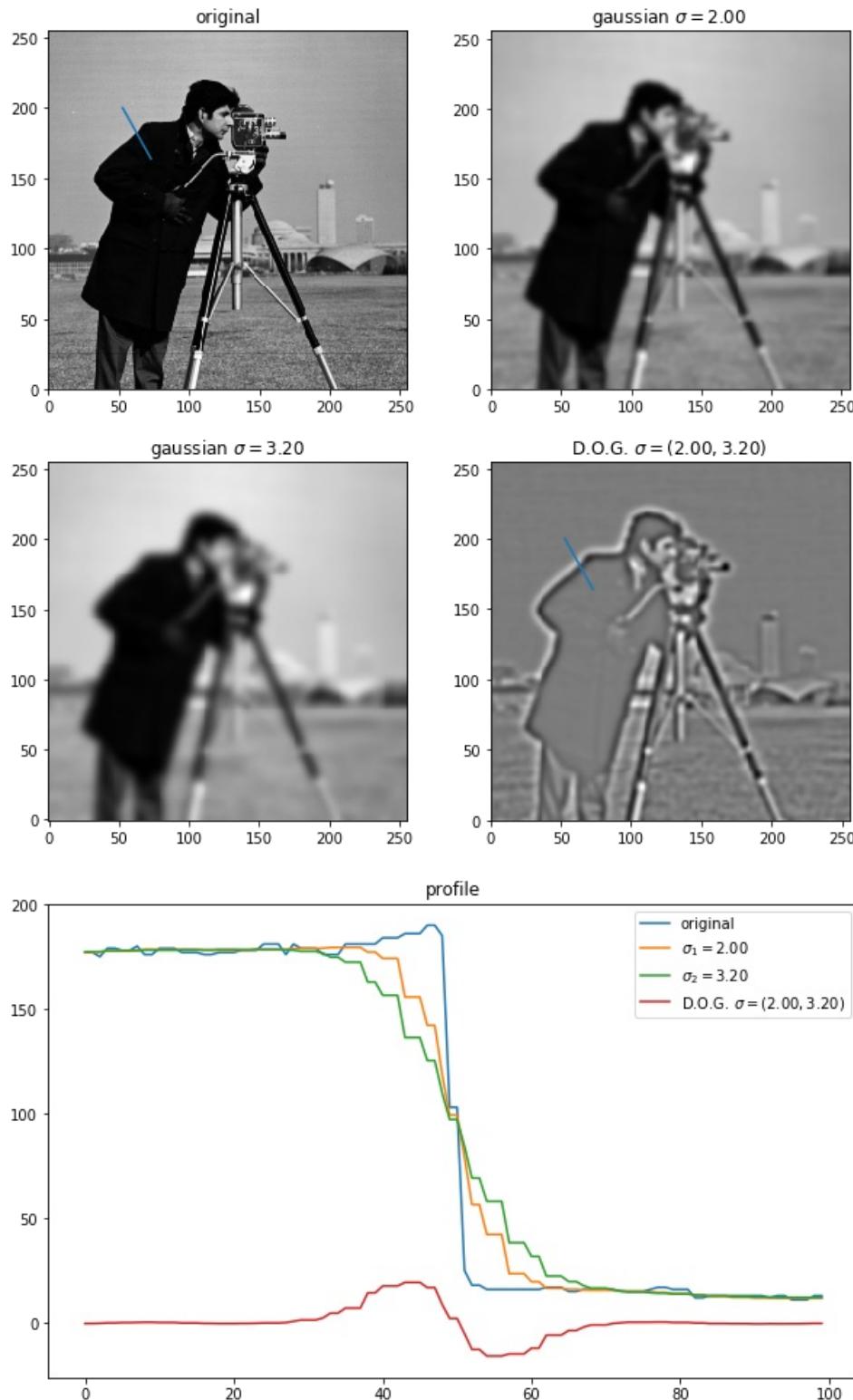
```
im = 1.*camera()[-1::-2,:,:2]

sigma1 = 2.
sigma2 = 1.6*sigma1
g1 = gaussian_filter(im,sigma1)
g2 = gaussian_filter(im,sigma2)

[x,y,p] = profile(im,(53,200),(73,164),100)
[x,y,p_s1] = profile(g1,(53,200),(73,164),100)
[x,y,p_s2] = profile(g2,(53,200),(73,164),100)
[x,y,p_s12] = profile(g1-g2,(53,200),(73,164),100)

plt.figure(figsize=[10,10])
plt.subplot(2,2,1)
plt.imshow(im,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.plot(x,y)
plt.gca().set_xlim(0,255)
plt.gca().set_ylim(0,255)
plt.title('original')
plt.subplot(2,2,2)
plt.imshow(g1,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('gaussian $\sigma=% .2f$ %sigma1')
plt.subplot(2,2,3)
plt.imshow(g2,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('gaussian $\sigma=% .2f$ %sigma2')
plt.subplot(2,2,4)
plt.imshow(1.*g1-g2,interpolation='nearest',cmap=cm.gray,origin='lower')
plt.title('D.O.G. $\sigma=(% .2f,% .2f)$' %(sigma1,sigma2));
plt.plot(x,y)
plt.gca().set_xlim(0,255)
plt.gca().set_ylim(0,255)

plt.figure(figsize=[10,6])
plt.plot(p,label='original')
plt.plot(p_s1,label='$\sigma_1=% .2f$ %sigma1')
plt.plot(p_s2,label='$\sigma_2=% .2f$ %sigma2')
plt.plot(p_s12,label='D.O.G. $\sigma=(% .2f,% .2f)$' %(sigma1,sigma2))
plt.title('profile')
plt.legend(loc=1);
```



