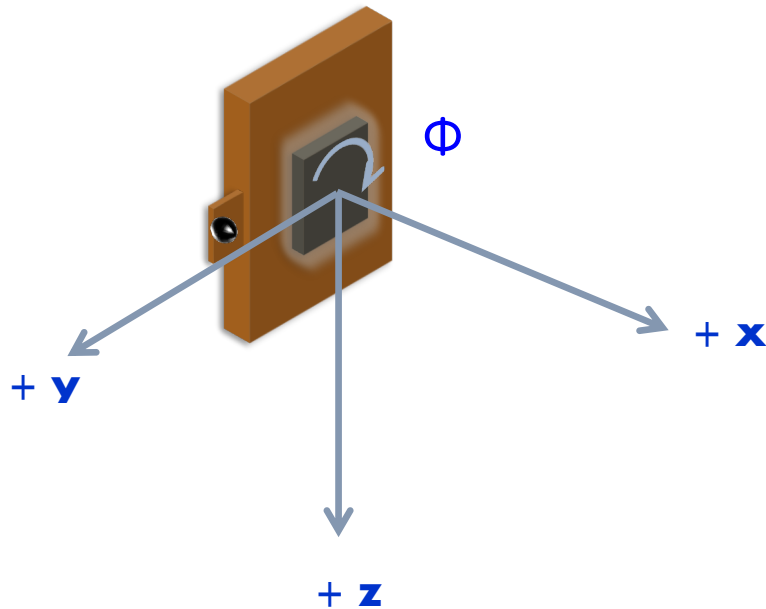


# I21 Diffcalc Guide

---



## Crystal orientation

Before moving in hkl space you must calculate a UB matrix by specifying the crystal's lattice parameters (which define the B matrix) and finding two reflections (from which the U matrix can be inferred); and, optionally for surface-diffraction experiments, determine how the surface of the crystal is oriented with respect to the phi axis.

### Start a new UB calculation

A *UB calculation* contains the description of the crystal-under-test, any saved reflections, reference angle direction, and a B & UB matrix pair if they have been calculated or manually specified. Starting a new UB calculation will clear all of these.

Before starting a UB-calculation, the `ub` command used to summarise the state of the current UB-calculation, will reflect that no UB-calculation has been started:

```
>>> ub
<<< No UB calculation started >>>
```

A new UB-calculation calculation may be started and lattice specified explicitly:

```
>>> newub 'example'
>>> setlat '1Acube' 1 1 1 90 90 90
```

or interactively:

```
>>> newub
calculation name: example
crystal name: 1Acube
crystal system
1) Triclinic
2) Monoclinic
3) Orthorhombic
4) Tetragonal
5) Rhombohedral
6) Hexagonal
7) Cubic
[1]: 7
   a[1]: 1
```

where a is unit cell basis vector in Angstroms for cubic crystal system.

The `ub` command will show the state of the current UB-calculation (and the current energy for reference):

```
>>> ub
UBCALC

  name:      example

REFERENCE

  n_hkl:      1.00000   0.00000   0.00000 <- set

SURFACE NORMAL

  n_phi:      0.00000   0.00000   1.00000 <- set

CRYSTAL

  name:      1Acube

  a, b, c:    1.00000   1.00000   1.00000
              90.00000   90.00000   90.00000   Cubic

  B matrix:   6.28319   0.00000   0.00000
              0.00000   6.28319   0.00000
              0.00000   0.00000   6.28319

UB MATRIX

  <<< none calculated >>>

REFLECTIONS

  <<< none specified >>>

CRYSTAL ORIENTATIONS

  <<< none specified >>>
```

## Load a UB calculation

To load the last used UB-calculation:

```
>>> lastub
Loading ub calculation: 'mono-Si'
```

To load a previous UB-calculation:

```
>>> listub
UB calculations in: /home/i21user/.diffcalc/i21

0) mono-Si          15 Feb 2017 (22:32)
1) i21-32          13 Feb 2017 (18:32)

>>> loadub 0
```

## Generate a U matrix from two lattice directions

Another approach to calculate a U matrix is to provide orientation of **two** crystal lattice directions using `addorient` command after aligning sample in laboratory frame of reference. The first lattice direction should be aligned along the selected direction in the laboratory frame. Best would be to provide the lattice direction normal to the surface of the sample. For the purpose of finding azimuthal orientation in U matrix calculation it is sufficient for the projection of the second lattice direction to be aligned to the given orientation in the laboratory frame in the plane perpendicular to the first lattice orientation.

