



Data Collection Worksheet

Please Note: The Data Collection Worksheet (DCW) is a tool to aid integration of a PhenX protocol into a study. The PhenX DCW is not designed to be a data collection instrument. Investigators will need to decide the best way to collect data for the PhenX protocol in their study. Variables captured in the DCW, along with variable names and unique PhenX variable identifiers, are included in the PhenX Data Dictionary (DD) files.

TA: 6:03

PAT: 2

Voxel size: 1.0 x 1.0 x 1.0 mm

Rel. SNR: 1.00

Properties

Prio Recon	Off
Before measurement	Off
After measurement	Off
Load to viewer	On
Inline movie	Off
Auto store images	On
Load to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Load to stamp segments	Off
Load images to graphic segments	Off
Auto open inline display	Off
Start measurement without further	On

preparation
 Wait for user to On
 start
 Start single
 measurements

Routine

Slab group 1
 Slab 1
 Dist. 50%
 Factor L0.0
 Position A14.2 H0.7
 Orientation S > C0.2
 Phase enc. A >> P
 dir. 0.00 deg
 Rotation
 Phase 0%
 oversampling
 Slice 0.0%
 oversampling
 Slices per slab 176
 FoV read 256 mm
 FoV phase 100.0%
 Slice thickness 1.00 mm
 TR 2530 ms
 TE 1 3.31 ms
 TE 2 6.99 ms
 Averages 1
 Concatenations 1
 Filter Prescan
 Coil elements Normalize
 HEA;HEP

Contrast

Magn. preparation Non-sel.
 preparation IR
 TI 1100 ms
 Flip angle 7.0 deg
 Fat suppr. None

Water suppr.	None
Averaging	Long term mode
Reconstruction	Magnitude
Measurements	1
Multiple series	Each measurement

Resolution

Base resolution	256
Phase resolution	100%
Slice resolution	100%
Phase partial Fourier	Off
Slice partial Fourier	Off
Interpolation	Off
PAT mode	GRAPPA
Accel. factor PE	2
Ref. lines PE	32
Matrix Coil Mode	Auto (Triple)
Reference scan mode	Integrated
Image Filter	Off
Distortion Corr.	Off
Unfiltered images	Off
Prescan Normalize	On
Normalize	Off
B1 filter	Off
Raw filter	Off
Elliptical filter	Off

Geometry

Multi-slice	Single shot mode
Series	Interleaved

System

Body	Off
HEP	On
HEA	On
Positioning mode	FIX
Table position	H
Table position	0 mm
MSMA	S - C - T
Sagittal	R >> L
Coronal	A >> P
Transversal	F >> H
Save uncombined	Off
Coil	Adaptive
Combine Mode	Combine
AutoAlign	Head > Brain
Auto Coil Select	Default
Shim mode	Standard
Adjust with body coil	Off
Confirm freq. adjustment	Off
Assume Silicone	Off
? Ref. amplitude 1H	0.000 V
Adjustment Tolerance	Auto
Adjust volume	
Position	L0.0

A14.2 H0.7

Orientation	S > C0.2
Rotation	0.00 deg
F >> H	256 mm
A >> P	256 mm
R >> L	176 mm

Physio

1st	None
Signal/Mode	
Dark blood	Off

Inline

Subtract	Off
Std-Dev-Sag	Off
Std-Dev-Cor	Off
Std-Dev-Tra	Off
Std-Dev-Time	Off
MIP-Sag	Off
MIP-Cor	Off
MIP-Tra	Off
MIP-Time	Off
Save original images	On

Sequence

Introduction	On
Dimension	3D
Elliptical scanning	Off
Asymmetric echo	Off
Contrasts	2
Bandwidth 1	195 Hz/Px
Bandwidth 2	651

	Hz/Px
Flow comp. 1No
Flow comp. 2No
Echo spacing10.1 ms
RF pulse typeFast
Gradient modeFast
ExcitationNon-sel.
RF spoilingOn
Readout polarityPositive
Readout trajectoryBipolar
Add. scale factor8.0
Gradient spoilingIntegral
Gradient moment factor3.0
Siemens reconstructionOn
Save raw k-space dataOff
AveragingRMS

Protocol source: <https://www.phenxtoolkit.org/protocols/view/660501>