

STIR User's Meeting

Reconstruction of PET data acquired
with the BrainPET using STIR

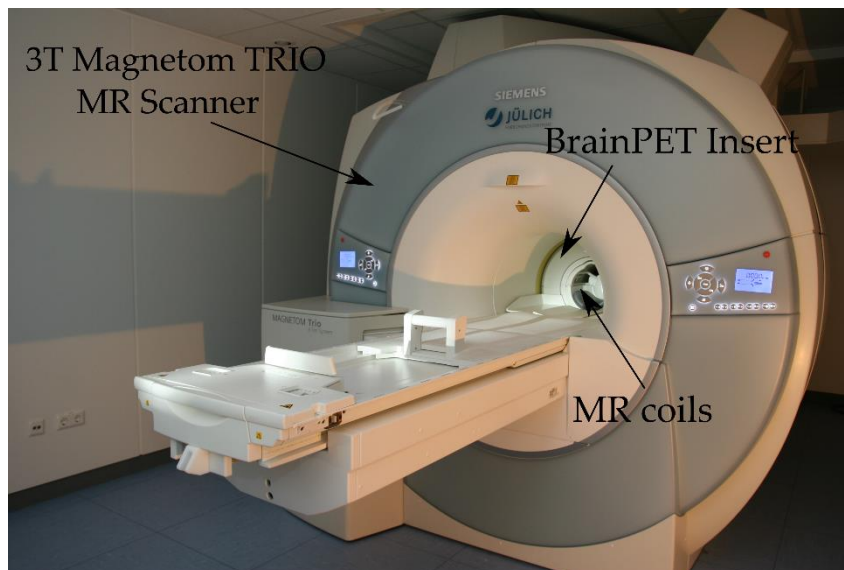
31. October 2013 | Liliana Caldeira

Outline

- 3T MR-BrainPET System
- Reading BrainPET data into STIR
- FBP reconstruction
- OSEM/OSMAPOSL reconstruction

