

Monte Python

the way back from cosmology to Fundamental Physics

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and

UC Berkeley



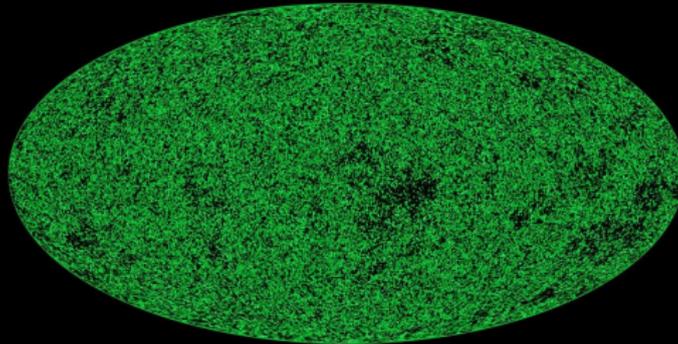
NORDITA



IFT School on Cosmology Tools

March 2017

Related Tools

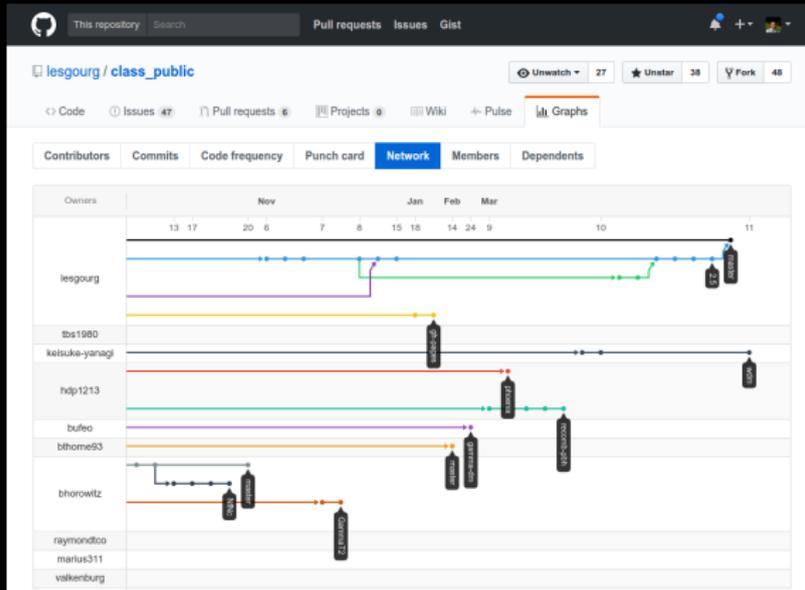


Version control and Python interfacing

PLANCK

Version control: GIT + github

- GIT: access all previous versions, restore, compare, branch...
- Github: website + interface
 - Network: users, branches intermediate versions



Version control: GIT + github

- GIT: access all previous versions, restore, compare, branch...
- Github: website + interface
 - Network: users, branches intermediate versions
 - Issues: troubleshooting, forum/improvements

The repository: Pull requests Issues Gist

lesgourg / class_public Unwatch 27 Unstar 38 Fork 46

Code Issues 47 Pull requests 6 Projects 0 Wiki Pulse Graphs

Filters Labels Milestones [New issue](#)

47 Open 83 Closed Author Labels Milestones Assignee Sort

- #143 How to add a new derived parameter in CLASS? opened 6 days ago by shouvikc 2
- #142 Default selection_magnification_bias opened 7 days ago by jesspetroo
- #141 dens-dens cls dependent on choice of nCl or nCl,sCl,ICl opened 23 days ago by hoyteb
- #140 Default Input Parameters opened 23 days ago by bubarakakikaka 1
- #139 Freeze value of age() in multiple copies. opened 27 days ago by mirandawebster
- #138 Dark energy perturbation for a variable equation of state in CLASS opened on Feb 10 by barump1985 3
- #136 growth factor for wCDM opened on Feb 1 by amoradinejad 5
- #132 late time interaction opened on Dec 6, 2016 by mahtaparsa
- #131 Non-constant equation of state opened on Dec 6, 2016 by damonge 2

Python interfacing with classy

- `classy` → use `CLASS` as a Python module
 - Required for MCMC (tomorrow!)
 - Useful for plotting

```
from classy import Class
import numpy as np
import matplotlib.pyplot as plt
cosmo = Class ()
cosmo.set ( {'output': 'tCl, pCl, lCl', 'lensing': 'yes'})
cosmo.compute ()
l = np.array ( range (2 ,2501) )
factor = l*(l +1) / (2*np.pi )
lensed_cl = cosmo.lensed_cl (2500)
#then just plot lensed_cl...
```

Python interfacing with classy

- classy → use CLASS as a Python module
 - Required for MCMC (tomorrow!)
 - Useful for plotting
- IPython → Interactive Python frontend
 - TAB auto-completion

```
miguel@Goedel:~$ ipython
Python 2.7.6 (default, Oct 26 2016, 20:30:19)
Type "copyright", "credits" or "license()" for more information.

IPython 5.1.0 -- An enhanced Interactive Python.
?                -> Introduction and overview of IPython's features.
%quickref        -> Quick reference.
help             -> Python's own help system.
object?         -> Details about 'object', use 'object??' for extra details.

In [1]: from classy import Class

In [2]: c = Class()

In [3]: c.
c.age                c.h                  c.Omega_nu
c.angular_distance  c.Hubble             c.pars
c.baryon_temperature c.ionization_fraction c.pk
c.compute            c.lensed_cl         c.raw_cl
c.density_cl        c.luminosity_distance c.rs_drag
c.empty              c.n_s                c.set
c.get_background     c.Neff              c.set_default
c.get_current_derived_parameters c.nonlinear_method  c.sigma8
c.get_perturbations  c.nonlinear_scale   c.state
c.get_pk             c.Omega0_m          c.struct_cleanup
c.get_primordial    c.Omega_b           c.T_cmb
c.get_thermodynamics c.omega_b           c.z_of_r
c.get_transfer       c.Omega_m
```

Python interfacing with classy

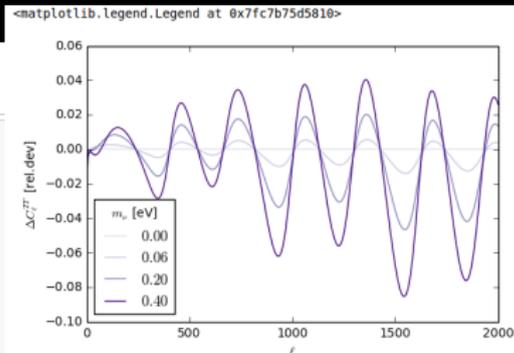
- classy → use CLASS as a Python module
 - Required for MCMC (tomorrow!)
 - Useful for plotting
- IPython → Interactive Python frontend
 - TAB auto-completion
- Jupyter → Notebook interface (Julia+Python+R)

Jupyter example

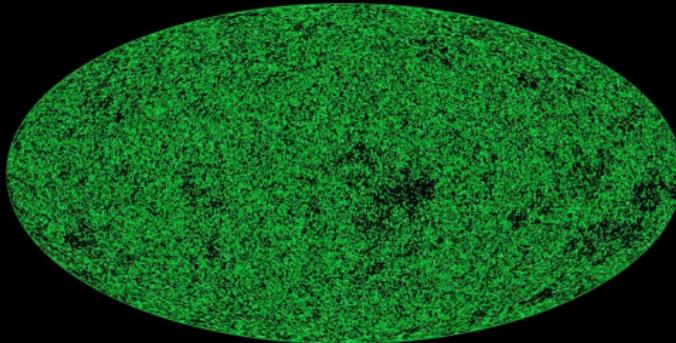
You can write \mathcal{L}_{FIT} and awesome equations like $m_\nu \gtrsim 60$ meV. And plot the effect of m_ν in few lines

```
cm = plt.get_cmap('Purples')
c = Class()
cl = {} #dictionary for output
for m in [0.0, 0.06, 0.2, 0.4]:
    c.set({'N_ncdm':1, 'N_ur':2.033, 'm_ncdm':m, 'output':'tcl'})
    c.compute()
    cl[m] = c.raw_cl(2000)
    plt.plot(cl[m]['ell'][2:], cl[m]['tt'][2:]/cl[0]['tt'][2:]-1.,
            color=cm((m+0.1)/0.5), label=r'%.2f$(m)')
    c.empty()

plt.xlabel(r'$\ell$')
plt.ylabel(r'$\Delta C_{\ell}^{\text{TT}}$ [rel.dev]')
plt.legend(loc='lower left', fontsize = 12, title= r'$m_{\nu}$ [eV]')
```



Monte Python



from cosmology back to fundamental physics

PLANCK

DISCLAIMER: Short time!

≲ 3h course ⇒ overview and basic usage

Learn further:

- MontePython slides by Sebastien Clesse (≪ 1h?)

https://lesgourg.github.io/class-tour/16.06.02_Lisbon_intro.pdf

- MontePython course (~ 5h)

<https://lesgourg.github.io/class-tour-Tokyo.html>

- Links to extra resources in exercise sheet

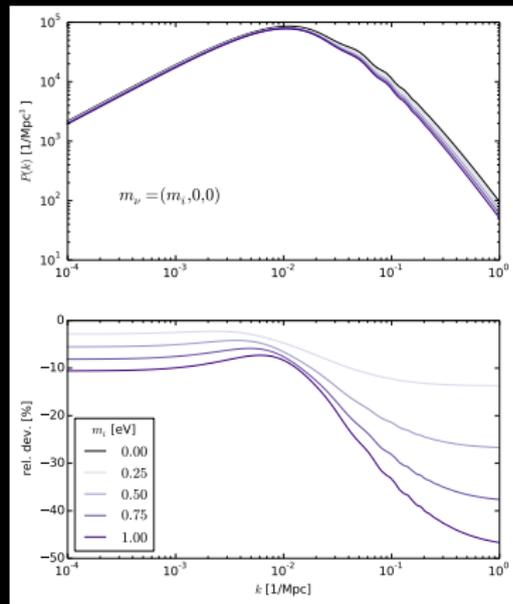
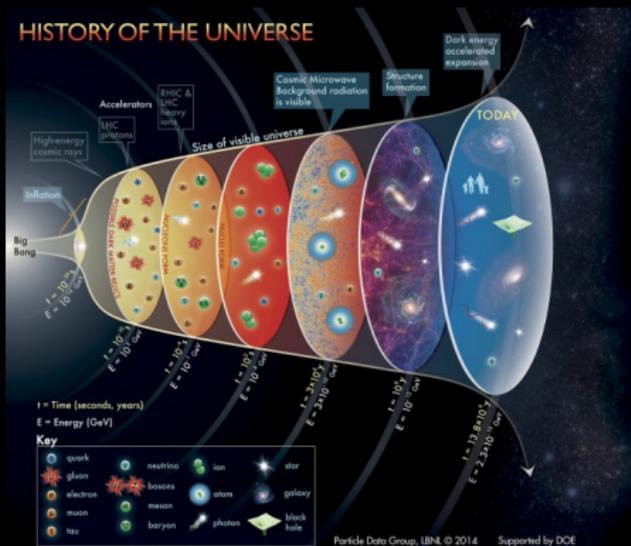
IMPORTANT DISCLAIMER:

↓ I'm mainly a user with little experience developing!

↑ Help from experts:

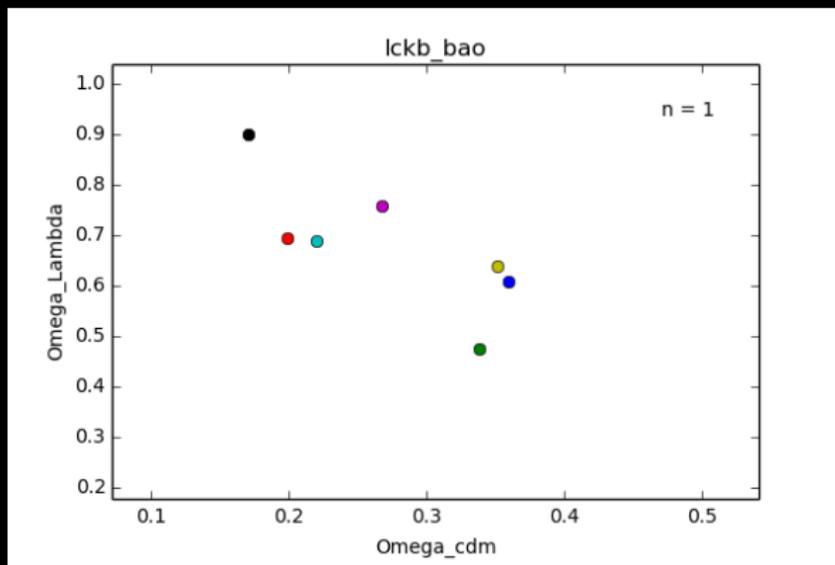
Thejs Brinckmann, Carlos Garcia, Deanna Hooper & Vivian Poulin

Fundamental physics and cosmology



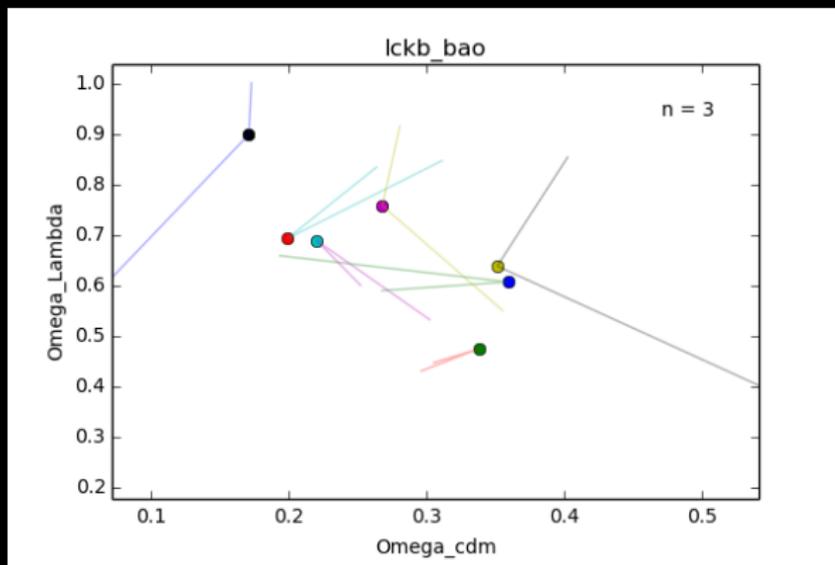
Initial conditions, Dark Matter, Neutrinos, Dark Energy, Gravity...

Scan space of parameters



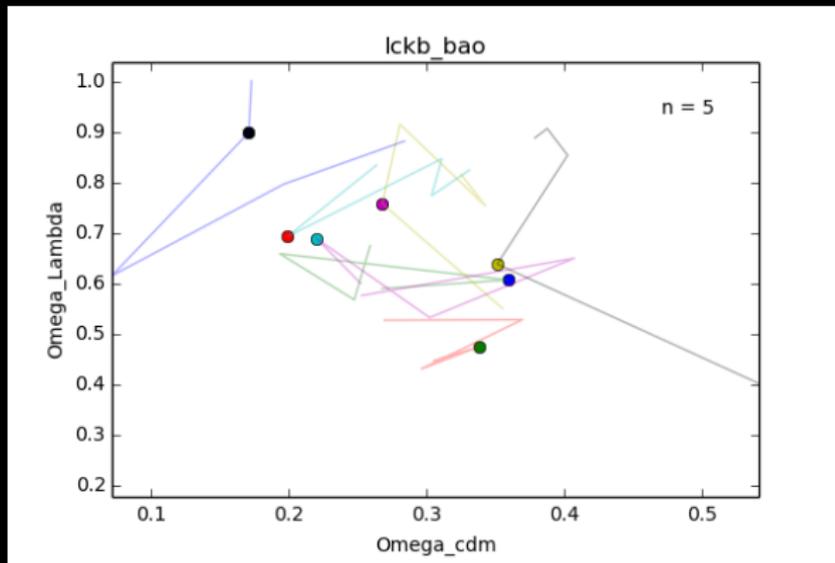
(recall lecture by Will Handley)

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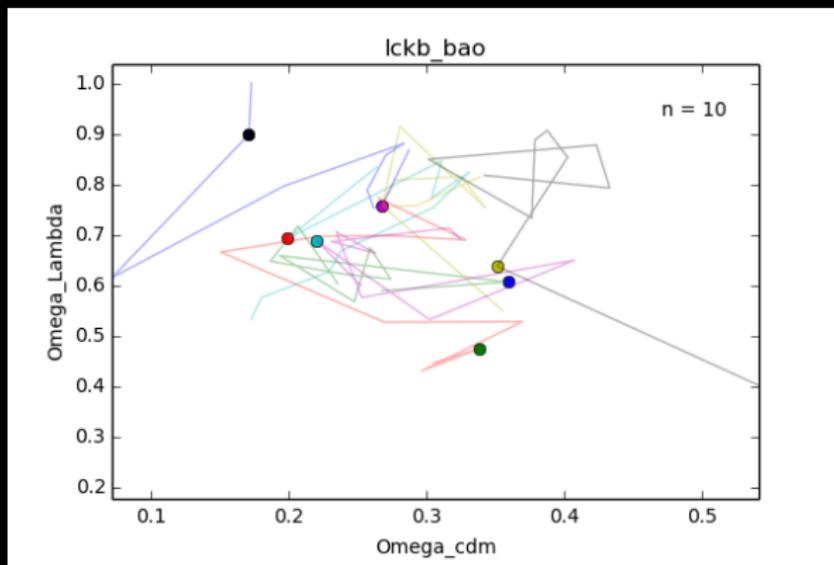
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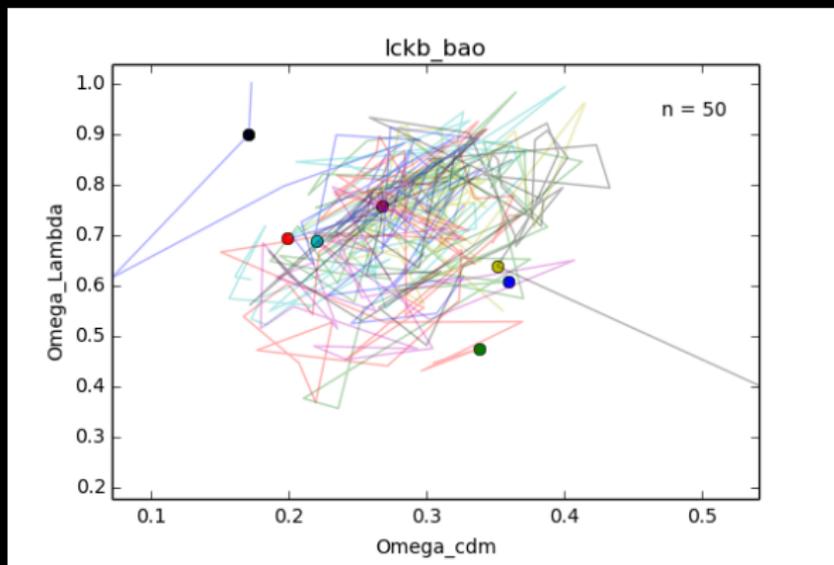
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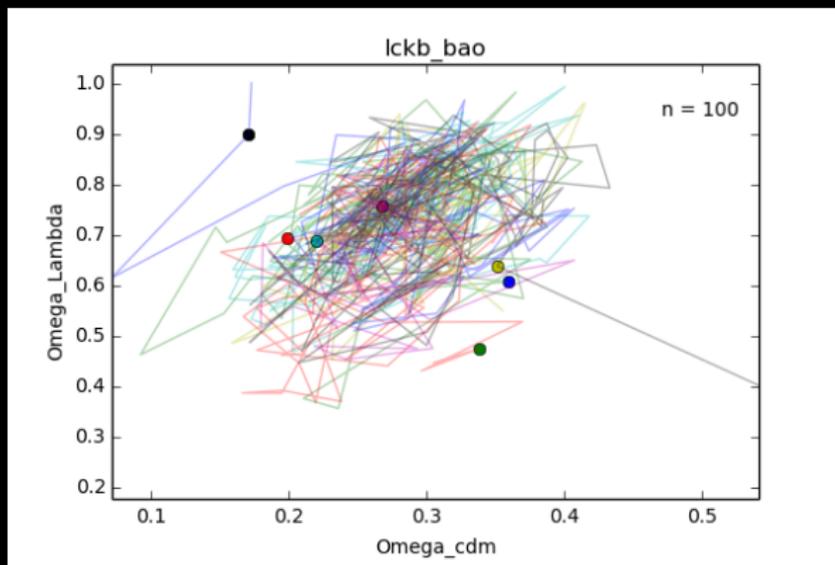
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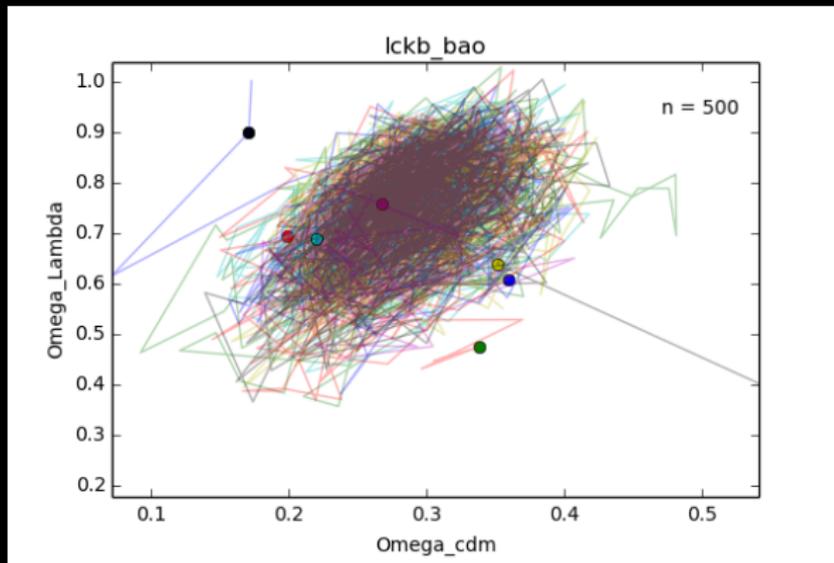
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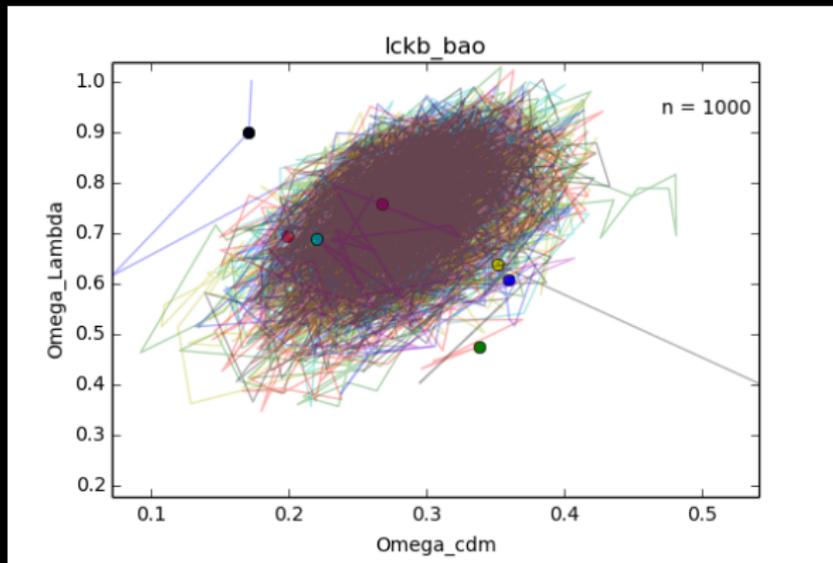
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MontePython

An Markov Chain Monte Carlo engine for parameter extraction:

Features

- Written in Python
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 - likelihoods for new experiments
 - features for sampling, plotting...

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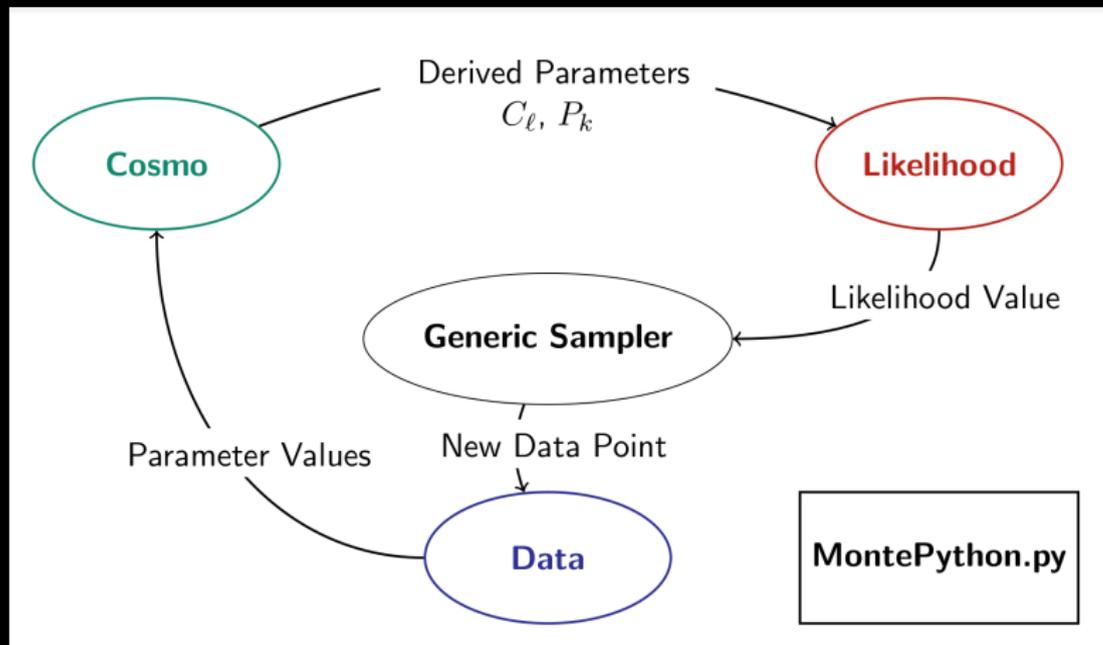
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- Uses CLASS through the *classy* wrapper
- Modular, easy to add
 - likelihoods for new experiments
 - features for sampling, plotting...
- Easy to use, intensively documented
- Parallelization is optional
 - simpler to install
 - runs in old/separate computers, short queue

Modular Structure



(from B. Audren's Monte Python slides)

Defining a run (model.param)

- List of experiments

```
data.experiments=['Planck_highl', 'Planck_lowl', 'Planck_lensing']
```

Collected in montepython/likelihoods:

```
miguel@Goedel:~/code/montepython_zuma/montepython/likelihoods$ ls
acbar          clik_wmap_full      __init__.py         quad
bao            clik_wmap_lowl     __init__.pyc       sdss_lrgDR4
bao_boss       cosmic_clocks_BC03 JLA                  sn
bao_boss_aniso cosmic_clocks_BC03_all JLA_simple          spt
bao_boss_aniso_gauss_approx cosmic_clocks_MaStro lowlike              spt_2500
bao_known_rs   da_rec              Planck_actspt       test_gaussian
bicep          euclid_lensing     Planck_highl        test_nuisance1
bicep2         euclid_pk           Planck_highl_lite   test_nuisance2
boomerang      fake_desi           Planck_highl_TTTEEE timedelay
cbi            fake_planck_bluebook Planck_lensing       WiggleZ
CFHTLens       gunn_peterson       Planck_lowl         WiggleZ_bao
CFHTLens_correlation hst                 Planck_SZ           wmap
clik_fake_planck igm_temperature     polarbear           wmap_9yr
```

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- Cosmological parameters

```
                # [mean, (bounds) , SIGMA, scale, type ]  
data.parameters['n_s']=[0.96, None, None, 0.008, 1 , 'cosmo']
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Fixed values:

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- Derived and Nuisance parameters

```
data.parameters['sigma8'] = [0, None, None, 0, 1, 'derived']  
data.parameters['A_cib_217'] = [61, 0, 200, 7, 1, 'nuisance']
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- MCMC parameters: data.N=10, data.write_step=5 ...

Running Monte Python (run)

Single chain:

```
python montepython/MontePython.py run \  
-p model.param \  
-o output_directory (...)
```

Parallel run (4 chains):

```
mpirun -nc 4 python (...)
```

Some options:

```
#!/bin/sh
Appending to an existing folder: using the log paran instead of
input/Lambda_bao_b paran
Running Monte Python v2.2.2

with CLASS v2.4.5

Testing likelihoods for:
-> bao_boss, bao_boss_aniso

#!/\ excluding isotropic CMASS measurement
Creating chains/lcdm_bao_b/2017-03-12_100_2.txt
Creating error file chains/lcdm_bao_b/error_log.txt

Deduced starting covariance matrix:
[['omega_b', 'Omega_cdm', 'Omega_k']
 [[ 0.  0.  0.]
 [ 0.  0.  0.]
 [ 0.  0.  0.]]]

# -logLkl 1e+02omega_b Omega_cdm Omega_k Omega_Lambda
3 11.9558 2.221160e+00 3.132032e-01 -1.692095e-02 6.553103e-01
1 12.7946 2.262336e+00 3.271086e-01 2.742679e-02 5.961614e-01
1 12.8483 2.256136e+00 2.494162e-01 1.271974e-01 5.742182e-01
3 11.0619 2.286885e+00 2.308487e-01 1.014554e-01 6.437860e-01
1 11.4992 2.275734e+00 2.188346e-01 5.439644e-02 6.851744e-01
1 11.1477 2.309970e+00 2.013242e-01 3.477642e-02 7.135600e-01
3 7.83276 2.297757e+00 2.751741e-01 9.733703e-02 5.774153e-01
14 6.96653 2.302347e+00 2.748064e-01 5.921469e-02 6.157454e-01
4 6.46171 2.372939e+00 2.523288e-01 2.276382e-02 6.731983e-01
13 6.95783 2.334726e+00 2.363992e-01 -3.071722e-02 7.434402e-01
4 8.883 2.350200e+00 2.572473e-01 -3.956319e-02 7.311015e-01
14 7.19446 2.380873e+00 2.218523e-01 -2.615132e-02 7.532173e-01
2 6.95177 2.359364e+00 2.629244e-01 7.590971e-02 6.097521e-01
6 7.18696 2.339764e+00 2.595452e-01 8.159985e-02 6.078676e-01
1 6.44771 2.353985e+00 2.392116e-01 3.775342e-02 6.717389e-01
4 8.22755 2.328585e+00 2.075475e-01 -2.491768e-03 7.442001e-01
1 7.89432 2.399661e+00 2.076928e-01 -2.671562e-02 7.669296e-01
6 7.21255 2.388188e+00 2.145373e-01 7.295908e-03 7.261260e-01
2 7.07413 2.397083e+00 2.685047e-01 6.233775e-02 6.169233e-01
1 7.37498 2.406336e+00 2.207678e-01 4.592795e-02 6.880866e-01
2 10.6892 2.439808e+00 1.946383e-01 5.075931e-02 7.014388e-01
3 11.7196 2.476394e+00 2.298767e-01 1.535277e-01 5.626362e-01
1 12.3995 2.441832e+00 3.239238e-01 2.005879e-01 4.222807e-01
3 10.2062 2.420720e+00 2.880782e-01 1.751385e-01 4.840349e-01
1 7.3809 2.439488e+00 2.699004e-01 7.482511e-02 6.021178e-01
3 7.25137 2.437977e+00 2.578438e-01 2.764830e-02 6.613929e-01
1 6.83719 2.459588e+00 2.214344e-01 4.108869e-02 6.839635e-01

# 100 steps done, acceptance rate: 0.26
```

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-o output_directory (...)
```

Parallel run (4 chains):

```
mpirun -nc 4 python (...)
```

Some options:

- `-N` → # points
- `-C` → covariance matrix
- `-r` → restart from last point of chain
- `--update` → update sampling + covariance

```
Appending to an existing folder: using the log paran instead of  
input/lambda_bao_b.paran  
Running Monte Python v2.2.2  
  
with CLASS v2.4.5  
  
Testing likelihoods for:  
-> bao_boss, bao_boss_aniso  
  
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All options explained

```
python montepython/MontePython.py info --help
```

Analyzing results (info)

Single model/experiment:

```
python montepython/MontePython.py info \  
output_directory (...)
```

Comparing several runs:

```
python montepython/MontePython.py info \  
output_1 output_2 output_3 (...)
```

Configuring the output/analysis

```
miguel@Goedel:~/code/montepython_zuma/chains_itp$ python ../montepython/MontePython.py info lckb_bao/  
Running Monte Python v2.2.2  
  
--> Finding global maximum of likelihood  
--> Removing burn-in  
--> Scanning file lckb_bao/2017-03-12_100000_3.txt : Removed 16  
6 non-markovian points, 0 points of burn-in, keep 10397 steps  
2017-03-12_10_1.txt : Removed 0  
non-markovian points, 2 points of burn-in, keep 1 steps  
2017-03-12_100000_4.txt : Removed 57  
non-markovian points, 0 points of burn-in, keep 10405 steps  
2017-03-12_100000_8.txt : Removed 0  
non-markovian points, 2 points of burn-in, keep 6783 steps  
2017-03-12_100000_7.txt : Removed 0  
non-markovian points, 4 points of burn-in, keep 13666 steps  
2017-03-12_100000_1.txt : Removed 92  
non-markovian points, 0 points of burn-in, keep 7975 steps  
2017-03-12_100000_6.txt : Removed 0  
non-markovian points, 4 points of burn-in, keep 20 steps  
2017-03-12_100000_5.txt : Removed 15  
5 non-markovian points, 0 points of burn-in, keep 11158 steps  
2017-03-12_10_2.txt : Removed 0  
non-markovian points, 2 points of burn-in, keep 1 steps  
--> Computing mean values  
--> Computing variance  
--> Computing convergence criterium (Gelman-Rubin)  
-> R-1 is 0.002068 for Omega_b  
0.001200 for Omega_cdm  
0.002191 for Omega_k  
0.002137 for Omega_Lambda  
-----  
-> Computing histograms for Omega_b  
-> Computing histograms for Omega_cdm  
-> Computing histograms for Omega_k  
-> Computing histograms for Omega_Lambda  
-----  
--> Saving figures to .pdf files  
--> Writing .info and .tex files
```

Analyzing results (info)

Single model/experiment:

```
python montepython/MontePython.py info \
    output_directory (...)
```

Comparing several runs:

```
python montepython/MontePython.py info \
    output_1 output_2 output_3 (...)
```

Configuring the output/analysis

- `--extra` → file with plot options
- `--bins` → # bins for posterior
- `--all` → plot every subplot separately
- `--no-mean` → only marginalized in 1D

```
miguel@Goedel:~/code/montepython_zuma/chains_itp$ python ../montepython/MontePython.py info lckb_bao/
Running Monte Python v2.2.2

--> Finding global maximum of likelihood
--> Removing burn-in
--> Scanning file lckb_bao/2017-03-12_100000_3.txt : Removed 16
6 non-markovian points, 0 points of burn-in, keep 10397 steps
2017-03-12_10_1.txt : Removed 0
non-markovian points, 2 points of burn-in, keep 1 steps
2017-03-12_100000_4.txt : Removed 57
non-markovian points, 0 points of burn-in, keep 10405 steps
2017-03-12_100000_8.txt : Removed 0
non-markovian points, 2 points of burn-in, keep 6783 steps
2017-03-12_100000_7.txt : Removed 0
non-markovian points, 4 points of burn-in, keep 13666 steps
2017-03-12_100000_1.txt : Removed 92
non-markovian points, 0 points of burn-in, keep 7975 steps
2017-03-12_100000_6.txt : Removed 0
non-markovian points, 4 points of burn-in, keep 20 steps
2017-03-12_100000_5.txt : Removed 15
5 non-markovian points, 0 points of burn-in, keep 11158 steps
2017-03-12_10_2.txt : Removed 0
non-markovian points, 2 points of burn-in, keep 1 steps
--> Computing mean values
--> Computing variance
--> Computing convergence criterium (Gelman-Rubin)
--> R-1 is 0.002068 for Omega_b
0.001280 for Omega_cdm
0.002191 for Omega_k
0.002137 for Omega_Lambda
-----
--> Computing histograms for Omega_b
--> Computing histograms for Omega_cdm
--> Computing histograms for Omega_k
--> Computing histograms for Omega_Lambda
-----
--> Saving figures to .pdf files
--> Writing .info and .tex files
```

All options explained

```
python montepython/MontePython.py info --help
```

A very minimal run

Write `lckb.param`:

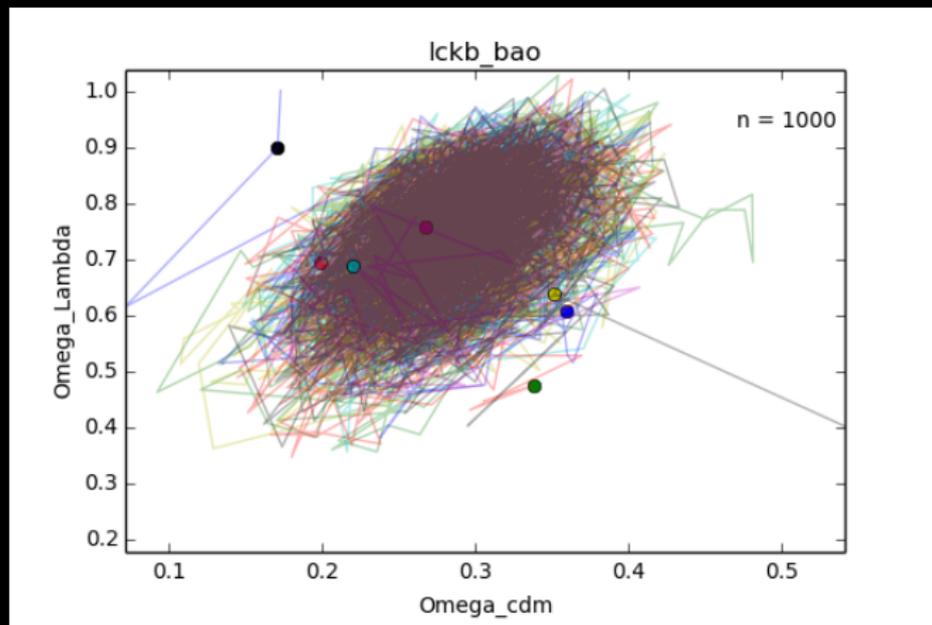
```
data.experiments=['bao_boss', 'bao_boss_aniso']
#Cosmo parameteress          [mean, min, max, sigma, scale, type]
data.parameters['Omega_b'] = [0.045, 0.01, None, 0.01, 1, 'cosmo']
data.parameters['Omega_cdm'] = [0.3, 0, None, 0.1, 1, 'cosmo']
data.parameters['Omega_k'] = [0.0, -0.5, 0.5, 0.1, 1, 'cosmo']
#Fixed parameters (sigma = 0)
data.parameters['H0'] = [67.8, None, None, 0, 1, 'cosmo']
data.cosmo_arguments['YHe'] = 0.24
#derived parameters
data.parameters['Omega_Lambda'] = [1, None, None, 0, 1, 'derived']
#mcmc parameters
data.N=10
data.write_step=5
```

Run ~ 7 chains with

```
python montepython/MontePython.py run -o chains/lckb_bao \
-p lckb_param --update 300 -N 100000
```

A very minimal run

The 7 chains explore the parameter space



Chains named yyyy-mm-dd_N_n.txt (date_points_id)

A very minimal run

Analyze:

```
python montepython/MontePython.py info chains/lckb_bao
```

- `lckb_bao.tex` → table with MCMC results

Param	best-fit	mean $\pm\sigma$	95% lower	95% upper
Ω_b	0.03595	$0.03977^{+0.0095}_{-0.015}$	0.01662	0.06547
Ω_{cdm}	0.2931	$0.2872^{+0.049}_{-0.048}$	0.1892	0.3847
Ω_k	-0.1183	$-0.08087^{+0.11}_{-0.14}$	-0.3182	0.1755
Ω_Λ	0.7891	$0.7538^{+0.12}_{-0.091}$	0.542	0.9564

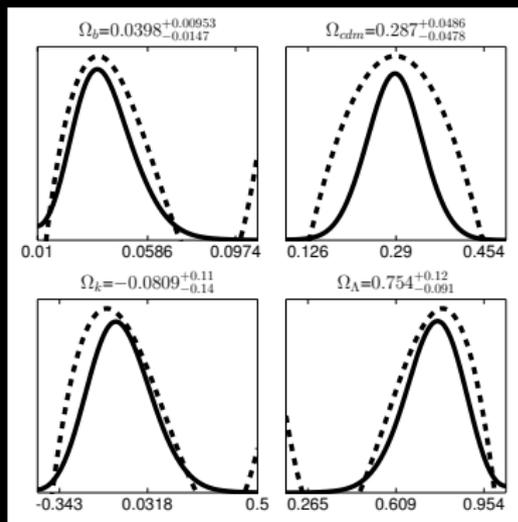
$-\ln \mathcal{L}_{\min} = 5.57269$, minimum $\chi^2 = 11.15$

- `lckb_bao.covmat` → covariance matrix
- `lckb_bao.bestfit` → best fit values

→ arguments for another run

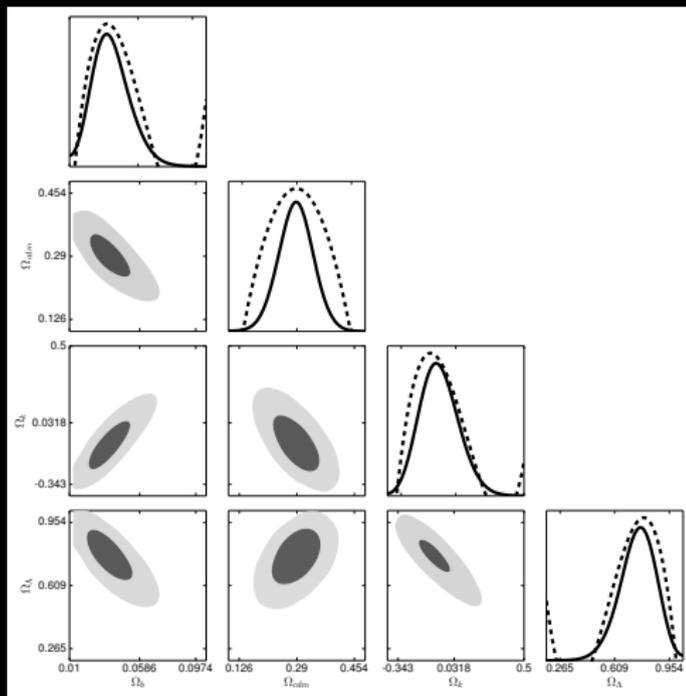
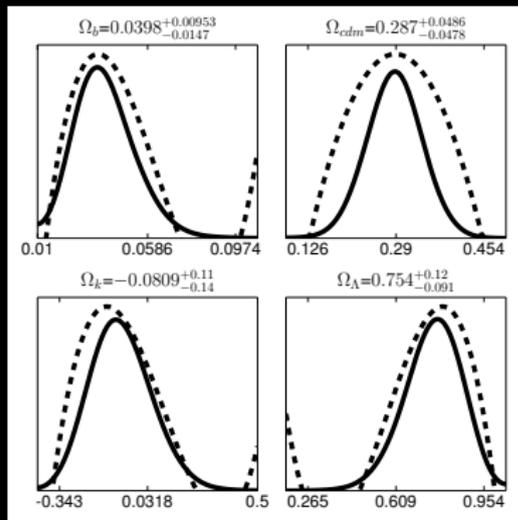
A very minimal run

In `lckb_bao/plots`:



A very minimal run

In `lckb_bao/plots`:

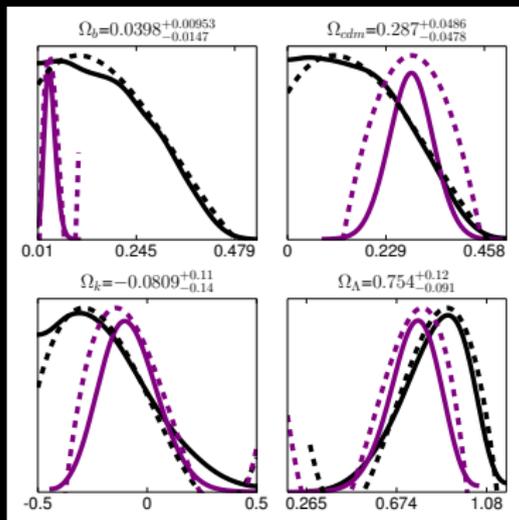


Comparing several runs

Run chains `lckb_sne` with `data.experiments=['sne']`

Analyze: `python ... info chains/lckb_sne chains/lckb_bao`

In `lckb_sne/plots`:

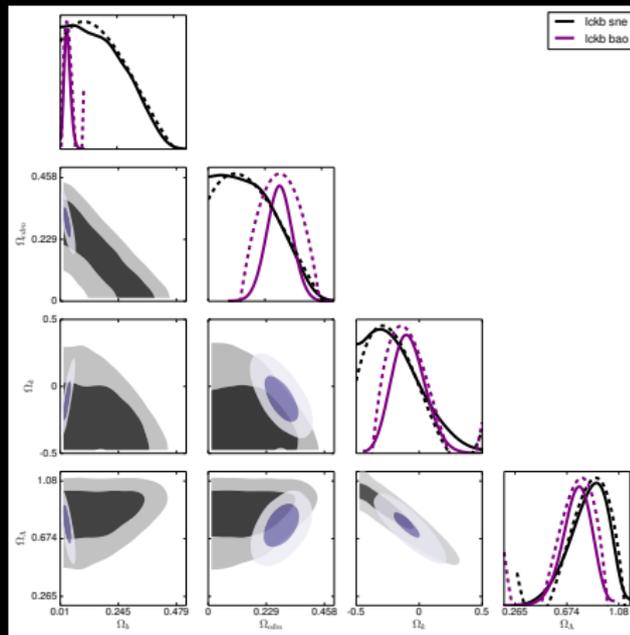
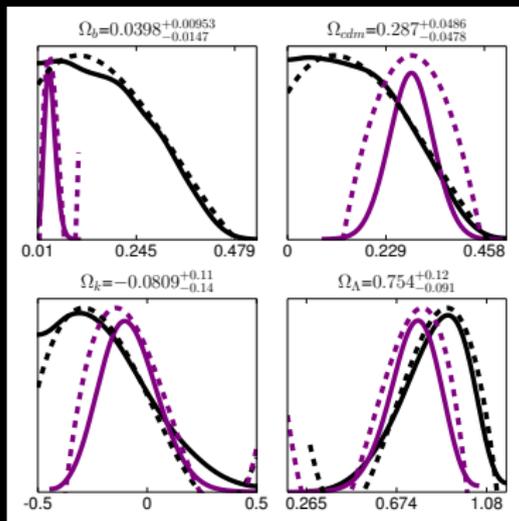


Comparing several runs

Run chains `lckb_sne` with `data.experiments=['sne']`

Analyze: `python ... info chains/lckb_sne chains/lckb_bao`

In `lckb_sne/plots`:



Experiments and Likelihoods

Simplest example ever: Prior on H_0 (1103.2976)

The Hubble Space Telescope measured $h_{obs} = 0.738 \pm 0.024$

$$\log(\mathcal{L}) = -\frac{1}{2} \frac{(h_{th} - h_{obs})^2}{\sigma_h^2}$$

Likelihood (montepython/likelihoods/hst/__init__):

```
from montepython.likelihood_class import Likelihood_prior
class hst(Likelihood_prior):
    def loglkl(self, cosmo, data):
        h = cosmo.h()
        loglkl = -0.5 * (h - self.h) ** 2 / (self.sigma ** 2)
        return loglkl
```

Data (montepython/likelihoods/hst/hst.data):

```
# Values for Hubble Space Telescope (following 1103.2976)
hst.h      = 0.738
hst.sigma  = 0.024
```

Likelihood rules

- Likelihoods in directory `montepython/likelihoods/l_name`
- Needed files: `__init__.py` and `l_name.data`
- `__init__.py` defines a class, inheriting from `Likelihood`
- Contains function `loglkl` $\rightarrow \log(\mathcal{L})$

Introducing your own Likelihoods

- Follow the above rules
- Inspire yourself with the examples
- \exists similar likelihood? \rightarrow you can inherit its methods!
- You can use additional python packages

(See also B. Audren's lecture on likelihoods)

Conclusions

- Brings all the power of CLASS to Python
- Easy to run chains and analyze likelihoods
- Many available experiments
- Advantages from object oriented features in python
 - Add likelihoods
 - Add samplers or other features
- This just scratches the surface, many more options!

(See also B. Audren's slides)

The hi_class academy

Coming soon!



www.hiclass-code.net

- Set of interrelated projects:
 - ✦ Theory & model building
 - ✦ Implementation and phenomenology
 - ✦ Compare with data
- Collaboration → Publishable results
 - ✦ Review of models
 - ✦ Observational constraints
- Stay tuned for more info!