

BSM phenomenology the computational way

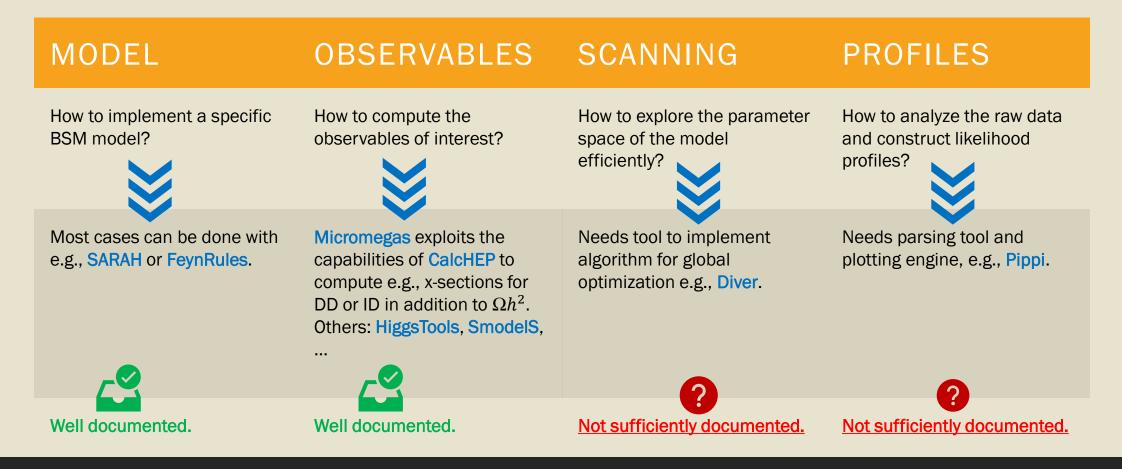
CATALINA ESPINOZA IF-UNAM / CONACYT

Taller "Más allá del Modelo Estándar y Astropartículas" | IFUNAM | 15 - marzo - 2023

### Summary

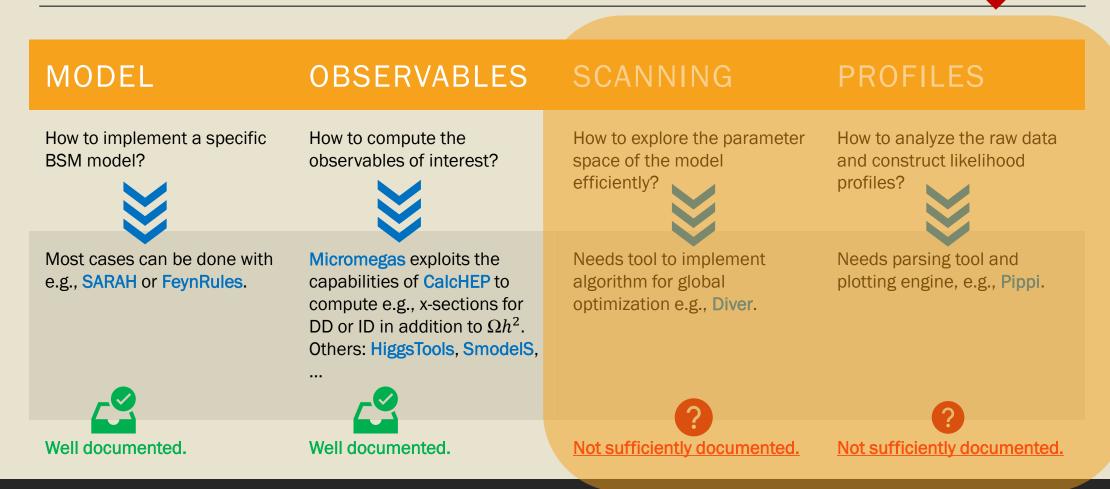
Introduction:	Will focus on BSM models of Particle Physics, but only on the part of scanning and construction of likelihood profiles of a given model.
Scanning the parameter space (PS):	Will describe a tool to explore the PS of a model computing a composite likelihood function with respect to the observables of interest.
Parsing the raw data and plotting the likelihood profiles:	Will describe a tool to generate the actual plots that constitute the main results of the numerical analysis.
Tutorial:	This is part of planned tutorials under development that will be available elsewhere. This is a summarized version of the tutorial on likelihood profiles, which will be much more detailed.

# Phenomenologing the computational way ...



# Phenomenologing the computational way ...

#### Will focus on this part!



### Simple BSM model example

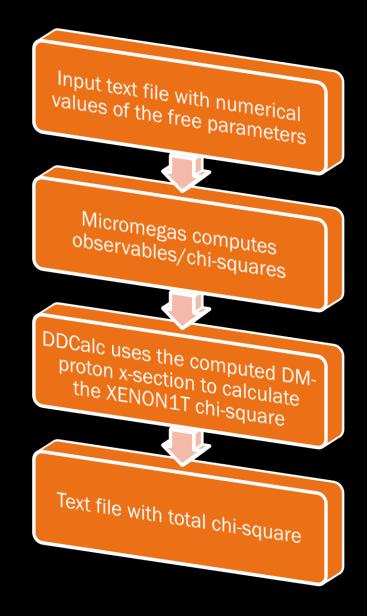
Scalar	Two SU(2) Higgs doublets H1 and H2 and one singlet $\phi$
sector	The singlet mixes with the real parts of the neutral components of the doublets into three CP-even physical scalars h1, h2 and h3
Dark	One majorana fermion coupled to the singlet: $y_{\Psi} \overline{\Psi}_R^C \Psi_R \phi$
sector	The mixing in the scalar sector induces couplings of the physical scalars with the DM and therefore we have effective DM-quark couplings and thus DM-nucleon scattering

### Observables

We calculate higgs mass:  $\chi_h^2 = (m_h - m_h^{PDG})^2 / \sigma_h^2$ the observables and from them the chi-square functions Higgs mass:  $\chi_{h}^2 = (\Omega h^2 - \Omega^{Planck} h^2)^2 / \sigma_{\Omega h}^2$ DM-proton x-sect:  $\chi_{DD}^2 = \dots$  (from DDCalc tool)

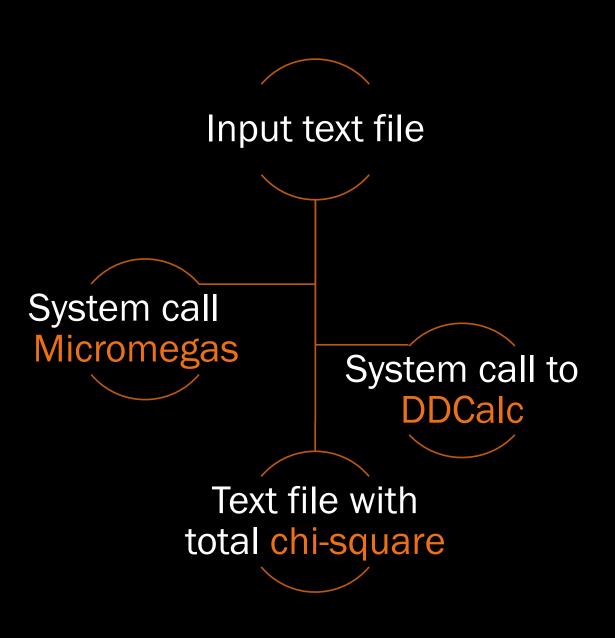
#### For example...

