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 (3)


 (1)


 (7)


Hospital (1)


 {B1860723-4F8F-476e-8075-D42C65706693} (1)


InBox (1)


 spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25:06.7


 SURVEY 00:33.8


 T1w 04:22.0

 T2w 05:11.5

 DWI 05:34.1

 GRE-MT 04:20.4

 GRE-T1 00:34.3

 GRE-ME 04:30.6

mk:@MSITStore:C:\Users\310073343\AppData\Local\Temp\Temp1\_spine\_generic\_phili... 9/29/2020

Hospital (1) | {B1860723-4F8F-476e-8075-D42C65706693} (1) |  
 InBox (1) | spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25.06.7 | SURVEY 00:33.8

INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	00:33.8	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	MS
Act. TR/TE (ms)	9.3 / 5.4	FOV RL (mm)	300	technique	FFE
ACQ matrix M x P	240 x 200	FH (mm)	300	Contrast enhancement	T1
ACQ voxel MPS (mm)	1.25 / 1.50 / 10.0	stack AP (mm)	32	Acquisition mode	cartesian
REC voxel MPS (mm)	0.94 / 0.94 / 10.0	ACQ voxel size RL (mm)	1.25	Fast Imaging mode	TFE
Scan percentage (%)	83.33334	FH (mm)	1.5	shot mode	single-shot
Packages	3	Slice thickness (mm)	10	TFE startup echoes	default
Min. slice gap (mm)	0	Recon voxel size RL (mm)	0.9375	profile order	linear
TFE factor	400	FH (mm)	0.9375	Echoes	1
TFE dur. shot / acq (ms)	3752.2 / 3715.1	Fold-over suppression	oversampling	partial echo	no
Act. WFS (pix) / BW (Hz)	0.817 / 531.5	H (mm)	150	shifted echo	no
Min. WFS (pix) / Max. BW (Hz)	0.817 / 531.5	F (mm)	150	TE	shortest
Local torso SAR	< 6 %	Reconstruction matrix	320	Flip angle (deg)	20
Whole body SAR / level	< 0.2 W/kg / normal	SENSE	no	TR	shortest
SED	0.0 kJ/kg	k-t BLAST	no	Halfscan	no
Coil Power	5 %	Stacks	3	Water-fat shift	minimum
Max B1+rms	0.55 uT	current	A	RF Shims	fixed
PNS / level	10 % / normal	type	parallel	Shim	default
dB/dt	15.4 T/s	slices	3	mDIXON	no
Sound Pressure Level (dB)	-0.8140956	slice gap	user defined	Fat suppression	no
Boil-off (hPa/h)	1.64134	gap (mm)	1	Water suppression	no
Max Boil-off (hPa/h)	1.646055	slice orientation	coronal	TFE prepulse	no
Boil-off (hPa)	0.01539666	fold-over direction	FH	MTC	no
<b>MOTION</b>		fat shift direction	L	T2prep	no
Cardiac synchronization	no	Stacks as packages	yes	Research prepulse	no
Heart rate > 250 bpm	no	Minimum number of packages	1	MDME	no
Respiratory compensation	no	Slice scan order	default	Diffusion mode	no
Navigator respiratory comp	no	Stack scan order	ascend	T1 mapping	no
Flow compensation	no	Move table per stack	no	Transmit channels	both
Temporal slice spacing	default	Stack alignment	no	SAR mode	low
fMRI echo stabilisation	no	Stack sort order	no	B1 mode	default
Motion smoothing	no	PlanAlign	yes	SAR allow first level	yes
NSA	1	REST slabs	0	PNS mode	low
MRE enable	no	Catheter tracking	no	Gradient mode	regular
<b>DYN/ANG</b>		Interactive positioning	no	SoftTone mode	yes
Angio / Contrast enh.	no	Allow table movement	no		
Quantitative flow	no	<b>OFFC/ANG</b>			
Manual start	no	Stacks	3		
Dynamic study	no	current	A		
Arterial Spin labeling	no	Stack Offc. AP (P=+mm)	0		
<b>POST/PROC</b>		RL (L=+mm)	0		
Preparation phases	auto	FH (H=+mm)	0		
Interactive F0	no	Ang. AP (deg)	0		
SmartPlan survey	no	RL (deg)	0		
B0 field map	no	FH (deg)	0		
B1 field map	no	Free rotatable	no		
MIP/MPR	no				
SWIp	no				
Images	M, no, no, no				
Autoview image	M				
Calculated images	no, no, no, no				
Reference tissue	Skeletal muscle				
Recon compression	No				
Preset window contrast	hard				
Reconstruction mode	real time				
Save raw data	no				
Hardcopy protocol	no				
Image filter	system default				
Uniformity correction	no				
Geometry correction	default				