Gaussian and Laplacian pyramids

In [13]:

```
from skimage import data
from skimage.transform import pyramid_gaussian,pyramid_laplacian

image = data.astronaut()
rows, cols, dim = image.shape
pyramid = tuple(pyramid_gaussian(image, downscale=2))

composite_image = np.zeros((rows, cols + int(cols/2), 3), dtype=np.double)
composite_image[:rows, :cols, :] = pyramid[0]

i_row = 0
for p in pyramid[1:]:
    n_rows, n_cols = p.shape[:2]
    composite_image[i_row:i_row + n_rows, cols:cols + n_cols] = p
    i_row += n_rows

plt.figure(figsize=[10,10])
plt.imshow(composite_image);
```

/home/olivier/miniconda3/envs/py3/lib/python3.7/site-packages/skimage/transform/_warps.py:23: UserWarning: The default multichannel argument (None) is deprecated. Please specify either True or False explicitly. multichannel will default to False starting with release 0.16.
warn('The default multichannel argument (None) is deprecated. Please '



In [14]:

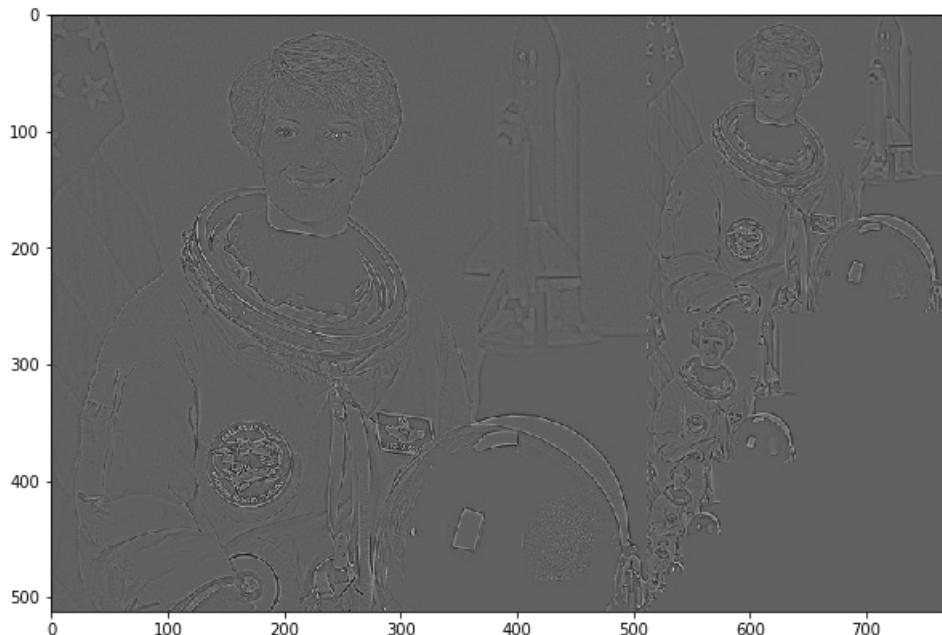
```
pyramid = tuple(pyramid_laplacian(image[:, :, 0], downscale=2))

composite_image = np.zeros((rows, cols + int(cols/2)), dtype=np.double)

composite_image[:rows, :cols] = pyramid[0]

i_row = 0
for p in pyramid[1:]:
    n_rows, n_cols = p.shape[:2]
    composite_image[i_row:i_row + n_rows, cols:cols + n_cols] = p
    i_row += n_rows

plt.figure(figsize=[10,10])
plt.imshow(composite_image,cmap=cm.gray);
```