- Use a text file to pass the numerical values of the free parameters (1 single point of parameter space) to micromegas
- Micromegas computes relic density, higgs mass and DMproton x-section and writes their values to a text file with also the chi-square from the simple formulas above
- DDCalc reads form the text file the DM-proton x-section and computes the chi-square relative to the XENON-1T experiment, writes the result to a text file



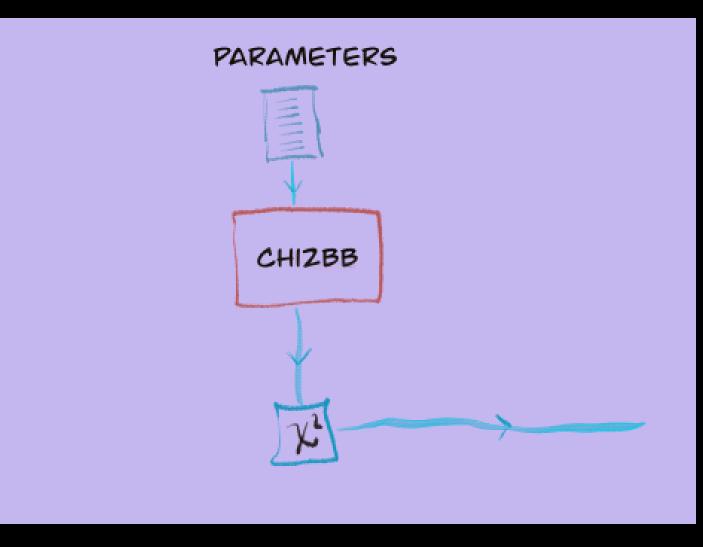
#### In practice ...

- From the scanning program we communicate with Micromegas and DDCalc through "system calls", for example in C:
- system("./my\_micromegas\_prog par\_file.txt")
- This executes Micromegas with input file 'par\_file.txt' previously created with the values of the free variables for 1 point of parameter space



#### We call this: Chi-square Black Box





### Exploring the parameter space

□ Next, we use a global optimization algorithm to explore the parameter space and find the minimum of the composite (total) chi-square function

□ Pictorially ... suppose we want to explore a 2-dimensional surface



# Number of points (NP)

- Choose NP points on the surface randomly
- Using the chi2BB, compute the heights for all of them
- Save to disk the coordinates, observables, heights of all points
- This is called a 'generation', with a population of NP points

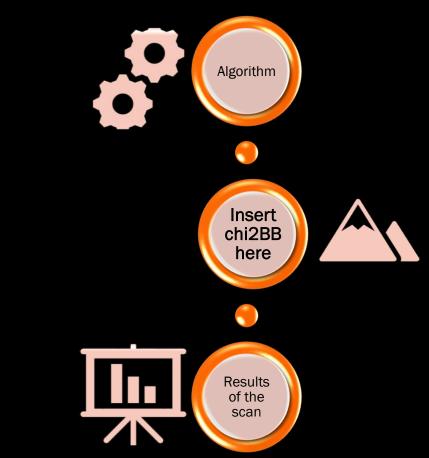


#### Clever algorithm

- Use some algorithm to 'move around' the points to new locations (a new generation) and repeat
- Stop repeating when certain
   'convergence criteria' is met (there is a high probability that the global minimum has been found)
- Clearly, the larger the 'population' is, the analysis is better
- Typically, NP=20,000 is recommended for dimension ~ 20 or less



The Global Optimisation Algorithm Black Box ➡ For practical purposes we don't need to know how the algorithm works: it is just a minimization black box



#### The 'Diver' package

We use this package authored by Pat Scott, which can be obtained from GitHub ...

🍪 🕠 GitHub - patscott/Di	iver: F× +			
< → c (	C A https://github.com/patscott/Diver		▣ ☆	V 🗈 🕈 🖑 H 🚳
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patscott / Diver     Public	े भी Pull requests 10 ⓒ Actions 田 P	rrojects 🛈 Security 🗠 Insights		لِ Notifications 💡 ۱
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		Lunfit_points option causing cae65fe on Dec 20, 2019	① 169 commits	Public Package for Differential Evolution
	🖿 build	Shifted main F code into root dir of repo.	8 years ago	🛱 Readme
	🖿 example_c	Updated readme	4 years ago	☆ 4 stars ⊙ 4 watching
	example_cpp	Minor changes to comments and status messages. This accepts pull	4 years ago	약 3 forks
	example_f	Added test of discard_unfit_points to Fortran example	4 years ago	
	🖿 include	Added option discard_unfit_points to extend max_acceptable_value	4 years ago	Releases
	🖿 lib	Shifted main F code into root dir of repo.	8 years ago	🛇 <b>6</b> tags
	src src	Fixed bugs with new discard_unfit_points option causing	4 years ago	
	🕒 .gitignore	Minor bugfix: exit -> cycle in routine for randomly generating new	9 years ago	Packages
	L HISTORY	Updated to v1.0.5	4 years ago	
	C README.md	Updated readme	4 years ago	
	🕒 ScannerBit.pdf	Updated ScannerBit/Diver paper to arxiv v2	6 years ago	Contributors <sub>3</sub>
	🕒 logo.pdf	Added logo	6 years ago	patscott Pat Scott
	ြ logo.svg	Added logo	6 years ago	jacopo-chevallard Jacopo Chevallard
	🗅 makefile	Fixed incorrect build dependencies for init module.	6 years ago	EthanCarragher Ethan Carragher
	i≣ README.md			
	Diver			Languages

#### The 'Diver' package

Click on 'Code' and then copy the address ...

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		Go to file	Code 🗸	
	Local	Codespaces		Public Package for Differential Evolution
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#### The 'Diver' package

From the 'Readme' we see that it compiles with a simple instruction ... Diver is written in Fortran2003. We originally wrote it with applications in mind, but there is nothing to prevent its use in other fields.

The code and its options are described in detail in the ScannerBit paper papers that use results or insights obtained with Diver should cite this p

 Martinez, McKay, Farmer, Scott, Roebber, Putze & Conrad 2017, Euro arXiv:1705.07959

#### Compilation

The Diver build system is not really complex enough to require autotools by hand to suit your system, or call it from another makefile or the comr

To build Diver as a static library, and build all examples, do

make

To instead build Diver as a shared library, do

ake libdiver.so

o build only the static library, do

#### Installing Diver

□ This is the complete set of instructions to install Diver

You may need to install also
 openmpi in case you don't have it,
 this is to execute the program
 using several cores in parallel

```
sudo apt-get -y install openmpi-bin
git clone https://github.com/patscott/Diver.git
cd Diver
make
cd Diver/example_c/
gedit example_c.c &
```

#### Examples

❑ The tool has an example of use, the same example is presented written in C, C++ and Fortran

■ We open the C source code of the example here ...

```
sudo apt-get -y install openmpi-bin
git clone https://github.com/patscott/Diver.git
cd Diver
make
cd Diver/example_c/
gedit example_c.c &
```

