# roofit\_example

September 6, 2017

## 1 RooFit Tutorial: Fit Cosmological Models to Type 1a Supernova Data

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#### 1.1 Introduction

In this tutorial, I show how to use the probability modeling package **RooFit**, released with the CERN data analysis package **ROOT**, to perform simple fits of cosmological models to Type 1a supernova data. See the **minuit\_example.ipynb** to get some background to this problem.

prerequisite: some familiarity with Python and ROOT is helpful.

## 1.2 Other Dependencies

This tutorial uses the github package **histutil**, which contains some simple ROOT-based utilities. To install this package do

git clone https://github.com/hbprosper/histutil.git and source the setup.sh script.

#### 1.3 PART 1: Create Probability Model

Note: if you want to make your plots interactive, do **%jsroot on**. But, it is off by default so that the LaTex annotation works correctly.

Welcome to JupyROOT 6.10/02

## 1.3.1 Model parameters

- ID: model identifier
- free: specifies whether parameter is free
- name: name of parameter
- guess: starting (or fixed) value of parameter
- step: step size during minimization

• min, max: parameter range

```
In [2]: #
                                 free, name, quess,
                                                     step,
                                                             min, max
       PARAMS = {'LCDM' :
                            [0, [(True,
                                        'OM',
                                                1, 1.e-3,
                                                                0, 10),
                                        'OL',
                                                                -10, 10),
                                 (True,
                                                0, 1.e-3,
                                (True,
                                        'HO',
                                               70, 1.e-2,
                                                                0, 200)]],
                 'phantom': [1, [(False, 'OM',
                                                1, 1.e-3,
                                                               0, 10),
                                (False, 'OL',
                                                0, 1.e-3,
                                                                -10, 10),
                                        'HO',
                                                                0, 200),
                                 (True,
                                                70, 1.e-2,
                                (True,
                                       'n',
                                               2, 1.e-3,
                                                                0,
                                                                     10)]]
               }
       # define ranges for redshifts and distance moduli
       ZMIN = 0.0
       ZMAX = 1.5
       MUMIN = 32.0
       MUMAX = 48.0
```

#### 1.3.2 Choose model

• MODEL = 'LCDM' or 'phantom'

```
In [3]: MODEL = 'phantom'
```

#### 1.3.3 Compile C++ classes Model and CosmicCode

- **Model** defines  $\Omega(a)$  for the cosmological models
- **CosmicCode** computes the distance modulus
- The model identifier, ID, is used later in some inline C++ code. Therefore, ID needs to be added to the ROOT namespace; that is, ROOT needs to know about it.

```
In [4]: def compileCode(modelname, modelparams):
    ROOT.gROOT.ProcessLine(open('../CosmicCode.cc').read())

# make sure model name is valid
    if not modelparams.has_key(modelname):
        sys.exit("** unknown model %s" % modelname)

# return model id and model parameters
    ID, params = modelparams[modelname]

# add ID to ROOT namespace
    ROOT.gROOT.ProcessLine('int ID = %d;' % ID)
    return params

In [5]: params = compileCode(MODEL, PARAMS)
```

#### 1.3.4 A useful bit of ROOT magic: mixing in a bit of C++

Here we create a bit of C++ code, which will be visible to both ROOT and Python. The object **code** and the function **distanceModulus** are used below in the definition of the probability model.

check that ROOT.distanceModulus is visible to Python

```
In [7]: print ROOT.distanceModulus(1, 0.3, 0.7, 70, 2)
43.9845970443
```

#### 1.3.5 Create workspace

Create an empty workspace called **Type1a**. This is useful for at least two reasons. First, you may wish to save the probability model you have created to a ROOT file. Second, it is much more convenient to use the workspace methods to build a model than to use the **RooFit** C++ classes directly.