Canny edge detection

In [15]:

```
from scipy import ndimage
from skimage import feature

# Generate noisy image of a square
im = np.zeros((128, 128))
im[32:-32, 32:-32] = 1

im = ndimage.rotate(im, 15, mode='constant')
im = ndimage.gaussian_filter(im, 4)
im += 0.2 * np.random.random(im.shape)

# Compute the Canny filter for two values of sigma
edges1 = feature.canny(im)
edges2 = feature.canny(im, sigma=3)

# display results
fig, (ax1, ax2, ax3) = plt.subplots(nrows=1, ncols=3, figsize=(8, 3))

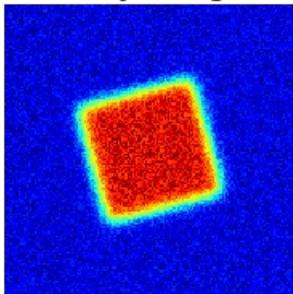
ax1.imshow(im, cmap=plt.cm.jet)
ax1.axis('off')
ax1.set_title('noisy image', fontsize=20)

ax2.imshow(edges1, cmap=plt.cm.gray)
ax2.axis('off')
ax2.set_title('Canny filter, $\sigma=1$', fontsize=20)

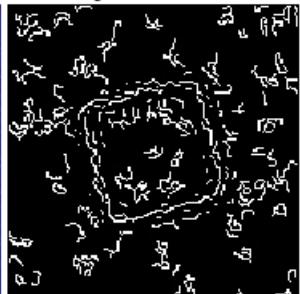
ax3.imshow(edges2, cmap=plt.cm.gray)
ax3.axis('off')
ax3.set_title('Canny filter, $\sigma=3$', fontsize=20)

fig.subplots_adjust(wspace=0.02, hspace=0.02, top=0.9,
                    bottom=0.02, left=0.02, right=0.98)
```

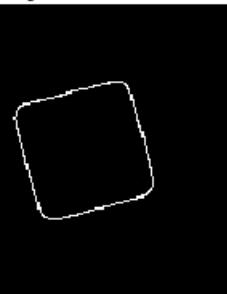
noisy image



Canny filter, $\sigma = 1$



Canny filter, $\sigma = 3$



In [16]:

```
from skimage.morphology import disk
import skimage.filters.rank as skr

# Generate noisy image of a square
im = np.zeros((128, 128))
im[32:-32, 32:-32] = 1

im = ndimage.rotate(im, 15, mode='constant')
im = ndimage.gaussian_filter(im, 4)
im += 0.2 * np.random.random(im.shape) - .1
im[im>1] = 1 #clip image
im[im<0] = 0 #clip image
im = (im*255).astype(np.uint8)
mgrad0 = skr.gradient(im,disk(1))
mgrad1 = skr.gradient(im,disk(3))

# display results
fig, (ax1, ax2, ax3) = plt.subplots(nrows=1, ncols=3, figsize=(8, 3))

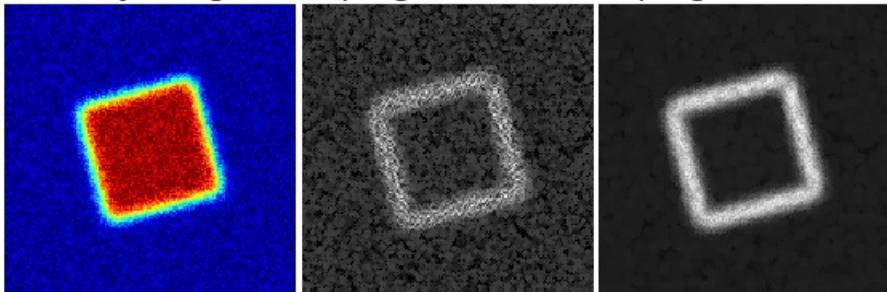
ax1.imshow(im, cmap=plt.cm.jet)
ax1.axis('off')
ax1.set_title('noisy image', fontsize=20)

ax2.imshow(mgrad0, cmap=plt.cm.gray)
ax2.axis('off')
ax2.set_title('morph.gradient, $r=1$', fontsize=20)

ax3.imshow(mgrad1, cmap=plt.cm.gray)
ax3.axis('off')
ax3.set_title('morph.gradient, $r=3$', fontsize=20)

fig.subplots_adjust(wspace=0.02, hspace=0.02, top=0.9,
                    bottom=0.02, left=0.02, right=0.98)
```