#### The C example

#### □ This is the entire program

```
The idea is to adapt this code to scan our model
```

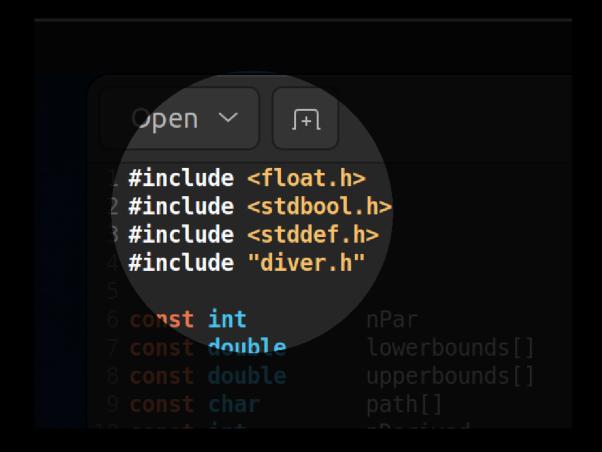
■ We can divide the code's parts in four blocks ...

```
#include <float.h>
#include <stdbool.h>
#include <stddef.h>
#include "diver.h"
const int
                  nPar
                                                                        // Dimensionality of the parameter space
                                        = 5;
const double
                  lowerbounds[]
const double
                  upperbounds[]
                                        = \{ 5., 50., 5., 50., 2. \};
                                                                        // Upper boundaries of parameter space
const char
                  path[]
                                        = "example c/output/example";
                                                                        // Path to save samples, resume files, etc
const int
                  nDerived
                                        = 0;
                                                                        // Number of derived quantities to output
                                        = 0;
                                                                        // Number of parameters that are to be treated as discrete
const int
const int
                  discrete[]
                                                                        // Indices of discrete parameters, Fortran style, i.e. starting
const bool
                  partitionDiscrete
                                       = false;
                                                                        // Split the population evenly amongst discrete parameters and
const int
                                        = 1;
const int
                  maxgen
                                       = 100;
const int
                  NP
                                       = 1000:
                                                                        // Population size (individuals per generation)
                                                                        // Size of the array indicating scale factors
const int
const double
                                       = \{0.6\};
                                                                        // Scale factor(s). Note that this must be entered as an array
const double
                                        = 0.9;
const double
                  lambda
                                        = 0.8;
const bool
                                                                        // Use current vector for mutation
                  current
const bool
                  expon
                                                                        // Use exponential crossover
                                                                        // Boundary constraint: 1=brick wall, 2=random re-initializatic
const int
                  bndry
const bool
                                                                        // Use self-adaptive choices for rand/1/bin parameters as per E
const bool
                  lambdajDE
                                                                        // Use self-adaptive rand-to-best/1/bin parameters; based on Br
const double
                  convthresh
                                        = 1.e-6;
                                                                        // Threshold for gen-level convergence: smoothed fractional imp
const int
                  convsteps
                                        = 10;
                                                                        // Number of steps to smooth over when checking convergence
const bool
                  removeDuplicates
                                                                        // Weed out duplicate vectors within a single generation
                                                                        // Calculate approximate log evidence and posterior weightings
                                        = false;
const bool
                  doBayesian
const double
                  maxNodePop
                                       = 1.9;
const double
                  Ztolerance
                                                                        // Input tolerance in log-evidence
const int
                                        = 100;
                                                                        // Save progress every savecount generations
                  savecount
const bool
                  resume
                                        = false:
                                                                        // Restart from a previous run
const bool
                  outputSamples
                                        = false:
                                                                        // Write output .raw and .sam (if nDerived != 0) files
                                                                        // Initialisation strategy: 0=one shot, 1=n-shot, 2=n-shot with
const int
                  init pop strategy
                                       = 0;
const bool
                  discard unfit points = false;
                                                                        // Recalculate any trial vector whose fitness is above max acce
                  max init attempts
const int
                                       = 10000;
const double
                  max acceptable val
                                                                        // Maximum fitness to accept for the initial generation if init
                                      = 1e6;
const int
                  seed
                                       = 1234567;
                                                                        // base seed for random number generation; non-positive or abse
const int
                                                                        // Output verbosity: 0=only error messages, 1=basic info, 2=civ
                  verbose
//Function to be minimized. Corresponds to -ln(Likelihood).
double gauss(double params[], const int param dim, int *fcall, bool *quit, const bool validvector, void** context)
 double result = 0.0;
  for (int i = 0; i<param dim; i++) result += params[i]*params[i] + 1.0;</pre>
  if (!validvector) result = DBL MAX;
  *fcall += 1:
  *quit = false;
  return result;
int main(int argc, char** argv)
 void* context = &gauss; //Not actually used in this example.
  cdiver(gauss, nPar, lowerbounds, upperbounds, path, nDerived, nDiscrete, discrete, partitionDiscrete,
         maxciv, maxgen, NP, nF, F, Cr, lambda, current, expon, bndry, jDE, lambdajDE, convthresh,
         convsteps, removeDuplicates, doBayesian, NULL, maxNodePop, Ztolerance, savecount, resume,
         outputSamples, init pop strategy, discard unfit points, max init attempts, max acceptable val, seed, context, verbose);
         //Note that prior, maxNodePop and Ztolerance are just ignored if doBayesian = false
                                                                                                                                   19
```

#### The first block

The first part is the 'include' block

Here you add libraries that you need, e.g. 'stdio.h' for handling files



## The fourth block

The last part is the 'main' function

This part does not need to be modified

■ Just make sure that the name of the chi2BB function (here 'gauss') is consistent with the third block as describe next ...

```
fieal( += 1;
     quit = false;
     return result;
55
56 int main(int argc, char** argv)
57 {
     void* context = &gauss;) //Not ac
     cdiver(gauss, nPar, lowerbounds, upper
            maxciv, maxgen, NP, nF, F, Cr,
            convsteps, removeDuplicates, do
            outputSamples, init pop strated
            //Note that prior, maxNodePop a
63
```

The third block is the function that is being minimized

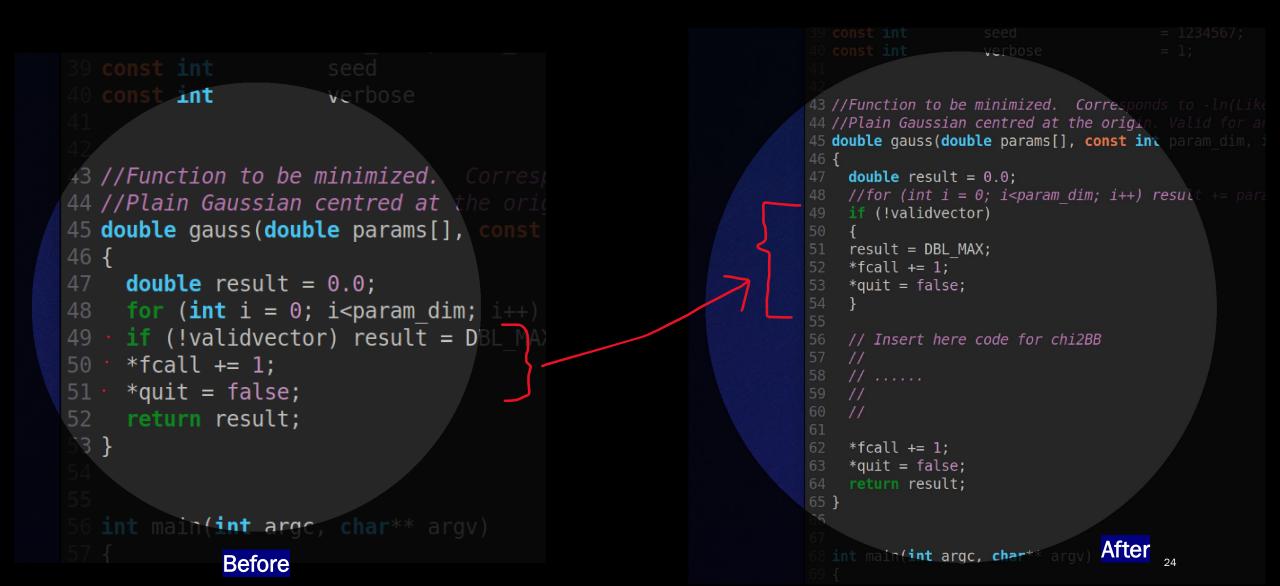
❑ We need to replace this with our chi2BB code in order to scan the parameter space of our model

```
int
                     verbose
43 //Function to be minimized. Corres
44 //Plain Gaussian centred at the onio
45 double gauss(double params[], const
46 {
     double result = 0.0;
     for (int i = 0; i<param dim; i+++)</pre>
48
     if (!validvector) result = DBL MAD
49
50
    *fcall += 1;
    *quit = false;
51
     return result;
  int main(int argc, char** argv)
```

Use to comment one line first ....