## 1.3.6 Create parameters

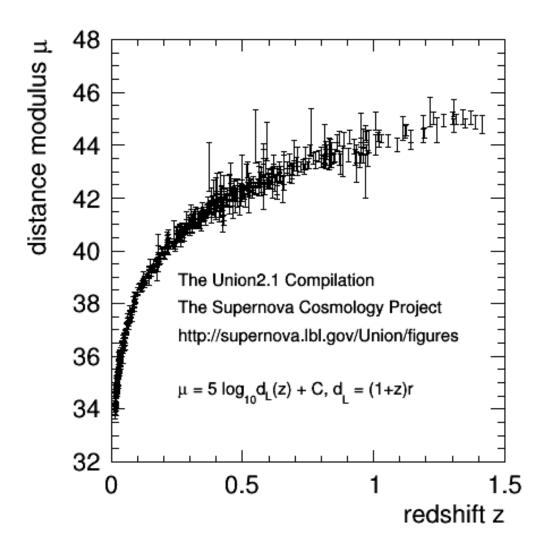
Create RooFit parameters using the workspace factory method.

```
In [9]: def createParameters(wspace, params,
                             zmin=ZMIN, zmax=ZMAX, mumin=MUMIN, mumax=MUMAX):
            # measured redshift
            wspace.factory('z[%f, %f]' % (zmin, zmax))
            # measured distance modulus
            wspace.factory('x[%f, %f]' % (mumin, mumax))
            # distance modulus uncertainty
            wspace.factory('dx[0, 2]')
            wspace.defineSet('set_data', 'z,x,dx')
            # cosmological parameters
            args = 'z'
            for (free, name, guess, step, xmin, xmax) in params:
                cmd = '%s[%f, %f, %f]' % (name, guess, xmin, xmax)
                wspace.factory(cmd)
                args += ',%s' % name
                if not free:
                    wspace.var(name).setConstant()
            wspace.defineSet('set_arguments', args)
            wspace.Print()
            return args
In [10]: arguments = createParameters(wspace, params)
        print 'arguments for distanceModulus: (%s)' % arguments
arguments for distanceModulus: (z,OM,OL,HO,n)
RooWorkspace(Type1a) Type1a contents
variables
-----
(HO,OL,OM,dx,n,x,z)
named sets
-----
set_arguments:(z,OM,OL,HO,n)
set_data:(z,x,dx)
1.3.7 Load Type 1a supernova data
In [11]: # -----
         # read Type Ia data
```

```
\# format: name, z, x, dx
# name: name of supernova
         measured redshift of supernova
# x +/- dx: measured distance modulus
# -----
def readData(wspace, filename):
   import os
   from array import array
   from string import split, atof
   if not os.path.exists(filename):
       sys.exit("** can't open file %s" % filename)
    # skip first 5 lines and convert 2nd through 4th
    # columns to floats
   data = map(lambda x: map(atof, x[1:-1]),
                  map(split,
                          open(filename).readlines()[5:]))
    # create a RooFit data set
   set_data = wspace.set('set_data')
   dataset = ROOT.RooDataSet('data', 'Type 1a data', set_data)
   print "number of observations: %d" % len(data)
   print "%5s\t%10s\t%10s +/- %-10s" % ('', 'z', 'x', 'dx')
    # and fill it with the supernova data
   # also copy data to arrays
   z = array('d')
   x = array('d')
   dz = array('d')
   dx = array('d')
   for ii, d in enumerate(data):
       z.append(d[0])
       x.append(d[1])
       dz.append(0)
       dx.append(d[2])
       set_data['z'].setVal(z[-1])
       set_data['x'].setVal(x[-1])
       set_data['dx'].setVal(dx[-1])
       dataset.add(set_data)
       if ii % 100 == 0:
           print "%5d\t%10.3f\t%10.4f +/- %-10.4f"%\
             (ii, z[-1], x[-1], dx[-1])
    # import data set into workspace
   getattr(wspace, 'import')(dataset)
```

```
wspace.Print()
             return (z, x, dz, dx)
In [12]: data = readData(wspace, '../SCPUnion2.1_mu_vs_z.txt')
number of observations: 580
                                        x +/- dx
                                  35.3466 +/- 0.2239
                  0.028
    0
  100
                  0.065
                                  37.3067 +/- 0.1628
  200
                  0.194
                                  39.9615 +/- 0.1264
  300
                  0.620
                                  43.2280 +/- 0.3903
  400
                  0.710
                                  43.0220 +/- 0.1843
  500
                  0.564
                                  42.3729 +/- 0.2920
RooWorkspace(Type1a) Type1a contents
variables
-----
(HO,OL,OM,dx,n,x,z)
datasets
-----
RooDataSet::data(z,x,dx)
named sets
-----
set_arguments:(z,OM,OL,HO,n)
set_data:(z,x,dx)
In [13]: setStyle()
         def plotData(data, code, zmin=ZMIN, zmax=ZMAX, mumin=MUMIN, mumax=MUMAX):
             from array import array
             print "plot data"
             z, x, dz, dx = data
             ndata = len(z)
             print "number of observations: %d" % ndata
             g = mkgraphErrors(z, x, dz, dx,
                               "redshift z",
                               "distance modulus #mu",
                               zmin, zmax,
                               ymin=mumin,
```

```
ymax=mumax,
                                color=ROOT.kBlack)
             ROOT.SetOwnership(g, 0)
             g.SetName('dataplot')
             g.SetTitle('')
             g.SetMarkerSize(0.2)
             c = ROOT.TCanvas("fig_data", "SN1a data", 500, 500)
             ROOT.SetOwnership(c, 0)
             c.cd()
             g.Draw("ap")
             xpos = 0.32
             ypos = 0.50
             textsize = 0.035
             scribe = Scribe(xpos, ypos, textsize)
             ROOT.SetOwnership(scribe, 0)
             scribe.write("The Union2.1 Compilation")
             scribe.write("The Supernova Cosmology Project")
             scribe.write("http://supernova.lbl.gov/Union/figures")
             scribe.write("")
             scribe.write("#mu = 5 \log_{10}d_{L}(z) + C, d_{L} = (1+z)r")
             c.SaveAs('.pdf')
             c.Draw()
             return g
In [14]: gd = plotData(data, ROOT.code)
plot data
number of observations: 580
Info in <TCanvas::Print>: pdf file ./fig_data.pdf has been created
```