3T MR-BrainPET System

3T Magnetom TRIO + BrainPET Insert

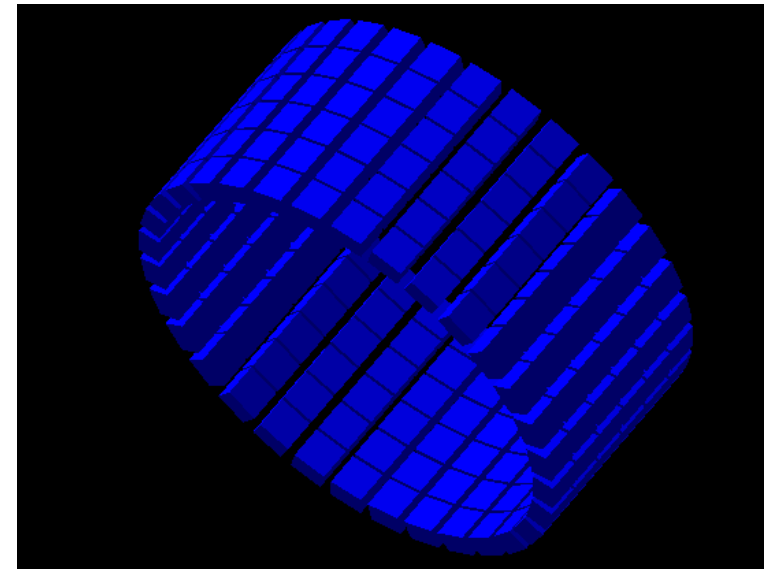


Complete System

BrainPET Insert

3T MR-BrainPET System

Detector Geometry



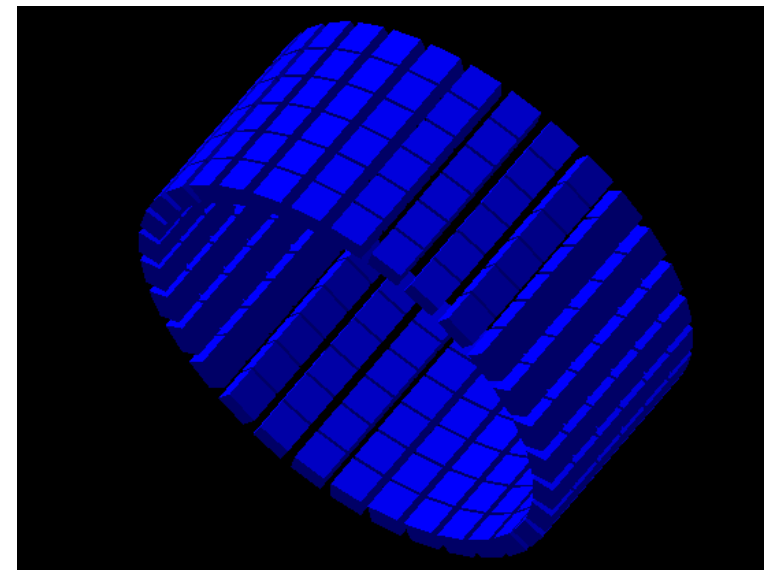
Detector Cassette

GATE model of BrainPET insert

3T MR-BrainPET System

BrainPET properties

| | |
|--------------------|----------------------------|
| Ring Diameter | 376 mm |
| Axial FOV | 192.5 mm |
| Crystal | 2.5x2.5x20 mm ³ |
| Crystal Block | 12x12 |
| # of Crystal Rings | 12x6=72 |
| # of Crystals/Ring | 12x32=384 |



GATE model of BrainPET insert

3T MR-BrainPET System

BrainPET properties including GAPS

| | |
|--------------------|----------------------------|
| Ring Diameter | 376 mm |
| Axial FOV | 192.5 mm |
| Crystal | 2.5x2.5x20 mm ³ |
| Crystal Block | 12x12 |
| # of Crystal Rings | 12x6=72 |
| # of Crystals/Ring | 12x32=384 |

| | |
|----------------------------|-----------------------------|
| Axial Gaps | 5 |
| Size Axial Gap | 2.5 mm |
| Transaxial Gaps | 32 |
| Size Transaxial Gap | 7.5 mm =3x2.5 mm |
| # of Crystal Rings | 12x6=72 +5 =77 |
| # of Crystals/Ring | 12x32=384 +32x3 =480 |

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Reading BrainPET data into STIR

Understanding Sinogram Structure

| | |
|--------------------|-----------------------------|
| Ring Diameter | 376 mm |
| Axial FOV | 192.5 mm |
| Crystal | 2.5x2.5x20 mm ³ |
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| # of Crystal Rings | 12x6=72 +5 =77 |
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- Sinogram Order
- Use simple data
- Use manip_projdata

Reading BrainPET data into STIR

Header File sinogram.hs (1/4)

```
!INTERFILE :=  
name of data file := sinogram.s  
originating system := Userdefined  
!GENERAL DATA :=  
!GENERAL IMAGE DATA :=  
!type of data := PET  
imagedata byte order := LITTLEENDIAN  
!PET STUDY (General) :=  
!PET data type := Emission  
applied corrections := {arc correction}  
!number format := float  
!number of bytes per pixel := 4
```

User Defined Scanner

Data Format

Reading BrainPET data into STIR

Header File sinogram.hs (2/4)

```
number of dimensions := 4
matrix axis label [4] := segment
!matrix size [4] := 15
matrix axis label [3] := view
!matrix size [3] := 192
matrix axis label [2] := axial coordinate
!matrix size [2] := {35,53,71,89,107,125,143,153,143,125,107,89,71,53,35}
matrix axis label [1] := tangential coordinate
!matrix size [1] := 256
minimum ring difference per segment :={-67,-58,-49,-40,-31,-22,-13,-4,5,
14,23,32,41,50,59}
maximum ring difference per segment :={-59,-50,-41,-32,-23,-14,-5,4,13,
22,31,40,49,58,67}
```

Sinogram Organisation:
Span 9 (5+4)

Reading BrainPET data into STIR

Header File sinogram.hs (3/4)

```
Scanner parameters:=  
Scanner type := Userdefined  
Number of rings           := 77  
Number of detectors per ring := 480  
Inner ring diameter (cm)   := 376  
Average depth of interaction (cm) := 0.7  
Distance between rings (cm) := 0.25  
Default bin size (cm)      := 0.125  
View offset (degrees)      := 0  
Maximum number of non-arc-corrected bins := 256  
Default number of arc-corrected bins    := 256
```