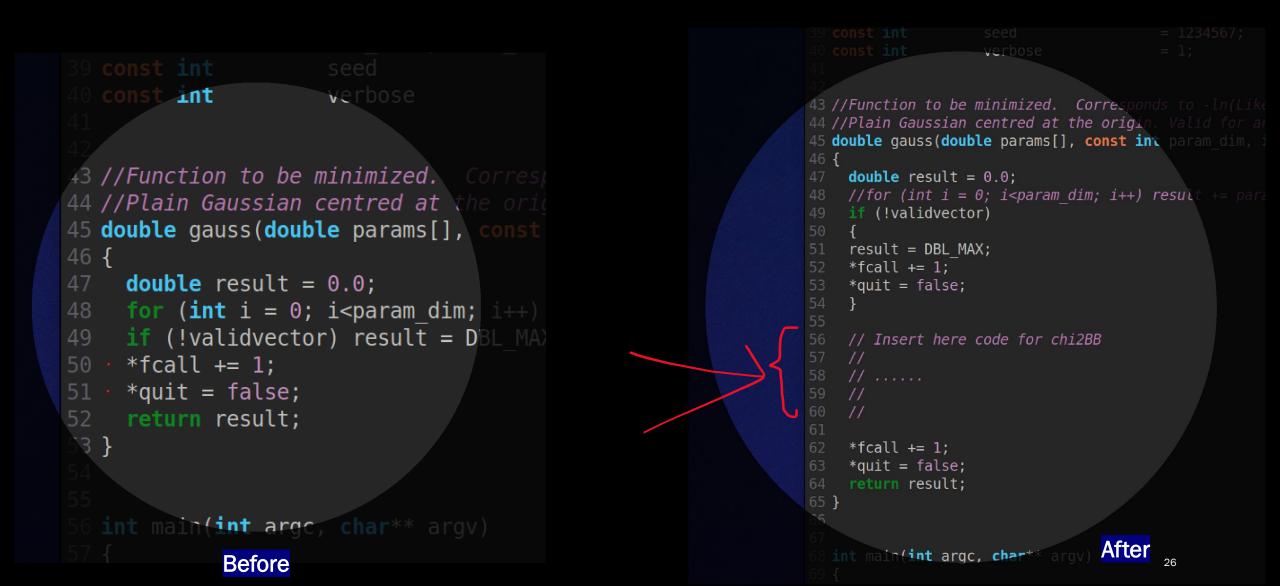
□ Then we group the next 3 lines in a single 'if' statement ...



□ Then we copy the 2 shown lines just before the 'return' statement ....



□ Finally, we write our code for the chi2BB in the space shown ...



But first let me describe the second block

In this block, global variables are defined in order to configure Diver

In most cases, almost all variables can be taken with the default values

I will describe those that need to be changed according to our model ...

Only the values in the 'pink' column must be changed ...

#include <float.h>
#include <stdbool.h>
#include <stddef.h>
#include "diver.h"

5	const	int	nPar	=	5;
	const	double	lowerbounds[]	=	{-5.,-50.,-5.,-50.,-2.};
8	const	double	upperbounds[]	=	{ 5., 50., 5., 50., 2.};
9	const	char	path[]	=	<pre>"example_c/output/example";</pre>
	const	int	nDerived	=	0;
	const		nDiscrete		0;
	const		discrete[]		{};
	const		partitionDiscrete	=	false;
	const		maxciv		1;
	const		maxgen	=	100;
	const		NP		1000;
	const		nF		1;
		double	F[]		{0.6};
		double	Cr		0.9;
		double	lambda		0.8;
	const		current		false;
	const		expon		false;
	const		bndry		3;
	const		jDE		true;
	const		lambdajDE		true;
		double	convthresh		1.e-6;
	const		convsteps		10;
	const		removeDuplicates		true;
	const		doBayesian		false;
		double	maxNodePop		1.9;
		double	Ztolerance		1.e-3;
	const		savecount		100; folso:
	const		resume		false;
	const const		outputSamples		false;
	const		<pre>init_pop_strategy discard unfit points</pre>		0; falso:
	const		<pre>discard_unfit_points max_init_attempts</pre>		
			<pre>max_init_attempts max_accontable_val</pre>		10000;
		double	<pre>max_acceptable_val sood</pre>		1e6;
	const		seed		1234567;
	const	INC	verbose	=	1;

//European to be minimized. Corresponds to -ln(Likelihood)

// M. // Ma

// Pop // Size

// Scal // Cros

// Mixir // Use c // Use e

// Bound // Use s

// Use s

// Thres // Numb // Weec

// Cal // Por

/ Ir

 nPar needs to be adjusted according to the number of free parameters in the model

The next 2 arrays have the values of the intervals in which your free parameters can take values, you choose the order of your parameters

 In this example, the first parameter can be varied between
 -5 and 5, the second between -50 and 50, etc.

2 #include					
3 #include					
4 #include	"diver.h				
	_		V		
o const int		nPar	= 5;	11	
7 const dou	ble	lowerbounds[]	= {-5.,-50.,-5.,-50.,-2.};		
8 const dou	ble	upperbounds[]	= { 5., 50., 5., 50., 2.};	11	<del>Upper</del> bo
9 const cha	r	path[]	<pre>= "example_c/output/example";</pre>		
10 const int		nDerived	= 0;		
11 const int		nDiscrete	= 0;	X	
12 const int		discrete[]	= {};	11	
13 const boo	ι	partitionDiscrete	= false;	11	Split th
14 const int		maxciv	= 1;	11	Maximum
15 const int		maxgen	= 100;	11	Maximum
16 const int		NP	= 1000;	11	Populati
17 const int		nF	= 1;	11	Size of
18 const dou	ble	F[]	= {0.6};	11	Scale fa
19 const dou	ble	Cr	= 0.9;	11	Crossove
20 const dou	ble	lambda	= 0.8;	11	Mixing
21 const boo	ι	current	= false;	11	Use curr
22 const boo		expon	= false;		Use expo
23 const int		bndry	= 3;		Boundary
24 const boo		jDE	= true;		Use self
25 const boo		lambdajDE	= true;		Use self
26 const dou		convthresh	= 1.e-6;		Threshol
27 const int		convsteps	= 10;		Number o
28 const boo		removeDuplicates	= true;		Weec out
29 const boo		doBayesian	= false;		Calculat
30 const dou		maxNodePop	= 1.9;		Populati
31 const dou		Ztolerance	= 1.e-3;		<b>Ir</b> put to
32 const int		savecount	= 100;		Save pro
33 const boo		resume	= false;	11	
34 const boo		outputSamples	= false;	11	
35 const int		<pre>init_pop_strategy</pre>	= 0;		
36 const boo		<pre>discard_unfit_points</pre>			
37 const int		<pre>max_init_attempts</pre>	= 10000;		
38 const dou		<pre>max_acceptable_val</pre>	= 1e6;		
39 const int		seed	= 1234567;		
const int		verbose	= 1;		

- Next, the directory where the data will be saved is stated under quotes
- In this example the directory is (relative to the directory where the executable is) example\_c/output/