## 1.3.8 Create probability model

We assume a Gaussian probability density function (pdf),

$$f(x | z, \sigma, \theta) = Gauss(x, \mu(z, \theta), \sigma),$$

where  $\theta$  are the parameters of the cosmological model. Neglecting correlations amongst the data, the likelihood of the data is

$$p(x \mid z, \sigma, \theta) = \prod_{i=1}^{N} f(x_i \mid z_i, \sigma_i, \theta),$$

with N=580. For a Bayesian calculation, we would, in addition, need to specify a prior density  $\pi(\theta)$  for the cosmological parameters. We assume that  $z_i$  and  $\sigma_i$  are known with negligible uncertainty.

```
In [15]: def createModel(wspace, arguments):
            #-----
            # create distance modulus expression
            # "mu".
            # note use of compiled C++ program
            # for expressions that are too
            # complicated to be written inline
            print "create model"
            set_arguments = wspace.set('set_arguments')
            cmd = 'distanceModulus(%s)' % arguments
            print cmd
            mu = ROOT.RooFormulaVar('mu', '#mu', cmd, ROOT.RooArgList(set_arguments))
            # import the "mu" expression into workspace
            getattr(wspace,'import')(mu)
            # finally, create probability model
            wspace.factory('Gaussian::model(x, mu, dx)')
            wspace.Print()
In [16]: createModel(wspace, arguments)
create model
distanceModulus(z,OM,OL,HO,n)
RooWorkspace(Type1a) Type1a contents
variables
(HO,OL,OM,dx,n,x,z)
p.d.f.s
RooGaussian::model[ x=x mean=mu sigma=dx ] = 0.601823
functions
RooFormulaVar::mu[actualVars=(z,OM,OL,HO,n) formula="distanceModulus(z,OM,OL,HO,n)"] = 42.7578
datasets
_____
RooDataSet::data(z,x,dx)
named sets
```

```
set_arguments:(z,OM,OL,HO,n)
set_data:(z,x,dx)
```

## 1.3.9 Save workspace to a ROOT file

```
In [17]: filename = '%s_workspace.root' % MODEL
    if wspace.writeToFile(filename) != 0:
        print "** problem saveing workspace to ROOT file %s" % filename
```

#### 1.4 PART 2: Perform Fit

Ordinarily, this part would be performed in a separate session, which would require loading the workspace back into memory as follows:

```
filename = '%s_workspace.root' % MODEL
rfile = TFile(filename)
if not rfile.IsOpen():
    exit("** can't open file %s" % filename)

name = 'Type1a'
wspace = rfile.Get(name)
if wspace == None:
    exit("** can't get workspace %s from file %s" % (name, filename))
```

