

herp

derp

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# Indhold

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# **1 Topogram**

## **1.1 Routine**

- mA: 35
- kV: 120
- Topogram length: 1024 mm
- Tube position: Top

## **1.2 Scan**

- mA: 35
- kV: 120
- Delay: 4s
- Topogram length: 1024 mm
- Direction: Craniocaudal
- Tube position: Top
- API (-1 betyder None, ellers skal tallet oversaettes (stoerre arbejde): -1
- Kernel: T80s sharp
- Window: Topogram Body

# **2 CT WB**

## **2.1 Routine**

- Eff. mAs: 225
- kV: 14
- CARE Dose4D: On
- CTDIvol: 15.1475mGy
- Scan time: 1.2000000 s

- Delay: 4.000 s
- Slice: 3.00 mm
- No. of images: Samme som i foerste recon, slet?(y/n)
- Tilt: 225 grader

## **2.2 Scan**

- Quality ref. mAs: 225
- Eff. mAs: 225
- kV: 120
- Scan time: 14.840 s
- Rotation time: 0.500 s
- Delay: 4.000 s
- Slice: 0.6 mm
- Pitch: 1.2000000
- Direction: Craniocaudal

## **2.3 Recon**

### **2.3.1 Recon 1**

- Series description: AC CT
- Slice: 3.00
- Kernel: "B19f PET very smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 1
- FoV: 700
- Center X: 0
- Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: AxialRJ
- Recon Axis: Axial
- Image order: Craniocaudal
- Recon increment: 3.000
- No. of images: 324

### **2.3.2 Recon 2**

- Series description: CT WB 3.0 B30f
- Slice: 3.00
- Kernel: "B30f medium smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 0
- FoV: 500
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: AxialRJ
- Recon Axis: Axial
- Image order: Craniocaudal
- Recon increment: 3.000
- No. of images: 324

### **2.3.3 Recon 3**

- Series description: CT WB 3.0 SPO cor
- Slice: 3.00
- Kernel: "B30f medium smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 0
- FoV: 1202
- Center X: 0
- Center Y: 0
- Mirroring: None
- Extended CT scale: Standard
- Recon job: VolumeRJ
- Recon Axis: Coronal
- Image order: AP
- Recon increment: 3.000
- No. of images: 267

### **2.3.4 Recon 4**

- Series description: CT WB 3.0 SPO sag
- Slice: 3.00
- Kernel: "B30f medium smooth"
- Window: "Abdomen"
- Extended FoV(1/0): 0
- FoV: 1202
- Center X: 0
- Center Y: 0

- Mirroring: None
- Extended CT scale: Standard
- Recon job: VolumeRJ
- Recon Axis: Sagittal
- Image order: LeftToRight
- Recon increment: 3.000
- No. of images: 267

### **3    Pause**

LoL, I dunno

## **4    PET WB**

### **4.1    Routine**

- Isotope: F-18
- Pharm.: FDG
- Inj. Dose: 400 MegaBequerels
- Inj. date (date/month - year): 15/06 - 2011
- Inj. time: 10:37:00
- Scan mode: Sinogram
- Scan range: Match CT FOV
- No. of beds: 6
- Scan duration/bed: 150 Seconds

## **4.2 Scan**

- Autoload: On
- Rebinner LUT: 0
- Scan output: Sinogram
- Sinogram mode: Trues
- Input trigger signal: None
- LLD (keV): 435
- ULD (keV): 650

## **4.3 Recon**

### **4.3.1 Recon 1**

- Series description: ET WB Image check
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: Iterative
- Iterations: 4
- Subsets: 8
- Image size: 168
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 4
- Offset X (mm): 0
- Offset Y (mm): 0
- Attenuation correction: (1,1)
- Scatter correction: 1
- Match CT slice location: 1
- Save intermediate data: 0



### 4.3.2 Recon 2

- Series description: ET WB Corr True X 2 m
- Recon range (bed): ups..
- Output image type: Corrected
- Recon method: TrueX
- Iterations: 3
- Subsets: 21
- Image size: 336
- Zoom: 1
- Filter: Gaussian
- FWHM (mm): 2
- Offset X (mm): 0
- Offset Y (mm): 0
- Attenuation correction: (1,1)
- Scatter correction: 1
- Match CT slice location: 1
- Save intermediate data: 0