noisy image morph.gradient, $r=1$ morph.gradient, $r=3$



Canny edge detection algorithm

1. image smoothing
2. gradient intensity detection
3. local non-maximum suppression
4. double border intensity threshold
5. weak edge suppression

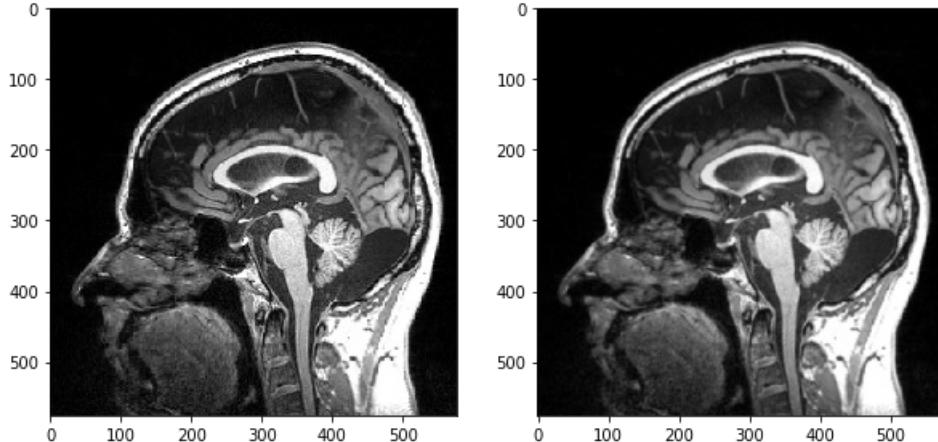
image smoothing

In [17]:

```
from skimage.io import imread
from skimage.filters import gaussian

ct = imread('https://upload.wikimedia.org/wikipedia/commons/5/5f/MRI_EGC_sagittal.png')
plt.figure(figsize=[10,5])
plt.subplot(1,2,1)
plt.imshow(ct);

smooth_ct = gaussian(ct[:, :, 0], 1.)
plt.subplot(1,2,2)
plt.imshow(smooth_ct, cmap=plt.cm.gray);
```