However, we shall just continue with the workspace already in memory.

```
In [18]: wspace.Print()

RooWorkspace(Type1a) Type1a contents

variables
-------
(H0,OL,OM,dx,n,x,z)

p.d.f.s
------
RooGaussian::model[ x=x mean=mu sigma=dx ] = 0.601823

functions
------
RooFormulaVar::mu[ actualVars=(z,OM,OL,HO,n) formula="distanceModulus(z,OM,OL,HO,n)" ] = 42.7578

datasets
-------
RooDataSet::data(z,x,dx)
```

```
named sets
-----
set_arguments:(z,OM,OL,HO,n)
set_data:(z,x,dx)
In [19]: def performFit(wspace, params, printLevel=-1):
          print "="*80
          print "performing fit..."
          swatch = ROOT.TStopwatch()
          swatch.Start()
          # save results of fit to results
          ROOT.RooMsgService.instance().setGlobalKillBelow(ROOT.RooFit.FATAL)
          r = wspace.pdf('model').fitTo(wspace.data('data'),
                                  ROOT.RooFit.Save(),
                                  ROOT.RooFit.PrintLevel(printLevel))
          print "real time: %10.3f s" % swatch.RealTime()
          print "="*80
          r.Print()
          print "="*80
          results = []
          npar = 0
          for p in params:
             free, name = p[:2]
             results.append((wspace.var(name).getVal(), wspace.var(name).getError()))
             if free:
                npar += 1
          return (results, npar)
In [20]: results, npar = performFit(wspace, params)
______
performing fit...
real time:
            3.049 s
______
_____
 RooFitResult: minimized FCN value: 1447.61, estimated distance to minimum: 8.69769e-10
            covariance matrix quality: Full, accurate covariance matrix
            Status: MINIMIZE=0 HESSE=0
   Floating Parameter FinalValue +/- Error
 _____
                    7.0425e+01 +/- 4.53e-01
               HO
                n 2.8213e+00 +/- 1.87e-01
```

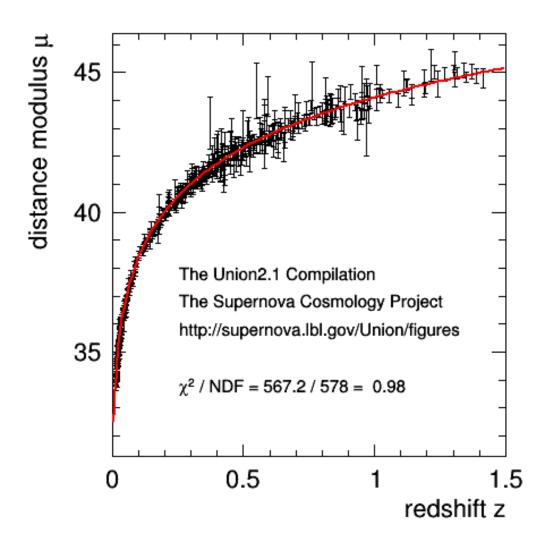
### 1.5 Plot Results

```
In [21]: def annotate(modelname, scribe, results, offset=0.01):
             if modelname == 'LCDM':
                 OM = tuple(results[0])
                 OL = tuple(results[1])
                 scribe.write("#LambdaCDM model")
                 scribe.write("")
                 scribe.write("#Omega(a) = #frac{#Omega_{M}}{a^{3}} + "
                       "#frac{(1 - #0mega_{M} - #0mega_{#Lambda})){a^{2}}"
                                " + #Omega_{#Lambda}",
                                offset)
                 scribe.write("")
                 scribe.write("\#0mega_{M} = \%5.2f \#pm \%-5.2f" % OM,
                                offset)
                 scribe.write("#0mega_{#Lambda} = %5.2f #pm %-5.2f " % OL,
                                offset)
             else:
                 n, dn = results[3]
                 x = 3.0/(2*n)
                 G = ROOT.TMath.Gamma(x)
                 T = G*ROOT.TMath.Sqrt(2.718)*2**x/n
                 scribe.write("phantom model")
                 scribe.write("")
                 scribe.write("#Omega(a) = #frac{#Omega_{M}}{a^{3}} + "
                            "#frac{e^{a^{n}-1} - \#Omega_{M}}{a^{3}}", offset)
                 scribe.write("")
                 scribe.write("H_{0}t = #sqrt{e} 2^{3/(2n)} #Gamma(3/(2n), a^{n}/2)/n", offset)
                 scribe.write("")
                 scribe.write("H_{0}t_{rip}) = #sqrt{e} 2^{3/(2n)} #Gamma(3/(2n))/n = %4.2f" % T,
                                offset)
                 scribe.write("")
                 scribe.write("where \#Gamma(s, x) = \#int_{0}^{x} t^{s-1} e^{-t} dt", offset)
                 scribe.write("and n = \frac{4.2f}{n}, offset)
In [22]: def plotModel(modelname, code, results, npar,
                       gd, data,
                       zmin=ZMIN, zmax=ZMAX, mumin=MUMIN, mumax=MUMAX):
             from array import array
             p = array('d')
             for value, error in results:
                 p.append(value)
             z, x, dz, dx = data
             ndata = len(z)
```

```
# compute chisq
chi2 = 0.0
for i in xrange(ndata):
    mu = code.distanceModulus(z[i], p)
    c = (x[i] - mu)/dx[i]
    chi2 += c*c
NDF = ndata - npar # number of degrees of freedom
# compute curve
nz = 100
zstep = (zmax - zmin) / nz
zz = array('d')
mu = array('d')
for ii in xrange(nz):
    zz.append( (ii+0.5)*zstep )
    mu.append( code.distanceModulus(zz[-1], p) )
g = mkgraph(zz, mu,
            "redshift z",
            "distance modulus #mu",
            zmin, zmax, color=ROOT.kRed, lwidth=2)
ROOT.SetOwnership(g, 0)
g.SetName('model')
g.SetTitle('')
c = ROOT.TCanvas("fig_%s_fit" % modelname, "SN1a model fit", 500, 500)
ROOT.SetOwnership(c, 0)
c.cd()
g.Draw('ac')
gd.Draw('psame')
g.Draw('csame')
xpos = 0.32
ypos = 0.50
textsize = 0.035
scribe = Scribe(xpos, ypos, textsize)
ROOT.SetOwnership(scribe, 0)
scribe.write("The Union2.1 Compilation")
scribe.write("The Supernova Cosmology Project")
scribe.write("http://supernova.lbl.gov/Union/figures")
scribe.write("")
scribe.write("#chi^{2} / NDF = \%5.1f / \%d = \%5.2f" % (chi2, NDF, chi2/NDF))
c.Draw()
c.SaveAs(".pdf")
```