Hospital (1) | {B1860723-4F8F-476e-8075-D42C65706693} (1) |  
 Inbox (1) | spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25.06.7 | T1w 04:22.0

INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	04:22.0	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	3D
Act. TR/TE (ms)	7.8 / 3.5	FOV FH (mm)	320	technique	FFE
ACQ matrix M x P	320 x 260	AP (mm)	260	Contrast enhancement	T1
ACQ voxel MPS (mm)	1.00 / 1.00 / 1.00	RL (mm)	192	Acquisition mode	cartesian
REC voxel MPS (mm)	1.00 / 1.00 / 1.00	ACQ voxel size FH (mm)	1	Fast Imaging mode	TFE
Scan percentage (%)	100	AP (mm)	1	3D non-selective	no
Act. slice gap (mm)	0	RL (mm)	1	shot mode	multishot
TFE shots	130	Recon voxel size FH (mm)	1	TFE factor	246
TFE dur. shot / acq (ms)	1968.1 / 1912.0	AP (mm)	1	3D free factor	no
Min. TI delay	996.3246	RL (mm)	1	startup echoes	default
Act. WFS (pix) / BW (Hz)	2.268 / 191.5	Fold-over suppression	no	shot interval (ms)	user defined
Min. WFS (pix) / Max. BW (Hz)	0.456 / 952.7	Slice oversampling	default	profile order	linear
Local torso SAR	< 16 %	RF select. FOS	no	turbo direction	Z
Whole body SAR / level	< 0.5 W/kg / normal	ENCASE enable	no	Echoes	1
SED	< 0.1 kJ/kg	Reconstruction matrix	320	partial echo	no
Coil Power	15 %	SENSE	yes	shifted echo	no
Max B1+rms	0.91 uT	P reduction (AP)	2	TE	shortest
PNS / level	74 % / normal	S reduction (RL)	1	Flip angle (deg)	9
dB/dt	117.8 T/s	k-t BLAST	no	TR	shortest
Sound Pressure Level (dB)	15.06801	Stacks	1	Halfscan	no
Boil-off (hPa/h)	7.042519	slices	192	Water-fat shift	maximum
Max Boil-off (hPa/h)	7.34327	slice orientation	sagittal	RF Shims	fixed
Boil-off (hPa)	0.5124765	fold-over direction	AP	Shim	auto
<b>CONFLICTS</b>		fat shift direction	F	mDIXON	no
AP (mm)	260	Multi-chunk	no	Fat suppression	no
ACQ voxel size FH (mm)	1	PlanAlign	no	Water suppression	no
AP (mm)	1	REST slabs	0	TFE prepulse	invert
Recon voxel size FH (mm)	1	Catheter tracking	no	slice selection	no
AP (mm)	1	Interactive positioning	no	delay	user defined
Reconstruction matrix	320	Allow table movement	no	(ms)	1000
slices	192	<b>OFFC/ANG</b>		PSIR	no
<b>MOTION</b>		Stacks	1	MTC	no
Cardiac synchronization	no	Stack Offc. AP (P=+mm)	4.959927	T2prep	no
Heart rate > 250 bpm	no	RL (L=+mm)	3.967938	Research prepulse	no
Respiratory compensation	no	FH (H=+mm)	47.61523	MDME	no
Navigator respiratory comp	no	Ang. AP (deg)	-2.273305	Diffusion mode	no
Flow compensation	no	RL (deg)	0	T1 mapping	no
fMRI echo stabilisation	no	FH (deg)	0	Transmit channels	both
Motion smoothing	no	Free rotatable	no	SAR mode	high
NSA	1	<b>POST/PROC</b>		B1 mode	default
MRE enable	no	Preparation phases	auto	SAR allow first level	yes
<b>DYN/ANG</b>		Interactive F0	no	PNS mode	moderate
Angio / Contrast enh.	no	SmartPlan survey	no	Gradient mode	maximum
Quantitative flow	no	B0 field map	no	SoftTone mode	no
CENTRA	no	B1 field map	no		
Manual start	no	MIP/MPR	no		
Dynamic study	no	SWIp	no		
Arterial Spin labeling	no	Images	M, no, no, no		
<b>POST/PROC</b>		Autoview image	M		
Preparation phases	auto	Calculated images	no, no, no, no		
Interactive F0	no	Reference tissue	White matter		
SmartPlan survey	no	Recon compression	No		
B0 field map	no	Preset window contrast	soft		
B1 field map	no	Reconstruction mode	immediate		
MIP/MPR	no	Save raw data	no		
SWIp	no	Hardcopy protocol	no		
Images	M, no, no, no	Image filter	weak		
Autoview image	M	Uniformity correction	no		
Calculated images	no, no, no, no	Geometry correction	default		
Reference tissue	White matter	Elliptical k-space shutter	default		
Recon compression	No				
Preset window contrast	soft				
Reconstruction mode	immediate				
Save raw data	no				
Hardcopy protocol	no				
Image filter	weak				
Uniformity correction	no				
Geometry correction	default				
Elliptical k-space shutter	default				