gradient intensity detection

$$\mathbf{G} = \sqrt{\mathbf{G}_x^2 + \mathbf{G}_y^2}$$

$$\Theta = \text{atan2}(\mathbf{G}_y, \mathbf{G}_x)$$

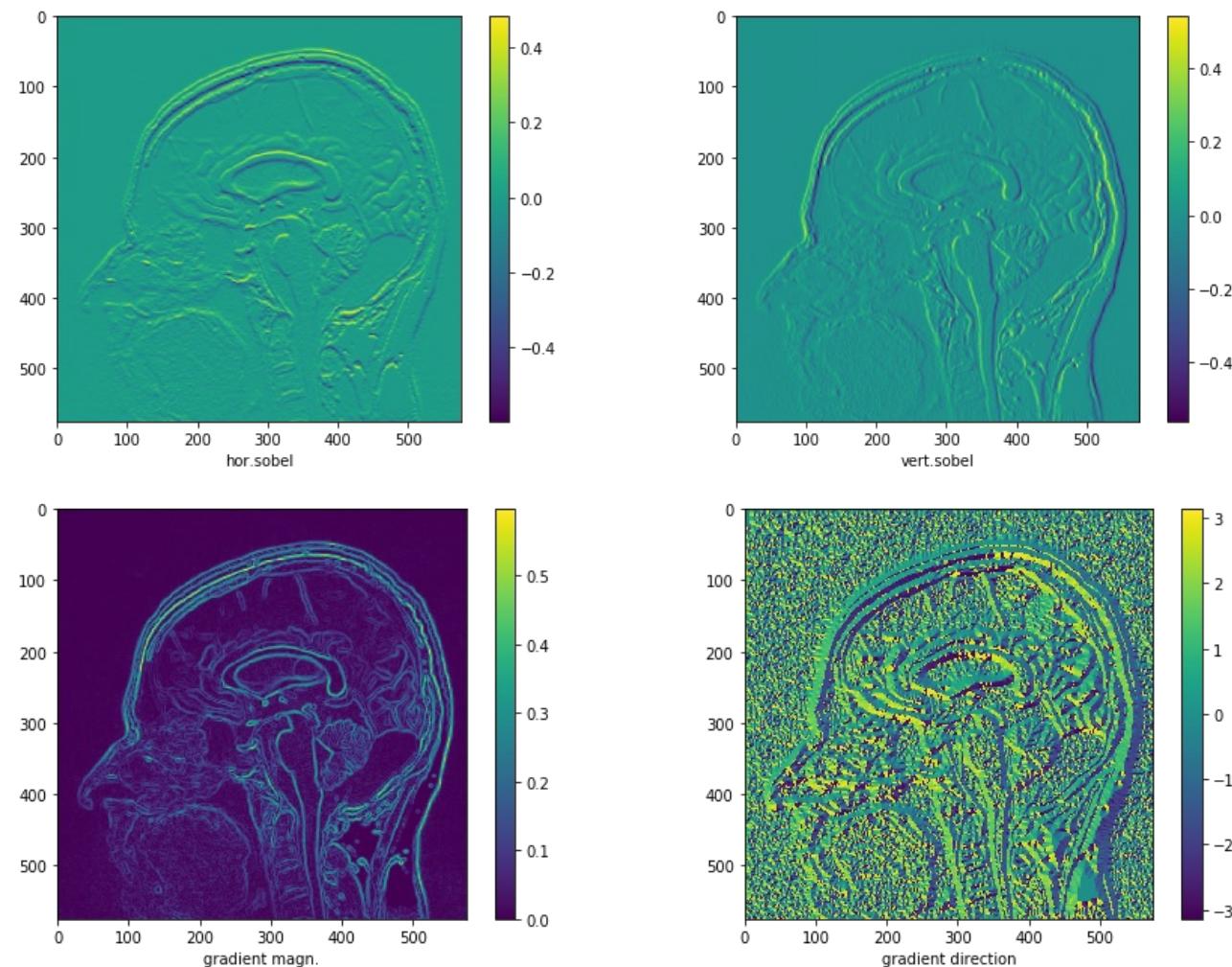
In [18]:

```
from skimage.filters import sobel_h,sobel_v
sh = sobel_h(smooth_ct)
sv = sobel_v(smooth_ct)
plt.figure(figsize=[15,5])
plt.subplot(1,2,1)
plt.imshow(sh)
plt.colorbar();
plt.xlabel('hor.sobel')
plt.subplot(1,2,2)
plt.imshow(sv)
plt.colorbar()
plt.xlabel('vert.sobel')

gm = np.sqrt(sh**2.+sv**2.)

angle = np.arctan2(sv,sh)

plt.figure(figsize=[15,5])
plt.subplot(1,2,1)
plt.imshow(gm)
plt.colorbar();
plt.xlabel('gradient magn.')
plt.subplot(1,2,2)
plt.imshow(angle)
plt.colorbar()
plt.xlabel('gradient direction');
```

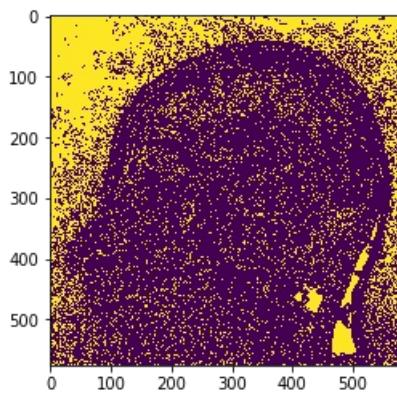


local non-maximum suppression

In [19]:

```
from skimage.morphology import disk
import skimage.filters.rank as skr

local_max = gm*255 >= skr.maximum((gm*255).astype(np.uint8),disk(1))
plt.imshow(local_max);
```