Find U matrix from two lattice directions:

```
>>> addorient [0 0 1] [1 0 0]
>>> addorient [1 0 0] [0 1 0]
Calculating UB matrix.
```

## Calculate a UB matrix

Unless a U or UB matrix has been manually specified, a new UB matrix will be calculated after the second reflection has been found, or whenever one of the first two reflections is changed.

Use command `orientub` to force the UB matrix to be calculated from the first two orientations.

UB matrix can be calculated from any combination of two reflections and/or orientations by providing corresponding reflection/orientation tags or numbers as an argument to `calcube`. In

case of using one reflection and one orientation it is recommended to use tags to avoid ambiguity.

If you have misidentified a reflection used for the orientation the resulting UB matrix will be incorrect. Always use the `checkub` command to check that the computed reflection indices agree with the estimated values:

```
>>> checkub
```

	ENERGY	H	K	L	H_COMP	K_COMP	L_COMP	TAG
1	12.3984	0.00	1.00	1.00	0.0000	1.0000	1.0000	
2	12.3984	0.00	0.00	1.00	0.0000	0.0000	1.0000	

## Calculate a U matrix from crystal mismatch

U matrix can be defined from crystal mismatch by using a rotation matrix calculated from a provided mismatch angle and axis. `setmismatch` command defines new U matrix by setting it to a rotation matrix calculated from the specified angle and axis parameters. `addmismatch` command applies the calculated rotation matrix to the existing U matrix, i.e. adds extra mismatch to the already existing one:

```
>>> setmismatch 5 [1 0 0]
n_phi: -0.00000 -0.08716 0.99619
n_hkl: 0.00000 0.00000 1.00000 <- set
normal:
  angle: 5.00000
  axis: 1.00000 -0.00000 0.00000
```

## Manually specify U matrix

Set U matrix manually (pretending sample is squarely mounted):

```
>>> setu [[1 0 0] [0 1 0] [0 0 1]]
Recalculating UB matrix.
NOTE: A new UB matrix will not be automatically calculated when the
orientation reflections are modified.
```

## Set the reference vector

The reference vector can be used to define azimuthal direction within the crystal with which we want to orient the incident or diffracted beam. Orientation of the reference vector w.r.t the incident and diffracted beam is indicated using `alpha` and `beta` angles.

By default in `i21` the reference vector is set parallel to the theta axis. That is, along the y-axis of the laboratory coordinate frame.

The `ub` command shows the current reference vector at the top its report (or it can be shown by calling `setnphi` or `setnhkl` with no args):

```
>>> ub
...
REFERENCE
```

```

n_phi:      0.00000  1.00000  0.00000
n_hkl:      1.00000  0.00000  0.00000 <- set
...

```

The <- set label here indicates that the reference vector is set in the reciprocal lattice space. In this case, therefore, its direction in the laboratory coordinate frame is inferred from the UB matrix.

To set the reference vector in the phi coordinate frame use:

```

>>> setnphi [0 1 0]
...

```

To set the reference vector in the crystal's reciprocal lattice space use:

```

>>> setnhkl [0 1 0]
...

```

## Constraining solutions for moving in hkl space

To get help and see current constraints:

```

>>> help con
...
>>> con
      REF              SAMP
-----
a_eq_b      th
alpha       chi
beta        phi
psi         mu_is_gam
bin_eq_bout bisect
betain      omega
betaout

!  1 more constraint required

Type 'help con' for instructions

```

## REFERENCE COLUMN:

- **alpha** - incident angle to reference vector
- **beta** - exit angle from reference vector
- **psi** - azimuthal rotation about scattering vector of reference vector (from scattering plane)
- **a\_eq\_b** - bisecting mode with alpha=beta. *Equivalent to psi=90*
- **betain** - incident angle to sample surface
- **betaout** - exit angle from sample surface

## SAMPLE COLUMN:

- **mu, eta, chi & phi** - physical settings
- **mu\_is\_gam** - force mu to follow gamma (results in a 5-circle geometry)
- **bisect** - bisecting mode with scattering vector in chi-circle plane
- **omega** - bisecting mode with omega angle between scattering vector and chi-circle plane

Diffcalc will report two other (un-constrainable) virtual angles:

- **theta** - half of  $2\theta$ , the angle through the diffracted beam bends
- **tau** - longitude of reference vector from scattering vector (in scattering plane)

## Configuring limits and cuts

Diffcalc uses motor limits set in GDA when used from GDA client running on a beamline. The standalone console version maintains its own limits on axes. These limits will be used when choosing solutions. If more than one detector solution exists Diffcalc will ask you to reduce the the limits until there is only one. However if more than one solution for the sample settings is available it will choose one that is closest to the current diffractometer orientation.