The files created with the data will have the same name but different extensions, in this example the name is 'example' #include <float.h>
#include <stdool.h>
#include <stddef.h>
#include "diver.h"

	const		nPar	= 5;		
		double	lowerbounds[]	= {-5.,-50.,-5.,-50.,-2.};		
		double	upperbounds[]	$= \{ 5., 50., 5., 50., 2. \};$	11	
	const	char 🔨	path[]	<pre>= "example_c/output/example";</pre>	$\leftarrow$	Path to
	const		nDerived	= 0;		
	const	int	nDiscrete	= 0;	$\sim$	
	const	int	discrete[]	= {};	11	
	const		partitionDiscrete	= false;	- //	Split t
	const		maxciv	= 1;	11	Maximum
	const		maxgen	= 100;		Maximum
16	const	int	NP	= 1000;	11	Populati
	const	int	nF	= 1;	- 77	Size
		double	F[]	= {0.6};		Scal
	const	double	Cr	= 0.9;	11	Crossov
	const	double	lambda	= 0.8;		Mixing
	const	bool	current	= false;	- //	Use c
	const	bool	expon	= false;	- 11	Use exp
	const	int	bndry	= 3;	11	Bound
	const	bool	jDE	= true;		Use s
	const	bool	lambdajDE	= true;	11	Use sel
26	const	double	convthresh	= 1.e-6;	- 11	Thresho
	const	int	convsteps	= 10;	- //	Number
	const	bool	removeDuplicates	= true;		Weec ou
29	const	bool	doBayesian	= false;	- 11	Calcula
	const	double	maxNodePop	= 1.9;	- 11	Populat
		double	Ztolerance	= 1.e-3;	11	Irput t
	const	int	savecount	= 100;	- 11	Save pr
	const	bool	resume	= false;	- //	
	const	bool	outputSamples	= false;	- 11	
	const	int	init_pop_strategy	= 0;		
	const		<pre>discard_unfit_points</pre>	= false;		
	const		<pre>max_init_attempts</pre>	= 10000;		
		double	<pre>max_acceptable_val</pre>	= 1e6;		
39	const		seed	= 1234567;		
	const	int	verbose	= 1;		

//Function to be minimized. Corresponds to -ln(Like)/hood

 Next, the variable nDerived refers to what we call observables, it is the number of observables that we want to save for each point

In our case, these are the Higgs mass, the relic density, the DM mass, the DM-proton x-section, 3 chi-square functions and their sum

Therefore, we should put nDerived = 8; #include <float.h>
#include <stdoool.h>
#include <stddef.h>
#include "diver.h"

	const		nPar	= 5;		
		double	lowerbounds[]	= {-5.,-50.,-5.,-50.,-2.};		
		double	upperbounds[]	$= \{ 5., 50., 5., 50., 2. \};$		
	const		path[]	<pre>= "example_c/output/example";</pre>		
	const		nDerived	= 0;		
	const		n <del>Discrete</del>	= 0;	$\mathcal{N}$	
	const	int	discrete[]	= {};	11	
	const		partitionDiscrete	= false;	11	Split th
	const		maxciv	= 1;		Maximum
	const		maxgen	= 100;	11	Maximum
	const		NP	= 1000;	11	Populati
	const		nF	= 1;		Size of
		double	F[]	= {0.6};		Scale in
		double	Cr	= 0.9;		Crossova
		double	lambda	= 0.8;		Mixing
	const		current	= false;		Use cum
	const		expon	= false;		Use exp
	const		bndry	= 3;		Boundary
	const		jDE	= true;		Use sela
	const		lambdajDE	= true;		Use sela
		double	convthresh	= 1.e-6;	11	Threshol
	const		convsteps	= 10;	!!.	Number
	const		removeDuplicates	= true;	<i>!</i> ,	Weed out
	const		doBayesian	= false;	<i>'</i> ,	Calculat
		double double	maxNodePop Ztolerance	= 1.9; = 1.e-3;	11	Populati
	const		savecount	= 100;	11	Input to Save pro
	const		resume	= false;	11	Pactart
	const		outputSamples	= false;	11	
	const		init pop strategy	= 0;		
	const		discard_unfit_points			
	const		max init attempts	= 10000;		
		double	<pre>max_acceptable_val</pre>	= 166;		
	const		seed	= 1234567;		
	const		verbose	= 1;		
	- one c	2				

Next, the variable maxgen should be put equal to a large number to ensure that a sufficiently large number of 'generations' are generated

If maxgen is small the calculation might be stopped before the convergence criteria is met

We recommend something like maxgen = 10000;

■ NP is the population; the recommended value is 20,000

"include "diver.h" = 5; const int nPar const double lowerbounds[] = {-5.,-50.,-5.,-50.,-2.}; 8 const double upperbounds[]  $= \{ 5., 50., 5., 50., 2. \};$ = "example c/output/example"; 9 const char path[] 10 const int nDerived = 0; 11 const int nDiscrete = 0; 12 const int discrete[]  $= \{\}$ : const bool = false; partitionDiscrete 14 const int maxciv = 1: // M. 15 const int = 100: // Ma maxgen NP 16 const int = 1000;// Por 17 const int nF = 1: // Size 18 const double F[]  $= \{0.6\};$ // Scal 19 const double Cr = 0.9;// Cros. 20 const double lambda = 0.8;// Mixiı 21 const bool current = false: 22 const bool = false; expon bndry 23 const int = 3: // Bound 24 const bool **jDE** = true; // Use s 25 const bool lambdaiDE = true; 26 const double convthresh = 1.e-6;// Thres 27 const int convsteps = 10;// Numb removeDuplicates 28 const bool = true; // Weec 29 const bool doBayesian = false: // Cal 30 const double maxNodePop = 1.9: // Por 31 const double Ztolerance = 1.e-3;// Ir 32 const int savecount = 100: 33 const bool = false; resume outputSamples = false; 34 const bool init pop strategy 35 const int = 0; discard unfit points = false; 36 const bool max init attempts 37 const int = 10000;max acceptable val 38 const double = 1e6:39 const int = 1234567;seed const int verbose = 1;

#include <stdbool.h>
#include <stddef.h>

 Next, the variable 'convthresh' is used for the convergence criteria, it is recommended to be set at the value 1e-4

Smaller values might take too much computing time, while greater values might lead to incomplete analysis of the parameter space #include <float.h>
#include <stdool.h>
#include <stddef.h>
#include "diver.h"

	const	int	nPar	=	5;		
	const	double	lowerbounds[]	=	{-5.,-50.,-5.,-50.,-2.};		
	const	double	upperbounds[]	=	{ 5., 50., 5., 50., 2.};		
	const	char	path[]	=	<pre>"example_c/output/example";</pre>		
	const	int	nDerived	=	0;		
	const	int	nDiscrete	=	0;	$\mathcal{N}$	
	const	int	discrete[]	=	{};	11	
	const	bool	partitionDiscrete	=	false;	11	Split
	const	int	maxciv	=	1;	11	Maxim
	const	int	maxgen	=	100;	11	Maxim
	const	int	NP	=	1000;	11	Popul
	const	int	nF		1;	//	Size
	const	double	F[]	=	{0.6};	11	Scal
	const	double	Cr	=	0.9;	11	Cros.
	const	double	lambda		0.8;	11	Mixir
	const	bool	current		false;	11	Use c
	const	bool	expon	=	false;	11	Use e
	const	int	bndry	=	3;	11	Bound
	const	bool	jDE	=	true;	//	Use s
	const	bool	lam <del>bdajDE</del>	=	true;	11	Use s
26	const	double 🧹	convthresh	=	1.e-6;	//	Three
	const	int	convsteps	=	10;	11	Numb
28	const	bool	removeDuplicates		true;	11	Weec
29	const	bool	doBayesian	=	false;	11	
30	const	double	maxNodePop	=	1.9;	//	Popula
31	const	double	Ztolerance	=	1.e-3;	11	Input
32	const	int	savecount		100;	11	<i>S</i> ave
	const		resume	=	false;	$\Pi$	
}4	const	bool	outputSamples	=	false;	$\Pi$	
	const		init_pop_strategy		Θ;		
	const		<pre>discard_unfit_points</pre>				
	const		<pre>max_init_attempts</pre>		10000;		
		double	<pre>max_acceptable_val</pre>		1e6;		
	const		seed		1234567;		
	const	int	verbose	=	1;		

//Function to be minimized. Corresponds to -ln(Like)

□ The variable savecount should always be equal to 1

This ensures that the observables are saved for all the points explored during the scan