Scanner Information

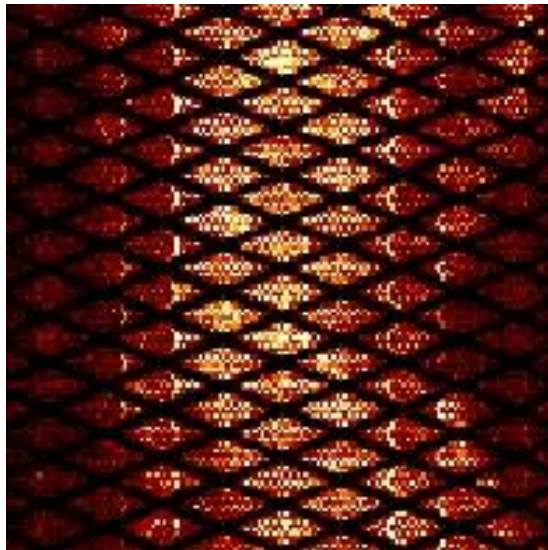
Reading BrainPET data into STIR

Header File sinogram.hs (4/4)

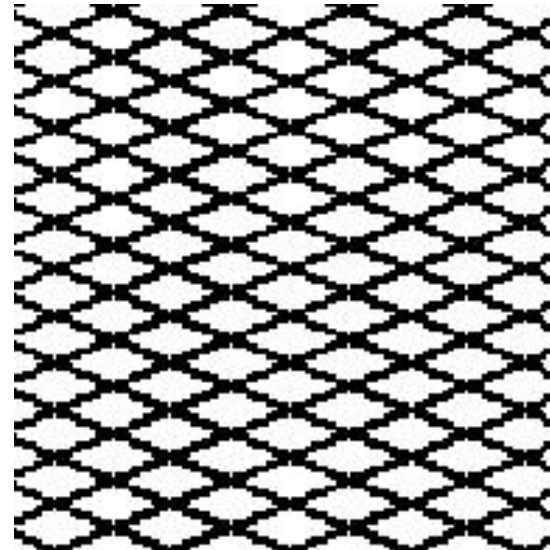
```
Number of blocks per bucket in transaxial direction      := 1
Number of blocks per bucket in axial direction          := 1
Number of crystals per block in axial direction         := 1
Number of crystals per block in transaxial direction    := 1
Number of detector layers                               := 1
Number of crystals per singles unit in axial direction  := 1
Number of crystals per singles unit in transaxial direction := 1
end scanner parameters:=
effective central bin size (cm) := 0.125
image scaling factor[1] := 1
data offset in bytes[1] := 0
number of time frames := 1
!END OF INTERFILE :=
```

Reading BrainPET data into STIR

Projection Data



BrainPET data (using manip_projdata)



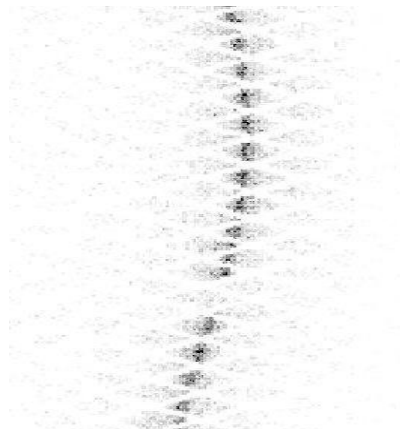
Gap Structure in Sinogram

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FBP reconstruction

Point Sources – Projection data



Example Figure from [3] H. Herzog et al. "Influence from high and ultra-high magnetic field on positron range measured with a 9.4T MR-BrainPET". IEEE NSS/MIC 2010.

FBP reconstruction

Point Sources – Parameter File (1/2)

FBP3DRP file.par

```
fbp3drpparameters :=  
input file := noisy_testsino.hs  
output filename prefix := output_noisy  
  
;;;;;;;;; output image parameters  
zoom := 1  
; defaults to cover whole FOV  
;xy output image size (in pixels) := 180  
maximum absolute segment number to process := 7  
;;;;;;;;; parameters for initial image  
num segments to combine with ssrb := -1
```