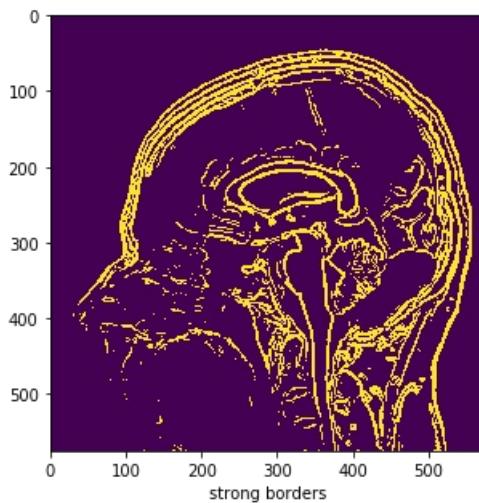
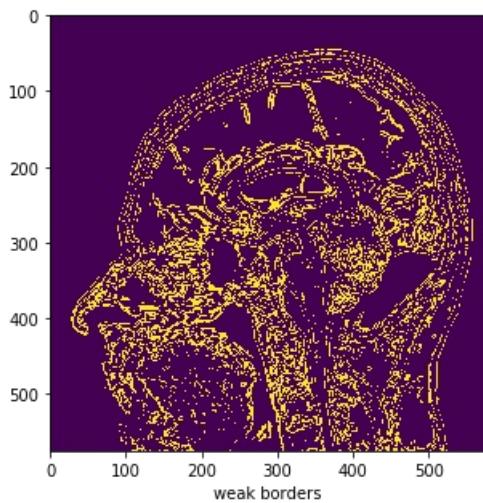
double border intensity threshold

e.g.

- weak border are $\geq 10\%$ of image maximum
- weak border are $\geq 20\%$ of image maximum

In [20]:

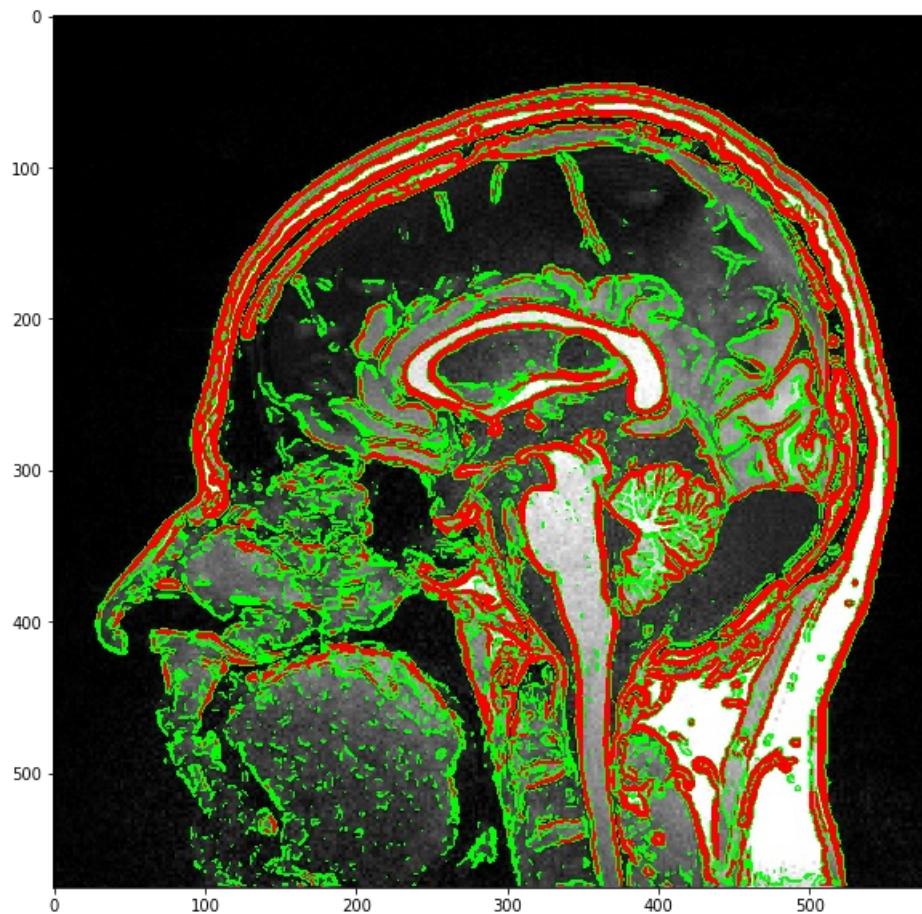
```
image_max = np.max(gm)
weak_borders = np.logical_and(.1*image_max <= gm, gm < .2*image_max)
strong_borders = gm >= .2*image_max
plt.figure(figsize=[15,5])
plt.subplot(1,2,1)
plt.imshow(weak_borders)
plt.xlabel('weak borders')
plt.subplot(1,2,2)
plt.imshow(strong_borders)
plt.xlabel('strong borders');
```