This is important to generate the contours of equal likelihood

#include <float.h>
#include <stdbool.h>
#include <stddef.h>
#include "diver.h"

	const	int	nPar	= 5;		
	const	double	lowerbounds[]	= {-5.,-50.,-5.,-50.,-2.};		
	const	double	upperbounds[]	$= \{ 5., 50., 5., 50., 2. \};$		
	const	char	path[]	<pre>= "example_c/output/example";</pre>		
	const	int	nDerived	= 0;		
	const		nDiscrete	= 0;	$\mathbf{N}$	
	const		discrete[]	= {};	11	
	const		partitionDiscrete	= false;		Split
	const		maxciv	= 1;		Maximu
	const		maxgen	= 100;		Maxamu
	const		NP	= 1000;		Popula
	const		nF	= 1;		Size
		double	F[]	= {0.6};		Scal
		double	Cr	= 0.9;		Cross
		double	lambda	= 0.8;		Mixin
	const const		current	= false;		Use c Use e
	const		expon bndry	= false; = 3;		Bound
	const		jDE	= 5; = true;		Use s
	const		lambdajDE	= true;		Use s
		double	convthresh	= 1.e-6;	11	Three
	const		convsteps	= 10;	11	Numb
	const		removeDuplicates	= true;		Weec
	const		doBayesian	= false;		Cal
		double	maxNodePop	= 1.9;		Popula
		double	7tolerance	= 1.e-3;		Input
	const	int 🤇	savecount	= 100;		Save r
	const	bool	resume	= false;	11	
	const	bool	outputSamples	= false;	11	
	const	int	<pre>init_pop_strategy</pre>	= 0;		
36	const	bool	discard_unfit_points	= false;		
	const		<pre>max_init_attempts</pre>	= 10000;		
		double	<pre>max_acceptable_val</pre>	= 1e6;		
39	const		seed	= 1234567;		
	const	int	verbose	= 1;		

Finally, the variable 'outputSamples' should be equal to 'true'

This ensures that the
 observables are saved to disk
 alongside the parameters for each
 point of parameter space explored

#include <float.h>
#include <stdool.h>
#include <stddef.h>
#include "diver.h"

5	const	int	nPar	= 5;			
	const	double	lowerbounds[]	= {-5.,-50.,-5.,	-50.,-2.};		
	const	double	upperbounds[]	= { 5., 50., 5.,	50., 2.};		
	const	char	path[]	<pre>= "example_c/out</pre>	put/example";		
	const	int	nDerived	= 0;			
	const	int	nDiscrete	= 0;		$\mathcal{N}$	
	const	int	discrete[]	= {};		11	
	const	bool	partitionDiscrete	= false;		11	Spl
	const	int	maxciv	= 1;		11	Mex
	const	int	maxgen	= 100;		11	Ma.
	const	int	NP	= 1000;		11	Рор
	const	int	nF	= 1;		11	Siz
	const	double	F[]	= {0.6};		11	Sca
	const	double	Cr	= 0.9;		11	Cro
	const	double	lambda	= 0.8;		11	Mix
	const	bool	current	= false;		11	Use
	const	bool	expon	= false;		11	Use
	const		bndry	= 3;		11	Bou
	const		jDE	= true;			Use
	const	bool	lambdajDE	= true;		11	Use
		double	convthresh	= 1.e-6;		11	Thr
	const		convsteps	= 10;		11	Num
	const	bool	removeDuplicates	= true;		11	Wee
	const		doBayesian	= false;		11	Cal
	const	double	maxNodePop	= 1.9;		11	Por
		double	Ztolerance	= 1.e-3;			Ir
	const		savecount	= 100;		11	Şav
	const		resume	= false;		11	
	const		outputSamples	= false; ——		11	
	const		<pre>init_pop_strategy</pre>	= 0;			
	const		<pre>discard_unfit_points</pre>				
	const		<pre>max_init_attempts</pre>	= 10000;			
		double	<pre>max_acceptable_val</pre>	= 1e6;			
	const		seed	= 1234567;			
	const	int	verbose	= 1;			

### Inserting the chi2BB code

Let us return to the third block

■ We will insert the code for the generation of the composite chisquare for a given point of parameter space

```
40 const int _____verbose
43 //Function to be minimized. Corresponds to -ln(
44 //Plain Gaussian centred at the origin. Valid fo
45 double gauss(double params[], const int param di
46 {
    double result = 0.0;
   //for (int i = 0; i<param dim; i++) result +=
    if (!validvector)
51 result = DBL MAX;
    *fcall += 1;
    *quit = false;
    // Insert here code for chi2BB
    11
    11 .....
     11
    11
    *fcall += 1:
     *quit = false;
     return result;
65 }
  S int main(int argc, char** argv)
                                                     35
```

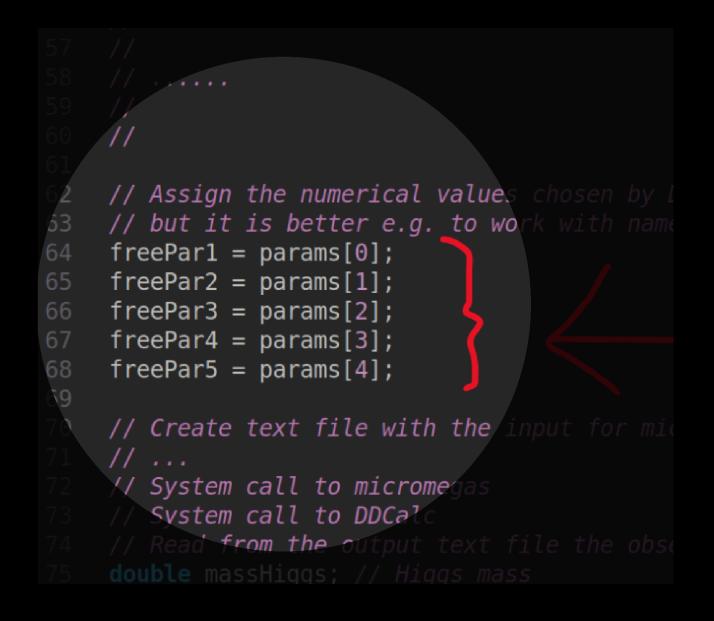
## Inserting the chi2BB code

This is the resulting code, except that we have not put explicitly the code for the generation of the text files nor the system calls

```
//Plain Gaussian centred at the origin. Valid for any number of dimensions. Minimum value is the number of dimension
double gauss(double params[], const int param dim, int *fcall, bool *quit, const bool validvector, void** context)
  double result = 0.0;
  //for (int i = 0; i<param dim; i++) result += params[i]*params[i] + 1.0;</pre>
   if (!validvector)
  result = DBL MAX;
  *fcall += 1:
  *quit = false;
  // Assign the numerical values chosen by Diver to variables of your choosing (this is optional,
  // but it is better e.g. to work with names like lambda1 for a coupling than params[0])
  freePar1 = params[0];
  freePar2 = params[1];
  freePar3 = params[2];
  freePar4 = params[3];
  freePar5 = params[4];
  // Create text file with the input for micromegas
  // System call to DDCalc
  double massHiggs; // Higgs mass
  double Omega2;
  double DMmass;
                    // DM mass
  double sigmaP;
                   // DM-proton x-sect
  double chi2Higgs; // chi-square higgs
  double chi2Relic; // chi-square relic
  double chi2XENON; // chi-square XENON1T
  fscanf(....,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON);
  double chiComposite = chi2Higgs + chi2Relic + chi2XENON;
  // save observables for this point in parameter space
  params[5] = massHiggs;
  params[11] = chi2XENON;
  params[12] = chiComposite;
  result
             = chiComposite;
  *fcall += 1;
  *quit = false;
  return result;
```

Diver <u>chooses values</u> for the free parameters and puts these values in the array 'params'; these values constitute 1 point in parameter space

In this example we have 5 free parameters, and we are assigning these values to new variables named freePar1, etc



