SIEMENS MAGNETOM Investigational_Device_7T syngo MR B17

VASO binary version 109

TA: 26:11 PAT: 2 Voxel size: 0.7×0.7×1.0 mm Rel. SNR: 1.00 USER: VASO_109

Properties		PAT mode	GRAPPA
Prio Recon	Off	Accel. factor PE Ref. lines PE	2 24
Before measurement		Accel. factor 3D	1
After measurement		Ref. lines 3D	8
Load to viewer	On	Reference scan mode	Separate
Inline movie	Off		
Auto store images	On	Prescan Normalize	Off
Load to stamp segments	Off	Raw filter	Off
Load images to graphic	Off	Elliptical filter	Off
segments		Hamming	Off
Auto open inline display	Off	Geometry	
Start measurement without	On	Multi-slice mode	Interleaved
further preparation		Series	Ascending
Wait for user to start	Off		
Start measurements	single	Special sat.	Parallel F
Routine		Gap	25.0 mm
Slab group 1		Thickness	100 mm
Slabs	1	Table position	ш
Dist. factor	50 %	Table position	H 0 mm
Position	R33.7 A7.8 H20.4	Table position Inline Composing	Off
Orientation	T > S35.9	Inline Composing	Oli
Phase enc. dir.	R >> L	System	
Rotation	60.00 deg	V32	Off
Phase oversampling	0 %	A32	On
Slice oversampling	0.0 %	Decite of a second	FIV
Slices per slab	12	Positioning mode	FIX
FoV read	32.7 mm	MSMA	S-C-T
FoV phase	300.0 %	Sagittal	R >> L
Slice thickness	0.99 mm	Coronal	A >> P
TR	1702.40 ms	Transversal	F >> H
TE	26 ms	Save uncombined	Off
Averages	1	Coil Combine Mode	Sum of Squares
Concatenations	1	AutoAlign	 D ()
Filter	None	Auto Coil Select	Default
Coil elements	A32	Shim mode	Standard
		Adjust with body coil	Off
Contrast		Confirm freq. adjustment	Off
Perfusion mode	Picore Q2TIPS	Assume Silicone	Off
TI2	750 ms	! Ref. amplitude 1H	220.000 V
TI1	50 ms	Adjustment Tolerance	Auto
TI1s	50 ms	Adjust volume	
Flip angle	34 deg	! Position	R28.2 A11.1 H12.7
Fat suppr.	Fat sat.	! Orientation	Transversal
Fat sat. mode	Strong	! Rotation	0.00 deg
Averaging mode	Long term	! R >> L	89 mm
Reconstruction	Magnitude	! A >> P	81 mm
Measurements	923	! F >> H	58 mm
Delay in TR	0 ms	Dhysia	
Multiple series	Off	Physio Physio	Name
		1st Signal/Mode	None
Perfusion mode	PICORE Q2T	BOLD	
Inversion time 1	50 ms	Motion correction	Off
Saturation stop time	50 ms	Spatial filter	Off
Inversion time 2	750.0 ms	· ·	
Flow limit	100.0 cm/s	Sequence	
Resolution		Introduction	On
	46	Dimension	3D
HASE RESOLUTION		Reordering	Linear
Base resolution	100 %		-
Phase resolution	100 % 100 %	Contrasts	1
Phase resolution Slice resolution	100 %	Bandwidth	1026 Hz/Px
Phase resolution Slice resolution Phase partial Fourier	100 % 6/8	Bandwidth Free echo spacing	1026 Hz/Px Off
Phase resolution Slice resolution	100 %	Bandwidth	1026 Hz/Px

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Gradient mode Normal Excitation Slab-sel. RF spoiling On	
Ampl 110 BWDTH 150 3.1kHz thickness 30 use Ernst angle Off Maxwell Correction Off log physio files Off FFT scale 1.00 dummy prepscan time 3 s z shim 0.00 mT/m*ms RF duration 1456 us RF BWTP 25.0 EFFECTIVE TR 20428 ms PatPartitions 12 EPI phase correction local PAT refscan mode Flash FlashRef BaseRes 46 FlashRef BW 107 Hz/px FlashRef TE 7000 us FlashRef FA 5 deg use CAIPI Off	

Additional Parameters:

- -> GRAPPA Kernel 2x3
- -> GRAPPA Regularization (NoisereductionI=5000)
 -> Partial Fourier Algorithm = POCS
- -> POCS iterations = 8