weak edge suppression

In [21]:

```
masked_ct = ct.copy()
masked_ct[weak_borders,:,:]=[0,255,0,255]
masked_ct[strong_borders,:,:]=[255,0,0,255]
plt.figure(figsize=[10,10])
plt.imshow(masked_ct);
```

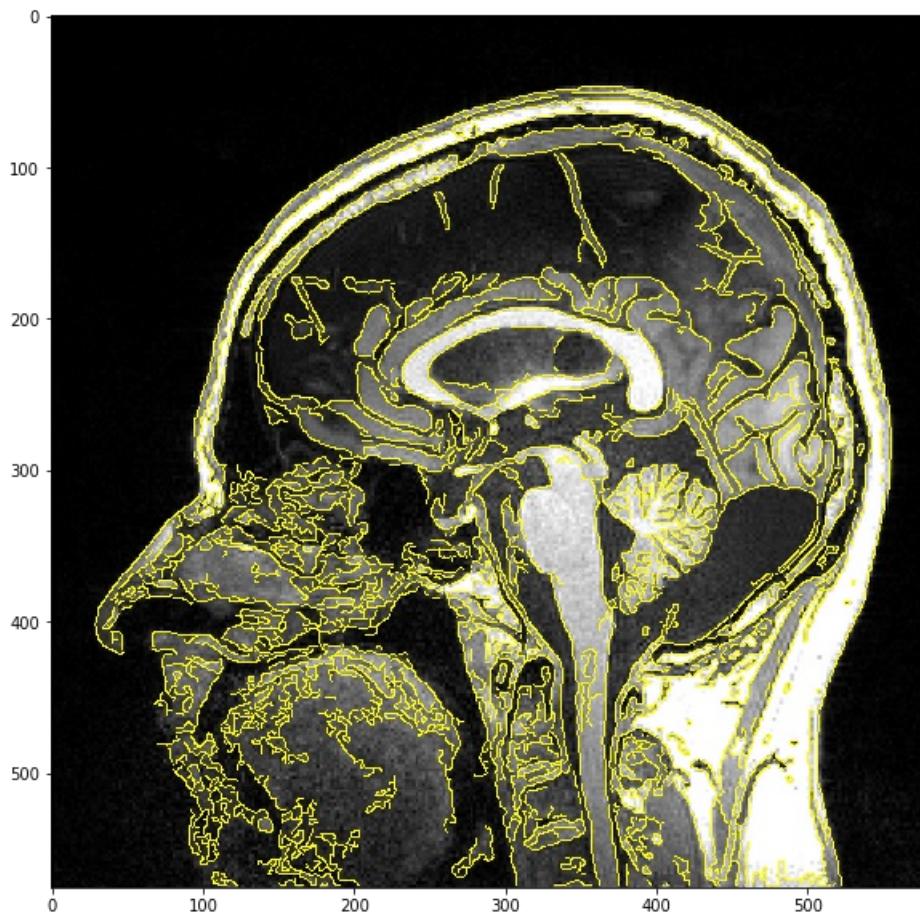


example of canny edge implementation (parameters may differ)

Canny edges overlayed on the original image

In [22]:

```
canny = feature.canny(ct[:, :, 0], low_threshold=.1*255, high_threshold=.4*255)
masked_ct = ct.copy()
masked_ct[canny, :] = [255, 255, 0, 255]
plt.figure(figsize=[10, 10])
plt.imshow(masked_ct);
```