FBP reconstruction

Point Sources – Parameter File (2/2)

```
;filter parameters, default to pure ramp
alpha parameter for ramp filter := 1
cut-off for ramp filter (in cycles) := 0.5
;;;;;;;;; parameters for Colsher filter
alpha parameter for colsher filter in axial direction := 1
cut-off for colsher filter in axial direction (in cycles) := 0.5
alpha parameter for colsher filter in planar direction := 1
cut-off for colsher filter in planar direction (in cycles) := 0.5
; define colsher on finer grid.
stretch factor for colsher filter definition in axial direction := 2
stretch factor for colsher filter definition in planar direction := 2
; allow less padding. DO NOT USE
transaxial extension for fft := 1
axial extension for fft := 1
;;;;;;;;; other parameters
save intermediate images := 0
display level := 0
end :=
```

FBP reconstruction

Point Sources – Reconstructed Images



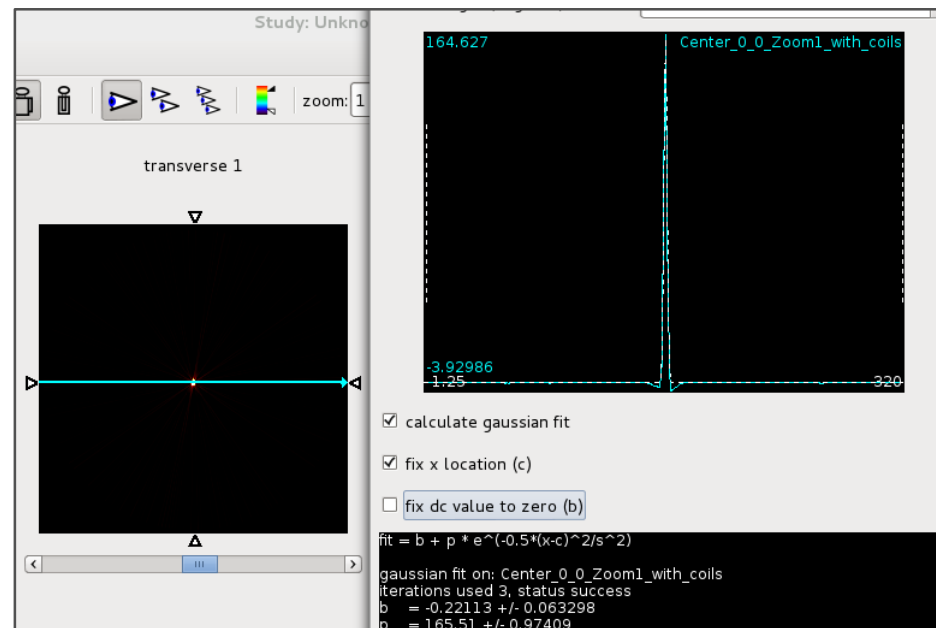
Transverse Image



Coronal Image

FBP reconstruction

Point Sources – Reconstructed Images



AMIDE

FBP reconstruction

Point Sources – Results in publications

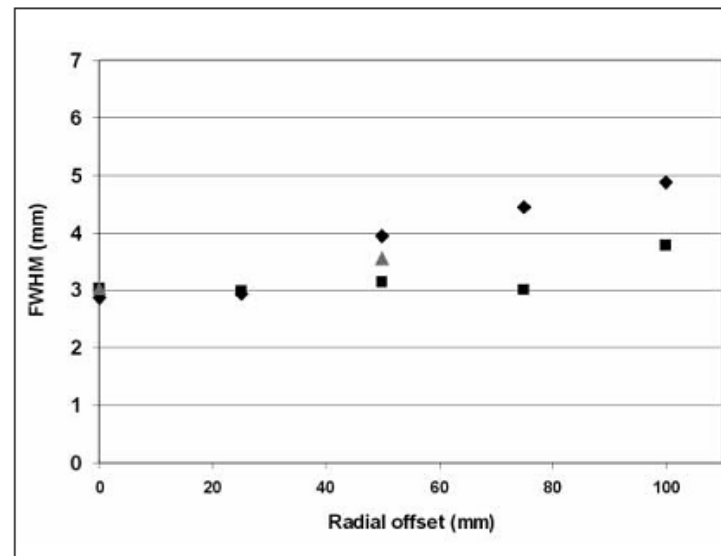


Fig. 2

Plots of tangential (■), radial (◆) and axial (Δ) image resolution expressed as full width at half maximum (FWHM) measured with line and point sources in air at different radial offsets.