Use the `hardware` command to see the current limits and cuts:

```
>>> hardware
      mu           (cut: -180.0)
      delta        (cut: -180.0)
      gam          (cut: -180.0)
      eta          (cut: -180.0)
      chi          (cut: -180.0)
      phi          (cut:   0.0)
```

Note: When auto sector/transforms are used,  
cuts are applied before checking limits.

To set the limits in standalone Diffcalc session:

```
>>> setmin delta -1
>>> setmax delta 145
```

To set a cut:

```
>>> setcut phi -180
```

This causes requests to move phi to be between the configured -180 and +360 degrees above this. i.e. it might dive to -10 degrees rather than 350.

## Configuring reciprocal space as function of selected detector

Due to the unique requirements of I21 RIXS spectrometer we have the possibility of using different detectors for diffcalc. For the aligning of the sample crystal axes using diffraction we can use the rotating photodiode that is located inside the sample vessel and whose motor is called `diffftth`. Once the alignment is completed and we move to the spectrometer to perform RIXS measurements, then we have the possibility to refer our geometry either to the collecting mirror `m5tth` or to any of the two collecting mirrors that we call `lowq` and `highq`.

To be able to select between this available geometries, we have a set of scannables `hkl_m5tth`, `hkl_lowq`, `hkl_highq` and `hkl_diffftth` that relate to different combinations of selected angles and geometries. They can be used as `hkl` scannable to do calculations and move diffractometer. The `hkl` can be also reassigned to one of these scannables for convenience using `usem5tth`, `uselowq`, `usehighq` and `usediffftth` commands.

## Moving in hkl space

Configure a mode, e.g. four-circle vertical:

```
>>> con psi 0
      psi : 0.0000
```

Simulate moving to a reflection:

```
>>> sim hkl [0 0 0.25]
_fourc would move to:
  delta : 76.3430
   th   : 38.1715
  chi   : 0.0000
  phi   : 0.0000

  alpha : -51.8285
  beta  : 51.8285
betain  : 38.1715
betaout : 38.1715
  naz   : 0.0000
  psi   : 0.0000
  qaz   : 0.0000
  tau   : 90.0000
  theta : 38.1715
  ttheta : 76.3430
```

Move to reflection:

```
>>> pos hkl [0 0 0.25]
hkl:      h: 0.00000 k: 0.00000 l: 0.25000
```

As we explained before since in i21 we have four different available geometries, it is important to make sure we are choosing the right one before moving to the reflection as the positions would be different for the different geometries.

```
>>> usediffftth
- setting hkl ---> hkl_diffftth
Loading ub calculation: 'manual'
```

```
WARNING: Ignoring constraint eta
WARNING: Ignoring constraint delta
INFO: difffcalc limits set in $difffcalc/startup/i21.py taken
from http://confluence.diamond.ac.uk/pages/viewpage.action?pageId=51413586
Current hardware limits set to:
  0.0 <= delta <= 180.0 (cut: -180.0)
  0.0 <= th <= 150.0 (cut: 0.0)
 -41.0 <= chi <= 36.0 (cut: -180.0)
-100.0 <= phi <= 100.0 (cut: -180.0)
Note: When auto sector/transforms are used,
      cuts are applied before checking limits.
```

```
>>> sim hkl [0 0 0.25]
_fourc would move to:
  delta : 76.3430
   th   : 38.1715
  chi   : 0.0000
  phi   : 0.0000

  alpha : -51.8285
  beta  : 51.8285
 betain : 38.1715
 betaout : 38.1715
   naz  : 0.0000
   psi  : 0.0000
   qaz  : 0.0000
   tau  : 90.0000
  theta : 38.1715
 ttheta : 76.3430
```