Our task is to tell Diver what value of the composite chi-square this point in parameter space has //Plain Gaussian centred at the origin. Valid for any number of dimensions. Minimum value is the number of dimension double gauss(double params[], const int param dim, int \*fcall, bool \*quit, const bool validvector, void\*\* context) double result = 0.0; //for (int i = 0; i<param dim; i++) result += params[i]\*params[i] + 1.0;</pre> (!validvector) result = DBL MAX; \*fcall += 1: \*quit = false; // Assign the numerical values chosen by Diver to variables of your choosing (this is optional, // but it is better e.g. to work with names like lambda1 for a coupling than params[0]) freePar1 = params[0]; freePar2 = params[1]; freePar3 = params[2]; freePar4 = params[3]; freePar5 = params[4]; // Create text file with the input for micromegas // System call to DDCalc double massHiggs; // Higgs mass double Omega2; **double** DMmass: // DM mass double sigmaP; // DM-proton x-sect double chi2Higgs; // chi-square higgs double chi2Relic; // chi-square relic double chi2XENON; // chi-square XENON1T fscanf(....,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON); **double** chiComposite = chi2Higgs + chi2Relic + chi2XENON; // save observables for this point in parameter space params[5] = massHiggs; params[11] = chi2XENON; params[12] = chiComposite; result = chiComposite; \*fcall += 1; \*quit = false; return result;

38

□ To do that we create a text file with the values of the free parameters

Next, we make a system call to micromegas passing this text file

Next, we make a system call to
 DDCalc which reads the DM-proton
 x-section calculated by micromegas
 and computes the XENON1T chi square

❑ Then we read from the output text file the values of the observables and the chi-squares and assign them to respective variables

```
freePar2 = params[1];
freePar3 = params[2];
freePar4 = params[3];
freePar5 = params[4];
   Create text file with the input for micromegas
 / System call to micromegas
// System call to DDCalc
// Read from the output text file the observables and chi-sou
double massHiggs; // Higgs mass
double Omega2; // relic
double DMmass; // DM mass
double sigmaP; // DM-proton x-sect
double chi2Higgs; // chi-square higgs
double chi2Relic; // chi-square relic
double chi2XENON; // chi-square XENON1T
fscanf(....,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2R/
```

// compopsite chi-square is the sum
oouble chiComposite = chi2Higgs + chi2Relic + chi2XENON;

// save observables for this point in parameter space
params[5] = massHiggs;

params[11] = chi2XENON;
params[12] = chiComposite;

The composite chi-square is just the sum of the 3 chi-squares computed

44 /,	/Function to be minimized. Corresponds to -ln(Likelihood). /Plain Gaussian centred at the origin. Valid for any number of dimensions. Minimum value is the number of dimens ouble gauss(double params[], const int param_dim, int *fcall, bool *quit, const bool validvector, void** context)
	<pre>double result = 0.0; //for (int i = 0; i<param_dim; +="" 1.0;<br="" i++)="" result="">if (!validvector) {</param_dim;></pre>
	result = DBL_MAX; *fcall += 1;
	*quit = false; }
	// Insert here code for chi2BB //
	// // Assign the numerical values chosen by Diver to variables of your choosing (this is optional,
	<pre>// but it is better e.g. to work with names like lambda1 for a coupling than params[0]) freePar1 = params[0]; freePar2 = params[1];</pre>
	freePar3 = params[2]; freePar4 = params[3];
	<pre>freePar5 = params[4]; // Create text file with the input for micromegas</pre>
	// // System call to micromegas // System call to DDCalc
	// Read from the output text file the observables and chi-squares, e.g. double massHiggs; // Higgs mass
	double Omega2;       // relic         double DMmass;       // DM mass         double sigmaP;       // DM-proton x-sect
	<pre>double chi2Higgs; // chi-square higgs double chi2Relic; // chi-square relic double chi2XENON; // chi-square XENON1T</pre>
82 <b>83</b>	<pre>fscanf(,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON);</pre>
	<pre>// compopsite chi-square is the sum double chiComposite = chi2Higgs + chi2Relic + chi2XENON;</pre>
	<pre>// save observables for this point in parameter space params[5] = massHiggs;</pre>
	 params[11] = chi2XENON;
	<pre>params[12] = chiComposite; // return the value of the composite chi-square for this point in parameter space</pre>
	result = chiComposite;
	<pre>*fcall += 1; *quit = false; return result;</pre>
L00 }	

□ The composite chi-square is just the sum of the 3 chi-squares computed // ...
// System call to micromegas
// System call to DDCalc
// Read from the output text file the observables and chi-squares, e.g.
double wassHiggs; // Higgs mass
double Omega2; // relic
double OMmass; // DM mass
double sigmaP; // DM-proton x-sect
double chi2Higgs; // chi-square higgs
couble chi2Relic; // chi-square relic
double chi2XENON; // chi-square xENON1T
fscanf(....,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON
// compopsite chi-square is the sum
double chiComposite = chi2Higgs + chi2Relic + chi2XENON;

```
// save observables for this point in parameter space
params[5] = massHiggs;
...
...
```

```
oarams[11] = chi2XENON;
oarams[12] = chiComposite;
```

// return the value of the composite chi-square for this point in parame
result = chiComposite;

\*fcall += 1;
\*quit = false;
return result;

Next, we must save the values of the observables and the chisquares

44 //	/Function to be minimized.  Corresponds to -ln(Likelihood). /Plain Gaussian centred at the origin. Valid for any number of dimensions.  Minimum value is the number of din puble gauss(double params[], const int param dim, int *fcall, bool *quit, const bool validvector, void** conte
46 {	
	double result = 0.0;
	//for (int i = 0; i <pre>param dim; i++) result += params[i]*params[i] + 1.0;</pre>
	if (!validvector)
	{
	result = DBL MAX;
	*fcall += 1;
	*quit = false;
	}
	// Insert here code for chi2BB
	·// ·····
	//
	// Assign the numerical values chosen by Diver to variables of your choosing (this is optional,
	// but it is better e.g. to work with names like lambda1 for a coupling than params[0])
	<pre>freePar1 = params[0];</pre>
	<pre>freePar2 = params[1];</pre>
	freePar3 = params[2];
	<pre>freePar4 = params[3];</pre>
	freePar5 = params[4];
	// Create text file with the input for micromegas
	// System call to micromegas
	// System call to DDCalc
	// Read from the output text file the observables and chi-squares, e.g.
	double massHiggs; // Higgs mass
	double Omega2; // relic
	double DMmass; // DM mass
	double sigmaP; // DM-proton x-sect
	double chi2Higgs; // chi-square higgs
	double chi2Relic; // chi-square relic
	double chi2XENON; // chi-square XENON1T
	fscanf(,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON);
83	// components shi caupacia the cum
	// compopsite chi-square is the sum
	<pre>double chiComposite = chi2Higgs + chi2Relic + chi2XENON;</pre>
	// save observables for this point in parameter space
87 88	params[5] = massHiggs;
91	params[11] = chi2XENON;
	<pre>params[11] = chi2xLNON, params[12] = chiComposite;</pre>
	// return the value of the composite chi-square for this point in parameter space
	result = chiComposite;
	*fcall += 1;
	*quit = false;
	return result;
L00 }	

xt)

■ These have to be saved in the same array 'params', but next to the free parameters, i. e. starting from params[5] onwards in this example

```
compopsite chi-square is the su
double chiComposite = chi2Higgs +
  save observables for this p
params[5] = massHiggs;
. . .
. . .
params[11] = chi2XENON;
params[12] = chiComposite;
 v return the value of the composit
result
           = chiComposite;
*fcall += 1:
```