Nuklearmedizin 2/2011

Figure from [1] H. Herzog et al. "High Resolution BrainPET combined with Simultaneous MRI". NuklearMedizin. 2011.

FBP reconstruction

Point Sources – Results in publications



*Sinograms of the ^{120}I point source measured at 0 T (left), 3 T, 7 T and 9.4 T (right):
Figure from [3] H. Herzog et al. "Influence from high and ultra-high magnetic field on
positron range measured with a 9.4 T MR-BrainPET". IEEE NSS/MIC 2010.*

FBP reconstruction

Point Sources – Results in publications

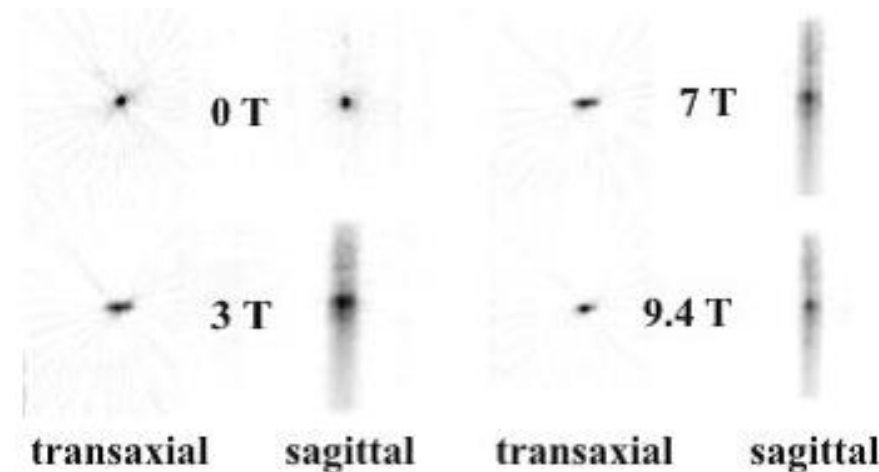


Fig. 7. Images of the point source filled with ^{120}I at different magnetic field strengths.

Figure from [3] H. Herzog et al. "Influence from high and ultra-high magnetic field on positron range measured with a 9.4T MR-BrainPET". IEEE NSS/MIC 2010.

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OSEM reconstruction

Patient data – Parameter File (1/2)

```
OSMAPOSLParameters :=  
objective function type:=  
PoissonLogLikelihoodWithLinearModelForMeanAndProjData  
  
PoissonLogLikelihoodWithLinearModelForMeanAndProjData Parameters:=  
input file := verynoisy_testsino.hs  
additive sinogram := 0  
  
projector pair type := Separate Projectors  
  Projector Pair Using Separate Projectors Parameters:=  
    Forward projector type := Ray Tracing  
      Forward Projector Using Ray Tracing Parameters:=  
        End Forward Projector Using Ray Tracing Parameters:=  
    Back projector type := Interpolation  
      Back Projector Using Interpolation Parameters:=  
        End Back Projector Using Interpolation Parameters:=  
    End Projector Pair Using Separate Projectors Parameters:=
```


OSEM reconstruction

Patient data – Parameter File (2/2)

```
recompute sensitivity := 1
sensitivity filename:=sensOSMAPOS�pair.hv
end PoissonLogLikelihoodWithLinearModelForMeanAndProjData Parameters:=

output filename prefix := veryoutputnoisyOSMAPOS�pair

number of subsets:= 1
number of subiterations:= 24
Save estimates at subiteration intervals:= 3

END :=
```

OSEM reconstruction

Patient data – Normalisation File

...

End Projector Pair Using Separate Projectors Parameters:=

```
Bin Normalisation type := From ProjData
Bin Normalisation From ProjData :=
normalisation projdata filename:= NORMaddnozeros.hs
End Bin Normalisation From ProjData:=
```

recompute sensitivity := 1

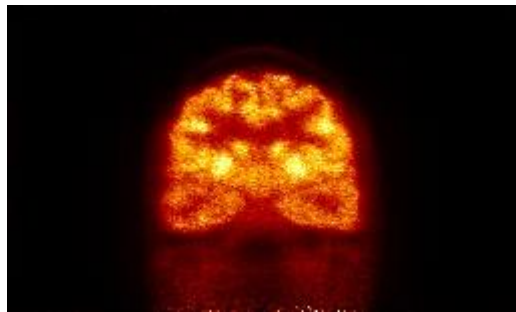
...