But if we choose a different geometrical configuration:

```
>>> usehighq
- setting hkl ---> hkl_highq
Loading ub calculation: 'manual'
WARNING: Ignoring constraint eta
WARNING: Ignoring constraint delta
INFO: difffcalc limits set in $difffcalc/startup/i21.py taken
from http://confluence.diamond.ac.uk/pages/viewpage.action?pageId=51413586
Current hardware limits set to:
  0.0 <= delta <= 180.0 (cut: -180.0)
  0.0 <= th <= 150.0 (cut: 0.0)
 -41.0 <= chi <= 36.0 (cut: -180.0)
-100.0 <= phi <= 100.0 (cut: -180.0)
Note: When auto sector/transforms are used,
      cuts are applied before checking limits.
```

```
>>> sim hkl [0 0 0.25]
_fourc would move to:
  delta : 72.3430
   th   : 38.1715
  chi   : 0.0000
  phi   : 0.0000

  alpha : -51.8285
```



```

beta : 51.8285
betain : 38.1715
betaout : 38.1715
naz : 0.0000
psi : 0.0000
qaz : 0.0000
tau : 90.0000
theta : 38.1715
ttheta : 76.3430

```

Note that the delta positions are different depending which detector we use as geometrical reference.

## Scanning energy keeping the Q constant. EfixQ

If we want to measure a energy dependence but keeping always the same geometry of a Q as a function of energy, first we need to align on the Q that we want to keep constant.

```

>>> energy
energy: 643.0000

>>> pos hkl [0 0 0.25]
hkl_diffth:h: -0.00000 k: -0.00000 l: 0.25000

>>> hkl

hkl:
hkl_diffth : -0.0000 -0.0000 0.2500

alpha : -51.8285
beta : 51.8285
betain : 38.1715
betaout : 38.1715
naz : 0.0000
psi : 0.0000
qaz : 0.0000
tau : 90.0000
theta : 38.1715
ttheta : 76.3430

_fourc:
delta : 76.3430
th : 38.1715
chi : 0.0000
phi : 0.0000

```

Then to scan the energy

```

>>> scan energy 636 660 1 ct 0.1 hkl [0 0 0.25] fourc
-----
en      h      k      l      delta      th      chi      phi      ct
-----
636.0000 0.00000 -0.00000 0.25000 77.3379 38.6689 0.0000 0.0000 0.32485
637.0000 0.00000 -0.00000 0.25000 77.1940 38.5970 0.0000 0.0000 0.32485
638.0000 -0.00000 -0.00000 0.25000 77.0507 38.5254 0.0000 0.0000 0.32485
639.0000 -0.00000 -0.00000 0.25000 76.9080 38.4540 0.0000 0.0000 0.32485
640.0000 0.00000 -0.00000 0.25000 76.7659 38.3829 0.0000 0.0000 0.32485

```

641.0000	-0.00000	-0.00000	0.25000	76.6244	38.3122	0.0000	0.0000	0.32485
642.0000	-0.00000	-0.00000	0.25000	76.4834	38.2417	0.0000	0.0000	0.32485
643.0000	-0.00000	-0.00000	0.25000	76.3430	38.1715	0.0000	0.0000	0.32485
644.0000	-0.00000	-0.00000	0.25000	76.2032	38.1016	0.0000	0.0000	0.32485
645.0000	-0.00000	-0.00000	0.25000	76.0640	38.0320	0.0000	0.0000	0.32485
646.0000	-0.00000	-0.00000	0.25000	75.9253	37.9626	0.0000	0.0000	0.32485
647.0000	0.00000	-0.00000	0.25000	75.7871	37.8936	0.0000	0.0000	0.32485
648.0000	-0.00000	-0.00000	0.25000	75.6496	37.8248	0.0000	0.0000	0.32485
649.0000	-0.00000	-0.00000	0.25000	75.5126	37.7563	0.0000	0.0000	0.32485
650.0000	-0.00000	-0.00000	0.25000	75.3761	37.6880	0.0000	0.0000	0.32485
651.0000	0.00000	-0.00000	0.25000	75.2402	37.6201	0.0000	0.0000	0.32485
652.0000	-0.00000	-0.00000	0.25000	75.1048	37.5524	0.0000	0.0000	0.32485
653.0000	-0.00000	-0.00000	0.25000	74.9699	37.4850	0.0000	0.0000	0.32485
654.0000	-0.00000	-0.00000	0.25000	74.8356	37.4178	0.0000	0.0000	0.32485
655.0000	0.00000	-0.00000	0.25000	74.7018	37.3509	0.0000	0.0000	0.32485
656.0000	-0.00000	-0.00000	0.25000	74.5686	37.2843	0.0000	0.0000	0.32485
657.0000	0.00000	-0.00000	0.25000	74.4358	37.2179	0.0000	0.0000	0.32485
658.0000	0.00000	-0.00000	0.25000	74.3036	37.1518	0.0000	0.0000	0.32485
659.0000	0.00000	-0.00000	0.25000	74.1719	37.0860	0.0000	0.0000	0.32485
660.0000	-0.00000	-0.00000	0.25000	74.0407	37.0204	0.0000	0.0000	0.32485