Hospital (1) | {B1860723-4F8F-476e-8075-D42C65706693} (1) |  
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INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	05:11.5	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	3D
Act. TR (ms)	3500	FOV FH (mm)	256	technique	SE
Act. TE (ms)	120	AP (mm)	236.8	Modified SE	no
ACQ matrix M x P	320 x 294	RL (mm)	50.4	Acquisition mode	cartesian
ACQ voxel MPS (mm)	0.80 / 0.81 / 0.80	ACQ voxel size FH (mm)	0.8	Fast Imaging mode	TSE
REC voxel MPS (mm)	0.80 / 0.80 / 0.80	AP (mm)	0.7996281	3D VIEW	no
Scan percentage (%)	99.32433	RL (mm)	0.8	shot mode	multishot
Act. slice gap (mm)	0	Recon voxel size FH (mm)	0.8	TSE factor	92
WFS (pix) / BW (Hz)	1.056 / 411.2	AP (mm)	0.8	3D free factor	no
TSE es / shot (ms)	6.2 / 578	RL (mm)	0.8	startup echoes	0
Min. TR (ms)	1733	Fold-over suppression	no	profile order	linear
Act. half scan factor	0.6278125	Slice oversampling	default	turbo direction	Y
Local torso SAR	< 50 %	RF select. FOS	no	DRIVE	yes
Whole body SAR / level	< 1.6 W/kg / normal	Reconstruction matrix	320	ultrashort	yes
SED	< 0.5 kJ/kg	SENSE	yes	shift	0
Coil Power	47 %	P reduction (AP)	2	fid reduction	default
Max B1+rms	1.61 uT	S reduction (RL)	1.5	3D non-selective	yes
PNS / level	68 % / normal	k-t BLAST	no	Echoes	1
dB/dt	102.5 T/s	Stacks	1	partial echo	no
Sound Pressure Level (dB)	21.71582	slices	63	TE	user defined
Boil-off (hPa/h)	0.8835939	slice orientation	sagittal	(ms)	120
Max Boil-off (hPa/h)	5.318677	fold-over direction	AP	Flip angle (deg)	90
Boil-off (hPa)	0.07645541	fat shift direction	F	Refocusing control	no
<b>MOTION</b>		Multi-chunk	no	TR	user defined
Cardiac synchronization	no	O-MAR	no	(ms)	3500
Heart rate > 250 bpm	no	PlanAlign	no	Halfscan	default
Respiratory compensation	no	REST slabs	0	Water-fat shift	maximum
Navigator respiratory comp	no	Catheter tracking	no	RF Shims	fixed
Flow compensation	no	Interactive positioning	no	Shim	default
Motion smoothing	no	Allow table movement	no	mDIXON	no
NSA	1	<b>OFFC/ANG</b>		Fat suppression	no
MRE enable	no	Stacks	1	Grad Rev Fat suppr	no
<b>DYN/ANG</b>		Stack Offc. AP (P=+mm)	4.959927	Water suppression	no
CENTRA	no	RL (L=+mm)	3.967938	BB pulse	no
Manual start	no	FH (H=+mm)	47.61523	MTC	no
Dynamic study	no	Ang. AP (deg)	-2.273305	Research prepulse	no
Arterial Spin labeling	no	RL (deg)	0	MDME	no
<b>POST/PROC</b>		FH (deg)	0	Zoom imaging	no
Preparation phases	auto	Free rotatable	no	Diffusion mode	no
Interactive F0	no			T1 mapping	no
SmartPlan survey	no			Transmit channels	both
B0 field map	no			SAR mode	high
B1 field map	no			B1 mode	default
MIP/MPR	no			SAR allow first level	yes
Images	M, no, no, no			PNS mode	moderate
Autoview image	M			Gradient mode	maximum
Calculated images	no, no, no, no			SoftTone mode	no
Reference tissue	Grey matter				
Recon compression	No				
Preset window contrast	soft				
Reconstruction mode	real time				
Save raw data	no				
Hardcopy protocol	no				
Image filter	weak				
Uniformity correction	no				
Geometry correction	default				
Elliptical k-space shutter	default				