Efficiencies Sinogram was available, but calculations were necessary for STIR:

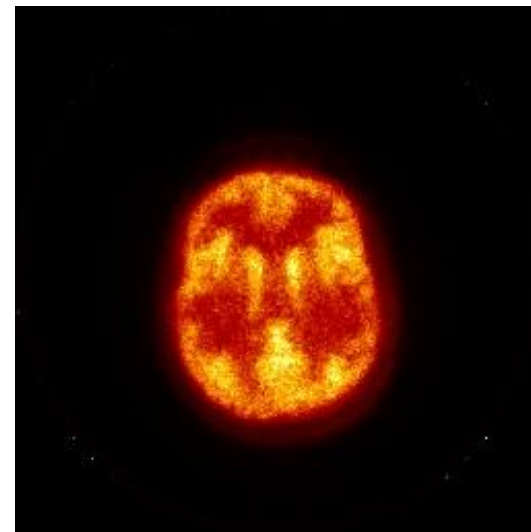
- 1) STIR uses multiplicative factors (1/efficiency)
- 2) Zero efficiency -> Very high multiplicative factor

OSEM reconstruction

Patient data – Reconstructed Images



Coronal Image



Transverse Image

OSEM reconstruction

Patient data – Parameter File using Median Root Prior

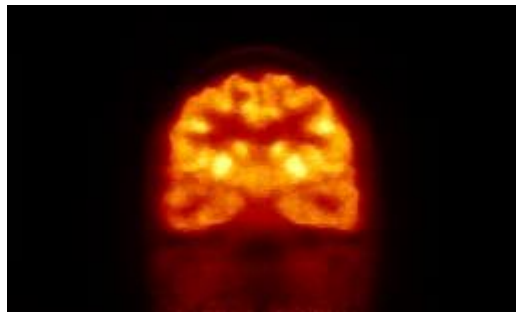
```
prior type := FilterRootPrior
FilterRootPrior Parameters :=
  penalisation factor := 1
  Filter type := Median
    Median Filter Parameters :=
      mask radius x := 1
      mask radius y := 1
      mask radius z := 1
    End Median Filter Parameters:=
END FilterRootPrior Parameters :=
```

| | | |
|-------------|-------------|-------------|
| λ_1 | λ_2 | λ_3 |
| λ_4 | λ_i | λ_5 |
| λ_6 | λ_7 | λ_8 |

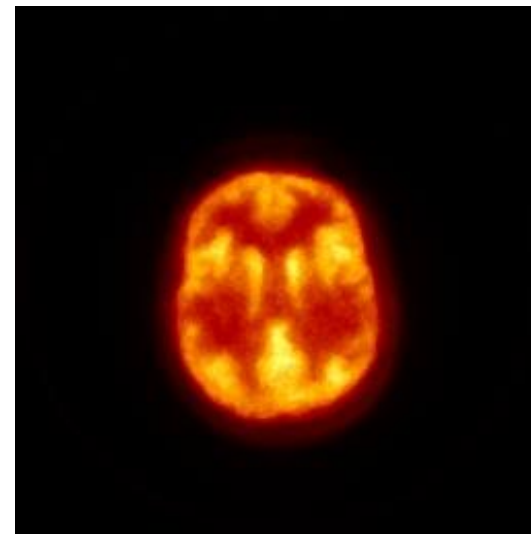
```
MAP model := multiplicative
```

OSEM reconstruction

Patient data – Reconstructed Images with Median Root Prior



Coronal Image



Transverse Image

Conclusion

- BrainPET data was read successfully into STIR
- FBP reconstruction for Point Source measurements
- OSEM/OSMAPOSL reconstruction for Patient data

References

- [1] H. Herzog et al. “High Resolution BrainPET combined with Simultaneous MRI”. NuklearMedizin. 2011.

- [2] L. Caldeira et al. “Reconstruction of PET Data Acquired with the BrainPET using STIR”. IEEE NSS/MIC 2012.

- [3] H. Herzog et al. “Influence from high and ultra-high magnetic field on positron range measured with a 9.4T MR-BrainPET”. IEEE NSS/MIC 2010.

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