Where  $hkl [0 0 0.25]$  is the  $Q$  that we want to keep fixed,  $ct$  is the counter that needs to be substituted by the names of the counters we want to use in each case. By including `fourc` at the end of the scan, GDA will print on the command line the values of `delta`, `th`, `chi` and `phi` for each point of the scan.

## Moving to different azimuthal angles keeping the Q constant.

Often when we align a particular Q direction for one sample we find that there are certain offsets in  $\theta$ ,  $\chi$  or  $\phi$  that need to be taken into account in order to be properly aligned.

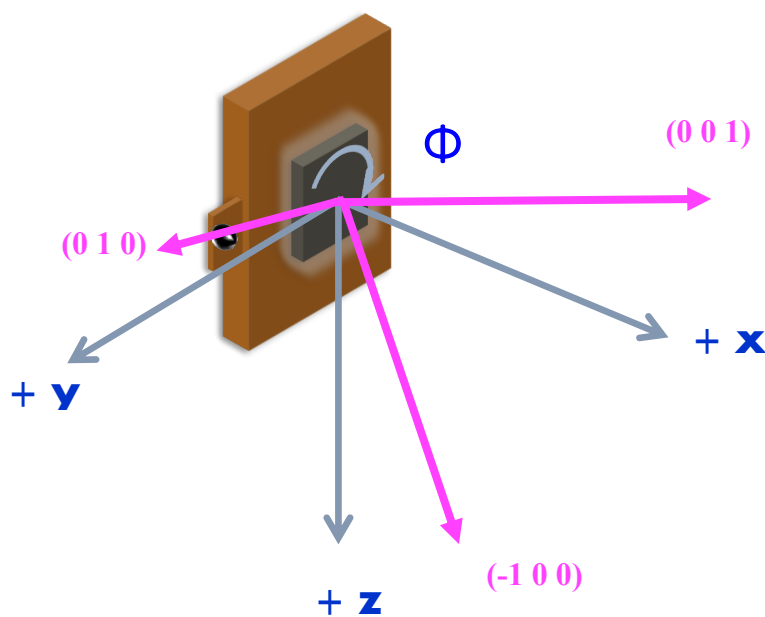
If we want to change geometry from one azimuthal angle to another keeping the specular geometry fixed, diffcalc can help us doing this while taking the offsets into account. For instance if we want to move from  $(\pi,0)$  geometry to  $(\pi,\pi)$  geometry.

Remember that

- **psi** - azimuthal rotation about scattering vector of reference vector (from scattering plane)

Often when we align the scattering plane we know the offsets in  $\phi$  only by the laue measurements as most of the times we cannot access any in plane reflection to refine it using diffraction.

Imaging a case as the one below when the scattering plane is forming an angle with the laboratory axes.



```
>>> newub
calculation name: uboffset
crystal name: SBMO
crystal system
1) Triclinic
2) Monoclinic
3) Orthorhombic
4) Tetragonal
5) Rhombohedral
6) Hexagonal
7) Cubic
[1]: 7
a[1]: 3.9
```

```
>>> addorient
h[0.0]: 0
k[0.0]: 0
l[0.0]: 1
x[0.0]: 1
y[0.0]: 0
z[0.0]: 0
current pos[y]: n
  delta[74.0407334683]: 0
  th[37.3675946909]: 0
  chi[10.0]: 10
  phi[5.0]: 5
tag: [001]
```

```
>>> In [121]: addorient
h[0.0]: 0
k[0.0]: 1
l[0.0]: 0
x[0.0]: 0
y[0.0]: 1
z[0.0]: 0
current pos[y]: n
  delta[74.0407334683]: 0
  th[37.3675946909]: 0
  chi[10.0]: 10
  phi[5.0]: 5
tag: [010]
Calculating UB matrix.
```