INFO PAGE	
Total scan duration	05:34.1
Rel. SNR	1
Act. TR (ms)	3429
Act. TE (ms)	70
ACQ matrix M x P	64 x 47
ACQ voxel MPS (mm)	0.88 / 0.89 / 5.00
REC voxel MPS (mm)	0.88 / 0.88 / 5.00
Scan percentage (%)	97.91666
Packages	2
Min. slice gap (mm)	0
Optimal slices	8
Max. slices	16
Diffusion gradient timing DELTA / delta (ms)	34.5 / 13.7
EPI factor	47
Entered heartrate	70 (< 78)
Trigger delay max. / act. (mm)	771.4 / 111.2
WFS (pix) / BW (Hz)	23.594 / 18.4
BW in EPI freq. dir. (Hz)	1505.5
SPIR offset act./default (Hz)	220 [220]
Local torso SAR	< 87 %
Whole body SAR / level	< 2.8 W/kg / 1st level
SED	< 0.9 kJ/kg
Cool Power	83 %
Max B1+rms	2.13 uT
PNS / level	80 % / normal
dB/dt	43.4 T/s
Sound Pressure Level (dB)	14.34447
Boil-off (hPa/h)	14.94871
Max Boil-off (hPa/h)	20.02934
Boil-off (hPa)	1.387528
MOTION	
Cardiac synchronization device	trigger PPU
R-R Window (%)	10, 20
Number of heart phases	single phase
arrhythmia rejection	no
no trig period (beats)	1
trigger delay	shortest
adaptive	no
cycled MS	no
Heart rate > 250 bpm	no
Slice following	no
REST grid	no
Respiratory compensation	no
Navigator respiratory comp	no
Flow compensation	no
Temporal slice spacing	minimal
NSA	1
MRE enable	no
DYN/ANG	
Manual start	no
Dynamic study	no
dyn stabilization	regular
Arterial Spin labeling	no
POST/PROC	
Preparation phases	full
Interactive F0	no
SmartPlan survey	no
B0 field map	no
B1 field map	no
MIP/MPR	no
Images	M, no, no, no
Autoview image	M
Calculated images	no, no, no, no
Reference tissue	Grey matter
Recon compression	No
Preset window contrast	soft
Reconstruction mode	immediate
Save raw data	no
Hardcopy protocol	no
Image filter	weak
Uniformity correction	no
Geometry correction	default

GEOMETRY	
Nucleus	H1
Uniformity	CLEAR
FOV RL (mm)	56
AP (mm)	42
FH (mm)	75
ACQ voxel size RL (mm)	0.875
AP (mm)	0.88
Slice thickness (mm)	5
Recon voxel size RL (mm)	0.875
AP (mm)	0.875
Fold-over suppression	zoom
Reconstruction matrix	64
SENSE	no
MB SENSE	no
k-t BLAST	no
Stacks	1
type	parallel
slices	15
slice gap	user defined
gap (mm)	0
slice orientation	transverse
flood-over direction	AP
fat shift direction	P
Minimum number of packages	2
Slice scan order	interleaved
PlanAlign	no
REST slabs	0
Catheter tracking	no
Interactive positioning	no
Allow table movement	no
OFFC/ANG	
Stacks	1
Stack Offc. AP (P=+mm)	4.959927
RL (L=+mm)	3.967938
FH (H=+mm)	47.61523
Ang. AP (deg)	-2.273305
RL (deg)	0
FH (deg)	0
Free rotatable	no
Shim Size AP (mm)	49.5138
RL (mm)	63.25952
FH (mm)	95
Offc. AP (P=+mm)	3.729211
RL (L=+mm)	5.026229
FH (H=+mm)	47.48476
Ang. AP (deg)	-2.272193
RL (deg)	0.1920719
FH (deg)	0.007537305

CONTRAST	
Scan type	Imaging
Scan mode	MS
technique	SE
Modified SE	no
Acquisition mode	cartesian
Fast Imaging mode	EPI
shot mode	single-shot
Echoes	1
partial echo	no
TE	shortest
Flip angle (deg)	90
TR (beats)	4
Halfscan	yes
factor	0.7450981
Water-fat shift	minimum
RF Shims	fixed
Shim	PB-volume
ShimAlign	no
mDIXON	no
Fat suppression	SPIR
strength	strong
frequency offset	user defined
offset (Hz)	220
Grad Rev Fat suppr	no
Water suppression	no
BB pulse	no
MTC	no
Research prepulse	no
MDME	no
Diffusion mode	DTI
sequence	SE
gradient duration	maximum
gradient overplus	no
directional resolution	opt 32
nr of b-factors	2
b-factor order	ascending
max b-factor	800
average high b	user defined
b-factor averages	(0) 5, (800) 1,, ..... .....
T1 mapping	no
Transmit channels	both
SAR mode	high
B1 mode	default
SAR allow first level	yes
PNS mode	moderate
Gradient mode	enhanced
SoftTone mode	no

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 InBox (1) | spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25.06.7 | GRE-MT 04:20.4

INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	04:20.4	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	3D
Act. TR/TE (ms)	57 / 2.0	FOV RL (mm)	230	technique	FFE
Dyn. scan time	02:10.1	AP (mm)	230	loop order	zy_order
Time to k0	01:05.0	FH (mm)	110	Contrast enhancement	T1
ACQ matrix M x P	256 x 255	ACQ voxel size RL (mm)	0.9	Acquisition mode	cartesian
ACQ voxel MPS (mm)	0.90 / 0.90 / 5.00	AP (mm)	0.8984375	Fast Imaging mode	none
REC voxel MPS (mm)	0.90 / 0.90 / 5.00	FH (mm)	5	3D non-selective	no
Scan percentage (%)	100	Recon voxel size RL (mm)	0.8984375	Echoes	1
Act. slice gap (mm)	0	AP (mm)	0.8984375	partial echo	yes
Act. WFS (pix) / BW (Hz)	1.663 / 261.1	FH (mm)	5	shifted echo	no
Min. WFS (pix) / Max. BW (Hz)	0.507 / 856.6	Fold-over suppression	no	TE	user defined
Min. TR/TE (ms)	51 / 1.54	Slice oversampling	user defined	(ms)	2
Local torso SAR	< 90 %	oversample factor	1.15	Flip angle (deg)	9
Whole body SAR / level	< 2.9 W/kg / 1st level	RF select. FOS	no	TR	user defined
SED	< 0.8 kJ/kg	ENCASE enable	no	(ms)	57
Coil Power	86 %	Reconstruction matrix	256	Halfscan	no
Max B1+rms	2.17 uT	SENSE	yes	Water-fat shift	user defined
PNS / level	54 % / normal	P reduction (AP)	2.1	(pixels)	1.66
dB/dt	89.0 T/s	S reduction (FH)	1	RF Shims	fixed
Sound Pressure Level (dB)	15.59372	k-t BLAST	no	Shim	PB-volume
Boil-off (hPa/h)	1.295321	Stacks	1	ShimAlign	no
Max Boil-off (hPa/h)	1.295321	slices	22	mDIXON	no
Boil-off (hPa)	0.09368623	slice orientation	transverse	Fat suppression	no
<b>MOTION</b>		fold-over direction	AP	Water suppression	no
Cardiac synchronization	no	fat shift direction	L	MTC	off resonance
Heart rate > 250 bpm	no	Multi-chunk	no	Research prepulse	no
Respiratory compensation	no	PlanAlign	no	MDME	no
Navigator respiratory comp	no	REST slabs	0	Diffusion mode	no
Flow compensation	yes	Catheter tracking	no	T1 mapping	no
fMRI echo stabilisation	no	Interactive positioning	no	Transmit channels	both
NSA	1	Allow table movement	no	SAR mode	high
MRE enable	no	<b>OFFC/ANG</b>		B1 mode	user defined
<b>DYN/ANG</b>		Stacks	1	amplitude (uT)	9.5
Angio / Contrast enh.	no	Stack Offc. AP (P=+mm)	4.959927	SAR allow first level	yes
Quantitative flow	no	RL (L=+mm)	3.967938	PNS mode	moderate
Manual start	no	FH (H=+mm)	47.61523	Gradient mode	maximum
Dynamic study	individual	Ang. AP (deg)	-2.273305	SoftTone mode	no
dyn scans	2	RL (deg)	0		
dyn scan times	shortest	FH (deg)	0		
fov time mode	default	Free rotatable	no		
dummy scans	0	Shim Size AP (mm)	49.5138		
immediate	no	RL (mm)	63.25952		
subtraction		FH (mm)	95		
fast next scan	no	Offc. AP (P=+mm)	3.729211		
synch. ext. device	no	RL (L=+mm)	5.026229		
MTC	yes	FH (H=+mm)	47.48476		
dyn stabilization	no	Ang. AP (deg)	-2.272193		
prospect. motion	no	RL (deg)	0.1920719		
corr.		FH (deg)	0.007537305		
Keyhole	no				
Arterial Spin labeling	no				
<b>POST/PROC</b>					
Preparation phases	full				
Interactive F0	no				
SmartPlan survey	no				
B0 field map	no				
B1 field map	no				
MIP/MPR	no				
SWIp	no				
Images	M, no, no, no				
Autoview image	M				
Calculated images	no, no, no, no				
Reference tissue	Grey matter				
Recon compression	No				
Preset window contrast	soft				
Reconstruction mode	real time				
reuse memory	no				
Save raw data	no				
Hardcopy protocol	no				
Image filter	weak				
Uniformity correction	no				
Geometry correction	default				
Elliptical k-space shutter	default				

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 InBox (1) | spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25.06.7 | GRE-T1 00:34.3

INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	00:34.3	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	3D
Act. TR/TE (ms)	15 / 2.0	FOV RL (mm)	230	technique	FFE
ACQ matrix M x P	256 x 255	AP (mm)	230	loop order	zy_order
ACQ voxel MPS (mm)	0.90 / 0.90 / 5.00	FH (mm)	110	Contrast enhancement	T1
REC voxel MPS (mm)	0.90 / 0.90 / 5.00	ACQ voxel size RL (mm)	0.9	Acquisition mode	cartesian
Scan percentage (%)	100	AP (mm)	0.8988745	Fast Imaging mode	none
Act. slice gap (mm)	0	FH (mm)	5	3D non-selective	no
Act. WFS (pix) / BW (Hz)	1.663 / 261.1	Recon voxel size RL (mm)	0.8984375	Echoes	1
Min. WFS (pix) / Max. BW (Hz)	0.507 / 856.6	AP (mm)	0.8984375	partial echo	yes
Min. TR/TE (ms)	5.1 / 1.59	FH (mm)	5	shifted echo	no
Local torso SAR	< 11 %	Fold-over suppression	no	TE	user defined
Whole body SAR / level	< 0.3 W/kg / normal	Slice oversampling	user defined	(ms)	2
SED	0.0 kJ/kg	oversample factor	1.15	Flip angle (deg)	15
Coil Power	10 %	RF select. FOS	no	TR	user defined
Max B1+rms	0.75 uT	ENCASE enable	no	(ms)	15
PNS / level	52 % / normal	Reconstruction matrix	256	Halfscan	no
dB/dt	77.5 T/s	SENSE	yes	Water-fat shift	user defined
Sound Pressure Level (dB)	15.97592	P reduction (AP)	2.1	(pixels)	1.66
Boil-off (hPa/h)	2.246981	S reduction (FH)	1	RF Shims	fixed
Max Boil-off (hPa/h)	2.246981	k-t BLAST	no	Shim	PB-volume
Boil-off (hPa)	0.02140249	Stacks	1	ShimAlign	no
MOTION		slices	22	mDIXON	no
Cardiac synchronization	no	slice orientation	transverse	Fat suppression	no
Heart rate > 250 bpm	no	fold-over direction	AP	Water suppression	no
Respiratory compensation	no	fat shift direction	L	MTC	no
Navigator respiratory comp	no	Multi-chunk	no	Research prepulse	no
Flow compensation	yes	PlanAlign	no	MDME	no
fMRI echo stabilisation	no	REST slabs	0	Diffusion mode	no
NSA	1	Catheter tracking	no	T1 mapping	no
MRE enable	no	Interactive positioning	no	Transmit channels	both
DYN/ANG		Allow table movement	no	SAR mode	high
Angio / Contrast enh.	no	OFFC/ANG		B1 mode	user defined
Quantitative flow	no	Stacks	1	amplitude (uT)	9.5
Manual start	no	Stack Offc. AP (P=+mm)	4.959927	SAR allow first level	yes
Dynamic study	no	RL (L=+mm)	3.967938	PNS mode	moderate
Arterial Spin labeling	no	FH (H=+mm)	47.61523	Gradient mode	maximum
POST/PROC		Ang. AP (deg)	-2.273305	SoftTone mode	no
Preparation phases	auto	RL (deg)	0		
Interactive F0	no	FH (deg)	0		
SmartPlan survey	no	Free rotatable	no		
B0 field map	no	Shim Size AP (mm)	49.5138		
B1 field map	no	RL (mm)	63.25952		
MIP/MPR	no	FH (mm)	95		
SWIp	no	Offc. AP (P=+mm)	3.729211		
Images	M, no, no, no	RL (L=+mm)	5.026229		
Autoview image	M	FH (H=+mm)	47.48476		
Calculated images	no, no, no, no	Ang. AP (deg)	-2.272193		
Reference tissue	Grey matter	RL (deg)	0.1920719		
Recon compression	No	FH (deg)	0.007537305		
Preset window contrast	soft				
Reconstruction mode	real time				
Save raw data	no				
Hardcopy protocol	no				
Image filter	weak				
Uniformity correction	no				
Geometry correction	default				
Elliptical k-space shutter	default				

Hospital (1) | {B1860723-4F8F-476e-8075-D42C65706693} (1) |  
 InBox (1) | spine\_generic\_philips\_R53\_2020\_09\_29 (7) 25.06.7 | GRE-ME 04:30.6

INFO PAGE		GEOMETRY		CONTRAST	
Total scan duration	04:30.6	Nucleus	H1	Scan type	Imaging
Rel. SNR	1	Uniformity	CLEAR	Scan mode	MS
Act. TR/TE1/delta TE (ms)	600 / 7.2 / 6.5	FOV RL (mm)	224	technique	FFE
ACQ matrix M x P	448 x 448	AP (mm)	224	Contrast enhancement	no
ACQ voxel MPS (mm)	0.50 / 0.50 / 5.00	FH (mm)	75	Acquisition mode	cartesian
REC voxel MPS (mm)	0.50 / 0.50 / 5.00	ACQ voxel size RL (mm)	0.5	Fast Imaging mode	none
Scan percentage (%)	100	AP (mm)	0.5	Echoes	3
Packages	1	Slice thickness (mm)	5	partial echo	no
Min. slice gap (mm)	0	Recon voxel size RL (mm)	0.5	shifted echo	no
Act. WFS (pix) / BW (Hz)	1.704 / 254.8	AP (mm)	0.5	TE first	user defined
Min. WFS (pix) / Max. BW (Hz)	0.911 / 476.9	Fold-over suppression	no	(ms)	7.2
Min. TR/TE1/delta TE (ms)	455 / 6.1 / 6.4	Reconstruction matrix	448	echospace	user defined
Local torso SAR	< 63 %	SENSE	yes	(ms)	6.5
Whole body SAR / level	< 2.0 W/kg / normal	P reduction (AP)	2	flyback	yes
SED	< 0.6 kJ/kg	k-t BLAST	no	Flip angle (deg)	30
Coil Power	61 %	Stacks	1	TR	user defined
Max B1+rms	1.82 uT	type	parallel	(ms)	600
PNS / level	79 % / normal	slices	15	Halfscan	no
dB/dt	102.7 T/s	slice gap	user defined	Water-fat shift	user defined
Sound Pressure Level (dB)	23.0589	gap (mm)	0	(pixels)	1.7
Boil-off (hPa/h)	10.58805	slice orientation	transverse	RF Shims	fixed
Max Boil-off (hPa/h)	10.58805	fold-over direction	AP	Shim	PB-volume
Boil-off (hPa)	0.7958682	fat shift direction	L	ShimAlign	no
<b>MOTION</b>		Minimum number of packages	1	mDIXON	no
Cardiac synchronization	no	Slice scan order	interleaved	Fat suppression	no
Heart rate > 250 bpm	no	PlanAlign	no	Water suppression	no
Respiratory compensation	no	REST slabs	1	MTX	no
Navigator respiratory comp	no	shared	no	Research prepulse	no
Flow compensation	yes	type	free	MDME	no
Temporal slice spacing	default	orientation	coronal	Diffusion mode	no
fMRI echo stabilisation	no	thickness (mm)	80	T1 mapping	no
NSA	2	power	1	Transmit channels	both
SMART	yes	Catheter tracking	no	SAR mode	high
MRE enable	no	Interactive positioning	no	B1 mode	default
<b>DYN/ANG</b>		Allow table movement	no	SAR allow first level	yes
Angio / Contrast enh.	no	<b>OFFC/ANG</b>		PNS mode	moderate
Quantitative flow	no	Stacks	1	Gradient mode	maximum
Manual start	no	Stack Offc. AP (P=+mm)	4.959927	SoftTone mode	no
Dynamic study	no	RL (L=+mm)	3.967938		
Arterial Spin labeling	no	FH (H=+mm)	47.61523		
<b>POST/PROC</b>		Ang. AP (deg)	-2.273305		
Preparation phases	full	RL (deg)	0		
Interactive F0	no	FH (deg)	0		
SmartPlan survey	no	Free rotatable	no		
B0 field map	no	Rest Offc. AP (P=+mm)	-95.04007		
B1 field map	no	RL (L=+mm)	3.967938		
MIP/MPR	no	FH (H=+mm)	47.61523		
SWIp	no	Ang. AP (deg)	-2.273305		
Images	M, no, no, no	RL (deg)	0		
Autoview image	M	FH (deg)	0		
Calculated images	no, no, no, no	Shim Size AP (mm)	49.5138		
Reference tissue	Grey matter	RL (mm)	63.25952		
Recon compression	No	FH (mm)	95		
Preset window contrast	soft	Offc. AP (P=+mm)	3.729211		
Reconstruction mode	real time	RL (L=+mm)	5.026229		
Save raw data	no	FH (H=+mm)	47.48476		
Hardcopy protocol	no	Ang. AP (deg)	-2.272193		
Image filter	weak	RL (deg)	0.1920719		
Uniformity correction	no	FH (deg)	0.007537305		
Geometry correction	default				