Finally, we return to Diver the value of the composite chi-square

//Plain Gaussian centred at the origin. Valid for any number of dimensions. Minimum value is the number of dimensior double gauss(double params[], const int param dim, int \*fcall, bool \*quit, const bool validvector, void\*\* context) double result = 0.0; //for (int i = 0; i<param dim; i++) result += params[i]\*params[i] + 1.0;</pre> if (!validvector) result = DBL MAX; \*fcall += 1: \*quit = false; // Assign the numerical values chosen by Diver to variables of your choosing (this is optional, // but it is better e.g. to work with names like lambda1 for a coupling than params[0]) freePar1 = params[0]; freePar2 = params[1]; freePar3 = params[2]; freePar4 = params[3]; freePar5 = params[4]; // Create text file with the input for micromegas // System call to DDCalc double massHiggs; // Higgs mass double Omega2; double DMmass; // DM-proton x-sect double sigmaP; double chi2Higgs; // chi-square higgs double chi2Relic; // chi-square relic double chi2XENON; // chi-square XENON1T fscanf(....,massHiggs,Omega2,DMmass,sigmaP,chi2Higgs,chi2Relic,chi2XENON); **double** chiComposite = chi2Higgs + chi2Relic + chi2XENON; // save observables for this point in parameter space params[5] = massHiggs; params[11] = chi2XENON; params[12] = chiComposite; result = chiComposite; \*fcall += 1; \*quit = false; return result;

□ Finally, we return to Diver the value of the composite chi-square

- This completes the code for the chi2BB
- Next, we compile the program with the 'make' instruction as before and execute it ...

```
ams[11] = chi2XENON;
 rams[12] = chiComposite;
// return the value of the com
result = chiComposite;
*fcall += 1;
*quit = false;
return result;
```

In this example we execute Diver with 4 cores,

'name\_of\_executable' is the program that results from compiling the code just shown

■ We redirect the output of the program to the text file 'log.txt' and we run in the 'background' mpiexec -np 4 name\_of\_executable >> log.txt &

git clone https://github.com/patscott/pippi.git
sudo apt-get -y install ctioga2

~/pippi/pippi q4.pip

❑ After a while (which can be of the order of ~1-2 months!) Diver generates a file with the results

□ To parse this file, we need the tool called 'pippi'

Pippi uses ctioga2 to plot so we also install this

❑ We call pippi by means of a configuration file, in this example the file called 'q4.pip', basically this files just defines which of observables we wish to plot

mpiexec -np 4 name\_of\_executable >> log.txt &

git clone https://github.com/patscott/pippi.git
 sudo apt-get -y install ctioga2

~/pippi/pippi q4.pip

This is an example of a pippi configuration file to plot the DMproton x-section as a function of the DM mass

But I will not enter into details of this here ....

key on 1D =

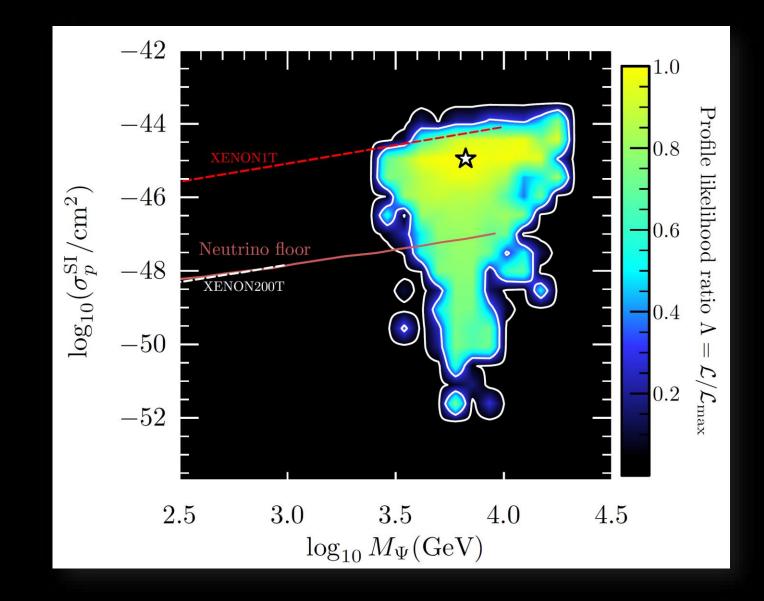
kev on 2D =

#### ---Common fields-----; EFN = empty for none;Filename of chain to operate on. For a group in an hdf5 file, something like 'myfile.hdf5:/group' main chain = './res/g4.sam' comparison chain = ;Filename of secondary chain to operate on, for comparison (EFN) do posterior pdf = F;Parse, script or plot posterior pdfs do profile like = T;Parse, script or plot chain into profile likelihoods oneD contour levels = ;Contour levels to plot in 1D plots (EFN; script operation ignores actual values) twoD contour levels = 68.395.4;Contour levels to plot in 2D plots (EFN; script operation ignores actual values) oneD\_plot\_quantities = ;Parameters/observables to construct 1D plots of (EFN) twoD plot quantities = {11 12}; ;Parameters/observable combinations to make 2D plots of (EFN) lot observables = ;Observables to make 2D plots of (EFN) ---Parsing----parse dir = 'parse diver' ;Directory to save parsing output in (EFN; default is to place parsed data in same folder as chains) cut on invalid observables = F ;Cut on any invalid entries in an hdf5 file, not just the likelihood (defaults to false, meaning cut on likelihood only) ;Default number of bins to sort samples into in each direction. default bins = 40 ; number of bins = ;Bins to use for specific observables (overrides default) interpolated resolution = 4000 ; ;Resolution of binwise interpolation for plotting (~300+ for publication) interpolation method = 'bilinear' ;Either bilinear (default) or spline (watch out for ringing in the latter case) chain type = other ;Algorithm used to generate chain (valid: MCMC, MultiNest, other) compute evidence = F;Compute and print evidence (only if chain type = MCMC) bf lnlike for profile like = ;Denominator of profile likelihood ratio (if empty, set to best fit determined from chain) use\_log\_scale = 11 12 ;Parameters/observables that need to be converted to a log scale for display (EFN) quantity\_rescalings = ;Scaling factors to apply to any pippi datastream indices (EFN) data ranges = ;Discard any points outside these ranges of parameters/observables 25:{-40.,1.} labels from file = ;Either a filename or blank (quantity labels is ignored if a filename is given) preamble = 'from preamble example import \*' ;A one-line python preamble to run before processing functional datastreams. assign to pippi datastream = ;Assign a python function or named hdf5 data stream to a particular pippi datastream index (use pippi probe to test; EFN) quantity labels = 1:'-lnlike' 4:'freePar1 5:'freePar2 6: freePar3 7:'freePar4 8: 'freePar5 9:'\$m h\$' 10:'\$\Omega h^2\$' 11:'\$\log {10} M \Psi \textrm{(GeV)}\$' 12:'\$\log {10} (\sigma p^{\rm SI} / \textrm{cm}^2)\$' 13:'\$\chi^2 h\$' 14:'\$\chi^2 {\Omega h^2}\$' \ 15:'\$\chi^2 {DD}\$' ;labels to apply to different pippi datastream indices (LaTeX for all but multiplicity, prior and likelihood) ---Scripting----script dir = 'scripts diver' ;Directory to save scripts in (EFN; default is to place scripts in same folder as parse output) ;1D plots that should include legends legend on 1D =legend locations 1D =;Locations of legends for 1D plots (subset of twoD plot quantities) ;Plot 1D distributions as histograms with no interpolation plot as histograms 1D = F

;2D plots that should include keys for best fit, posterior mean and/or reference pt key locations 1D = ;Locations of keys for 2D plots ;2D plots that should include legends legend on 2D = ;legend locations 2D = ;;Locations of legends for 2D plots (single location or partial list that will be filled in with 'bl') :2D plots that should include kevs for best fit, posterior mean and/or reference pt

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Suffice to say that it generates this plot of the DM-proton xsection as a function of the DM mass



#### Subtleties

There are certain subtleties with Diver and Pippi which are hard to explain in a talk

We will address these topics with a lot more of detail in the corresponding tutorials

 We shall also cover Micromegas,
 DDCalc and others in respective tutorials



Thank you for your attention!