In [23]: plotModel(MODEL, ROOT.code, results, npar, gd, data)

Info in <TCanvas::Print>: pdf file ./fig\_phantom\_fit.pdf has been created



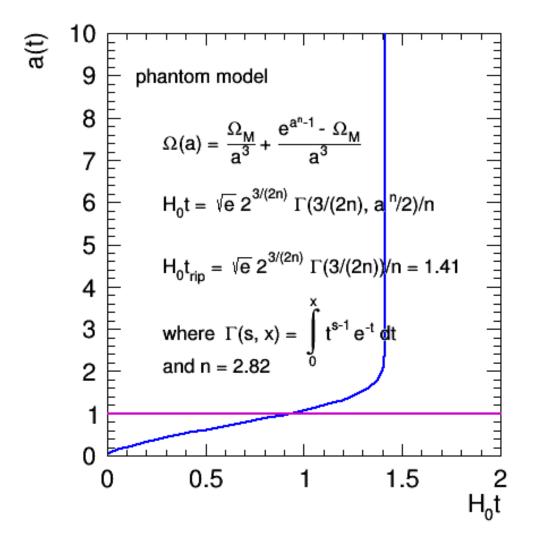
```
In [24]: def plotScaleFactor(modelname, code, results, amax=10, tmax=2):
    from array import array

p = array('d')
    for value, error in results:
        p.append(value)

# create a vs H0 t plot
a = array('d'); a.fromlist(code.N*[0])
t = array('d'); t.fromlist(code.N*[0])
```

```
code.scaleFactor(amax, p, t, a)
             g = mkgraph(t, a,
                         "H_{0}t}", "a(t)",
                         0, tmax, color=ROOT.kBlue, lwidth=2)
             ROOT.SetOwnership(g, 0)
             g.SetName('scaleFactor')
             # create horizontal line at a = 1
             x = array('d'); x.append(0); x.append(tmax)
             y = array('d'); y.append(1); y.append(1)
             glineH = mkgraph(x, y, '', '', 0, tmax,
                              color=ROOT.kMagenta+1, lwidth=2)
             ROOT.SetOwnership(glineH, 0)
             glineH.SetName('line')
             c = ROOT.TCanvas("fig_%s_scaleFactor" % modelname,
                              "SN1a scalefactor",
                              500, 500)
             ROOT.SetOwnership(c, 0)
             c.cd()
             htmp = mkhist1('htmp', 'H_{0}t', 'a(t)',
                                50, 0, tmax, ymin=0, ymax=10)
             ROOT.SetOwnership(htmp, 0)
             htmp.Draw()
             g.Draw('csame')
             glineH.Draw('csame')
             offset = 0.05
             xpos = 0.25
             ypos = 0.87
             scribe = Scribe(xpos, ypos)
             ROOT.SetOwnership(scribe, 0)
             annotate(modelname, scribe, results, offset)
             c.Draw()
             c.SaveAs(".pdf")
In [25]: plotScaleFactor(MODEL, ROOT.code, results)
```

Info in <TCanvas::Print>: pdf file ./fig\_phantom\_scaleFactor.pdf has been created



In []: