

## 3D IMAGE (VOLUME) ORIENTATION AND LOCATION IN SPACE:

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There are 3 different methods by which continuous coordinates can

attached to voxels. The discussion below emphasizes 3D volumes, and

the continuous coordinates are referred to as  $(x,y,z)$ . The voxel

index coordinates (i.e., the array indexes) are referred to as  $(i,j,k)$ ,

with valid ranges:

$i = 0 \dots \text{dim}[1]-1$   
 $j = 0 \dots \text{dim}[2]-1$  (if  
 $\text{dim}[0] \geq 2$ )

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k = 0 .. dim[3]-1 (if  
dim[0] >= 3)
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The (x,y,z) coordinates refer to the CENTER of a voxel. In methods

2 and 3, the (x,y,z) axes refer to a subject-based coordinate system,

with

+x = Right +y = Anterior  
+z = Superior.

This is a right-handed coordinate system. However, the exact direction

these axes point with respect to the subject depends on qform\_code

(Method 2) and sform\_code (Method 3).

N.B.: The i index varies most rapidly, j index next, k index

slowest.

Thus, voxel (i,j,k) is stored starting at location  
 $(i + j \cdot \text{dim}[1] + k \cdot \text{dim}[1] \cdot \text{dim}[2]) \cdot (\text{bitpix}/8)$   
into the dataset array.

N.B.: The ANALYZE 7.5 coordinate system is  
+x = Left +y = Anterior  
+z = Superior  
which is a left-handed coordinate system. This backwardness is  
too difficult to tolerate, so this NIFTI-1 standard specifies the  
coordinate order which is most common in functional neuroimaging.

N.B.: The 3 methods below all

give the locations of the voxel centers

in the  $(x,y,z)$  coordinate system. In many cases, programs will wish

to display image data on some other grid. In such a case, the program

will need to convert its desired  $(x,y,z)$  values into  $(i,j,k)$  values

in order to extract (or interpolate) the image data. This operation

would be done with the inverse transformation to those described below.

**N.B.: Method 2 uses a factor 'qfac' which is either -1 or 1; qfac is**

**stored in the otherwise**

unused pixdim[0]. If  
pixdim[0]=0.0 (which  
should not occur), we take  
qfac=1. Of course, pixdim[0] is  
only used  
when reading a NIFTI-1  
header, not when reading an  
ANALYZE 7.5 header.

In NIFTI-1 files, dimensions  
1,2,3 are for space, dimension 4  
is for time,

and dimension 5 is for  
storing multiple values at each  
spatiotemporal

voxel. Some examples:

- A typical whole-brain  
FMRI experiment's time series:

- dim[0] = 4

- dim[1] = 64

pixdim[1] = 3.75 xyz\_t\_units =  
NIFTI\_UNITS\_MM

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        - dim[2] = 64
pixdim[2] = 3.75
NIFTI_UNITS_SEC
        - dim[3] = 20
pixdim[3] = 5.0
        - dim[4] = 120
pixdim[4] = 2.0
    - A typical T1-weighted
      anatomical volume:
        - dim[0] = 3
        - dim[1] = 256
pixdim[1] = 1.0   xyz_t_units =
NIFTI_UNITS_MM
        - dim[2] = 256
pixdim[2] = 1.0
        - dim[3] = 128
pixdim[3] = 1.1
    - A single slice EPI time
      series:
        - dim[0] = 4
        - dim[1] = 64
pixdim[1] = 3.75 xyz_t_units =

```

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NIFTI_UNITS_MM
    - dim[2] = 64
pixdim[2] = 3.75
NIFTI_UNITS_SEC
    - dim[3] = 1
pixdim[3] = 5.0
    - dim[4] = 1200
pixdim[4] = 0.2
    - A 3-vector stored at each
point in a 3D volume:
    - dim[0] = 5
    - dim[1] = 256
pixdim[1] = 1.0    xyzt_units =
NIFTI_UNITS_MM
    - dim[2] = 256
pixdim[2] = 1.0
    - dim[3] = 128
pixdim[3] = 1.1
    - dim[4] = 1
pixdim[4] = 0.0
    - dim[5] = 3
intent_code =

```

NIFTI\_INTENT\_VECTOR

- A single time series with  
a 3x3 matrix at each point:

- dim[0] = 5

- dim[1] = 1

xyzt\_units = NIFTI\_UNITS\_SEC

- dim[2] = 1

- dim[3] = 1

- dim[4] = 1200

pixdim[4] = 0.2

- dim[5] = 9

intent\_code =

NIFTI\_INTENT\_GENMATRIX

- intent\_p1 = intent\_p2

= 3.0 (indicates matrix  
dimensions)

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METHOD 2 (used when qform\_code >  
0, which should be the "normal"  
case):

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The (x,y,z) coordinates are  
given by the pixdim[] scales, a  
rotation

matrix, and a shift. This

method is intended to represent

"scanner-anatomical"

coordinates, which are often

embedded in the

image header (e.g., DICOM

fields (0020,0032), (0020,0037),

(0028,0030),

and (0018,0050)), and

represent the nominal

orientation and location of

the data. This method can

also be used to represent

"aligned"

coordinates, which would  
typically result from some post-  
acquisition

alignment of the volume to a  
standard orientation (e.g., the  
same

subject on another day, or a  
rigid rotation to true  
anatomical

orientation from the tilted position of the subject in the scanner).

The formula for  $(x,y,z)$  in terms of header parameters and  $(i,j,k)$  is:

$$\begin{bmatrix} x \\ \text{pixdim}[1] * i \\ \text{qoffset}_x \end{bmatrix} = \begin{bmatrix} R11 & R12 & R13 \\ R21 & R22 & R23 \end{bmatrix}$$

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[          pixdim[2] * j ] +
[ qoffset_y ]

[ z ]    [ R31 R32 R33 ]

[ qfac * pixdim[3] * k ]
[ qoffset_z ]

```

The `qoffset_*` shifts are in  
 the NIFTI-1 header. Note that  
 the center  
 of the  $(i,j,k)=(0,0,0)$  voxel  
 (first value in the dataset

array) is

just

$(x, y, z) = (qoffset\_x, qoffset\_y, qoffset\_z)$ .

The rotation matrix  $R$  is calculated from the quatern\_\* parameters.

This calculation is described below.

The scaling factor  $q_{\text{fac}}$  is either 1 or -1. The rotation matrix  $R$

defined by the quaternion parameters is "proper" (has determinant 1).

This may not fit the needs of the data; for example, if the image

grid is

$i$  increases from Left-to-

Right

j increases from Anterior-  
to-Posterior

k increases from Inferior-  
to-Superior

Then  $(i,j,k)$  is a left-handed  
triple. In this example, if  
 $qfac=1,$

the R matrix would have to be

$$\begin{bmatrix} 1 & 0 & 0 \end{bmatrix}$$



$$\begin{bmatrix} 0 & -1 & 0 \end{bmatrix}$$
 which is  
"improper" (determinant = -1).

$$\begin{bmatrix} 0 & 0 & 1 \end{bmatrix}$$

If we set  $qfac=-1$ , then the R  
matrix would be

$$\begin{bmatrix} 1 & 0 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 0 & -1 & 0 \end{bmatrix}$$
 which is  
proper.

$$\begin{bmatrix} 0 & 0 & -1 \end{bmatrix}$$

This R matrix is represented  
by quaternion  $[a,b,c,d] =$   
 $[0,1,0,0]$

(which encodes a 180 degree  
rotation about the x-axis).