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\\Study Protocols **BRAIN** Stroke Investigate - E161686 v1b AAHead Scout 32ch-head-coil CVR_pace_2.5mm_iso_v1_TEST_SCAN CVR_pace_2.5mm_iso_v1 AAHead_Scout_32ch-head-coil t1_mprage_sag_p3_iso_Munich t2_3D_FLAIR v2 t2_space_v4 PD_VIBE_v1a SWI_v6c t2_blade_dark-fluid_tra_v2b DTI_custom_v1_AP DTI_custom_v1_rev localizer **TOF** PC_Carotids (C2-C3)localizer PC_SACSF localizer vessel_scout PC_Sinus t1_fl2d_sag_localiser PC_Aqueduct AAHead_Scout_32ch-head-coil IR-SPGR_TI_600_v4 IR-SPGR_TI_1500_v4 IR-SPGR_IROff_2_v4 IR-SPGR_IROff_5_v4 IR-SPGR_IROff_12_v4 t1_vibe_sag_DCE_2mm

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\AAHead_Scout_32ch-head-coil

TA: 0:14 PM: REF Voxel size: 1.6×1.6×1.6 mmPAT: 3 Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	On
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
TE	1.37 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3.15 ms
TE	1.37 ms
Flip angle	8 deg

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Resolution - Common

FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	69 %
Phase partial Fourier	6/8
Slice partial Fourier	6/8
Trajectory	Cartesian

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	3
Ref. lines PE	24
Accel. factor 3D	1

Resolution - iPAT

Reference scan mode	Integrated	
Resolution - Filter Image		
Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

B1 filter

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
Multi-slice mode	Sequential
Series	Ascending
Concatenations	1

Geometry - AutoAlign

_	
Slab group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off

Coil Select Mode	Default

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
A >> P R >> L F >> H	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Flip angle	8 deg
Measurements	1
Time to center	6.2 s

Inline - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
	- · ·
MapIt	None
Flip angle	8 deg
1 ' "	5
Measurements	1

Inline - MapIt

Contrasts	1
TR	3.15 ms
TE	1.37 ms

Sequence - Part 1

Introduction	On
Dimension	3D
Asymmetric echo	Weak
Contrasts	1
Multi-slice mode	Sequential
Bandwidth	540 Hz/Px

Sequence - Part 2

RF pulse type	Fast
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On

Mode	Off

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\CVR_pace_2.5mm_iso_v1_TEST_SCAN

TA: 0:12 PM: REF Voxel size: 2.5×2.5×2.5 mmPAT: 2 Rel. SNR: 1.00 : epfid

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	On
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	On
Start measurements	Single measurement

Routine

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
TR	3000 ms
TE	30.0 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3000 ms
TE	30.0 ms
MTC	Off
Flip angle	90 deg
Fat suppr.	Fat sat.

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Delay in TR	0 ms
Multiple series	Off

Resolution - Common

FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
Base resolution	94
Phase resolution	100 %
Phase partial Fourier	Off
Interpolation	Off

Resolution - iPAT

Г		
ı	Accel. mode	GRAPPA
ı	Accel. factor PE	2
ı	Ref. lines PE	24

Resolution - iPAT

Reference scan mode	EPI/separate	
Resolution - Filter Image		
Distortion Corr.	Off	
Prescan Normalize	On	

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
Hamming	Off

Geometry - Common

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
TR	3000 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	Fat sat.
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

-,	•
Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlian	Head > Brain

Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	235 mm
R >> L	235 mm
F >> H	130 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Di Olimii modo	11401 01111

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	3000 ms
Concatenations	1

BOLD

GLM Statistics	Off
Dynamic t-maps	Off
Ignore meas. at start	0
Ignore after transition	0
Model transition states	On
Temp. highpass filter	On
Threshold	4.00
Paradigm size	3
Meas[1]	Baseline
Meas[2]	Baseline
Meas[3]	Active
Motion correction	Off
Spatial filter	Off
Measurements	1
Delay in TR	0 ms
Multiple series	Off

Sequence - Part 1

Introduction	Off
Multi-slice mode	Interleaved
Free echo spacing	Off
Echo spacing	0.52 ms
Bandwidth	2312 Hz/Px

Sequence - Part 2

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Performance

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\CVR_pace_2.5mm_iso_v1

TA: 12:30 PM: REF Voxel size: 2.5×2.5×2.5 mmPAT: 2 Rel. SNR: 1.00 : epfid

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	On
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	On
Start measurements	Single measurement

Routine

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
TR	3000 ms
TE	30.0 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3000 ms	
TE MTC	30.0 ms	
MTC	Off	
Flip angle	90 deg	
Fat suppr.	Fat sat.	

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	247
Delay in TR	0 ms
Multiple series	Off

Resolution - Common

FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
Base resolution	94
Phase resolution	100 %
Phase partial Fourier	Off
Interpolation	Off

Resolution - iPAT

ſ	Accel. mode	GRAPPA
	Accel. factor PE	2
	Ref. lines PE	24

Resolution - iPAT

Reference scan mode	EPI/separate
Resolution - Filter Image	
Distortion Corr.	Off
Prescan Normalize	On

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
Hamming	Off

Geometry - Common

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
FoV read	235 mm
FoV phase	100.0 %
Slice thickness	2.5 mm
TR	3000 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	Fat sat.
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

-,	
Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlian	Head > Brain

ř.	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	235 mm
R >> L	235 mm
F >> H	130 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	3000 ms
Concatenations	1

BOLD

GLM Statistics	Off
Dynamic t-maps	Off
Ignore meas. at start	0
Ignore after transition	0
Model transition states	On
Temp. highpass filter	On
Threshold	4.00
Paradigm size	3
Meas[1]	Baseline
Meas[2]	Baseline
Meas[3]	Active
Motion correction	Off
Spatial filter	Off
Measurements	247
Delay in TR	0 ms
Multiple series	Off

Sequence - Part 1

Introduction	Off
Multi-slice mode	Interleaved
Free echo spacing	Off
Echo spacing	0.52 ms
Bandwidth	2312 Hz/Px

Sequence - Part 2

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Performance

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\AAHead_Scout_32ch-head-coil

TA: 0:14 PM: REF Voxel size: 1.6×1.6×1.6 mmPAT: 3 Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	On
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
TE	1.37 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3.15 ms
TE	1.37 ms
Flip angle	8 deg

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Resolution - Common

FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	69 %
Phase partial Fourier	6/8
Slice partial Fourier	6/8
Trajectory	Cartesian

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	3
Ref. lines PE	24
Accel. factor 3D	1

Resolution - iPAT

Reference scan mode	Integrated	
Resolution - Filter Image		

Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
Multi-slice mode	Sequential
Series	Ascending
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off

Coil Select Mode	Default

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
A >> P R >> L F >> H	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Flip angle	8 deg
Measurements	1
Time to center	6.2 s

Inline - Inline

Subtract	Off	
Measurements	1	
StdDev	Off	
Save original images	On	

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	8 deg
Measurements	1

Inline - MapIt

Contrasts	1
TR	3.15 ms
TE	1.37 ms

Sequence - Part 1

Introduction	On
Dimension	3D
Asymmetric echo	Weak
Contrasts	1
Multi-slice mode	Sequential
Bandwidth	540 Hz/Px

Sequence - Part 2

RF pulse type	Fast
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On

Mode	Off

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t1_mprage_sag_p3_iso_Munich

TA: 3:45 PM: FIX Voxel size: 1.0×1.0×1.0 mmPAT: 3 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
TR	2500.0 ms
TE	4.37 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	2500.0 ms
TE	4.37 ms
Magn. preparation	Non-sel. IR
TI	1100 ms
Flip angle	7 deg
Fat suppr.	Water excit. fast
Fat suppr. Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
Base resolution	256
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	7/8
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA	
Accel. factor PE	3	
Ref. lines PE	24	
Accel. factor 3D	1	
Reference scan mode	Integrated	

Resolution - Filter Image

Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
TR	2500.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Basis
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Sagittal
Rotation	0.00 deg
A >> P	256 mm
A >> P F >> H	256 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	2500.0 ms
Concatenations	1

Physio - Cardiac

Magn. preparation	Non-sel. IR
TI	1100 ms
Fat suppr.	Water excit. fast
Dark blood	Off
FoV read	256 mm
FoV phase	100.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	7 deg
Measurements	1
TR	2500.0 ms
TE	4.37 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	11.1 ms
Bandwidth	140 Hz/Px

Sequence - Part 2

RF pulse type	Fast
Gradient mode	Fast
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	Off
Turbo factor	192

Mode	Off
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t2_3D_FLAIR v2

TA: 5:57 PM: REF Voxel size: 1.0×1.0×1.0 mmPAT: 3 Rel. SNR: 1.00 : spcir

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
TR	5000 ms
TE	388 ms
Averages	1.0
Concatenations	1
Filter	Raw filter, Prescan
	Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	5000 ms
TE	388 ms
MTC	Off
Magn. preparation	Non-sel. T2-IR
TI 1	1800 ms
Fat suppr.	Fat sat.
Fat sat. mode	Strong
Blood suppr.	Off
Restore magn.	Off

Contrast - Dynamic

Averages	1.0
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
Base resolution	256
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Allowed
Slice partial Fourier	Off

Resolution - Common

Interpolation	Off	

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	3
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	On
Elliptical filter	Off

Geometry - Common

•	
Slab group	1
Slabs	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.00 mm
TR	5000 ms
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Fat suppr.	Fat sat.
Fat sat. mode	Strong
Restore magn.	Off
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off	
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Geometry - Tim Planning Suite

Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Basis
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Sagittal
Rotation	0.00 deg
A >> P F >> H	256 mm
F >> H	256 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
Trigger delay	0 ms
TR	5000 ms
Concatenations	1

Physio - Cardiac

Magn. preparation	Non-sel. T2-IR
TI 1	1800 ms
Fat suppr.	Fat sat.
Dark blood	Off
FoV read	256 mm
FoV phase	100.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag MIP-Cor MIP-Tra MIP-Time	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Sequence - Part 1

Introduction	On
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Flow comp.	No
Echo spacing	3.46 ms
Adiabatic-mode	Off
Bandwidth	751 Hz/Px

Sequence - Part 2

Echo train duration	865 ms
RF pulse type	Normal
Gradient mode	Fast
Excitation	Non-sel.
Flip angle mode	T2 var
Turbo factor	278

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t2_space_v4

TA: 3:42 PM: REF Voxel size: 0.9×0.9×0.9 mmPAT: 4 Rel. SNR: 1.00 : spcR

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	18.2 %
Slices per slab	176
FoV read	240 mm
FoV phase	100.0 %
Slice thickness	0.90 mm
TR	3200 ms
TE	408 ms
Averages	1.4
Concatenations	1
Filter	Raw filter, Prescan
	Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3200 ms
TE	408 ms
MTC	Off
Magn. preparation	None
Fat suppr.	None
Blood suppr.	Off
Restore magn.	On

Contrast - Dynamic

Averages	1.4
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	240 mm
FoV phase	100.0 %
Slice thickness	0.90 mm
Base resolution	256
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Allowed
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA	
Accel. factor PE	2	
Ref. lines PE	24	
Accel. factor 3D	2	
Ref. lines 3D	24	
Reference scan mode	Integrated	

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	On
Elliptical filter	Off

Geometry - Common

Slab group	1
Slabs	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
Slice oversampling	18.2 %
Slices per slab	176
FoV read	240 mm
FoV phase	100.0 %
Slice thickness	0.90 mm
TR	3200 ms
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Initial Position	Isocenter
L	0.0 mm
Р	0.0 mm
Н	0.0 mm
Initial Rotation	90.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	None
Restore magn.	On
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Performance
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slab-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
Trigger delay	0 ms
TR	3200 ms
Concatenations	1

Physio - Cardiac

Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	240 mm
FoV phase	100.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract Off

Inline - Common

Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Sequence - Part 1

Introduction	On
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Flow comp.	No
Echo spacing	3.61 ms
Adiabatic-mode	Off
Bandwidth	723 Hz/Px

Sequence - Part 2

Echo train duration	910 ms
RF pulse type	Normal
Gradient mode	Fast
Excitation	Slab-sel.
Flip angle mode	T2 var
Turbo factor	282

Allowed delay	30 s	
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\PD_VIBE_v1a

TA: 1:57 PM: FIX Voxel size: 1.0×1.0×1.0 mmPAT: 3 Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	Off
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	L0.0 A5.1 F8.3 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.0 mm
TR	6.04 ms
TE	2.44 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP;SP1

Contrast - Common

TR	6.04 ms
TE	2.44 ms
Flip angle	2.0 deg
Fat suppr. Water suppr.	None
Water suppr.	None
Dixon	Off

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.0 mm
Base resolution	256
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Trajectory	Cartesian
View sharing	Off

Resolution - Common

Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	3
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
POCS	Off

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	L0.0 A5.1 F8.3 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	192
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	1.0 mm
TR	6.04 ms
Multi-slice mode	Sequential
Series	Ascending
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	L0.0 A5.1 F8.3 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	Head > Basis
Initial Position	L0.0 A5.1 F8.3
R	0.0 mm
A	5.1 mm
F	8.3 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Fat suppr.	None
Water suppr.	None
Dixon	Off
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol Off

Geometry - Tim Planning Suite

Tal	ole position	Н
Tal	ole position	0 mm
Inli	ne Composing	Off

System - Miscellaneous

Positioning mode	FIX
	, .
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
Coil Focus	Flat
AutoAlign	Head > Basis
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.0 A5.1 F8.3 mm
Orientation	Sagittal
Rotation	0.00 deg
A >> P F >> H R >> L	256 mm
F >> H	256 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

View sharing	Off	
Flip angle	2.0 deg	
Measurements	1	
Burn time-to-center	Off	
Temporal interpolation	1	
3D centric reordering	Off	
Time to center	58.9 s	

Inline - Inline

Subtract	Off
Measurements	1
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Sag MIP-Cor MIP-Tra MIP-Time	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Soft Tissue

Wash - In	Off	
Wash - Out	Off	
TTP	Off	
PEI	Off	
TTP PEI MIP - time	Off	
Measurements	1	

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	2.0 deg
Measurements	1
Contrasts	1
TR	6.04 ms
TE	2.44 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Asymmetric echo	Weak
Contrasts	1
Optimization	Min. TE
Multi-slice mode	Sequential
Bandwidth	200 Hz/Px

Sequence - Part 2

RF pulse type	Fast
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On

Mode	Off
Allowed delay	0 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\SWI_v6c

TA: 4:02 PM: FIX Voxel size: 0.6×0.6×3.0 mmPAT: 2 Rel. SNR: 1.00 : swi_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	L0.0 A2.3 H2.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	23.1 %
Slices per slab	52
FoV read	240 mm
FoV phase	81.3 %
Slice thickness	3.00 mm
TR	28.0 ms
TE	20.00 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	28.0 ms
TE	20.00 ms
MTC	Off
Magn. preparation	None
Flip angle	9 deg
Fat suppr.	None
Water suppr.	None
SWI	On

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magn./Phase
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	240 mm
FoV phase	81.3 %
Slice thickness	3.00 mm
Base resolution	384
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off

Resolution - Common

Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	L0.0 A2.3 H2.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
Slice oversampling	23.1 %
Slices per slab	52
FoV read	240 mm
FoV phase	81.3 %
Slice thickness	3.00 mm
TR	28.0 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	L0.0 A2.3 H2.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Initial Position	L0.0 A2.3 H2.2
L	0.0 mm
Α	2.3 mm
Н	2.2 mm
Initial Rotation	89.61 deg
Initial Orientation	Transversal

Geometry - Saturation

Saturation mode	Standard
Fat suppr.	None
Water suppr.	None
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н

Geometry - Tim Planning Suite

Table position	0 mm
Inline Composing	Off

System - Miscellaneous

D 20 1	FIV
Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.0 A2.3 H2.2 mm
Orientation	Transversal
Rotation	89.61 deg
R >> L	195 mm
A >> P	240 mm
F >> H	156 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slab-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	28.0 ms
Concatenations	1
Segments	1

Physio - Cardiac

Tagging	None
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	240 mm
FoV phase	81.3 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off	
MIP-Cor MIP-Tra MIP-Time	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	1

Inline - Composing

Inline Composing	Off	
Distortion Corr.	Off	

Inline - MapIt

Save original images	On
Maplt	None
Flip angle	9 deg
Measurements	1
Contrasts	1
TR	28.0 ms
TE	20.00 ms

Sequence - Part 1

Introduction	On
Dimension	3D
Elliptical scanning	On
Phase stabilisation	Off
Asymmetric echo	Off
Contrasts	1
Flow comp.	Yes
Multi-slice mode	Interleaved
Bandwidth	120 Hz/Px

Sequence - Part 2

Segments	1
Acoustic noise reduction	None
RF pulse type	Fast
Gradient mode	Normal
Excitation	Slab-sel.
RF spoiling	On

Mode	Off
Allowed delay	30 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t2_blade_dark-fluid_tra_v2b

TA: 3:22 PM: REF Voxel size: 0.9×0.9×3.0 mmPAT: 2 Rel. SNR: 1.00 : tirB_rr

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Phase oversampling	0.0 %
FoV read	240 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
TR	10000.0 ms
TE	124 ms
Averages	1
Concatenations	2
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

Contrast - Common		
TR	10000.0 ms	
TE	124 ms	
TD	0.0 ms	
MTC	Off	
Magn. preparation	Slice-sel. IR	
TI	2606 ms	
Flip angle	130 deg	
Fat suppr.	Fat sat.	
Fat sat. mode	Strong	
Water suppr.	None	
Restore magn.	Off	
Freeze suppressed tissue	On	

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	240 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
Base resolution	256
BLADE coverage	100.0 %
Trajectory	BLADE

Resolution - Common

Interpolation	Off
•	

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	8
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Slice group	1
Slices	52
Dist. factor	0 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
FoV read	240 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
TR	10000.0 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	2

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	Head > Brain
Initial Position	Isocenter
L	0.0 mm
Р	0.0 mm
Н	0.0 mm
Initial Rotation	90.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	Fat sat.
Fat sat. mode	Strong
Water suppr.	None
Restore magn.	Off
Special sat.	Parallel F
Gap	10 mm
Thickness	70 mm

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	90.00 deg
R >> L	240 mm
R >> L A >> P F >> H	240 mm
F >> H	156 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	10000.0 ms
Concatenations	2

Physio - Cardiac

Magn. preparation	Slice-sel. IR
ті	2606 ms
Fat suppr.	Fat sat.
Dark blood	Off
FoV read	240 mm
FoV phase	100.0 %
BLADE coverage	100.0 %
Trajectory	BLADE

Physio - PACE

Resp. control	Off
Concatenations	2

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag MIP-Cor MIP-Tra MIP-Time	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off	
Distortion Corr.	Off	

Sequence - Part 1

Introduction	On
Dimension	2D
Compensate T2 decay	Off
Contrasts	1
Flow comp.	Read
Multi-slice mode	Interleaved
Free echo spacing	Off
Echo spacing	8.26 ms
Bandwidth	362 Hz/Px

Sequence - Part 2

Define	Turbo factor
Echo trains per slice	9
Phase correction	Automatic
Acoustic noise reduction	None
RF pulse type	Low SAR
Gradient mode	Fast
Hyperecho	Off
WARP	Off
Motion correction	On
Red. EC sensitivity	Off
Turbo factor	28

Mode	Min flip angle
Min flip angle	130 deg
Allowed delay	30 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\DTI_custom_v1_AP

TA: 11:12 PM: REF Voxel size: 2.0×2.0×2.0 mmPAT: 4 Rel. SNR: 1.00 : epse

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	72
Dist. factor	0 %
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	
Phase oversampling	0 %
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
TR	4300 ms
TE	74.0 ms
Averages	1
Concatenations	1
Filter	Raw filter, Prescan
	Normalize
Coil elements	HEA;HEP

Contrast - Common

300 ms
1.0 ms
ff
one
at sat.
rong

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Delay in TR	0 ms
Multiple series	Off

Resolution - Common

FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
Base resolution	128
Phase resolution	100 %
Phase partial Fourier	7/8
Interpolation	Off

Resolution - iPAT

Accel. mode Slice accel.

Resolution - iPAT

Accel. factor PE	2
Ref. lines PE	40
Accel. factor slice	2
Reference scan mode	EPI/separate

Resolution - Filter Image

Distortion Corr.	Off
Prescan Normalize	On
Dynamic Field Corr.	Off

Resolution - Filter Rawdata

Raw filter	On
Elliptical filter	Off

Geometry - Common

Slice group	1
Slices	72
Dist. factor	0 %
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	A >> P
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
TR	4300 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	
Initial Position	L0.7 P3.0 H31.9
L	0.7 mm
P	3.0 mm
Н	31.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	Fat sat.
Fat sat. mode	Strong
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm

MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	H >> F
Coil Combine Mode	Adaptive Combine
Matrix Optimization	Performance
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Rotation	0.00 deg
A >> P	256 mm
R >> L	256 mm
F >> H	144 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Standard

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	4300 ms
Concatenations	1

Physio - PACE

Resp. control	Off
Concatenations	1

Diff - Neuro

Diffusion mode	Free
Diff. directions	151
Diffusion Scheme	Monopolar
Diff. weightings	1
b-value	2000 s/mm ²
b-value	1
Diff. weighted images	On
Trace weighted images	Off
ADC maps	Off
FA maps	Off
Mosaic	Off
Tensor	Off
Noise level	40

Diff - Body

Diffusion mode	Free
Diff. directions	151
Diffusion Scheme	Monopolar
Diff. weightings	1
b-value	2000 s/mm ²
b-value	1
Diff. weighted images	On
Trace weighted images	Off
ADC maps	Off
Exponential ADC Maps	Off
FA maps	Off
Invert Gray Scale	Off
Calculated Image	Off
b-Value >=	0 s/mm²
Noise level	40

Diff - Composing

Inline Composing	Off	
Distortion Corr.	Off	

Sequence - Part 1

Introduction	On
Optimization	None
Multi-slice mode	Interleaved
Free echo spacing	Off
Echo spacing	0.78 ms
Bandwidth	1446 Hz/Px

Sequence - Part 2

EPI factor	128
RF pulse type	Normal
Gradient mode	Performance
Excitation	Standard

Sequence - pTX Pulses

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\DTI_custom_v1_rev

TA: 0:35 PM: REF Voxel size: 2.0×2.0×2.0 mmPAT: 4 Rel. SNR: 1.00 : epse

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	72
Dist. factor	0 %
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	P >> A
AutoAlign	
Phase oversampling	0 %
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
TR	4300 ms
TE	74.0 ms
Concatenations	1
Filter	Raw filter, Prescan
	Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	4300 ms
TE	74.0 ms
MTC	Off
Magn. preparation	None
Fat suppr.	Fat sat.
Fat sat. mode	Strong

Contrast - Dynamic

Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Delay in TR	0 ms
Multiple series	Off

Resolution - Common

FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
Base resolution	128
Phase resolution	100 %
Phase partial Fourier	7/8
Interpolation	Off

Resolution - iPAT

Accel. mode	Slice accel.
Accel. factor PE	2
Ref. lines PE	40

Resolution - iPAT

Accel. factor slice	2
Reference scan mode	EPI/separate

Resolution - Filter Image

Distortion Corr.	Off
Prescan Normalize	On
Dynamic Field Corr.	Off

Resolution - Filter Rawdata

Raw filter	On	
Elliptical filter	Off	

Geometry - Common

Slice group	1
Slices	72
Dist. factor	0 %
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	P >> A
FoV read	256 mm
FoV phase	100.0 %
Slice thickness	2.0 mm
TR	4300 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Phase enc. dir.	P >> A
AutoAlign	
Initial Position	L0.7 P3.0 H31.9
L	0.7 mm
Р	3.0 mm
Н	31.9 mm
Initial Rotation	180.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	Fat sat.
Fat sat. mode	Strong
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	REF
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L

Coronal	A >> P
Transversal	H >> F
Coil Combine Mode	Adaptive Combine
Matrix Optimization	Performance
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.7 P3.0 H31.9 mm
Orientation	Transversal
Rotation	180.00 deg
A >> P	256 mm
A >> P R >> L F >> H	256 mm
	144 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Standard

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	4300 ms
Concatenations	1

Physio - PACE

Resp. control	Off	
Concatenations	1	

Diff - Neuro

Diffusion mode	MDDW
Diff. directions	6
Diffusion Scheme	Monopolar
Diff. weightings	1
b-value	0 s/mm²
b-value	3
Diff. weighted images	On
Trace weighted images	Off
ADC maps	Off
FA maps	Off
Mosaic	Off
Tensor	Off
Noise level	40

Diff - Body

Diffusion mode	MDDW
Dillusion mode	IVIDDVV

Diff - Body

Diff. directions	6
Diffusion Scheme	Monopolar
Diff. weightings	1
b-value	0 s/mm ²
b-value	3
Diff. weighted images	On
Trace weighted images	Off
ADC maps	Off
Exponential ADC Maps	Off
FA maps	Off
Invert Gray Scale	Off
Calculated Image	Off
b-Value >=	0 s/mm ²
Noise level	40

Diff - Composing

Inline Composing	Off
Distortion Corr.	Off

Sequence - Part 1

Introduction	On
Optimization	None
Multi-slice mode	Interleaved
Free echo spacing	Off
Echo spacing	0.78 ms
Bandwidth	1446 Hz/Px

Sequence - Part 2

EPI factor	128
RF pulse type	Normal
Gradient mode	Performance
Excitation	Standard

Sequence - pTX Pulses

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\localizer

TA: 0:11 PM: ISO Voxel size: 1.2×1.2×10.0 mmPAT: Off Rel. SNR: 1.00 : fl_rr

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
· ·	Off
Wait for user to start	Oli
Start measurements	Single measurement

Routine

Slice group	1
Slices	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice group	2
Slices	5
Dist. factor	300 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Slices	2
Dist. factor	70 %
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	300 mm
FoV phase	100.0 %
Slice thickness	10.0 mm
TR	9.7 ms
TE	3.80 ms
Averages	1
Concatenations	8
Filter	Distortion Corr.(2D),
	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	9.7 ms
TE	3.80 ms
TD	0 ms
MTC	Off
Magn. preparation	None
Flip angle	40 deg
Fat suppr.	None
Water suppr.	None
SWI	Off

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Contrast - Dynamic

Multiple series

Resolution - Common		
FoV read	300 mm	
FoV phase	100.0 %	
Slice thickness	10.0 mm	
Base resolution	256	
Phase resolution	50 %	
Phase partial Fourier	Off	
Interpolation	Off	

Each measurement

Resolution - iPAT

ĺ	PAT mode	None
	1 7 (1 111000	140110

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Geometry - Common	
Slice group	1
Slices	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice group	2
Slices	5
Dist. factor	300 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Slices	2
Dist. factor	70 %
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
FoV read	300 mm
FoV phase	100.0 %
Slice thickness	10.0 mm
TR	9.7 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	8

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P

Geometry - AutoAlign

2
Isocenter
Transversal
A >> P
3
L0.0 A30.0 H0.0 mm
Coronal
R >> L
Isocenter
0.0 mm
0.0 mm
0.0 mm
0.00 deg
Sagittal

Geometry - Saturation

Saturation mode	Standard
Fat suppr.	None
Water suppr.	None
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

_ 	
Positioning mode	ISO
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	Default

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

ř	
Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	9.7 ms
Concatenations	8
Segments	1

Physio - Cardiac

Tagging	None
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	300 mm
FoV phase	100.0 %
Phase resolution	50 %

Physio - PACE

Resp. control	Off
Concatenations	8

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	1

Inline - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	40 deg
Measurements	1
Contrasts	1
TR	9.7 ms
TE	3.80 ms

SIEMENS MAGNETOM Prisma

Sequence - Part 1

Introduction	On
Dimension	2D
Phase stabilisation	Off
Asymmetric echo	Allowed
Contrasts	1
Flow comp.	Read
Multi-slice mode	Sequential
Bandwidth	320 Hz/Px

Sequence - Part 2

Segments	1
Acoustic noise reduction	None
RF pulse type	Normal
Gradient mode	Normal
Excitation	Slice-sel.
RF spoiling	On

Mode	Off	
Allowed delay	0 s	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\TOF

TA: 2:45 PM: ISO Voxel size: 0.5×0.5×0.8 mmPAT: 2 Rel. SNR: 1.00 : fl_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	6
Dist. factor	-20.83 %
Position	R4.6 A0.9 F54.5 mm
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	
Phase oversampling	0 %
Slice oversampling	16.7 %
Slices per slab	24
FoV read	200 mm
FoV phase	75.5 %
Slice thickness	0.80 mm
TR	20.0 ms
TE	3.51 ms
Averages	1
Concatenations	6
Filter	Distortion Corr.(2D), Prescan Normalize,
	Elliptical filter
Coil elements	HE3,4;NE1,2;SP1

Contrast - Common

TR TE	20.0 ms
TE	3.51 ms
TD	0.000 ms
MTC	Off
Flip angle	20 deg
Fat suppr. Water suppr.	None
Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Resolution - Common

FoV read	200 mm
FoV phase	75.5 %
Slice thickness	0.80 mm
Base resolution	384
Phase resolution	70 %
Slice resolution	50 %
Phase partial Fourier	7/8
Slice partial Fourier	7/8
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	On
POCS	Off

Geometry - Common

•	
Slab group	1
Slabs	6
Dist. factor	-20.83 %
Position	R4.6 A0.9 F54.5 mm
Orientation	Transversal
Phase enc. dir.	A >> P
Slice oversampling	16.7 %
Slices per slab	24
FoV read	200 mm
FoV phase	75.5 %
Slice thickness	0.80 mm
TR	20.0 ms
Multi-slice mode	Sequential
Series	Descending
Concatenations	6

Geometry - AutoAlign

Slab group	1
Position	R4.6 A0.9 F54.5 mm
Orientation	Transversal
Phase enc. dir.	A >> P
AutoAlign	
Initial Position	R4.6 A0.9 F54.5
R	4.6 mm
A	0.9 mm
F	54.5 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Fat suppr.	None
Fat suppr. Water suppr.	None
Special sat.	Tracking H
Gap	10 mm
Gap Thickness	40 mm

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
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Geometry - Tim Planning Suite

Table position	F
Table position	55 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	ISO
Table position	F
Table position	55 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Save uncombined	Off
Matrix Optimization	Off
Coil Focus	Flat
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	20.0 ms
Concatenations	6

Physio - Cardiac

Fat suppr.	None
Dark blood	Off
FoV read	200 mm
FoV phase	75.5 %
Phase resolution	70 %

Angio - Common

TONE ramp	80 %
Flow direction	F >> H

Angio - Common

Flip angle	20 deg
MTC	Off
Measurements	1
3D centric reordering	On

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Angio - MIP

MIP-Sag	On
MIP-Cor	On
MIP-Tra	Off
MIP-Sag MIP-Cor MIP-Tra MIP-Time	Off
Save original images	On

Angio - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Sequence - Part 1

Introduction	On
Dimension	3D
Elliptical scanning	Off
Asymmetric echo	Allowed
Contrasts	1
Flow comp.	Yes
Multi-slice mode	Sequential
Bandwidth	250 Hz/Px

Sequence - Part 2

Gradient mode	Fast
RF spoiling	On

Mode Off

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\PC_Carotids

TA: 1:39 PM: ISO Voxel size: 1.0×1.0×5.0 mmPAT: 2 Rel. SNR: 1.00 : fl_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	1
Dist. factor	20 %
Position	R4.9 A18.6 F16.5 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	19.60 ms
TE	5.82 ms
Averages	1
Concatenations	1
Filter	Distortion Corr.(2D),
	Prescan Normalize
Coil elements	HE3,4;NE1,2

Contrast - Common

TR	19.60 ms
TE	5.82 ms
TD	0 ms
Flip angle	12 deg
Wrap-up Magn.	None

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Off

Resolution - Common

FoV read	160 mm	
FoV phase	100.0 %	
Slice thickness	5.0 mm	
Base resolution	160	
Phase resolution	100 %	
Phase partial Fourier	Off	
Trajectory	Cartesian	
View sharing	Off	
Interpolation	Off	

Resolution - iPAT

PAT mode	GRAPPA
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Resolution - iPAT

Accel. factor PE	2
Ref. lines PE	32
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off	
Distortion Corr.	On	
Mode	2D	
Unfiltered images	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
POCS	Off

Geometry - Common

Slice group	1
Slices	1
Dist. factor	20 %
Position	R4.9 A18.6 F16.5 mm
Orientation	Transversal
Phase enc. dir.	R >> L
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	19.60 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	R4.9 A18.6 F16.5 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	R4.9 A18.6 F16.5
R	4.9 mm
A	18.6 mm
F	16.5 mm
Initial Rotation	90.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Wrap-up Magn.	None
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	F
Table position	17 mm
Inline Composing	Off

Positioning mode	ISO
Table position	F
Table position	17 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

R4.9 A18.6 F16.5 mm
Transversal
90.00 deg
160 mm
160 mm
5 mm
Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	Pulse/Retro
Average cycle	No Signal ms
Average cycle	No Signal ms
Calculated phases	32
TR	19.60 ms
Concatenations	1
Segments	1
Arrhythmia detection	None

Physio - PACE

Resp. control	Off
Concatenations	1

Angio - Common

7.11.9.0 CO.III.I.O.I.	
Flow mode	Single dir.
Encodings	1
Velocity enc.	70 cm/s
Direction	Through plane
Rephased images	On
Magnitude images	On

Angio - Common

Magnitude sum	Off	
Phase images	On	

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Angio - Cardiac

Inline Evaluation	Off
TE	5.82 ms
TR	19.60 ms
Save original images	On

Angio - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Angio - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Sequence - Part 1

Introduction	On
Dimension	2D
Reordering	Linear
Asymmetric echo	Weak
Flow comp.	Yes
Optimization	Min. TE TR
Multi-slice mode	Sequential
Echo spacing	9.8 ms
Sequence type	Gre
Bandwidth	202 Hz/Px

Sequence - Part 2

Define	Segments
Segments	1
RF pulse type	Normal
Gradient mode	Fast
Excitation	Slice-sel.
Flip angle mode	Constant
RF spoiling	On
Phase Enc. Rewinder	On
Cine	On

Mode	Off
Allowed delay	0 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\localizer (C2-C3)

TA: 0:11 PM: ISO Voxel size: 1.2×1.2×10.0 mmPAT: Off Rel. SNR: 1.00 : fl_rr

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group 1 Slices 1 Dist. factor 20 % Position Isocenter Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize Coil elements		
Dist. factor 20 % Position Isocenter Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slice group	1
Position Orientation Phase enc. dir. Sagittal Phase enc. dir. Slice group Slices Dist. factor Orientation Phase enc. dir. Slice group Slice group Slice group Slice group Slices Dist. factor Orientation Phase enc. dir. Slices Dist. factor Position Coronal Phase enc. dir. A >> P Slice group Slices Dist. factor Fosition Coronal Phase enc. dir. AutoAlign Phase oversampling FoV read Slice thickness To mm Transversal Do mm Transversal Do mm Transversal Do mm Transversal Distortion Corr.(2D), Prescan Normalize	Slices	1
Orientation Phase enc. dir. Sagittal Phase enc. dir. A >> P Slice group 2 Slices Dist. factor Orientation Phase enc. dir. Slice group 3 Slices Dist. factor Position Unientation Phase enc. dir. Slices Dist. factor Position Coronal Phase enc. dir. A >> P Slices Dist. factor Position Orientation Coronal Phase enc. dir. AutoAlign Phase oversampling FoV read Slice thickness To.0 mm TR TE 3.80 ms Averages Concatenations Filter Distortion Corr.(2D), Prescan Normalize	Dist. factor	20 %
Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Position	Isocenter
Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Orientation	Sagittal
Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Phase enc. dir.	A >> P
Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slice group	2
Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slices	5
Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Dist. factor	300 %
Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Position	Isocenter
Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Orientation	Transversal
Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Phase enc. dir.	A >> P
Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slice group	3
Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slices	2
Orientation Coronal Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Dist. factor	70 %
Phase enc. dir. R >> L AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Position	L0.0 A30.0 H0.0 mm
AutoAlign Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Orientation	Coronal
Phase oversampling 0 % FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Phase enc. dir.	R >> L
FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	AutoAlign	
FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Phase oversampling	0 %
Slice thickness 10.0 mm TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	FoV read	300 mm
TR 9.7 ms TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	FoV phase	100.0 %
TE 3.80 ms Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	Slice thickness	10.0 mm
Averages 1 Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	TR	9.7 ms
Concatenations 8 Filter Distortion Corr.(2D), Prescan Normalize	TE	3.80 ms
Filter Distortion Corr.(2D), Prescan Normalize	Averages	1
Prescan Normalize	Concatenations	8
	Filter	
Coil elements HE1-4;NE1,2		
	Coil elements	HE1-4;NE1,2

Contrast - Common

TR	9.7 ms	
TE	3.80 ms	
TD	0 ms	
MTC	Off	
Magn. preparation	None	
Flip angle	40 deg	
Fat suppr.	None	
Water suppr.	None	
SWI	Off	

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Contrast - Dynamic

Multiple series	Each measurement	
Resolution - Common		
FoV read	300 mm	
FoV phase	100.0 %	
Slice thickness	10.0 mm	
Base resolution	256	
Phase resolution	50 %	
Phase partial Fourier	Off	
Interpolation	Off	

Resolution - iPAT

ĺ	PAT mode	None
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Resolution - Filter Image

Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Geometry - Common	
Slice group	1
Slices	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice group	2
Slices	5
Dist. factor	300 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Slices	2
Dist. factor	70 %
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
FoV read	300 mm
FoV phase	100.0 %
Slice thickness	10.0 mm
TR	9.7 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	8

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P

Geometry - AutoAlign

Slice group	2
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	Isocenter
L	0.0 mm
Р	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Saturation mode	Standard
Fat suppr.	None
Water suppr.	None
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	ISO
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	9.7 ms
Concatenations	8
Segments	1

Physio - Cardiac

Tagging	None
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	300 mm
FoV phase	100.0 %
Phase resolution	50 %

Physio - PACE

Resp. control	Off
Concatenations	8

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	1

Inline - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	40 deg
Measurements	1
Contrasts	1
TR	9.7 ms
TF	3.80 ms

SIEMENS MAGNETOM Prisma

Sequence - Part 1

Introduction	On
Dimension	2D
Phase stabilisation	Off
Asymmetric echo	Allowed
Contrasts	1
Flow comp.	Read
Multi-slice mode	Sequential
Bandwidth	320 Hz/Px

Sequence - Part 2

Segments	1
Acoustic noise reduction	None
RF pulse type	Normal
Gradient mode	Normal
Excitation	Slice-sel.
RF spoiling	On

Mode	Off
Allowed delay	0 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\PC_SACSF

TA: 1:55 PM: ISO Voxel size: 0.8×0.8×5.0 mmPAT: 2 Rel. SNR: 1.00 : fl_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	1
Dist. factor	20 %
Position	L0.1 A7.3 F11.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	25.18 ms
TE	8.45 ms
Averages	1
Concatenations	1
Filter	Distortion Corr.(2D),
	Prescan Normalize, Elliptical filter
Coil elements	HE3,4;NE1,2

Contrast - Common

TR	25.18 ms
TE	8.45 ms
TD	0 ms
Flip angle	12 deg
Wrap-up Magn.	None

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Off

Resolution - Common

FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
Base resolution	192
Phase resolution	100 %
Phase partial Fourier	Off
Trajectory	Cartesian
View sharing	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	32
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	On	
POCS	Off	

Geometry - Common

Slice group	1
Slices	1
Dist. factor	20 %
Position	L0.1 A7.3 F11.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	25.18 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slice group	1
Position	L0.1 A7.3 F11.2 mm
Orientation	Transversal
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	L0.1 A7.3 F11.2
L	0.1 mm
Α	7.3 mm
F	11.2 mm
Initial Rotation	90.00 deg
Initial Orientation	Transversal

Geometry - Saturation

Wrap-up Magn.	None
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	F
Table position	11 mm
Inline Composing	Off

Positioning mode	ISO
Table position	F
Table position	11 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.1 A7.3 F11.2 mm
Orientation	Transversal
Rotation	90.00 deg
R >> L	160 mm
A >> P	160 mm
R >> L A >> P F >> H	5 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	Pulse/Retro
Average cycle	No Signal ms
Average cycle	No Signal ms
Calculated phases	32
TR	25.18 ms
Concatenations	1
Segments	1
Arrhythmia detection	None

Physio - PACE

Resp. control	Off
Concatenations	1

Angio - Common

g.c - c	
Flow mode	Single dir.
Encodings	1
Velocity enc.	6 cm/s
Direction	Through plane
Rephased images	On
Magnitude images	On

Angio - Common

Magnitude sum	Off	
Phase images	On	

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Angio - Cardiac

Inline Evaluation	Off
TE	8.45 ms
TR	25.18 ms
Save original images	On

Angio - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Angio - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Sequence - Part 1

Introduction	On
Dimension	2D
Reordering	Linear
Asymmetric echo	Weak
Flow comp.	Yes
Optimization	Min. TE TR
Multi-slice mode	Sequential
Echo spacing	12.6 ms
Sequence type	Gre
Bandwidth	200 Hz/Px

Sequence - Part 2

Define	Segments
Segments	1
RF pulse type	Normal
Gradient mode	Fast*
Excitation	Slice-sel.
Flip angle mode	Constant
RF spoiling	On
Phase Enc. Rewinder	On
Cine	On

Mode	Off
Allowed delay	0 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\localizer

TA: 0:11 PM: ISO Voxel size: 1.2×1.2×10.0 mmPAT: Off Rel. SNR: 1.00 : fl_rr

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice group	2
Slices	5
Dist. factor	300 %
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Slices	2
Dist. factor	70 %
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	300 mm
FoV phase	100.0 %
Slice thickness	10.0 mm
TR	9.7 ms
TE	3.80 ms
Averages	1
Concatenations	8
Filter	Distortion Corr.(2D),
	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	9.7 ms
TE	3.80 ms
TD	0 ms
MTC	Off
Magn. preparation	None
Flip angle	40 deg
Fat suppr.	None
Water suppr.	None
SWI	Off

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Contrast - Dynamic

Multiple series

Resolution - Common		
FoV read	300 mm	
FoV phase	100.0 %	
Slice thickness	10.0 mm	
Base resolution	256	
Phase resolution	50 %	
Phase partial Fourier	Off	
Interpolation	Off	

Each measurement

Resolution - iPAT

ĺ	PAT mode	None
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Resolution - Filter Image

Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Slice group 1 Slices 1 Dist. factor 20 % Position Isocenter Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved Concatenations 8		
Dist. factor 20 % Position Isocenter Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slice group	1
Position Isocenter Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slices	1
Orientation Sagittal Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Dist. factor	20 %
Phase enc. dir. A >> P Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Position	Isocenter
Slice group 2 Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Orientation	Sagittal
Slices 5 Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Phase enc. dir.	A >> P
Dist. factor 300 % Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slice group	2
Position Isocenter Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slices	5
Orientation Transversal Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Dist. factor	300 %
Phase enc. dir. A >> P Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Position	Isocenter
Slice group 3 Slices 2 Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Orientation	Transversal
Slices 2	Phase enc. dir.	A >> P
Dist. factor 70 % Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slice group	3
Position L0.0 A30.0 H0.0 mm Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Slices	2
Orientation Coronal Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Dist. factor	70 %
Phase enc. dir. R >> L FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Position	L0.0 A30.0 H0.0 mm
FoV read 300 mm FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Orientation	Coronal
FoV phase 100.0 % Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	Phase enc. dir.	R >> L
Slice thickness 10.0 mm TR 9.7 ms Multi-slice mode Sequential Series Interleaved	FoV read	300 mm
TR 9.7 ms Multi-slice mode Sequential Series Interleaved	FoV phase	100.0 %
Multi-slice mode Sequential Series Interleaved	Slice thickness	10.0 mm
Series Interleaved	TR	9.7 ms
	Multi-slice mode	Sequential
Concatenations 8	Series	Interleaved
Contactions	Concatenations	8

Geometry - AutoAlign

Slice group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P

Geometry - AutoAlign

Slice group	2
Position	Isocenter
Orientation	Transversal
Phase enc. dir.	A >> P
Slice group	3
Position	L0.0 A30.0 H0.0 mm
Orientation	Coronal
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	Isocenter
L	0.0 mm
Р	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Saturation mode	Standard
Fat suppr.	None
Water suppr.	None
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	ISO
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	Default

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	9.7 ms
Concatenations	8
Segments	1

Physio - Cardiac

Tagging	None
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	300 mm
FoV phase	100.0 %
Phase resolution	50 %

Physio - PACE

Resp. control	Off
Concatenations	8

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	1

Inline - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Inline - MapIt

Save original images	On	
MapIt	None	
Flip angle	40 deg	
Measurements	1	
Contrasts	1	
TR	9.7 ms	
TE	3.80 ms	

SIEMENS MAGNETOM Prisma

Sequence - Part 1

Introduction	On
Dimension	2D
Phase stabilisation	Off
Asymmetric echo	Allowed
Contrasts	1
Flow comp.	Read
Multi-slice mode	Sequential
Bandwidth	320 Hz/Px

Sequence - Part 2

Segments	1
Acoustic noise reduction	None
RF pulse type	Normal
Gradient mode	Normal
Excitation	Slice-sel.
RF spoiling	On

Mode	Off	
Allowed delay	0 s	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\vessel_scout

TA: 0:25 PM: FIX Voxel size: 1.4×1.4×5.0 mmPAT: 2 Rel. SNR: 1.00 : pc

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Olak amana	4
Slab group	1
Slabs	1
Dist. factor	20 %
Position	R5.1 A9.0 F14.7 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	26
FoV read	350 mm
FoV phase	59.4 %
Slice thickness	5.00 mm
TR	27.65 ms
TE	5.51 ms
Averages	1
Concatenations	1
Filter	Distortion Corr.(2D),
	Elliptical filter, Image Filter
Coil elements	HEA;HEP

Contrast - Common

TR	27.65 ms
TE	5.51 ms
Flip angle	12 deg

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	350 mm
FoV phase	59.4 %
Slice thickness	5.00 mm
Base resolution	256
Phase resolution	65 %
Slice resolution	54 %
Phase partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode GRAPPA

Resolution - iPAT

Accel. factor PE	2
Ref. lines PE	30
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	On
Intensity	Medium
Edge Enhancement	3
Smoothing	3
Unfiltered images	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	On	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	R5.1 A9.0 F14.7 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	26
FoV read	350 mm
FoV phase	59.4 %
Slice thickness	5.00 mm
TR	27.65 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R5.1 A9.0 F14.7 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	
Initial Position	R5.1 A9.0 F14.7
R	5.1 mm
Α	9.0 mm
F	14.7 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Isocenter
Transversal
0.00 deg
263 mm
350 mm
350 mm
Off

System - pTx Volumes

B1 Shim mode	TrueForm

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	27.65 ms
Concatenations	1
Segments	1

Angio - Common

Flow mode	Free
Encodings	2
Velocity enc. 1	40 cm/s
Velocity enc. 2	40 cm/s
Direction 1	F >> H
Direction 2	Through plane
Rephased images	On
Magnitude images	Off
Magnitude sum	On
Phase images	Off

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off

Angio - Inline

Save original images	On	
Angio - MIP		
MIP-Sag	On	
MIP-Cor	On	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Angio - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Asymmetric echo	Weak
Contrasts	1
Flow comp.	No
Multi-slice mode	Sequential
Bandwidth	300 Hz/Px

Sequence - Part 2

Segments	1
RF pulse type	Fast
Gradient mode	Fast
RF spoiling	Off

Mode	Off	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\PC_Sinus

TA: 2:11 PM: FIX Voxel size: 0.7×0.7×5.0 mmPAT: 2 Rel. SNR: 1.00 : fl_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	<u>·</u>
Dist. factor	20 %
Position	R6.3 P58.2 H77.5 mm
Orientation	C > T16.3 > S2.6
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	21.70 ms
TE	6.59 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	21.70 ms
TR TE TD	6.59 ms
TD	0 ms
Flip angle Wrap-up Magn.	12 deg
Wrap-up Magn.	None

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Off

Resolution - Common

FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
Base resolution	224
Phase resolution	100 %
Phase partial Fourier	Off
Trajectory	Cartesian
View sharing	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2

Resolution - iPAT

Ref. lines PE	32
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

F	Raw filter	Off
E	Elliptical filter	Off
F	POCS	Off

Geometry - Common

Slice group	1
Slices	1
Dist. factor	20 %
Position	R6.3 P58.2 H77.5 mm
Orientation	C > T16.3 > S2.6
Phase enc. dir.	R >> L
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	21.70 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Occinicity AutoAngii	
Slice group	1
Position	R6.3 P58.2 H77.5 mm
Orientation	C > T16.3 > S2.6
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	R6.3 P58.2 H77.5
R	6.3 mm
Р	58.2 mm
Н	77.5 mm
Initial Rotation	0.00 deg
Initial Orientation	C > T
C > T	16.3
> S	2.6

Geometry - Saturation

Wrap-up Magn.	None	
Special sat.	None	

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	H
. abio pooliion	• •
Table position	0 mm
. abio pooliion	•
Inline Composing	Off
minic composing	Oli

Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

R6.3 P58.2 H77.5 mm
C > T16.3 > S2.6
0.00 deg
160 mm
160 mm
5 mm
Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	Pulse/Retro
Average cycle	No Signal ms
Average cycle	No Signal ms
Calculated phases	32
TR	21.70 ms
Concatenations	1
Segments	1
Arrhythmia detection	None

Physio - PACE

Resp. control	Off
Concatenations	1

Angio - Common

7 til glo - Goliminon	
Flow mode	Single dir.
Encodings	1
Velocity enc.	50 cm/s
Direction	Through plane
Rephased images	On
Magnitude images	On

Angio - Common

Magnitude sum	Off	
Phase images	On	

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Angio - Cardiac

Inline Evaluation	Off
TE	6.59 ms
TR	21.70 ms
Save original images	On

Angio - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Angio - Composing

Inline Composing	Off
Distortion Corr.	Off

Sequence - Part 1

Introduction	On
Dimension	2D
Reordering	Linear
Asymmetric echo	Weak
Flow comp.	Yes
Optimization	Min. TE TR
Multi-slice mode	Sequential
Echo spacing	10.9 ms
Sequence type	Gre
Bandwidth	203 Hz/Px

Sequence - Part 2

Define	Segments
Segments	1
RF pulse type	Normal
Gradient mode	Fast*
Excitation	Slice-sel.
Flip angle mode	Constant
RF spoiling	On
Phase Enc. Rewinder	On
Cine	On

Mode	Off	
Allowed delay	0 e	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t1_fl2d_sag_localiser

TA: 0:51 PM: REF Voxel size: 0.7×0.7×3.0 mmPAT: Off Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	Off
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	20
Dist. factor	0 %
Position	L0.9 P5.3 H70.4 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
AutoAlign	
Phase oversampling	30 %
FoV read	220 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
TR	150.0 ms
TE	2.46 ms
Averages	1
Concatenations	1
Filter	Distortion Corr.(2D),
	Prescan Normalize,
	Elliptical filter
Coil elements	HEA;HEP

Contrast - Common

TR	150.0 ms
TE	2.46 ms
MTC	Off
Magn. preparation	None
Flip angle	70 deg
Fat suppr.	None
Water suppr.	None
SWI	Off

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	220 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
Base resolution	320
Phase resolution	80 %
Phase partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	None
Resolution - Filter Ima	ge
Image Filter	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	On	

Off

Geometry - Common

B1 filter

Slice group	1
Slices	20
Dist. factor	0 %
Position	L0.9 P5.3 H70.4 mm
Orientation	Sagittal
Phase enc. dir.	A >> P
FoV read	220 mm
FoV phase	100.0 %
Slice thickness	3.0 mm
TR	150.0 ms
Multi-slice mode	Interleaved
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

1
L0.9 P5.3 H70.4 mm
Sagittal
A >> P
L0.9 P5.3 H70.4
0.9 mm
5.3 mm
70.4 mm
0.00 deg
Sagittal

Geometry - Saturation

Saturation mode	Standard
Fat suppr.	None
Water suppr.	None
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	REF
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

-	
Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	150.0 ms
Concatenations	1
Segments	1

Physio - Cardiac

Tagging	None
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	220 mm
FoV phase	100.0 %
Phase resolution	80 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off

Inline - Common

Liver registration	Off	
Save original images	On	

Inline - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	1

Inline - Composing

Inline Composing	Off
Distortion Corr.	On
Mode	2D
Unfiltered images	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	70 deg
Measurements	1
Contrasts	1
TR	150.0 ms
TE	2.46 ms

Sequence - Part 1

Introduction	On
Dimension	2D
Phase stabilisation	Off
Asymmetric echo	Allowed
Contrasts	1
Flow comp.	No
Multi-slice mode	Interleaved
Bandwidth	320 Hz/Px

Sequence - Part 2

Segments	1
Acoustic noise reduction	None
RF pulse type	Fast
Gradient mode	Performance
Excitation	Slice-sel.
RF spoiling	On

Mode	Off
Allowed delay	60 s

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\PC_Aqueduct

TA: 2:27 PM: FIX Voxel size: 0.6×0.6×5.0 mmPAT: 2 Rel. SNR: 1.00 : fl_r

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
l' '	0"
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slice group	1
Slices	1
Dist. factor	20 %
Position	L0.6 P22.4 H72.9 mm
Orientation	T > C21.4 > S-4.8
Phase enc. dir.	R >> L
AutoAlign	
Phase oversampling	0 %
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	22.94 ms
TE	7.06 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

22.94 ms
7.06 ms
0 ms
12 deg
None

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1
Multiple series	Off

Resolution - Common

FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
Base resolution	256
Phase resolution	100 %
Phase partial Fourier	Off
Trajectory	Cartesian
View sharing	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2

Resolution - iPAT

Ref. lines PE	32
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
POCS	Off

Geometry - Common

Slice group	1
Slices	1
Dist. factor	20 %
Position	L0.6 P22.4 H72.9 mm
Orientation	T > C21.4 > S-4.8
Phase enc. dir.	R >> L
FoV read	160 mm
FoV phase	100.0 %
Slice thickness	5.0 mm
TR	22.94 ms
Multi-slice mode	Sequential
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Geometry - AutoAngn	
Slice group	1
Position	L0.6 P22.4 H72.9 mm
Orientation	T > C21.4 > S-4.8
Phase enc. dir.	R >> L
AutoAlign	
Initial Position	L0.6 P22.4 H72.9
L	0.6 mm
P	22.4 mm
Н	72.9 mm
Initial Rotation	90.20 deg
Initial Orientation	T > C
T > C	21.4
> S	-4.8

Geometry - Saturation

Wrap-up Magn.	None
Special sat.	None

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	H
. abio pooliion	• •
Table position	0 mm
. abio pooliion	•
Inline Composing	Off
minic composing	Oli

Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Sum of Squares
Matrix Optimization	Off
AutoAlign	
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	L0.6 P22.4 H72.9 mm
Orientation	T > C21.4 > S-4.8
Rotation	90.20 deg
R >> L	160 mm
A >> P	160 mm
A >> P F >> H	5 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Slice-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	High
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	Pulse/Retro
Average cycle	No Signal ms
Average cycle	No Signal ms
Calculated phases	32
TR	22.94 ms
Concatenations	1
Segments	1
Arrhythmia detection	None

Physio - PACE

Resp. control	Off
Concatenations	1

Angio - Common

7.11.9.0 CO.11.11.01.1	
Flow mode	Single dir.
Encodings	1
Velocity enc.	10 cm/s
Direction	Through plane
Rephased images	On
Magnitude images	On

Angio - Common

Magnitude sum	Off	
Phase images	On	

Angio - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Angio - Cardiac

Inline Evaluation	Off
TE	7.06 ms
TR	22.94 ms
Save original images	On

Angio - MIP

MIP-Sag	Off	
MIP-Cor	Off	
MIP-Tra	Off	
MIP-Time	Off	
Save original images	On	

Angio - Composing

Inline Composing	Off
Distortion Corr.	Off

Sequence - Part 1

Introduction	On
Dimension	2D
Reordering	Linear
Asymmetric echo	Weak
Flow comp.	Yes
Optimization	Min. TE TR
Multi-slice mode	Sequential
Echo spacing	11.5 ms
Sequence type	Gre
Bandwidth	201 Hz/Px

Sequence - Part 2

Define	Segments
Segments	1
RF pulse type	Normal
Gradient mode	Fast*
Excitation	Slice-sel.
Flip angle mode	Constant
RF spoiling	On
Phase Enc. Rewinder	On
Cine	On

Mode	Off	
Allowed delay	Λe	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\AAHead_Scout_32ch-head-coil

TA: 0:14 PM: FIX Voxel size: 1.6×1.6×1.6 mmPAT: 3 Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	On
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
TE	1.37 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3.15 ms
TE	1.37 ms
Flip angle	8 deg

Contrast - Dynamic

Averages	1
Averaging mode	Short term
Reconstruction	Magnitude
Measurements	1

Resolution - Common

FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	69 %
Phase partial Fourier	6/8
Slice partial Fourier	6/8
Trajectory	Cartesian

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	3
Ref. lines PE	24
Accel. factor 3D	1

Resolution - iPAT

Reference scan mode	Integrated	
Resolution - Filter Image		

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	128
FoV read	260 mm
FoV phase	100.0 %
Slice thickness	1.6 mm
TR	3.15 ms
Multi-slice mode	Sequential
Series	Ascending
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	Isocenter
Orientation	Sagittal
Phase enc. dir.	A >> P
Initial Position	Isocenter
L	0.0 mm
P	0.0 mm
Н	0.0 mm
Initial Rotation	0.00 deg
Initial Orientation	Transversal

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off

Coil Select Mode	Default

System - Adjustments

B0 Shim mode	Tune up
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	Isocenter
Orientation	Transversal
Rotation	0.00 deg
A >> P	263 mm
R >> L	350 mm
F >> H	350 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	1.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Flip angle	8 deg
Measurements	1
Time to center	6.2 s

Inline - Inline

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
	- · ·
MapIt	None
Flip angle	8 deg
1 ' "	5
Measurements	1

Inline - MapIt

Contrasts	1
TR	3.15 ms
TE	1.37 ms

Sequence - Part 1

Introduction	On
Dimension	3D
Asymmetric echo	Weak
Contrasts	1
Multi-slice mode	Sequential
Bandwidth	540 Hz/Px

Sequence - Part 2

RF pulse type	Fast
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On

Mode	Off

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\IR-SPGR_TI_600_v4

TA: 1:55 PM: FIX Voxel size: 1.2×1.2×1.2 mmPAT: 2 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	Off
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	1040.0 ms
TE	1.82 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	1040.0 ms
TE	1.82 ms
Magn. preparation	Non-sel. IR
TI	600 ms
Flip angle	5 deg
Fat suppr.	None
Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	1040.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
A >> P	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	3.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	1040.0 ms
Concatenations	1

Physio - Cardiac

Magn. preparation	Non-sel. IR
ТІ	600 ms
Fat suppr.	None
Dark blood	Off
FoV read	192 mm
FoV phase	125.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off	
Concatenations	1	

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
	÷
MapIt	None
Flip angle	5 deg
Measurements	1
TR	1040.0 ms
TE	1.82 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	5.4 ms
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On
Turbo factor	160

Mode	Off
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\IR-SPGR_TI_1500_v4

TA: 3:35 PM: FIX Voxel size: 1.2×1.2×1.2 mmPAT: 2 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	1940.0 ms
TE	1.82 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	1940.0 ms
TE	1.82 ms
Magn. preparation	Non-sel. IR
TI	1500 ms
Flip angle	5 deg
Fat suppr.	None
Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	24
Accel. factor 3D	1
Reference scan mode	Integrated

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	1940.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
A >> P	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	3.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	1940.0 ms
Concatenations	1

Physio - Cardiac

Magn. preparation	Non-sel. IR
TI	1500 ms
Fat suppr.	None
Dark blood	Off
FoV read	192 mm
FoV phase	125.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off	
Concatenations	1	

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	5 deg
Measurements	1
TR	1940.0 ms
TE	1.82 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	5.4 ms
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On
Turbo factor	160

Mode	Off
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\IR-SPGR_IROff_2_v4

TA: 1:36 PM: FIX Voxel size: 1.2×1.2×1.2 mmPAT: 2 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	On
preparation	0"
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
TE	1.82 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR TE	865.0 ms	
TE	1.82 ms	
Magn. preparation	None	
Flip angle	2 deg	
Fat suppr.	None	
Water suppr.	None	

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA	
Accel. factor PE	2	
Ref. lines PE	24	
Accel. factor 3D	1	
Reference scan mode	Integrated	

Resolution - Filter Image

Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
F >> H A >> P R >> L	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	3.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	865.0 ms
Concatenations	1

Physio - Cardiac

•	
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	192 mm
FoV phase	125.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	2 deg
Measurements	1
TR	865.0 ms
TE	1.82 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	5.4 ms
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On
Turbo factor	160

Mode	Off
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\IR-SPGR_IROff_5_v4

TA: 1:36 PM: FIX Voxel size: 1.2×1.2×1.2 mmPAT: 2 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
TE	1.82 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR TE	865.0 ms
TE	1.82 ms
Magn. preparation	None
Flip angle	5 deg
Fat suppr.	None
Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA	
Accel. factor PE	2	
Ref. lines PE	24	
Accel. factor 3D	1	
Reference scan mode	Integrated	

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
A >> P	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	3.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	865.0 ms
Concatenations	1

Physio - Cardiac

•	
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	192 mm
FoV phase	125.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off	
Distortion Corr.	Off	

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	5 deg
Measurements	1
TR	865.0 ms
TE	1.82 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	5.4 ms
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On
Turbo factor	160

Mode	Off
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\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\IR-SPGR_IROff_12_v4

TA: 1:36 PM: FIX Voxel size: 1.2×1.2×1.2 mmPAT: 2 Rel. SNR: 1.00 : tfl

Properties

Prio recon	Off
Load images to viewer	On
Inline movie	Off
Auto store images	On
Load images to stamp segments	On
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further	r On
preparation	
Wait for user to start	Off
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
TE	1.82 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR TE	865.0 ms
TE	1.82 ms
Magn. preparation	None
Flip angle	12 deg
Fat suppr.	None
Water suppr.	None

Contrast - Dynamic

Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
Base resolution	160
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off

Resolution - iPAT

PAT mode	GRAPPA	
Accel. factor PE	2	
Ref. lines PE	24	
Accel. factor 3D	1	
Reference scan mode	Integrated	

Resolution - Filter Image

Image Filter	Off	
Distortion Corr.	Off	
Prescan Normalize	On	
Unfiltered images	Off	
Normalize	Off	
B1 filter	Off	

Resolution - Filter Rawdata

Raw filter	Off	
Elliptical filter	Off	

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	50 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	160
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	1.20 mm
TR	865.0 ms
Multi-slice mode	Single shot
Series	Interleaved
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Navigator

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

Positioning mode	FIX
Table position	Н
Table position	0 mm

MSMA	S-C-T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
A >> P	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	3.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - Signal1

1st Signal/Mode	None
TR	865.0 ms
Concatenations	1

Physio - Cardiac

•	
Magn. preparation	None
Fat suppr.	None
Dark blood	Off
FoV read	192 mm
FoV phase	125.0 %
Phase resolution	100 %

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

Subtract	Off
Measurements	1
StdDev	Off
Save original images	On

Inline - MIP

MIP-Sag	Off
MIP-Cor	Off
MIP-Sag MIP-Cor MIP-Tra MIP-Time	Off
MIP-Time	Off
Save original images	On

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	12 deg
Measurements	1
TR	865.0 ms
TE	1.82 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Reordering	Linear
Asymmetric echo	Off
Flow comp.	No
Multi-slice mode	Single shot
Echo spacing	5.4 ms
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On
Turbo factor	160

Mode	Off	

\\Study Protocols\BRAIN\Stroke\Investigate - E161686 v1b\t1_vibe_sag_DCE_2mm

TA: 21:08 PM: FIX Voxel size: 2.0×2.0×2.0 mmPAT: Off Rel. SNR: 1.00 : fl

Properties

Prio recon	Off
Load images to viewer	Off
Inline movie	Off
Auto store images	On
Load images to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Auto close inline display	Off
Start measurement without further preparation	On
Wait for user to start	On
Start measurements	Single measurement

Routine

Slab group	1
Slabs	1
Dist. factor	20 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Phase oversampling	0 %
Slice oversampling	0.0 %
Slices per slab	96
FoV read	192 mm
FoV phase	125.0 %
Slice thickness	2.0 mm
TR	3.44 ms
TE	1.68 ms
Averages	1
Concatenations	1
Filter	Prescan Normalize
Coil elements	HEA;HEP

Contrast - Common

TR	3.44 ms
TE	1.68 ms
Flip angle	15.0 deg
Fat suppr.	None
Water suppr.	None
Dixon	Off

Contrast - Dynamic

Contrast - Dynamic	
Averages	1
Averaging mode	Long term
Reconstruction	Magnitude
Measurements	32
Pause after meas. 1	0.0 s
Pause after meas. 2	0.0 s
Pause after meas. 3	0.0 s
Pause after meas. 4	0.0 s
Pause after meas. 5	0.0 s
Pause after meas. 6	0.0 s
Pause after meas. 7	0.0 s
Pause after meas. 8	0.0 s
Pause after meas. 9	0.0 s
Pause after meas. 10	0.0 s
Pause after meas. 11	0.0 s
Pause after meas. 12	0.0 s
Pause after meas. 13	0.0 s

Contrast - Dynamic

Pause after meas. 14	0.0 s
Pause after meas. 15	0.0 s
Pause after meas. 16	0.0 s
Pause after meas. 17	0.0 s
Pause after meas. 18	0.0 s
Pause after meas. 19	0.0 s
Pause after meas. 20	0.0 s
Pause after meas. 21	0.0 s
Pause after meas. 22	0.0 s
Pause after meas. 23	0.0 s
Pause after meas. 24	0.0 s
Pause after meas. 25	0.0 s
Pause after meas. 26	0.0 s
Pause after meas. 27	0.0 s
Pause after meas. 28	0.0 s
Pause after meas. 29	0.0 s
Pause after meas. 30	0.0 s
Pause after meas. 31	0.0 s
Multiple series	Off

Resolution - Common

FoV read	192 mm
FoV phase	125.0 %
Slice thickness	2.0 mm
Base resolution	96
Phase resolution	100 %
Slice resolution	100 %
Phase partial Fourier	Off
Slice partial Fourier	Off
Trajectory	Cartesian
View sharing	Off
Interpolation	Off

Resolution - iPAT

PAT mode	None

Resolution - Filter Image

Image Filter	Off
Distortion Corr.	Off
Prescan Normalize	On
Unfiltered images	Off
Normalize	Off
B1 filter	Off

Resolution - Filter Rawdata

Raw filter	Off
Elliptical filter	Off
POCS	Off

Geometry - Common

Slab group	1
Slabs	1
Dist. factor	20 %
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
Slice oversampling	0.0 %
Slices per slab	96
FoV read	192 mm
FoV phase	125.0 %

Geometry - Common

Slice thickness	2.0 mm
TR	3.44 ms
Multi-slice mode	Sequential
Series	Ascending
Concatenations	1

Geometry - AutoAlign

Slab group	1
Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Phase enc. dir.	A >> P
AutoAlign	Head > Brain
Initial Position	L0.0 A3.9 F2.9
L	0.0 mm
A	3.9 mm
F	2.9 mm
Initial Rotation	0.00 deg
Initial Orientation	Sagittal

Geometry - Saturation

Fat suppr.	None
Water suppr.	None
Dixon	Off
Special sat.	None

Geometry - Tim Planning Suite

Set-n-Go Protocol	Off
Table position	Н
Table position	0 mm
Inline Composing	Off

System - Miscellaneous

Positioning mode	FIX
Table position	Н
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Coil Combine Mode	Adaptive Combine
Save uncombined	Off
Matrix Optimization	Off
AutoAlign	Head > Brain
Coil Select Mode	On - AutoCoilSelect

System - Adjustments

B0 Shim mode	Standard
B1 Shim mode	TrueForm
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Dominant Fat	Off
Assume Silicone	Off
Adjustment Tolerance	Auto

System - Adjust Volume

Position	R8.0 A13.6 H6.2 mm
Orientation	S > T-4.2 > C-2.3
Rotation	97.97 deg
F >> H	192 mm
A >> P	240 mm
R >> L	192 mm
Reset	Off

System - pTx Volumes

B1 Shim mode	TrueForm
Excitation	Non-sel.

System - Tx/Rx

Frequency 1H	123.244448 MHz
Correction factor	1
Gain	Low
Img. Scale Cor.	6.000
Reset	Off
? Ref. amplitude 1H	0.000 V

Physio - PACE

Resp. control	Off
Concatenations	1

Inline - Common

View sharing	Off
Flip angle	15.0 deg
Measurements	32
Pause after meas. 1	0.0 s
Pause after meas. 2	0.0 s
Pause after meas. 3	0.0 s
Pause after meas. 4	0.0 s
Pause after meas. 5	0.0 s
Pause after meas. 6	0.0 s
Pause after meas. 7	0.0 s
Pause after meas. 8	0.0 s
Pause after meas. 9	0.0 s
Pause after meas. 10	0.0 s
Pause after meas. 11	0.0 s
Pause after meas. 12	0.0 s
Pause after meas. 13	0.0 s
Pause after meas. 14	0.0 s
Pause after meas. 15	0.0 s
Pause after meas. 16	0.0 s
Pause after meas. 17	0.0 s
Pause after meas. 18	0.0 s
Pause after meas. 19	0.0 s
Pause after meas. 20	0.0 s
Pause after meas. 21	0.0 s
Pause after meas. 22	0.0 s
Pause after meas. 23	0.0 s
Pause after meas. 24	0.0 s
Pause after meas. 25	0.0 s
Pause after meas. 26	0.0 s
Pause after meas. 27	0.0 s
Pause after meas. 28	0.0 s
Pause after meas. 29	0.0 s
Pause after meas. 30	0.0 s
Pause after meas. 31	0.0 s
Burn time-to-center	Off
Temporal interpolation	1
3D centric reordering	Off
Time to center	20.0 s

Inline - Inline

Subtract	Off
Measurements	32
StdDev	Off
Liver registration	Off
Save original images	On

Inline - MIP

MIP-Sag	Off	
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Inline - MIP

MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Inline - Soft Tissue

Wash - In	Off
Wash - Out	Off
TTP	Off
PEI	Off
MIP - time	Off
Measurements	32
Pause after meas. 1	0.0 s
Pause after meas. 2	0.0 s
Pause after meas. 3	0.0 s
Pause after meas. 4	0.0 s
Pause after meas. 5	0.0 s
Pause after meas. 6	0.0 s
Pause after meas. 7	0.0 s
Pause after meas. 8	0.0 s
Pause after meas. 9	0.0 s
Pause after meas. 10	0.0 s
Pause after meas. 11	0.0 s
Pause after meas. 12	0.0 s
Pause after meas. 13	0.0 s
Pause after meas. 14	0.0 s
Pause after meas. 15	0.0 s
Pause after meas. 16	0.0 s
Pause after meas. 17	0.0 s
Pause after meas. 18	0.0 s
Pause after meas. 19	0.0 s
Pause after meas. 20	0.0 s
Pause after meas. 21	0.0 s
Pause after meas. 22	0.0 s
Pause after meas. 23	0.0 s
Pause after meas. 24	0.0 s
Pause after meas. 25	0.0 s
Pause after meas. 26	0.0 s
Pause after meas. 27	0.0 s
Pause after meas. 28	0.0 s
Pause after meas. 29	0.0 s
Pause after meas. 30	0.0 s
Pause after meas. 31	0.0 s

Inline - Composing

Inline Composing	Off
Distortion Corr.	Off

Inline - MapIt

Save original images	On
MapIt	None
Flip angle	15.0 deg
Measurements	32
Contrasts	1
TR	3.44 ms
TE	1.68 ms

Sequence - Part 1

Introduction	Off
Dimension	3D
Elliptical scanning	Off
Asymmetric echo	Off
Contrasts	1
Optimization	Min. TE

Sequence - Part 1

Multi-slice mode	Sequential
Bandwidth	520 Hz/Px

Sequence - Part 2

RF pulse type	Normal
Gradient mode	Normal
Excitation	Non-sel.
RF spoiling	On
Incr. Gradient spoiling	On

Mode	Off
Allowed delay	0 s