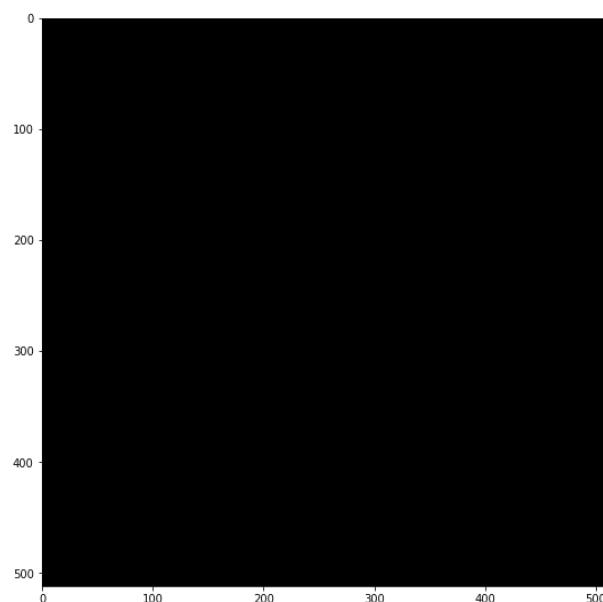


Comparison between canny edges and sobel edges

In [23]:

```
im = camera()
canny = feature.canny(im)*255

plt.figure(figsize=[20, 10])
plt.subplot(1, 2, 1)
plt.imshow(im)
plt.subplot(1, 2, 2)
plt.imshow(canny, cmap=plt.cm.gray);
```

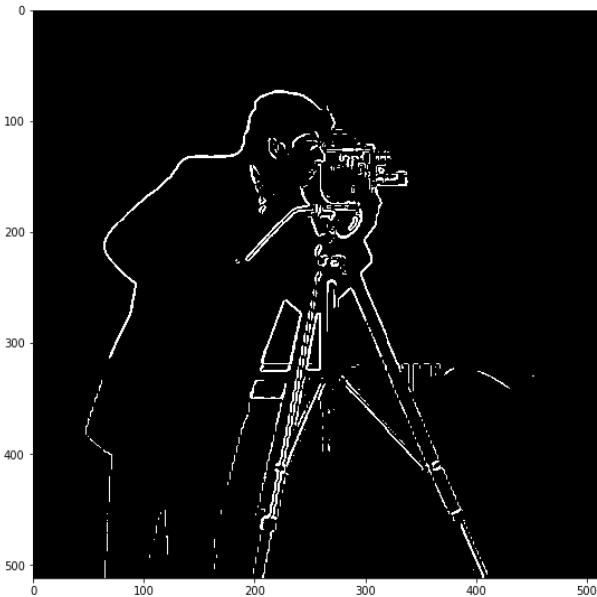
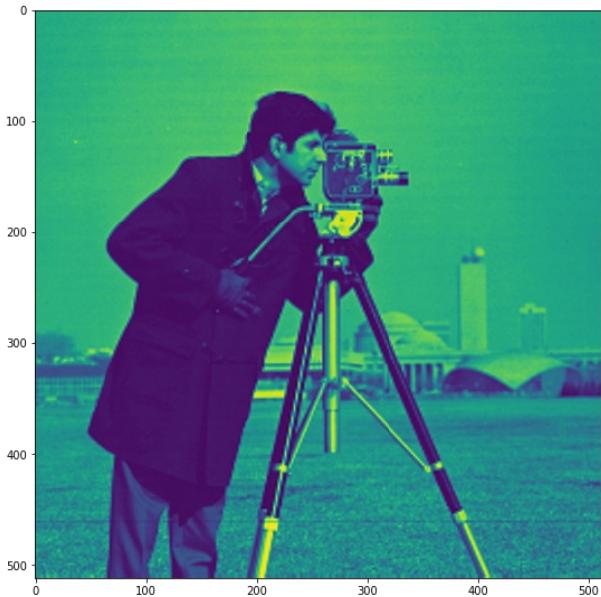


In [24]:

```
im = camera().astype(np.float)
_,_,fsobel = sobel(im)

norm = 255*fsobel/np.max(fsobel)

plt.figure(figsize=[20,10])
plt.subplot(1,2,1)
plt.imshow(im)
plt.subplot(1,2,2)
plt.imshow(norm>100,cmap=plt.cm.gray);
```



In []: