PY410 / 505 Computational Physics 1

Salvatore Rappoccio

Code for this lecture

- Code will be found in
 - -<u>https://github.com/ubsuny/CompPhys/tree/main/</u> DataAnalysis/Fitting

Some documentation for you for today's class

 Statistics derivation is best described in the Particle Data Group :

-<u>http://pdg.lbl.gov/2013/reviews/rpp2012-rev-statistics.pdf</u>

- You can also check the Numerical Recipes if you want (although not necessary, strictly speaking):
 - -<u>http://apps.nrbook.com/empanel/index.html</u>

First Example : Hubble's Law

- As a first physics application, we will study <u>Hubble's Law</u>, and learn how to perform a least squares fit to Edwin Hubble's measurements on extra-galactic nebulae given in his <u>1929 article</u>.
- This is a linear fit, which will give us some intuition about fitting in general

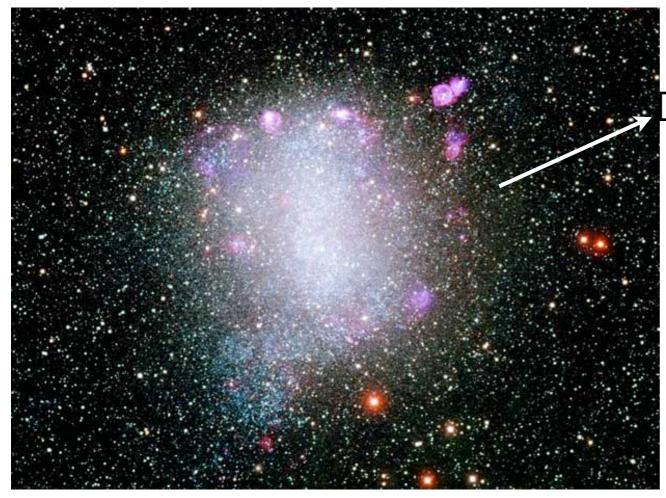
- Galaxies are collections of hundreds of billions of stars –Very enormous!
- Here's a picture from the Hubble telescope released in 2012 showing lots of different galaxies from the Hubble extreme Deep Field
- <u>http://en.wikipedia.org/wiki/</u> <u>Hubble_Extreme_Deep_Field</u>
- This is a photo of an event 13.2 billion years ago, just after the universe underwent inflation!





• And for a comical perspective :

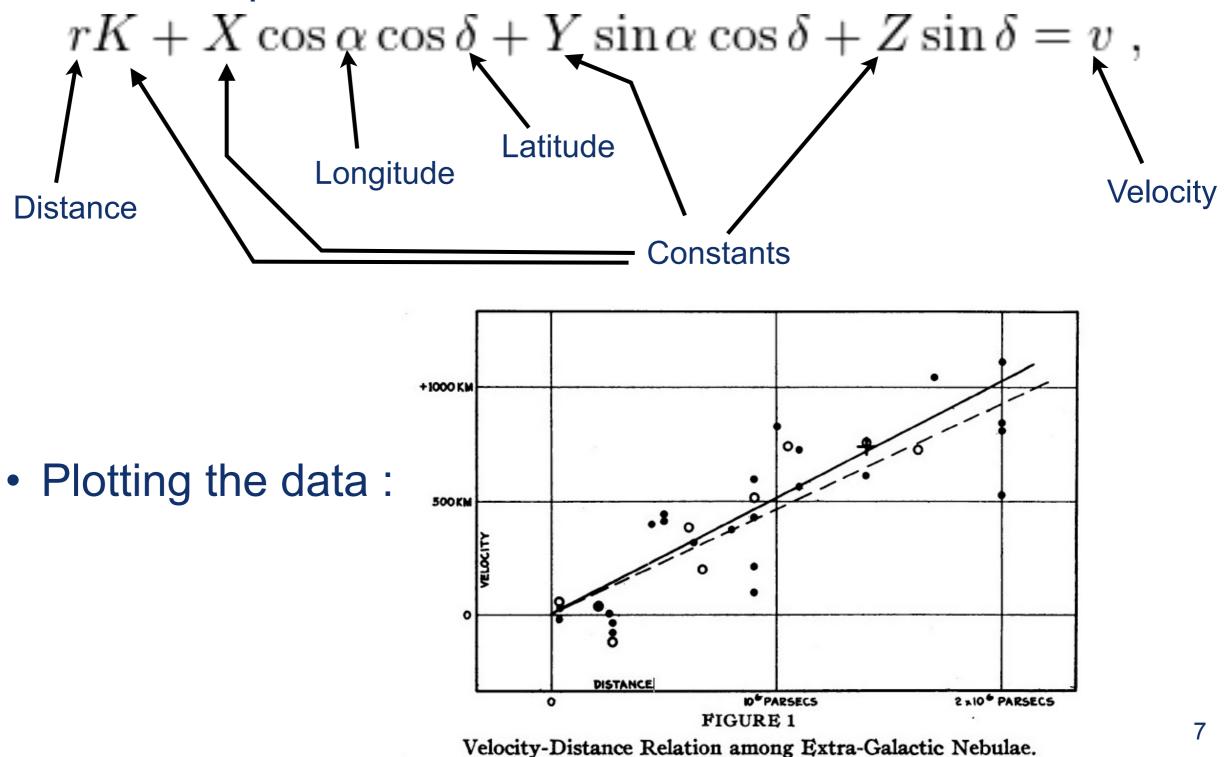
- We're going to analyze the data from the original 1929 paper
- Local Group Galaxy NGC 6822 (Barnard's Galaxy)
- r = 0.214 Mpc (1 Mega-parsec = 3.086x10¹⁹ km) moving towards us with speed v = 130 km/s.



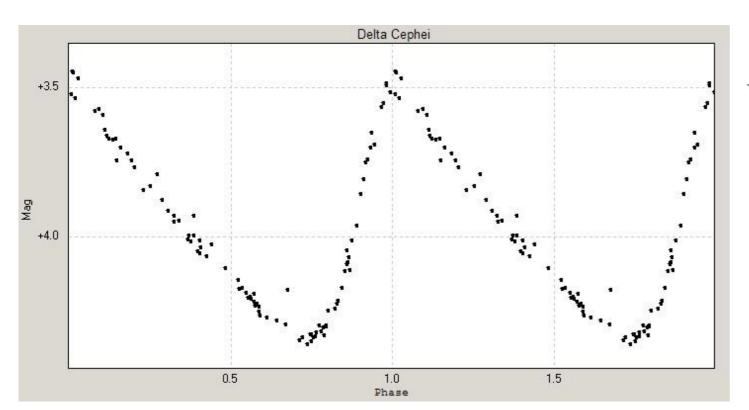
NEBULAE WHOSE DISTANCES HAVE BEEN ESTIMATED FROM STARS INVOLVED OR FROM MEAN LUMINOSITIES IN A CLUSTER					
OBJECT	MEAN LU	MINOSITIES IN	A CLUSTER	<i>m</i> ,	M,
S. Mag.		0.032	+ 170	1.5	-16.0
L. Mag		0.034	+ 290	0.5	17.2
N. G. C. 6822		0.214	- 130	9.0	12.7
598		0.263	- 70	7.0	15.1
221		0.275	- 185	8.8	13.4
224		0.275	- 220	5.0	17.2
5457	17.0	0.45	+ 200	9.9	13.3
4736	17.3	0.5	+ 290	8.4	15.1
5194	17.3	0.5	+ 270	7.4	16.1
4449	17.8	0.63	+ 200	9.5	14.5
4214	18.3	0.8	+ 300	11.3	13.2
3031	18.5	0.9	- 30	8.3	16.4
3627	18.5	0.9	+ 650	9.1	15.7
4826	18.5	0.9	+ 150	9.0	15.7
5236	18.5	0.9	+ 500	10.4	14.4
1068	18.7	1.0	+ 920	9.1	15.9
5055	19.0	1.1	+450	9.6	15.6
7331	19.0	1.1	+ 500	10.4	14.8
4258	19.5	1.4	+ 500	8.7	17.0
4151	20.0	1.7	+ 960	12.0	14.2
4382		2.0	+ 500	10.0	16.5
4472		2.0	+ 850	8.8	17.7
4486		2.0	+ 800	9.7	16.8
4649		2.0	+1090	9.5	17.0

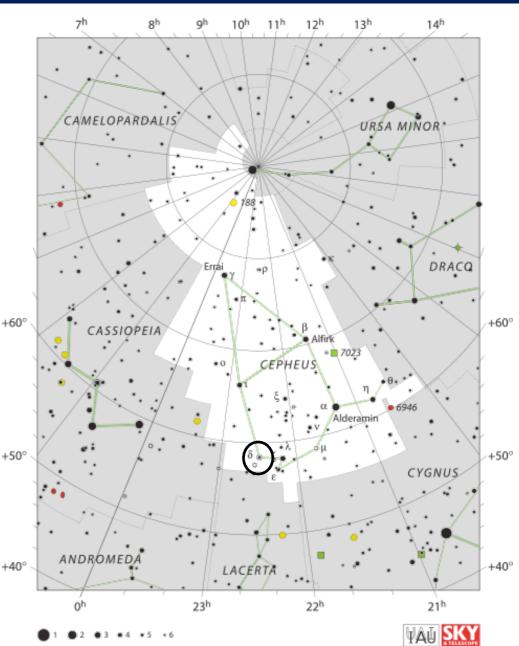
TABLE 1

• Hubble used this equation to determine a linear relationship :



- How do we get the luminosity for distant objects?
- Use Cepheid Variables!
 - -http://en.wikipedia.org/wiki/Cepheid_variable
- Luminosity of the star can be estimated from its period!
 - -Period is very easy to measure
 - -Convert to luminosity
- For instance : Delta Cephei:
 - -<u>http://en.wikipedia.org/wiki/Delta_Cephei</u>



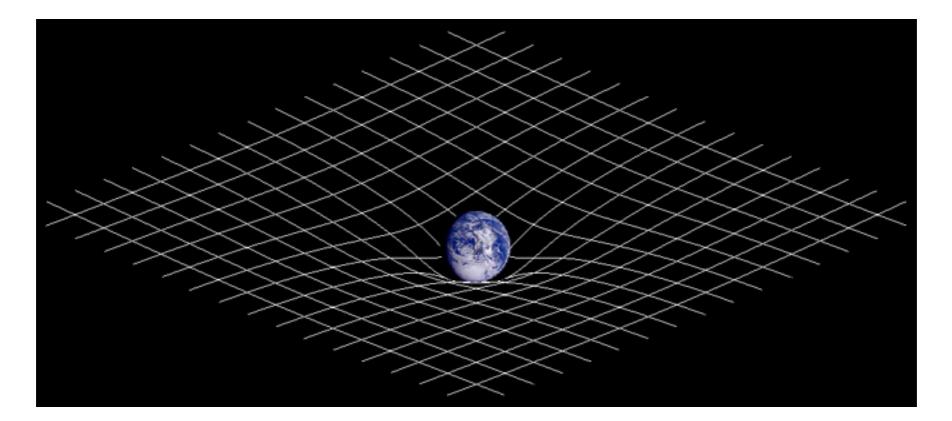


- Why do we expect that the further the distance of the galaxy, the faster they should be moving away from us?
- A priori, no reason
- It just happens to be so in our universe!

• So, now for a bit of general relativity and cosmology

General Relativity

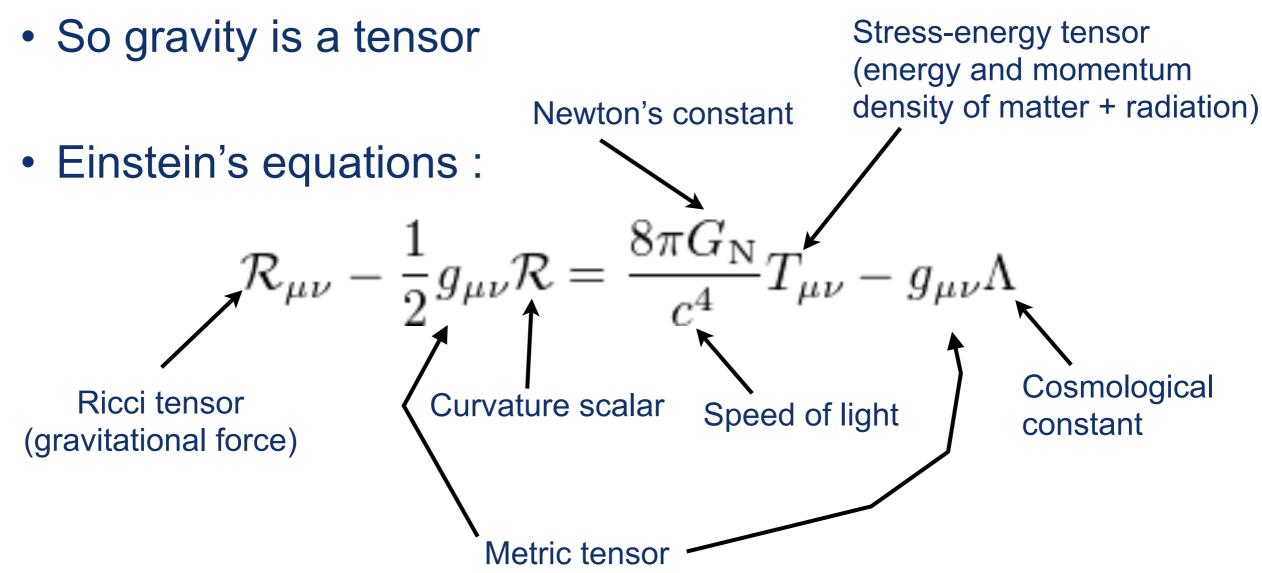
- <u>http://en.wikipedia.org/wiki/General_relativity</u>
- Relates gravity to the curvature of space-time!



Objects with mass or energy distort space-time, and this induces a gravitational field

General Relativity

Space-time is a tensor



General Relativity

- That's a huge set of nonlinear partial differential equations, and can be arbitrarily complicated ($T_{\mu\nu}$ has no constraint to its format)
- A few simple cases can be derived :
 - -If spacetime is homogeneous and isotropic, this is the Robertson-Walker metric : Cosmological scale factor

$$ds^2 \equiv \sum_{\mu=0}^3 \sum_{\nu=0}^3 g_{\mu\nu} dx^{\mu} dx^{\nu} = c^2 dt^2 - R^2(t) \left[\frac{dr^2}{1 - kr^2} + r^2 (d\theta^2 + \sin^2 \theta d\phi^2) \right] \; ,$$

–Assuming that the matter+radiation behave like a uniform perfect fluid with density ρ and pressure p, this is the Friedmann-Lamaitre equations:

$$H^{2} \equiv \left(\frac{\dot{R}}{R}\right)^{2} = \frac{8\pi G_{\rm N}\rho}{3} - \frac{kc^{2}}{R^{2}} + \frac{\Lambda c^{2}}{3}, \qquad \frac{\ddot{R}}{R} = -\frac{4\pi G_{\rm N}}{3}(\rho + 3p) + \frac{\Lambda c^{2}}{3}.$$

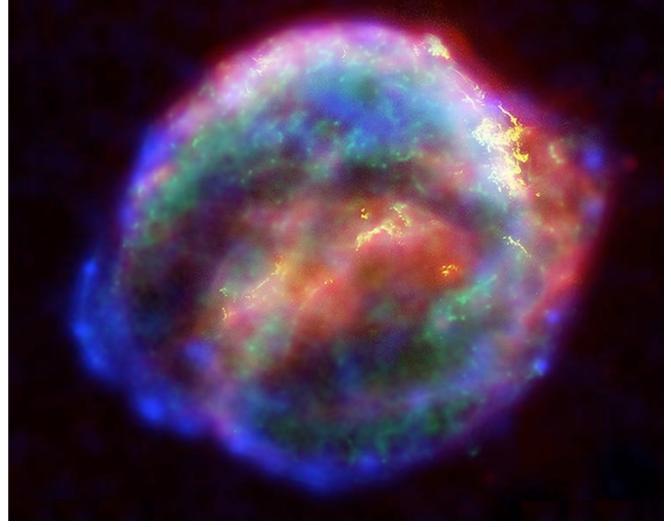
Hubble parameter : H(t0) = 72 km/s/Mpc at present time

- Supernovae occur when a star exhausts its hydrogen fuel, and blows off the outer shell
- It reduces in size, but the Pauli exclusion principle prevents collapse

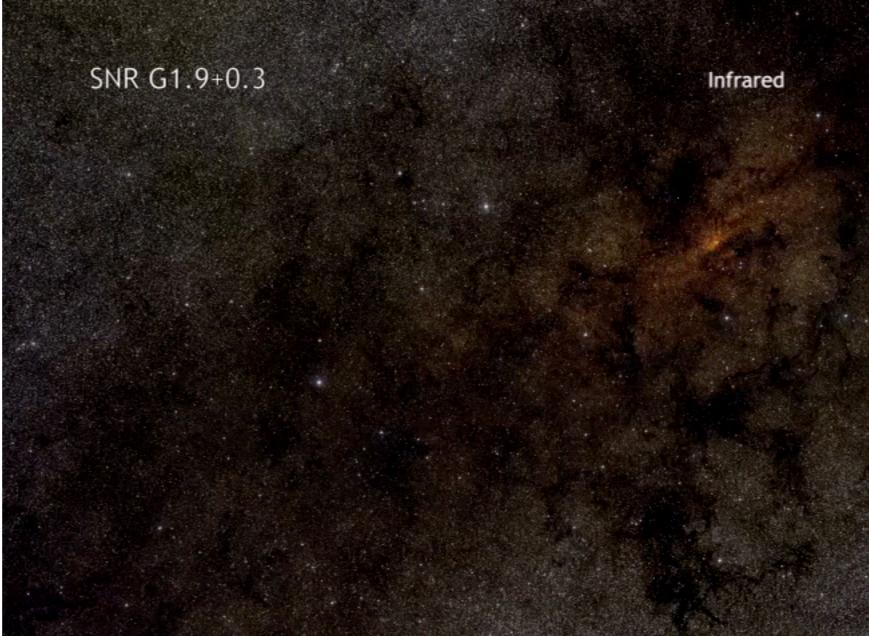
-White dwarf

- White dwarf then accretes material from nearby stars
- The core explodes in a thermonuclear event
- That's the supernova!
- This emits light at specific frequencies, which can be used to estimate the distance!

SN 1604 (discovered by Johannes Kepler)



- Supernovae occur when a star exhausts its hydrogen fuel, and blows off the outer shell
- It reduces in size, but the Pauli exclusion principle prevents collapse —White dwarf
- White dwarf then accretes material from nearby stars
- The core explodes in a thermonuclear event
- That's the supernova!
- This emits light at specific frequencies, which can be used to estimate the distance!



http://chandra.harvard.edu/photo/2013/g19/g19_BU_sm_web.mov

- PDG's Review of big bang cosmology has a nice set of data
 :
- <u>http://pdg.lbl.gov/2012/reviews/rpp2012-rev-bbang-</u> <u>cosmology.pdf</u>
- Brightness is measured by absolute magnitude "M"
- Apparent magnitude is "m"
- M is equal to m at 10 pc
- r is distance in pc
- Distance modulus is :

$$\mu \equiv m - M = 5 \log_{10} r - 5 \; ,$$

• Luminosity distance is :

$$D_L = 10^{\frac{(m-M)}{5}+1}$$

The distance modulus is approximately (for distant SN's)

$$\mu = 25 + 5\log_{10}\left(\frac{cz}{H_0}\right) + 1.086(1-q_0)z + \dots$$

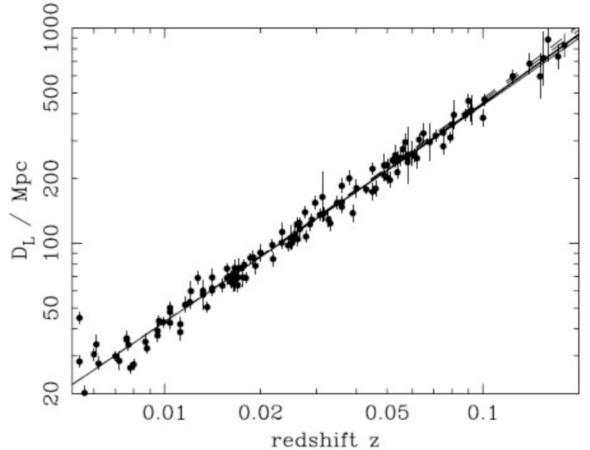
• Combine with G.R. doppler shift :

$$z+1 = rac{
u_1}{
u_2} = rac{R_2}{R_1} \simeq 1 + rac{
u_{12}}{c}$$

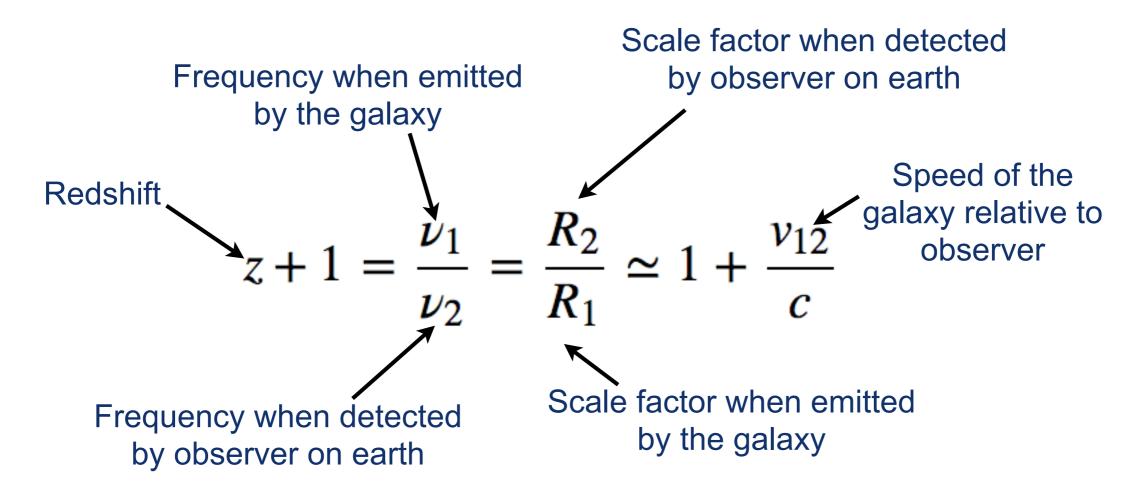
- We conclude that faster objects
 have more redshift!
- There is a linear relationship between brightness and redshift for supernovae!



(Careful : we will use the distance modulus and not luminosity distance since that's what we have data for) 16



- Specifics don't matter here, but we just want to state the relation of redshifts of galaxies to their velocities
- Obtained from G.R. doppler shift!

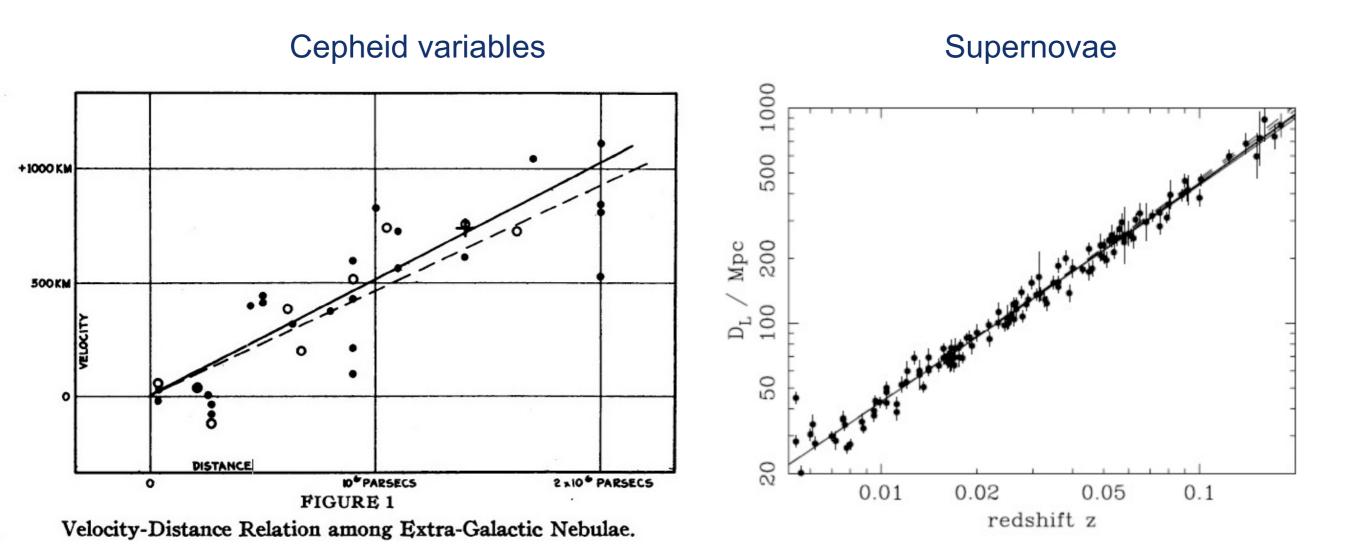


I don't expect you to be able to derive this, but we'll just fit the data

Recall : Development

- Step 1 : Write the algorithm down on paper
- Step 2 :
 - -If you don't understand everything : goto step 1
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- Step 4 : continue
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 - -else : continue
- Step 5 : write code
- Step 6 : check code with unit tests
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 - -Check "fail" criterion
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- Step 7 : Publish!

- Want to fit a line to a bunch of points
- Let's think for a bit about what this means and how we should expect to implement it



• Think about the simplest case : 1 point.

-What happens here?

• Think about the simplest case : 1 point.

-What happens here?

- Nothing! You can't fit a line to a point.
 - So, this should be checked before we attempt to fit anything

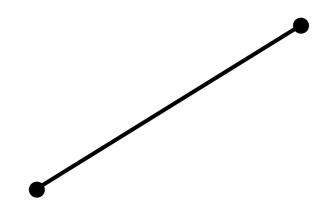
• OK, next simplest case : 2 points

-What happens here?

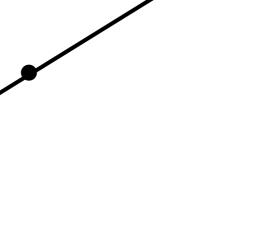
• OK, next simplest case : 2 points

-What happens here?

• That's easy too : there's no fit, you just draw a line



- What about 3 points?
 –Now this gets interesting!
- There's degeneracy that you can exploit
- If the three points are colinear, this works

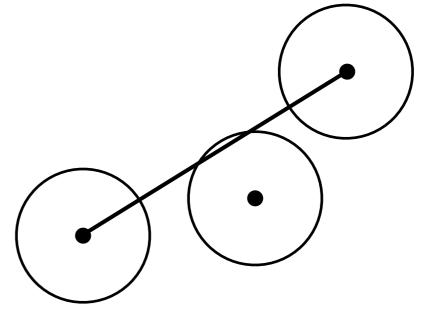


- What about 3 points?
 –Now this gets interesting!
- There's degeneracy that you can exploit
- If the three points are colinear, this works
- Otherwise, it doesn't!

Cannot draw a line to connect these three!

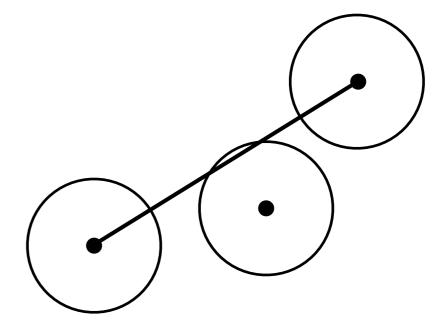
- So what do we do? We can't just give up!
- Very important aspect to remember :
 - -These points are not points : they are actually ellipses!
 - -They come with uncertainties!

 The uncertainties can come from two sources : –Statistical sampling



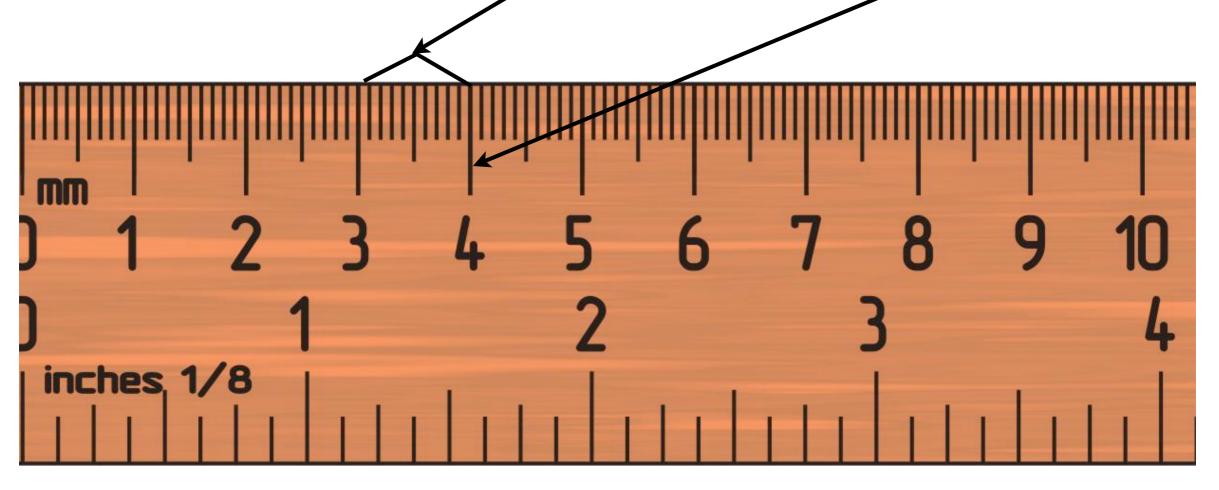
-Systematic effects

- The uncertainties can come from two sources :
 - -Statistical sampling
 - From variations in repeated trials
 - Mathematically "well-behaved"
 - Easy to estimate



- -Systematic effects
 - Intrinsic uncertainty from non-deterministic sources
 - Not mathematically "well-behaved"
 - Difficult to estimate

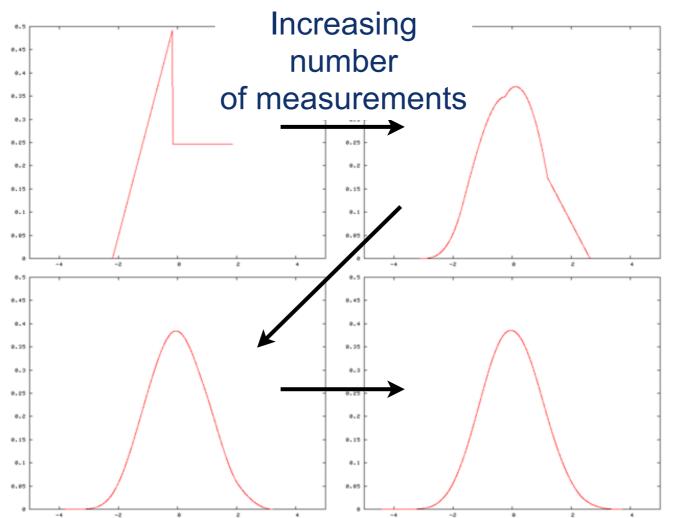
- Example : measuring distance with a ruler
 - -Systematic limitation : ruler has finite width of the lines, and finite number of lines to measure!



-Statistical variation : you can try to repeat the same measurement over and over to get a better estimate of the "true" value

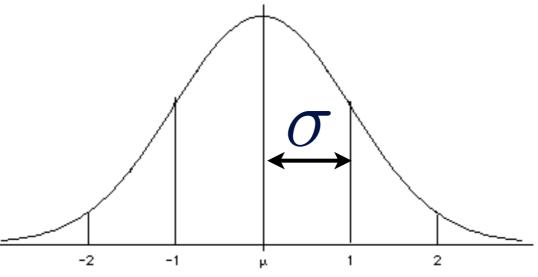
- Easy one first : statistical uncertainties
- Problem stated :
 - –We have a true value $\widetilde{\mathcal{X}}$
 - -We have several measured values $x_0, x_1, \ldots x_{n-1}$
 - -How well can we estimate \tilde{x} given $x_0, x_1, \ldots x_{n-1}$?

- Central limit theorem!
 - -http://en.wikipedia.org/wiki/Central_limit_theorem
- If your measurements are uncorrelated :
 - -As you make more measurements, they follow a Gaussian (or "normal", or bell-shaped curve) distribution
 - -http://en.wikipedia.org/wiki/Normal_distribution



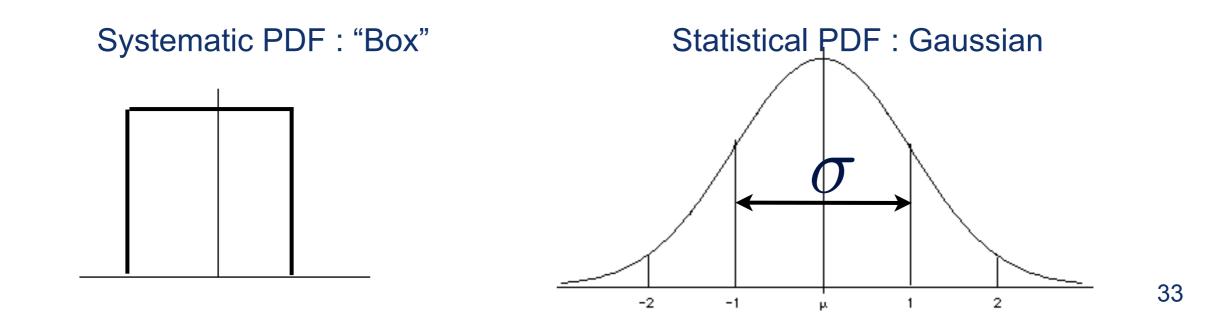
Called a "probability distribution function" (or PDF)

- So, the distribution of values will follow a Gaussian distribution for statistical uncertainties
- We usually quote the "sigma" (σ) of the Gaussian as the uncertainty band



- What about systematic effects?
 - -If you repeat the trial again and again, what happens?
 - –It's a systematic effect, so you actually get basically the same thing!

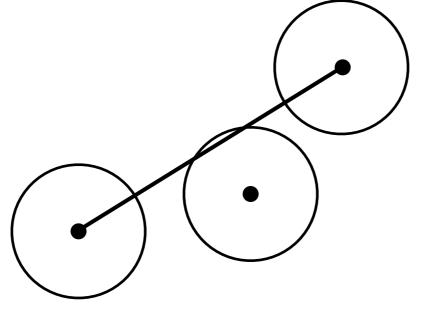
- Systematic uncertainties are very hard to estimate
- Typically we (as scientists) "reckon" them in some way
 - -Control samples
 - -Minimum resolution of your device
 - -And so on
- So, the probability distribution function for these are basically FLAT —You don't know where it is, but it's somewhere within that range



- You now say : "Sal, I thought you were supposed to teach me about programming? I haven't seen a line of code yet!"
- "Ah, my good students," I reply. "But remember we must understand what we're doing!"

- So why am I telling you all of this?
 - The systematic effects don't follow a mathematical formalism
 - -The statistical effects do follow a mathematical formalism
 - –So, we usually just pretend that systematic effects are like statistical effects, and assume Gaussian uncertainties too!

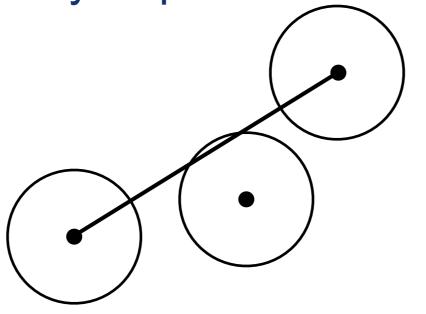
- So, back to linear fits!
- What the heck are these circles?
 - -They're the uncertainties!



- We'll just pretend that they're statistical
 - -Scientists usually pretend they're Gaussian anyway!

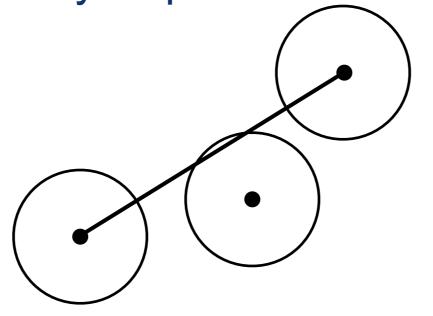
So now, what do we actually want to do?

We want to draw the line that intersects all of the possible uncertainty ellipses :



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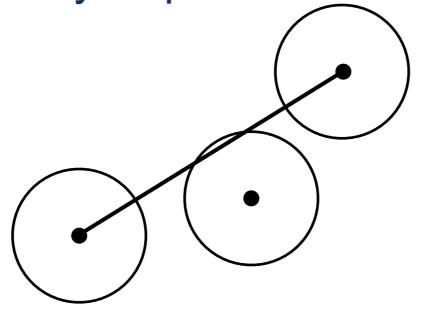
We want to draw the line that intersects all of the possible uncertainty ellipses :



BAD : we're ignoring the third point entirely! So what to do?

So now, what do we actually want to do?

We want to draw the line that intersects all of the possible uncertainty ellipses :

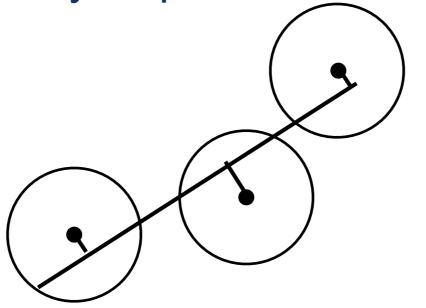


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Minimize the least-squares distance!

So now, what do we actually want to do?

We want to draw the line that intersects all of the possible uncertainty ellipses :



Minimize the least-squares distance!

- So what exactly do we want to compute, and what do we want to minimize?
- Assume the data are described by

$$y(x) = a + bx$$

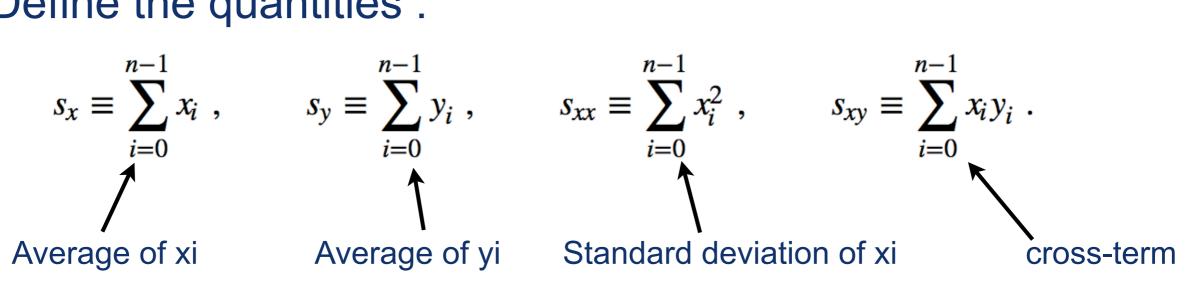
- Presume first that all of the uncertainties are exactly the same
- Then you just have to minimize the distance :

$$f(a,b) \equiv \sum_{i=0}^{n-1} \left(y_i - a - b x_i \right)^2$$

• At the minimum, the derivatives are zero:

$$\frac{\partial f}{\partial a} = -2\sum_{i=0}^{n-1} (y_i - a - bx_i) = 0 , \quad \text{and} \quad \frac{\partial f}{\partial b} = -2\sum_{i=0}^{n-1} x_i (y_i - a - bx_i) = 0$$

- Can solve these simultaneously for the two unknowns a and b
 - -Two equations, two unknowns!
- Define the quantities :



• With this definition, can compute a and b :

$$a = \frac{s_{xx}s_y - s_x s_{xy}}{ns_{xx} - s_x^2} , \qquad b = \frac{ns_{xy} - s_x s_y}{ns_{xx} - s_x^2}$$

- -This algorithm is discussed in <u>Section 15.2: Fitting data</u> to a straight line of <u>Numerical Recipes</u>.
- -(you can access a certain number of pages per month for free... but in any case, you don't need this if you don't want to use it)

- But! We're not quite done.
- What are the uncertainties on a and b?
- Actually, what we want is the uncertainty per degree of freedom of the fit
- Degrees of freedom is : number of data points - number of constraints
- For a linear fit we have two constraints
- Variance is therefore :

$$\sigma^2 \equiv \frac{f(a,b)}{\nu} = \frac{1}{n-2} \sum_{i=0}^{n-1} (y_i - y(x_i))^2 .$$

- What if the uncertainties are not all equal?
- The same principle applies, but instead of minimizing the mean-squared distance :

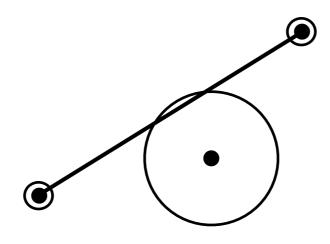
$$f(a,b) \equiv \sum_{i=0}^{n-1} \left(y_i - a - b x_i \right)^2$$

 You instead minimize the "chi-squared" which is the distance divided by the uncertainty :

$$\chi^2(a,b) \equiv \sum_{i=0}^{n-1} \left(\frac{y_i - a - bx_i}{\sigma_i} \right)^2$$

Note : This is only strictly true for GAUSSIAN uncertainties!

- How does this help?
- It "ignores" values with large uncertainties :



- The two points on the ends are very precise
- The third point in the middle is not precise
- Good point to check for a unit test!

- So how does this get modified?
- Parameters and uncertainties are :

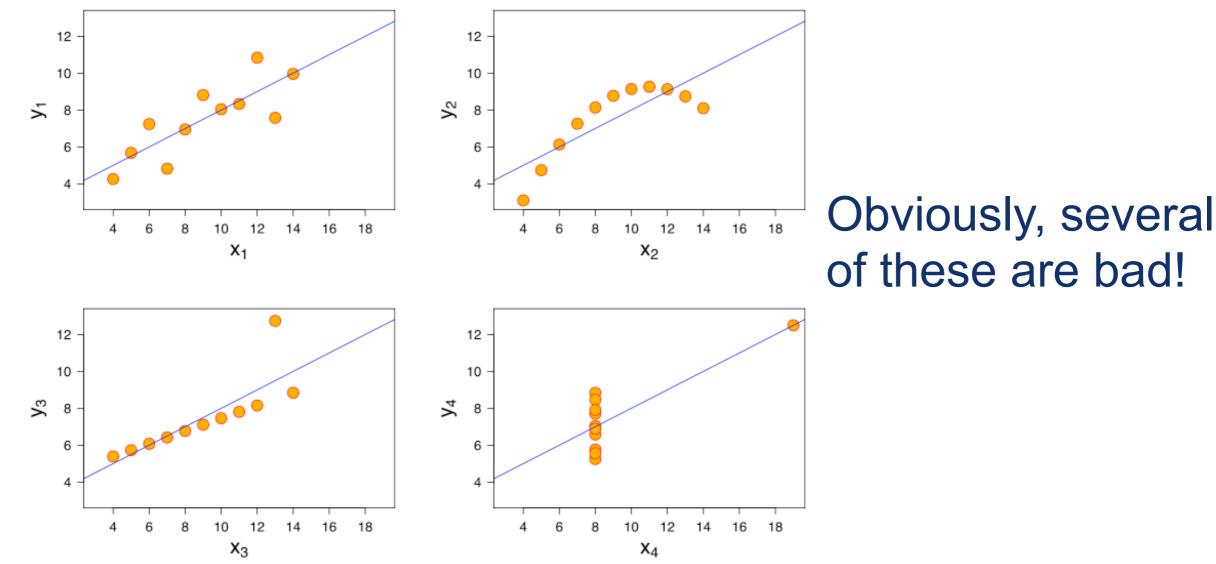
$$b = \frac{1}{S_{tt}} \sum_{i=0}^{n-1} \frac{t_i y_i}{\sigma_i} , \qquad a = \frac{S_y - S_x b}{S} \qquad \qquad \sigma_a^2 = \frac{1}{S} \left(1 + \frac{S_x^2}{SS_{tt}} \right) , \qquad \sigma_b^2 = \frac{1}{S_{tt}}$$

• Where :

$$t_i = \frac{1}{\sigma_i} \left(x_i - \frac{S_x}{S} \right) , \qquad S_{tt} = \sum_{i=0}^{n-1} t_i^2 ,$$

$$S = \sum_{i=0}^{n-1} \frac{1}{\sigma_i^2}, \qquad S_x = \sum_{i=0}^{n-1} \frac{x_i}{\sigma_i^2}, \qquad S_y = \sum_{i=0}^{n-1} \frac{y_i}{\sigma_i^2}.$$

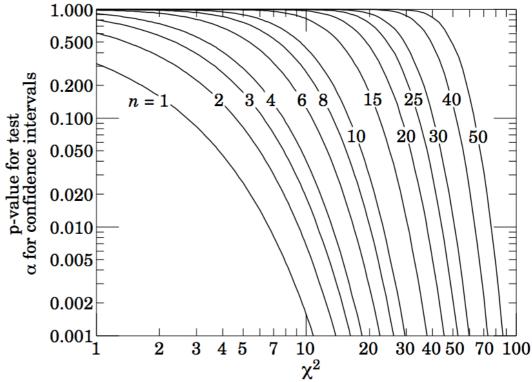
- With uncertainties on the inputs, you can compute "goodness of fit"
- Why do you care?
- All of these have the same fit :



- The chi-squared per degree of freedom is what we're looking for here $\chi^2/\text{d.o.f} \equiv \frac{1}{n-2} \sum_{i=0}^{n-1} \left(\frac{y_i a bx_i}{\sigma_i}\right)^2 \approx 1 \ .$
- Roughly speaking, it's the number of "standard deviations" that you're "off" in the fit

About 1 S.D. off :

 If you've estimated your uncertainties correctly, it should follow a distribution of values called the "chi-squared" distribution :



- **Figure 36.1:** One minus the χ^2 cumulative distribution, $1 F(\chi^2; n)$, for *n* degrees of freedom. This gives the *p*-value for the χ^2 goodness-of-fit test as well as one minus the coverage probability for confidence regions (see Sec. 36.3.2.4).
- <u>http://en.wikipedia.org/wiki/Chi-squared_distribution</u>
- <u>http://pdg.lbl.gov/2013/reviews/rpp2012-rev-statistics.pdf</u>

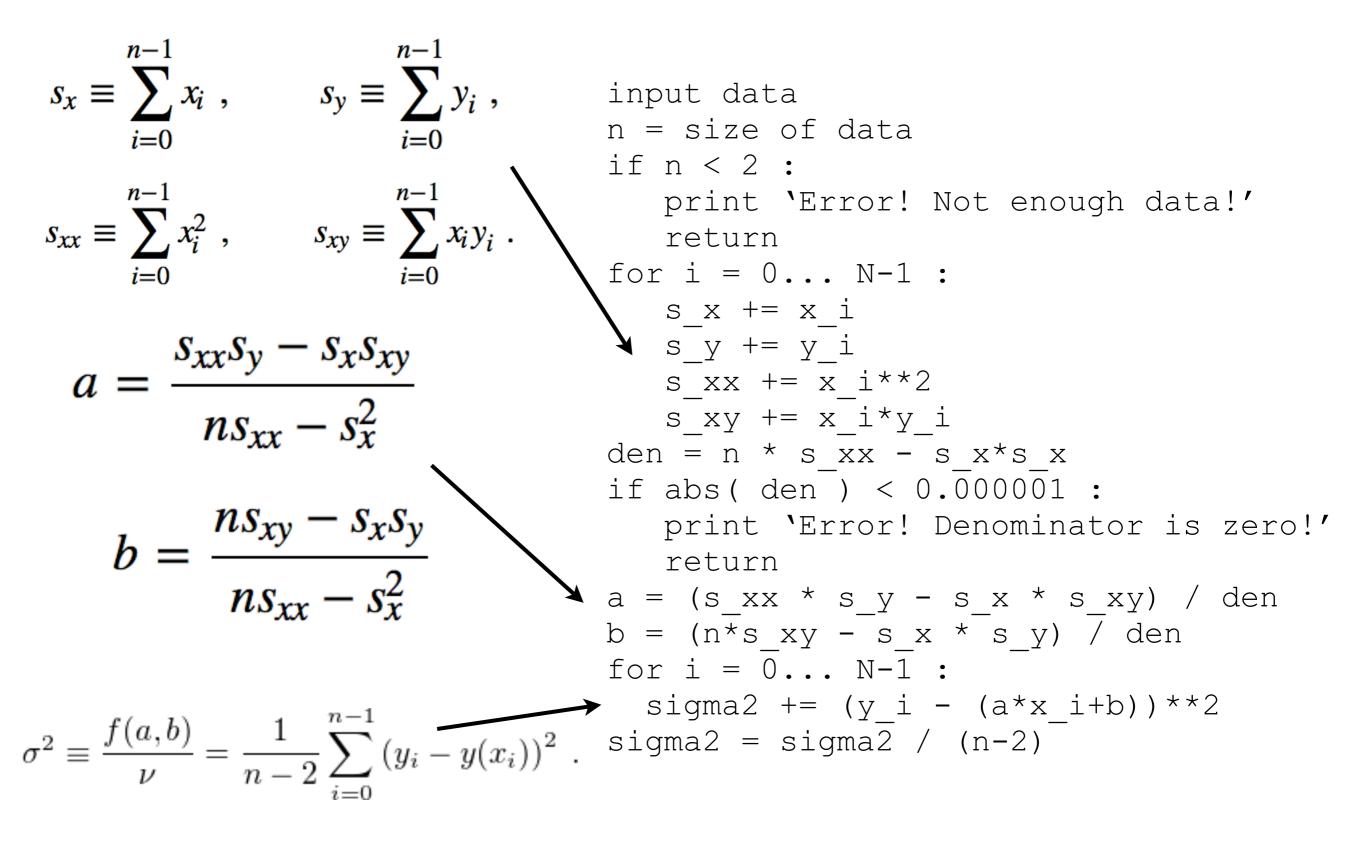
- OK : moment of truth
- We have two cases : with, and without uncertainties
- Let's do without first, it's easier

- Step 1 : Write the algorithm down on paper
- Step 2 :

-If you don't understand everything : goto step 1

- -else : continue
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Pseudocode : No uncertainties



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Write code : No uncertainties

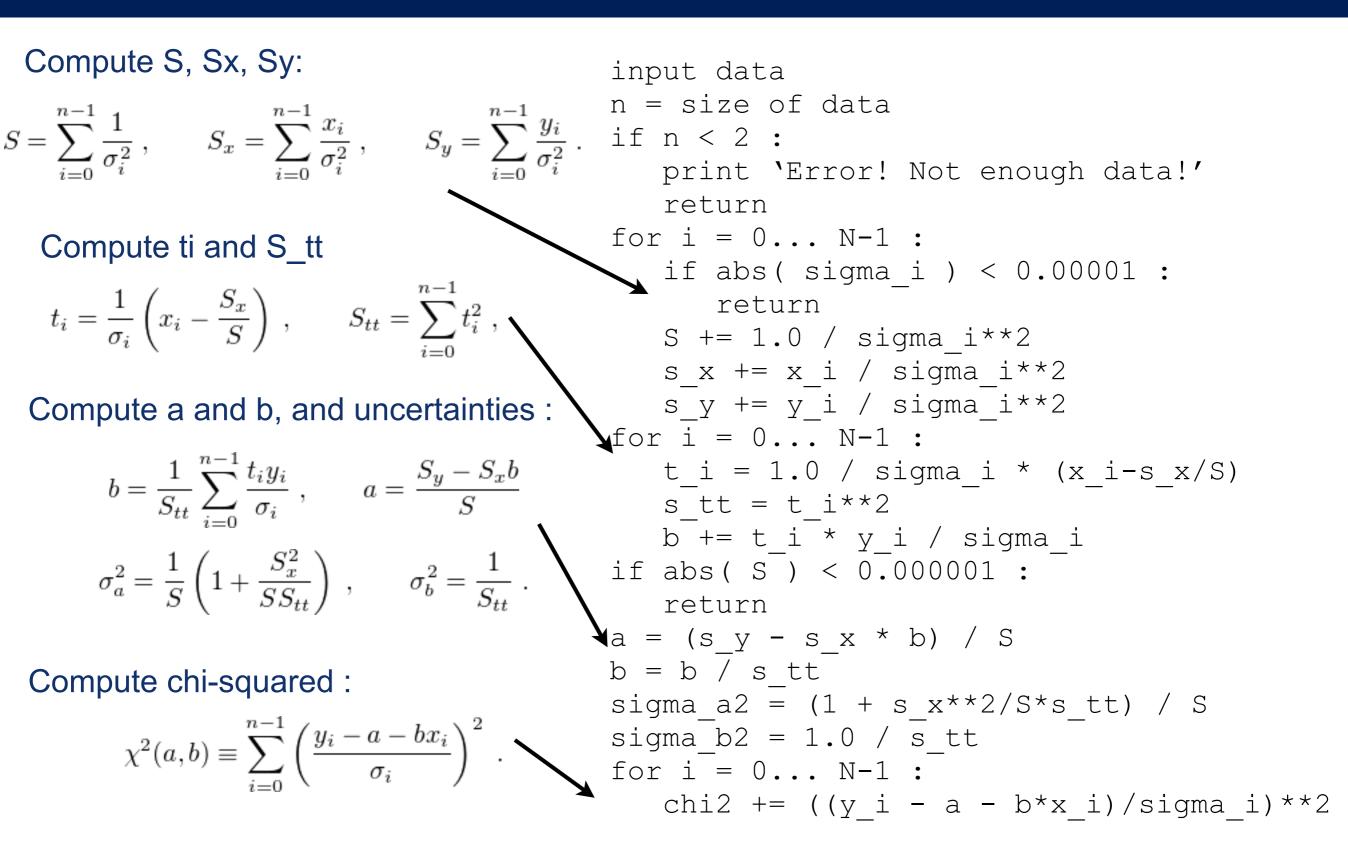
```
python
             Pseudocode
                                                     n = len(x) # number of galaxies
input data
                                                      if n <= 2 :
n = size of data
                                                         print ('Error! Need at least two data points!')
if n < 2:
                                                         exit()
   print 'Error! Not enough data!'
                                                      # Compute all of the stat. variables we need
   return
for i = 0... N-1 :
                                                      s x = np.sum(x)
                                                      s y = np.sum(y)
   s x += x i
                                                      s xx = np.sum(x**2)
   s y += y_i
                                                     s xy = np.sum(x*y)
   s xx += x i**2
                                                     denom = n * s xx - s x**2
   s xy += x i*y i
                                                     if abs( denom ) < 0.000001 :
den = n * s xx - s x*s x
                                                         print ('Error! Denomominator is zero!')
                                                         exit()
if abs(den) < 0.00001:
   print 'Error! Denominator is zero!'
                                                      # Compute y-intercept and slope
   return
                                                     a = (s xx * s y - s x * s xy) / denom
a = (s_x x * s_y - s x * s xy) / den
                                                     b = (n*s xy - s x * s y) / denom
b = (n \cdot s xy - s x \cdot s y) / den
                                                     # Compute uncertainties
for i = 0... N-1:
                                                      if n > 2:
  sigma2 += (y i - (a*x i+b))**2
                                                           sigma = np.sqrt(np.sum((y - (a+b*x))**2) / (n-2))
sigma2 = sigma2 / (n-2)
                                                           sigma a = np.sqrt(sigma**2 * s xx / denom)
                                                           sigma b = np.sqrt(sigma**2 * n / denom)
                                                      else :
                                                           sigma = 0.
                                                           sigma a = 0.
                                                           sigma b = 0.
                                                      return [a, b, sigma, sigma_a, sigma_b]
```

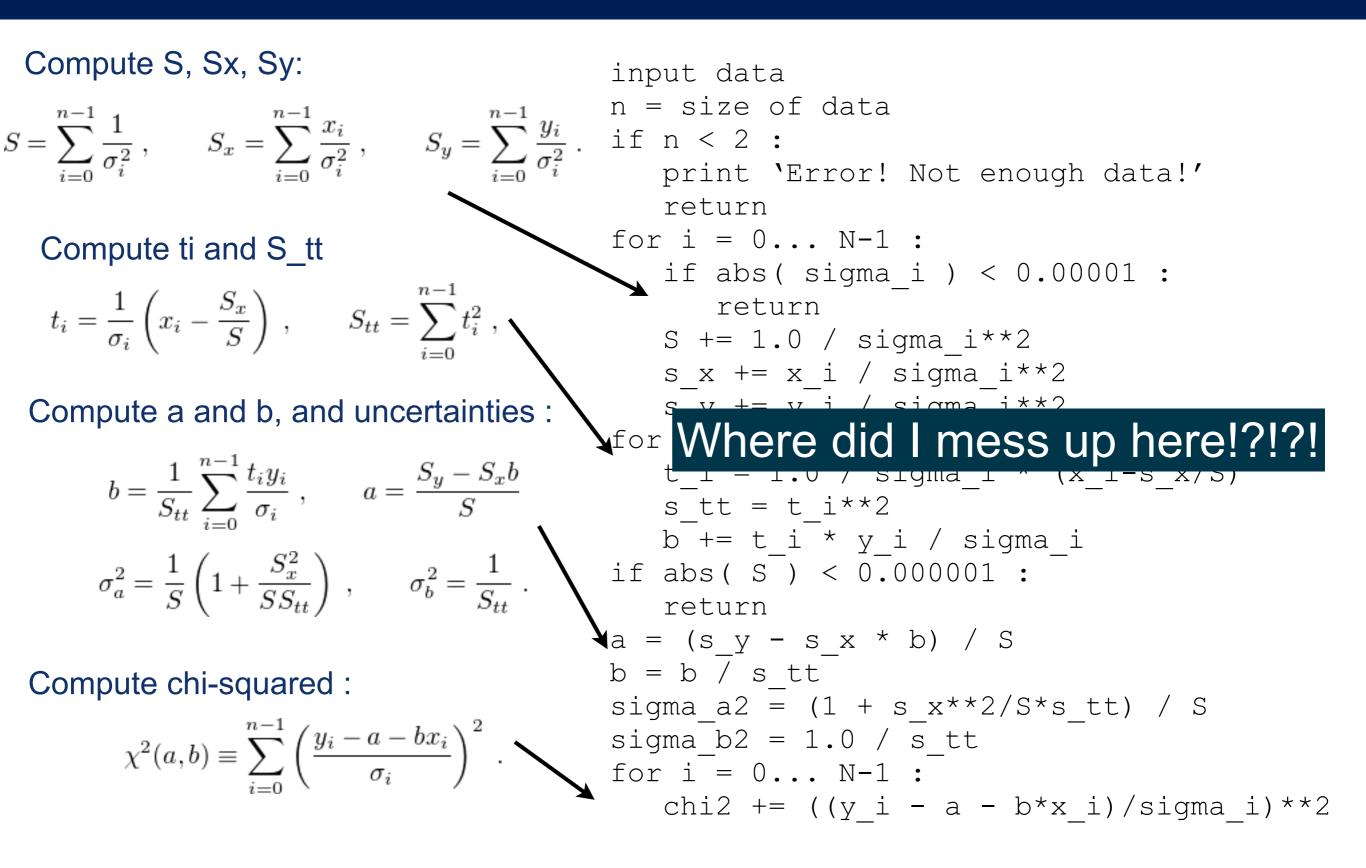
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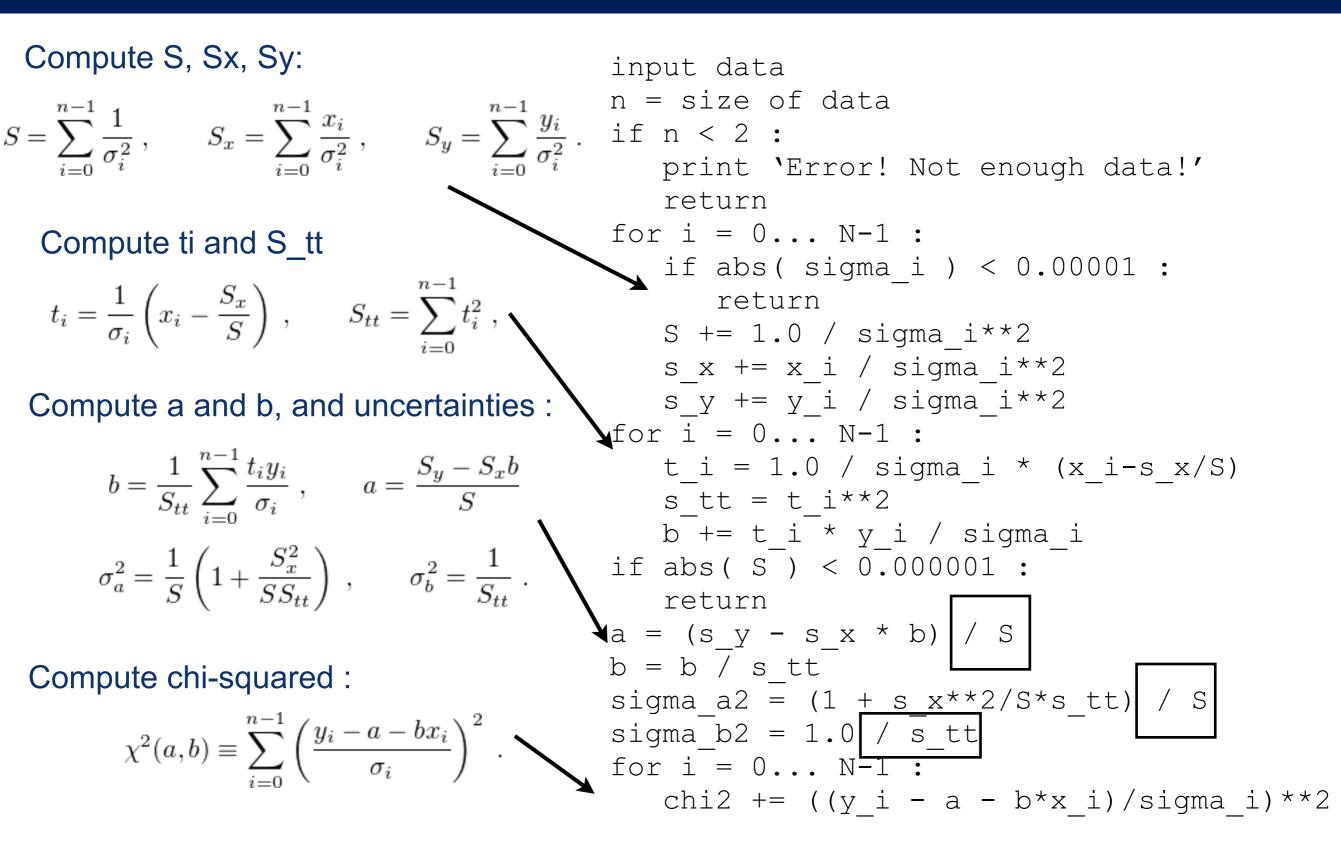
 Check "pass" criterion
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- Step 7 : Publish!

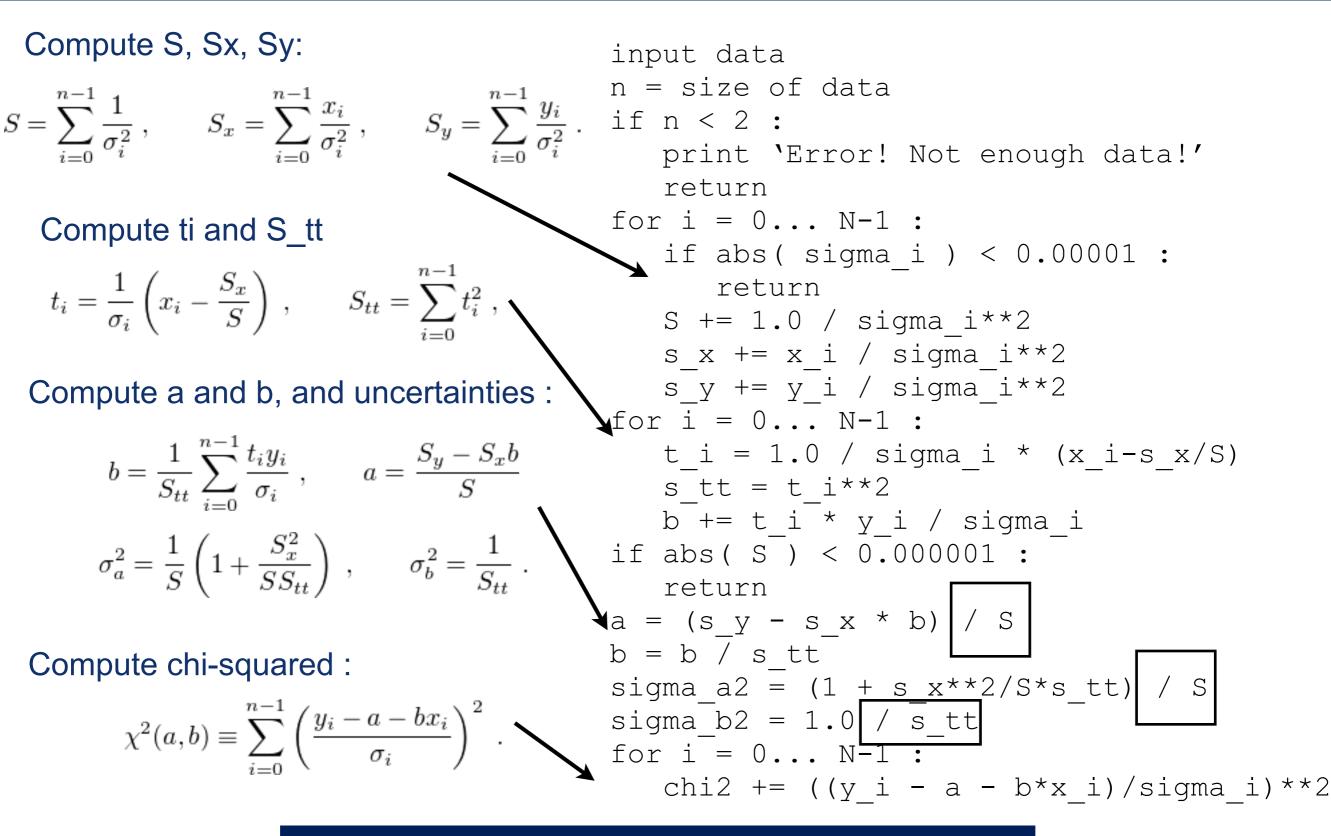
Hands-on!

Effectively the same as without uncertainties, with a few minor modifications









Unlikely to happen, but good to be paranoid anyway!

Write code : With uncertainties

Pseudocode

```
input data
n = size of data
if n < 2:
   print 'Error! Not enough data!'
   return
for i = 0... N-1 :
   if abs( sigma i ) < 0.00001 :
      return
   S += 1.0 / sigma i**2
   s x += x i / sigma i**2
   s y += y i / sigma i**2
for i = 0... N-1 :
  t i = 1.0 / sigma i * (x i-s x/S)
   s tt = t i^{*2}
  b += t i * y i / sigma i
if abs(S) < 0.00001:
   return
a = (s y - s x * b) / S
b = b / s tt
sigma a2 = (1 + s x^{*2}/S^{*s} tt) / S
sigma b2 = 1.0 / s tt
for i = 0 \dots N-1 :
   chi2 += ((y i - a - b*x i)/sigma i)**2
```

python

```
import numpy as np
def chi square fit(x, y, err):
    n = len(x)
    if n < 2:
        print ('Error! Need at least 2 data points!')
        exit()
    S = np.sum(1/err**2)
    if abs(S) < 0.00001 :
        print ('Error! Denominator S is too small!')
        exit()
    S x = np.sum(x/err**2)
   S y = np.sum(y/err**2)
    t = (x - S_x/S) / err
    S tt = np.sum(t**2)
    if abs(S tt) < 0.00001 :
        print ('Error! Denominator S is too small!')
        exit()
    b = np.sum(t*y/err) / S tt
    a = (Sy - Sx * b) / S
    sigma a2 = (1 + S x**2/S/S tt) / S
    sigma b2 = 1/S tt
    if sigma a2 < 0.0 or sigma b2 < 0.0 :
        print ('Error! About to pass a negative to sqrt')
        exit()
    sigma a = np.sqrt(sigma a2)
    sigma b = np.sqrt(sigma b2)
    chi square = np.sum((y - a - b*x) / err)**2)
    return(a, b, sigma a, sigma b, chi square)
```

Write code : With uncertainties

```
void chi square fit ( // makes a linear chi-square fit
 const vector<double>& x, // vector of x values - input
 const vector<double>& y, // vector of y values - input
 const vector<double>& err, // vector of y error values - input
 double& a,
                           // fitted intercept - output
                          // fitted slope - output
 double& b,
                        // estimated error in intercept - output
 double& sigma a,
 double& sigma b,
                          // estimated error in slope - output
 double& chi square)
                          // minimized value of chi-square sum - output
 int n = x.size();
 assert(n \ge 2);
 double S = 0, S_x = 0, S_y = 0;
 for (int i = 0; i < n; i++) {</pre>
   assert ( fabs(err[i]) >= 0.000001 );
   S += 1 / err[i] / err[i];
   S_x += x[i] / err[i] / err[i];
   S_y += y[i] / err[i] / err[i];
 ♠
 vector<double> t(n);
 for (int i = 0; i < n; i++)</pre>
   t[i] = (x[i] - S_x/S) / err[i];
 double S tt = 0;
 for (int i = 0; i < n; i++)</pre>
   S_tt += t[i] * t[i];
 b = 0;
 for (int i = 0; i < n; i++)</pre>
  b += t[i] * y[i] / err[i];
 assert( fabs(S tt) > 0.00001);
 b /= S tt;
 assert(fabs(S) > 0.00001);
 a = (S y - S x * b) / S;
 sigma a = sqrt((1 + S x * S x / S / S tt) / S);
 sigma_b = sqrt(1 / S_tt);
 chi square = 0;
 for (int i = 0; i < n; i++) {</pre>
   double diff = (y[i] - a - b * x[i]) / err[i];
   chi square += diff * diff;
  }
```

C++ tip : assert yourself!

- If you have a condition that your algorithm must fulfill, you can use a few C++ mechanisms to handle this.
- Python handles exceptions on its own.
- C++ behavior there is undefined, and compiler dependent!
- So, you can use a simple "assert(condition)" to make sure it's true
- Alternatively you can use exception handling but that can get a little hairy. I won't cover it in this class much (if at all).

Hands-on!

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Fitting curves

- What if we want to fit something besides a line?
- Well, there are a few cases :

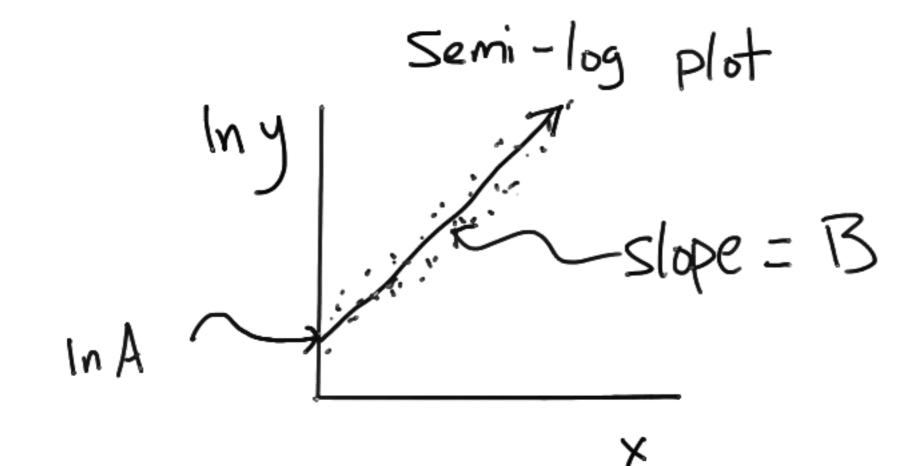
example!

- Does change of variables in x and y make it a line?
 Everything else
- In the first case, it's actually easiest to just fit a line again —In fact, you're technically doing this in your homework

 In the second case, it's also not much harder conceptually, but is more computationally intensive

First case : transform to linear

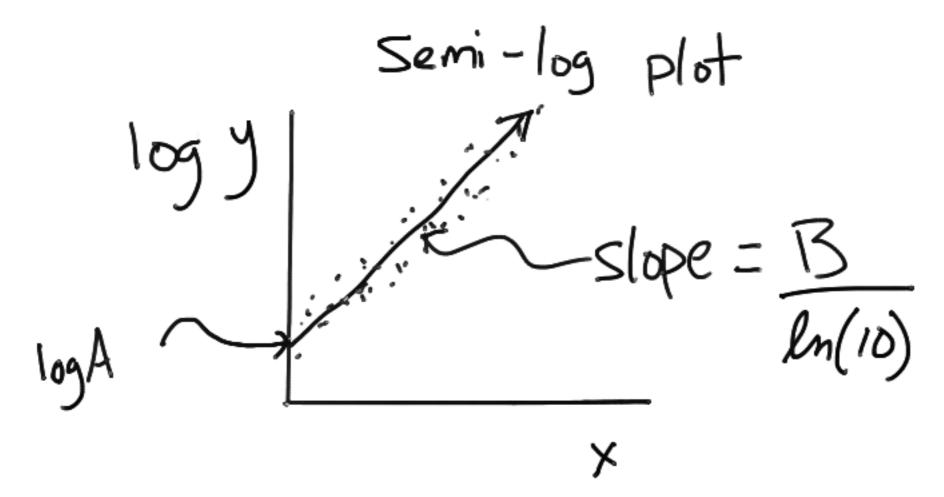
- Say we have a model like $y = A e^{Bx}$
- What do we do?
- Take the logarithm of both sides! $\ln y = \ln A + Bx$



- OK, but what if I do a base-10 logarithm instead of natural?
- Not a problem, just transform!

$$\log_{b} (x) = \frac{\log_{d} (x)}{\log_{d} (b)}$$

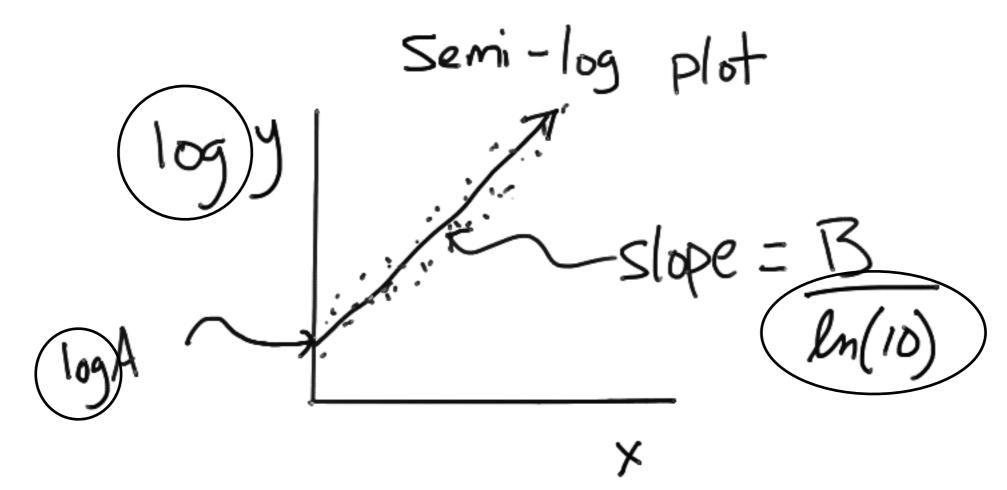
• So in this case :



- OK, but what if I do a base-10 logarithm instead of natural?
- Not a problem, just transform!

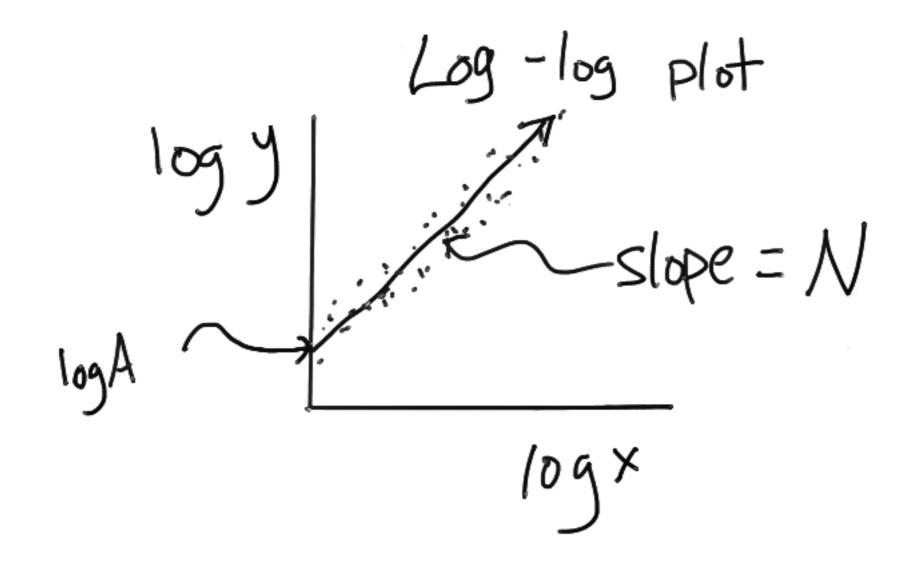
$$\log_{b} (x) = \frac{\log_{d} (x)}{\log_{d} (b)}$$

• So in this case :



- Now try $y = A x^N$
- Take the logarithm of both sides :

$$\log y = \log A + N \log x$$



- What about the uncertainties?
- We did a change of variables :

$$y' = f(y)$$

• So, propagating the uncertainties, we get :

$$\sigma_{y'}^2 = \left(\frac{\partial f}{\partial y}\right)^2 \sigma_y^2$$

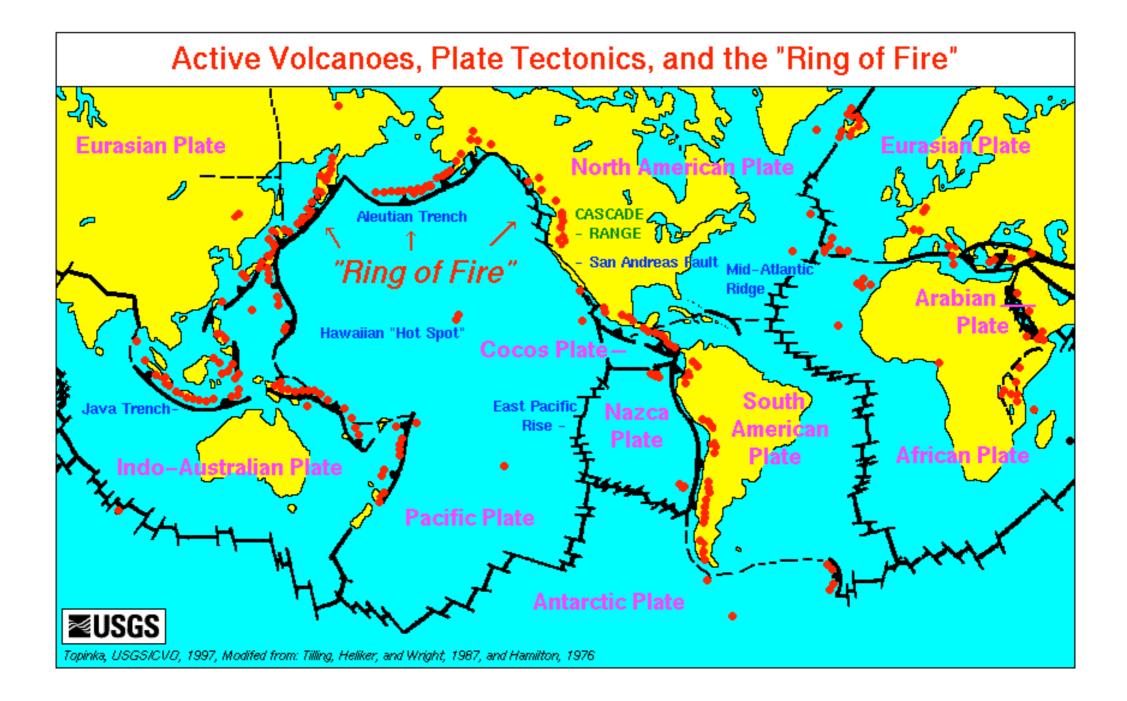
- So you just have to remember this in the chi-squared minimization
- Example :

$$y' = \ln y = \ln A + Bx$$
 $\sigma_{y'} = \frac{\sigma_y}{|y|}$

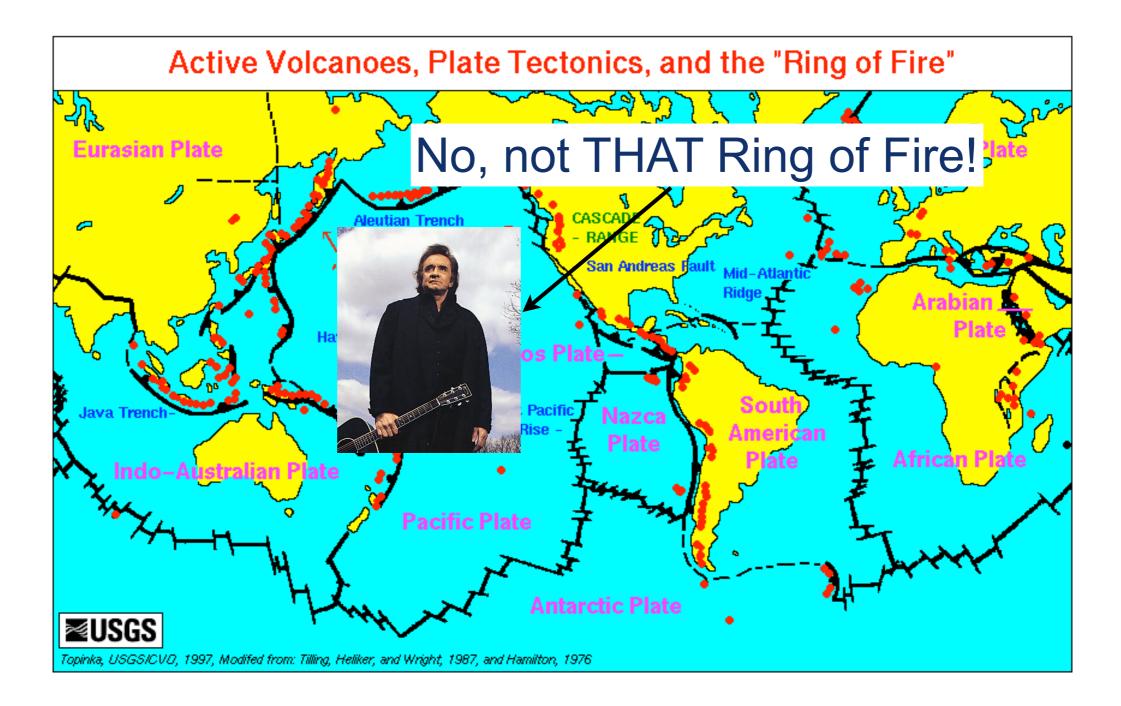
(Note : In Assignment 1 you're already given $\sigma_{y'}$ so you don't have to worry! ₇₆

- Modulo that, it's already a "solved problem"
- You should be doing this in your homeworks already!

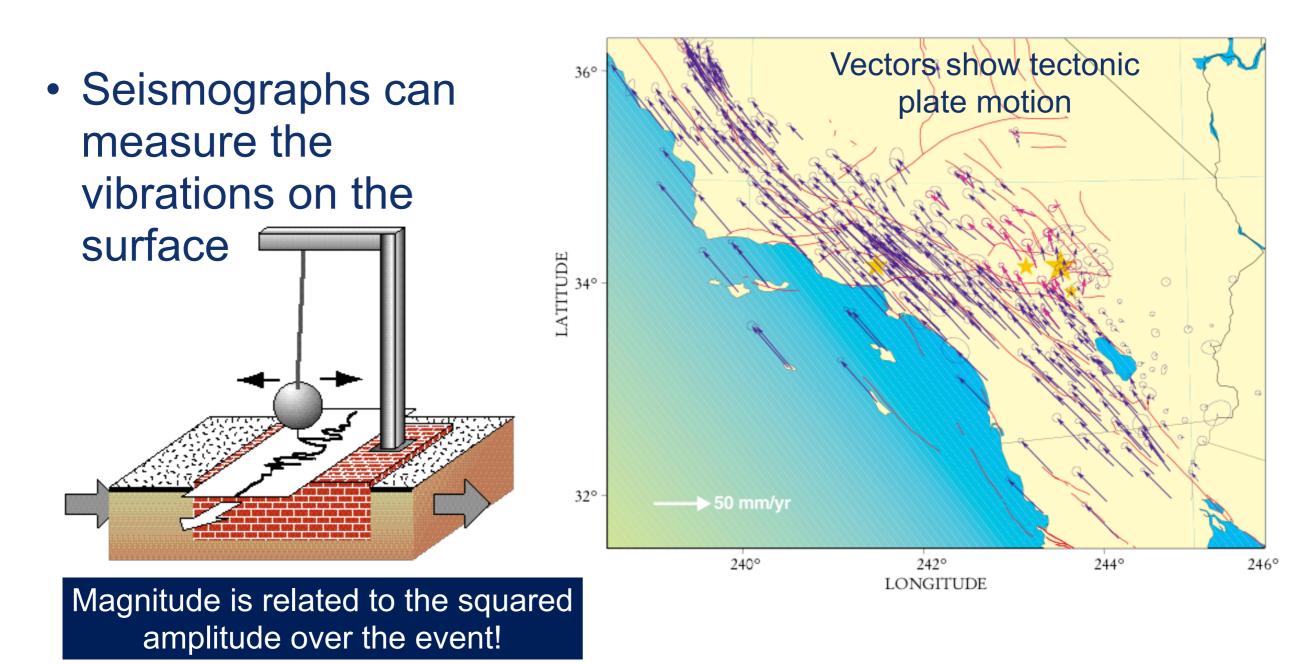
 Earthquakes occur when tectonic plates of the earth move relative to one another



 Earthquakes occur when tectonic plates of the earth move relative to one another

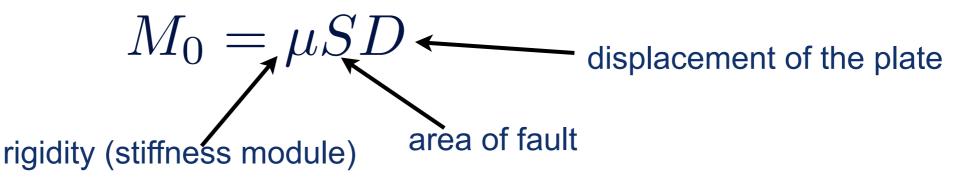


- When they rub against each other, can get stuck!
- Builds pressure, then slips, releasing a lot of energy



H. Kanamori and E.E. Brodsky, <u>The Physics of Earthquakes</u>, Physics Today **54**, 34-40 (2001) <u>http://www.colorado.edu/physics/phys2900/homepages/Marianne.Hogan/graphs.html</u>

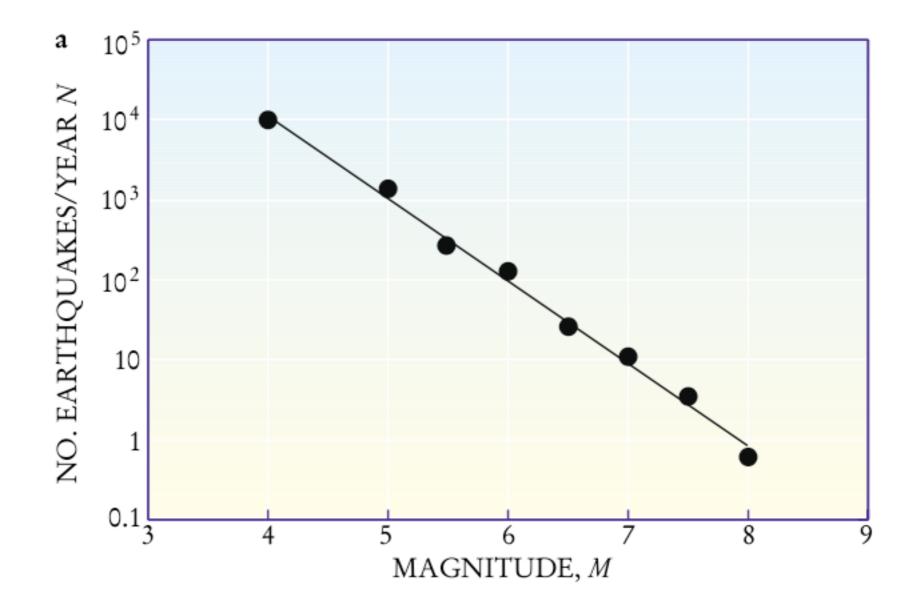
- The "Richter scale" was developed by Richter in the 1930's
- Relates the LOCAL magnitude scale $_{M_L}$
 - -Defined by the amount of amplitude variation on a seismograph
- Replaced in the 1970's by the MOMENT magnitude scale



- Gutenberg-Richter Law Model :
 - –Frequency (N) of earthquakes of magnitude (M) : defined as number of events with magnitude >= M
 - -Empirical model :

$$\log N = a - bM$$
$$M \sim \log M_0$$

 Frequency vs magnitude plot of earthquakes between 1904 and 2000 :



• Get data from :

-<u>http://earthquake.usgs.gov/earthquakes/eqarchives/epic/</u>

- Many formats for the data file :
 - -Map & List
 - -CSV (comma-separated values)
 - -KML (google-based geographical data representation)
 - -QuakeML (XML for earthquakes)
 - -GeoJSON (JSON for earthquakes)
- We'll go for CSV :

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• Example :

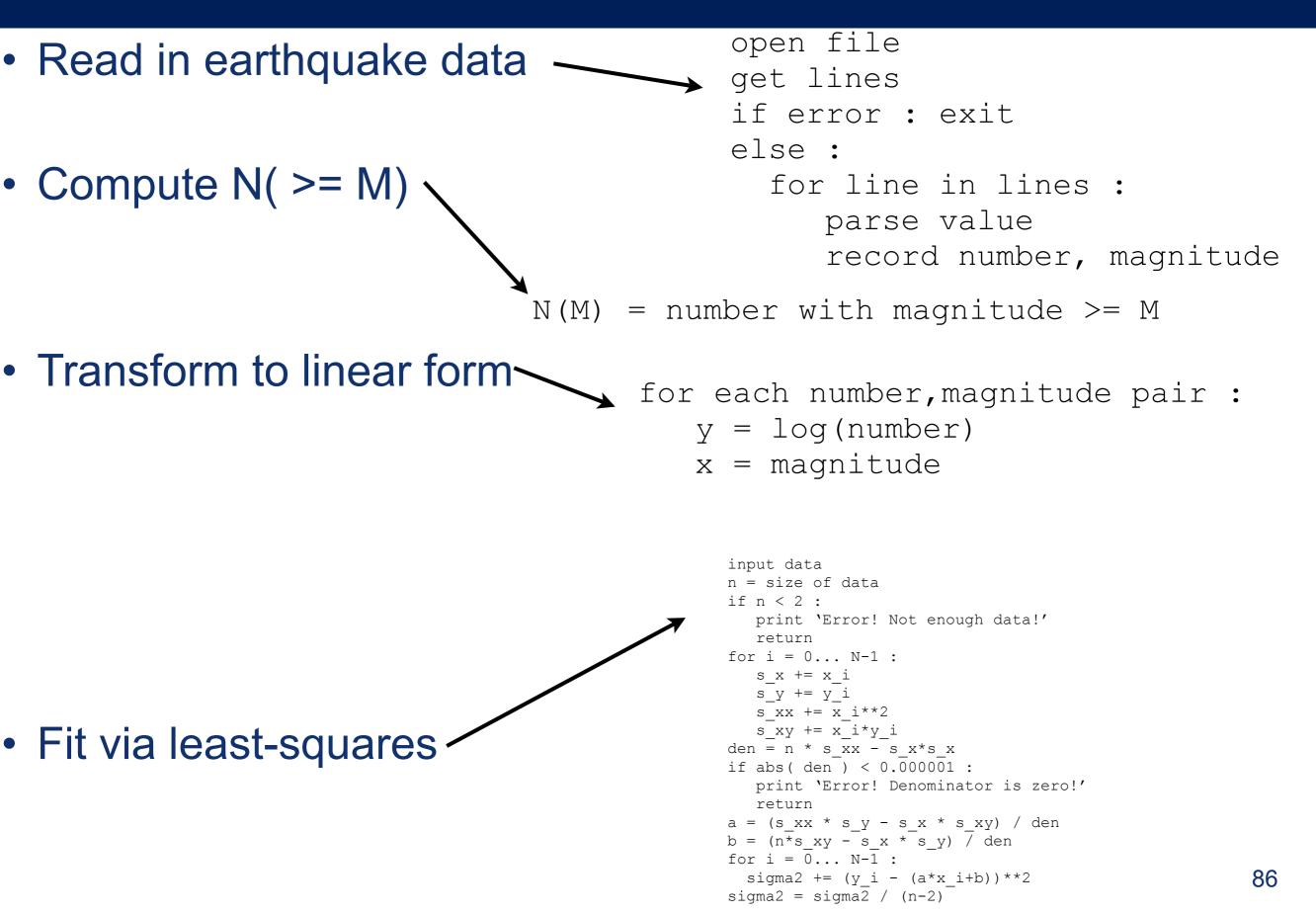
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 The data we'll fit : all earthquakes in southern California from 1973 until today

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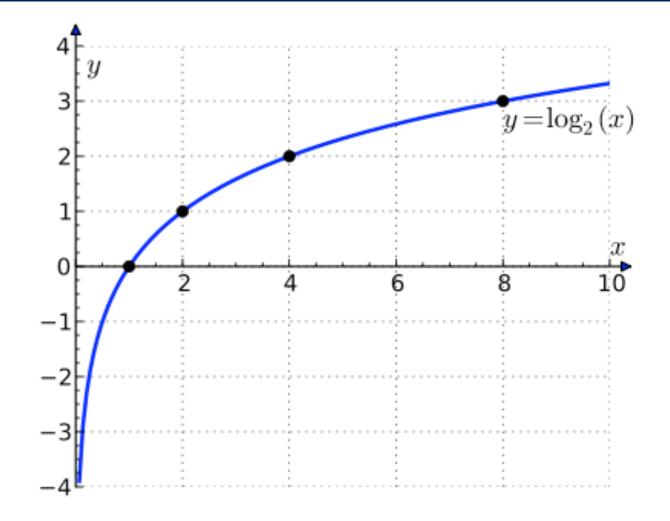
The big picture



Numerical issues : logarithms

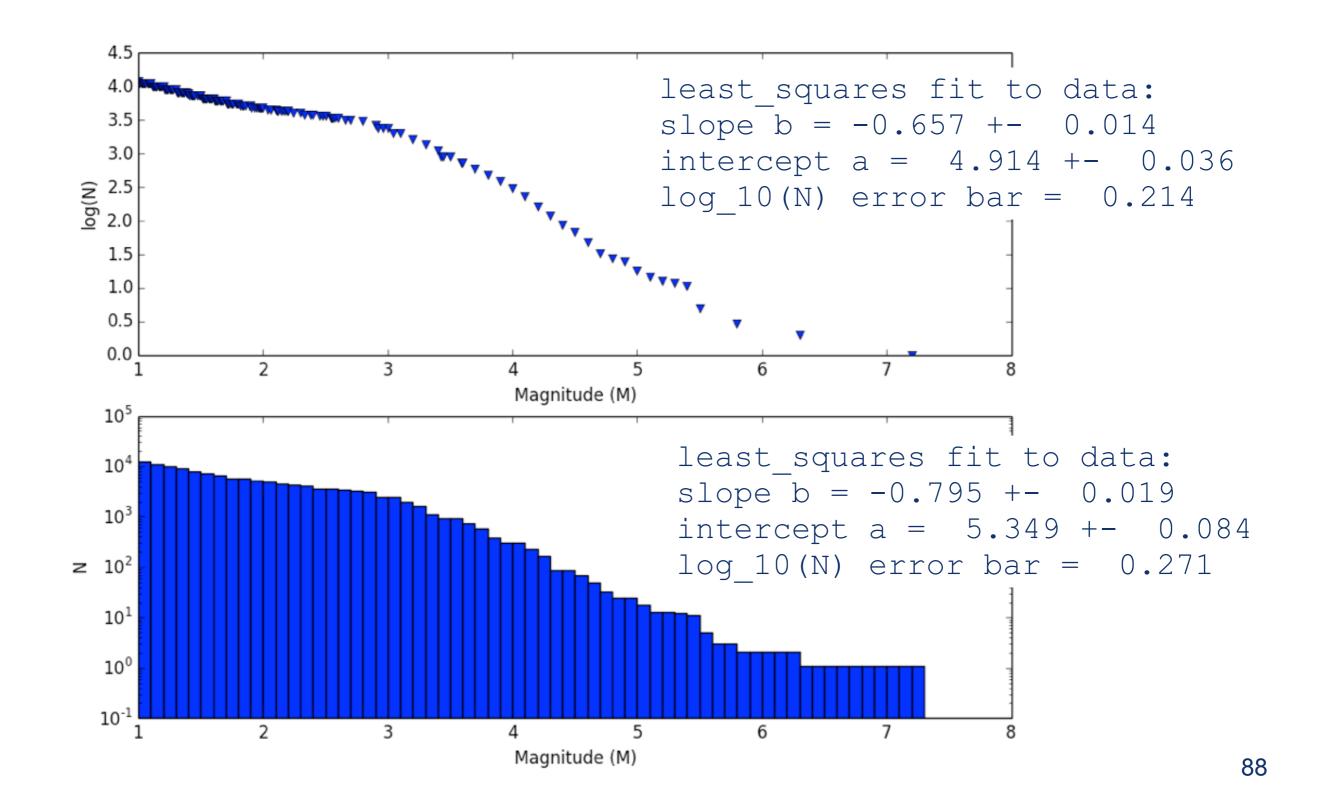
- Logarithms cannot be <= 0
- Always need to check this!
- Other than that, very nice because it transforms multiplication into addition!

$$\log x \times y = \log x + \log y$$



Binning!

Be sure to be careful about fitting binned data!



- For general curve-fitting, it's not conceptually more difficult
- However, it is computationally more difficult

• Define the function and its parameters as :

$$\{a, b\} \rightarrow \vec{a}$$
 $y = y(x; \vec{a})$
"y is a function of x, with parameters a"

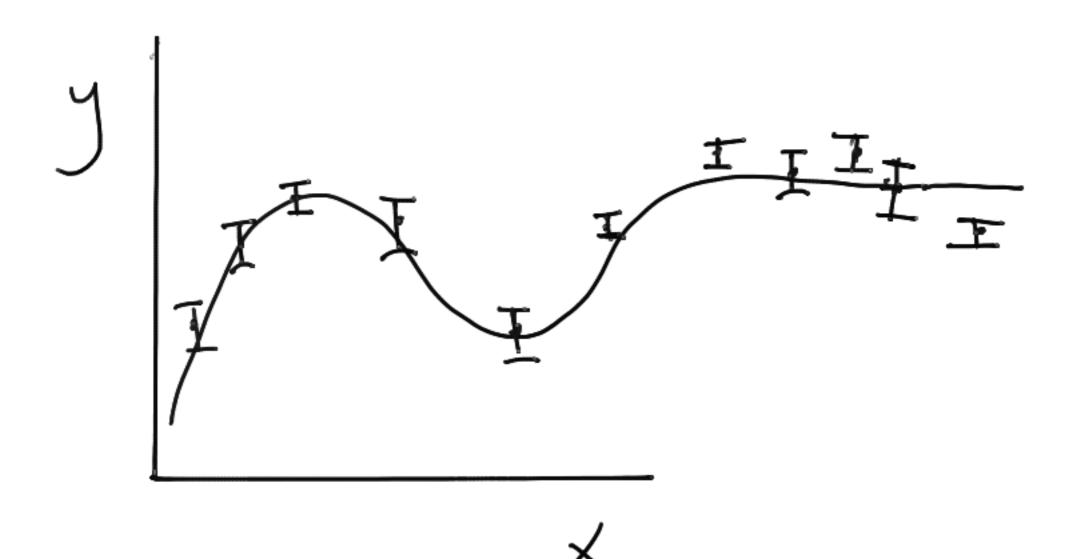
• Rewrite our chi-squred expression :

$$\chi^2(\vec{a}) \equiv \sum_{i=0}^{n-1} \left(\frac{y_i - y(x; \vec{a})}{\sigma_i} \right)^2$$

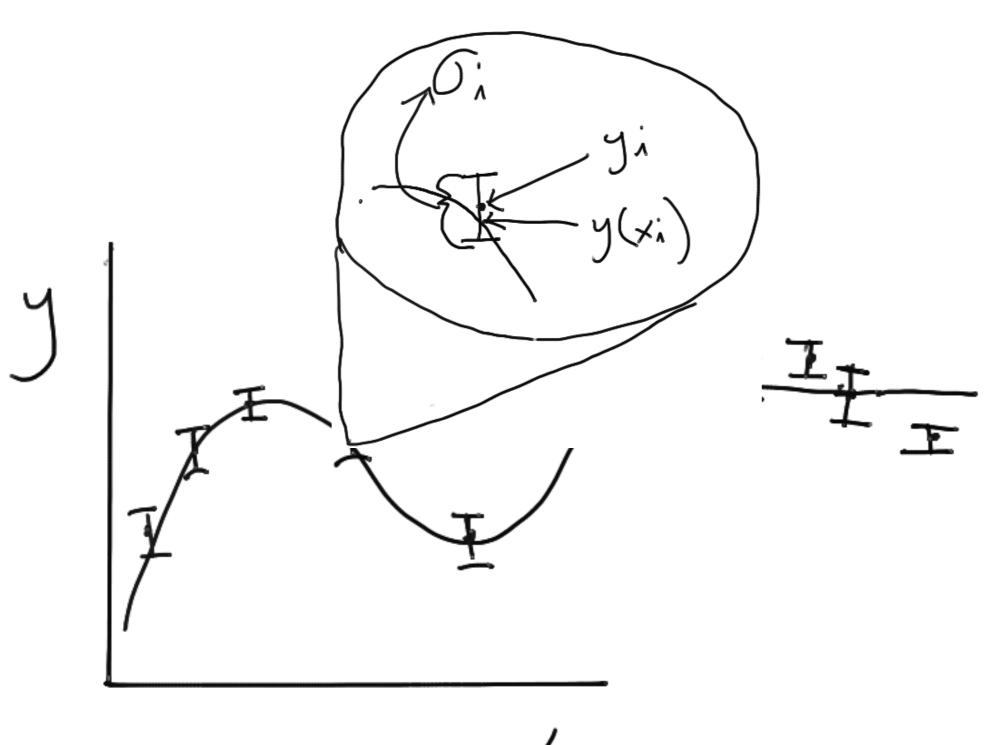
• Now this actually should be obvious!

• This completely generalizes to nonlinear y(x)

• Still just minimizing the distance within the uncertainties



• Still just minimizing the distance within the uncertainties



• Same strategy as before : minimize the chi2!

$$\chi^2(\vec{a}) \equiv \sum_{i=0}^{n-1} \left(\frac{y_i - y(x; \vec{a})}{\sigma_i} \right)^2$$

- So, let's say that y(x) is some expansion of functions Y_k(x) : $y(x; \vec{a}) = \sum_{k=0}^{m-1} a_k Y_k(x)$
- Then the chi2 is :

$$\chi^2(x,\vec{a}) = \sum_{i=0}^{n-1} \frac{1}{\sigma_i^2} \left[y_i - \sum_{k=0}^{m-1} a_k Y_k(x) \right]^2$$

• We minimize :

$$\frac{\partial \chi^2}{\partial a_j} = \frac{\partial}{\partial a_j} \sum_{i=0}^{n-1} \frac{1}{\sigma_i^2} \left[y_i - \sum_{k=0}^{m-1} a_k Y_k(x) \right]^2 = 0$$

• Taking the derivative :

$$2a_j \sum_{i=0}^{n-1} \frac{1}{\sigma_i^2} Y_j(x) \left[y_i - \sum_{k=0}^{m-1} a_k Y_k(x) \right] = 0$$

 The 2*a_j cancels. Then we multiply the sum through, and bring over the second term, so we get :

$$\sum_{i=0}^{n-1} \sum_{k=0}^{m-1} \frac{Y_j(x_i)Y_k(x_i)}{\sigma_i^2} a_k = \sum_{i=0}^{n-1} \frac{Y_j(x_i)y_i}{\sigma_i^2}$$

• This is a matrix equation, so we define the "design matrix" :

$$A_{ij} = \frac{Y_j(x_i)}{\sigma_i}$$

$$\mathbf{A} = \begin{bmatrix} Y_1(x_1)/\sigma_1 & Y_2(x_1)/\sigma_1 & \dots \\ Y_1(x_2)/\sigma_2 & Y_2(x_2)/\sigma_2 & \dots \\ \dots & \dots & \dots & \dots \end{bmatrix}$$

• Then our chi2 minimization becomes :

$$(\mathbf{A}^T \mathbf{A})\vec{a} = \mathbf{A}^T \vec{b}$$

• SO :

$$\vec{a} = (\mathbf{A}^T \mathbf{A})^{-1} \mathbf{A}^T \vec{b}$$

• If we define the "correlation matrix" :

$$\mathbf{C} = (\mathbf{A}^T \mathbf{A})^{-1}$$

• Then the uncertainty on a_j is :

$$\sigma_{a_j} = \sqrt{C_{jj}}$$

• As a first example, let's look at polynomial fits

$$y = \sum_{k=0}^{m-1} a_k x^k$$

- Slight generalization of the linear fit we did previously
- General solution is to minimize the chi2 :

$$\chi^2(\vec{a}) \equiv \sum_{i=0}^{n-1} \left(\frac{y_i - y(x; \vec{a})}{\sigma_i} \right)^2$$

• In this case :

$$\chi^{2}(\vec{a}) = \sum_{i=0}^{n-1} \left(\frac{y_{i} - \sum_{j=0}^{M} a_{j} x^{j}}{\sigma_{i}} \right)^{2}$$

• Our design matrix is therefore :

 $A_{ij} = x_i^j / \sigma_i$

 Caveat : This oftentimes is ill-formed, so don't go too crazy here. Typically we do quadratic, cubic, quartic, but above that it strains credibility.

• Will return to this after we do some linear algebra!