Here we have included the offsets we know from the Laue measurements:  $\phi = 5$  deg. and  $\chi = 10$  deg. while defining the orientation reflections that will create our ub matrix.

```
>>> ub
UBCALC

  name:          uboffset

REFERNCE

  n_phi:         0.17365  -0.08583  -0.98106
  n_hkl:         1.00000   0.00000   0.00000 <- set

SURFACE NORMAL

  n_phi:         1.00000   0.00000   0.00000 <- set
  n_hkl:         0.17365   0.00000   0.98481

CRYSTAL

  name:          SBMO

  a, b, c:       3.90000   3.90000   3.90000
                90.00000  90.00000  90.00000  Cubic

  B matrix:      0.00000   0.00000   1.61107
                0.00000   1.61107   0.00000
                -1.61107   0.00000   0.00000

UB MATRIX
```

```
U matrix:  0.98481  0.00000  -0.17365
           0.01513  0.99619  0.08583
           0.17299  -0.08716  0.98106
```

```
miscut:
  angle: 10.00000
  axis:  0.00000  -0.99619  0.08716
```

```
UB matrix:  0.27976  0.00000  1.58660
            -0.13828  1.60494  0.02438
            -1.58056  -0.14041  0.27870
```

#### REFLECTIONS

```
<<< none specified >>>
```

#### CRYSTAL ORIENTATIONS

	H	K	L	X	Y	Z	DELTA	TH	CHI	PHI	TAG
1	0.00	0.00	1.00	1.00	0.00	0.00	0.0000	0.0000	10.0000	5.0000	[001]
2	0.00	1.00	0.00	0.00	1.00	0.00	0.0000	0.0000	10.0000	5.0000	[010]

We want to define the  $(\pi, 0)$  position when the plane of scattering coincides with the plane contained by the [001] and [010] directions. In this case since there is an offset between this plane and the  $\text{phi}=0$  position, we need to consider as a reference for the azimuthal rotation the [010] crystal axes and not the laboratory reference. For doing this:

```
>>> setnhkl [0 1 0]
      n_phi: -0.00000  0.99619  -0.08716
      n_hkl:  0.00000  1.00000  0.00000 <- set
```

Then we constrain the azimuthal angle so that the  $(\pi, 0)$  plane coincides with the plane of scattering. Based on how we have defined the ub matrix, in this case it would be  $\text{psi} = 0$ .

```
>>> con psi 0
      psi : 0.0000

>>> pos en 643
      en:  643.0000

>>> sim hkl [0 0 0.25]
_fourc would move to:
      delta :  76.3430
      th :   38.1715
      chi :   10.0000
      phi :    5.0000

      alpha : -51.8285
      beta :  51.8285
      betain : 37.4904
      betaout : 37.4904
      naz : -0.0000
      psi :  0.0000
      qaz :  0.0000
      tau :  90.0000
      theta : 38.1715
      ttheta : 76.3430
```

```
>>> pos hkl [0 0 0.25]
hkl_diffth:h: -0.00000 k: -0.00000 l: 0.25000
```

This will move the motors to the  $(\pi, 0)$  plane taking into account all offsets.

If now we want to move to the  $(\pi, \pi)$  scattering plane, then we just need to constrain the azimuthal angle for 45 deg.

```
>>> con psi 45
psi : 45.0000

>>> sim hkl [0 0 0.25]
_fourc would move to:
  delta : 76.3430
   th   : 31.0644
   chi  :  7.0530
   phi  : 50.4385

  alpha : -33.7729
  beta  : 33.7729
betain  : 30.8036
betaout : 44.8425
  naz   : -58.2832
  psi   : 45.0000
  qaz   :  0.0000
  tau   : 90.0000
  theta : 38.1715
  ttheta : 76.3430
```

```
>>> pos hkl [0 0 0.25]
hkl_diffth:h: -0.00000 k: -0.00000 l: 0.25000
```

This will move the motors to the  $(\pi, \pi)$  plane taking into account all offsets.

Analogously we can use this system to move to any other azimuthal angle we want considering all the offsets.