Diffusion MRI (dMRI) Sequence Settings

In this article, we will review the settings of the diffusion (dMRI) MRI sequence. Spin Echo (SE) Echo Planer Imaging (EPI) single shot (SS) sequence is mainly used for dMRI acquisition by the PBS researchers. We have also installed the same sequence, but highly customizable WIP by CMRR also. Thus, we have the generic Siemens' version and the versatile CMRR version of the same sequence. With CMRR dMRI sequence, we can acquire diffusion data with high accuracy b-values and multi-shell data.



Figure 1: SE-EPI SS sequence diagram

The SE-EPI-SS diffusion sequence is ultra-fast and can acquire the whole brain within one repetition time (TR) for a given diffusion direction. We will lay out important parameters in the sequence and show how someone would see it on the sequence card at the console computer.

Number of slices, FOV

In general, field of view (FOV) and number of slices are chosen to cover the interested region of the brain. Researchers usually acquire the whole brain. The number of slices depends on the slice thickness and the slice gap (if any) to cover the interested region. In the case of simultaneous slice acquisition, the number of slices must be divisible to an integer by the multi-band acceleration factor (MB or SMA).

TA: 8:01 PM: FIX	PAT: 2 Vox	el size: 1.7×1.7×1.7mm Re	I. SNR: 0.71 : epse
Routine Contrast Re	solution Geometry	System Physio Diff	Sequence
	Common AutoAlign	Saturation Navigator	
Slice group	1 -	FoV re	ad 240 📜 mm
Slices	81 📜	FoV pha	ise 100.0 📜 %
Dist. factor	0 📜 %	Slice thickne	ess 1.70 📮 mm
Position Isocent	er		TR 4110 🗧 ms
Orientation Transve	ersal 🔹	Multi-slice mode	Interleaved
Phase enc. dir.	A >> P •	Series	Interleaved -
Phase oversampling	0 📜 %	Multi-band accel. fac	tor 3 ÷

Figure 2: Geometry - slices, slice gap (Dist. factor), slice thickness, FOV

Echo time (TE) and Repetition Time (TR)

Echo Time (TE) is set to the minimum for dMRI scans. Value of TR depends on the number of slices and the simultaneous multiband factor (SMA/MB). TR can also be set to the minimum value the sequence

can take. The SMA factor is equal to how many slices the sequence acquires simultaneously during each TR.

TA: 7:20	PM: FIX	PAT: (Off Vox	el size: 1.7>	<1.7×1.7mm Re	I. SNR: 1.00	: epse
Routine	Contrast	Resolution	Geometry	System	Physio Diff	Sequence	
			Common	Dynami	ic		
		TR 41	10 📫 ms		Fat suppr.	Fat sat.	
		TE 88.	40 📜 ms	Grad	l. rev. fat suppr.	Disabled	
		MT					
Magn.	preparation	None					
	Flip angle		90 📜 deg				

Figure 3: TE, TR,

Flip Angle

For the most experiments, the flip angle is set to 90 degrees for the excitation pulse and 180 degrees for the refocusing pulse.

TA: 7:20	PM: FIX	PAT: C)ff ∨ox	el size: 1.7	×1.7×1.7m	m Re	I. SNR: 1.00	: epse
Routine (Contrast	Resolution	Geometry	System	Physio	Diff	Sequence	
			Common	n Dynam	ic			
		TR 41'	10 📫 ms		Fat s	uppr.	Fat sat.	
		TE 88.4	40 📜 ms	Grad	d. rev. fat s	uppr.	Disabled	
		МТО						
Magn. pi	reparation	None						
F	lip angle	•	90 📜 deg					

Figure 4: Flip Angle

Acceleration Techniques: GRAPPA, multiband factor (MB/SMA) and partial Fourier factor

GRAPPA is the in-plane acceleration factor; GRAPPA of 2 will reduce scan time almost by half. Multiband factor (SMA) is the slice acceleration factor; SMA of 4 will reduce scan time almost by ¼. Partial Fourier cuts back some of the k-space lines along Phase Encoding direction to accelerate acquisition; if used, most studies use factor of 6/8. We can apply all these acceleration techniques simultaneously if needed with a compromise to the image quality. Most studies use GRAPPA of 2 with SMA of 2 or 4. Some researchers just use SMA of 8 only. With SMA, some researchers also save reference scans for post-processing.

TA: 7:20) PM: FIX	PAT:	Off Vox	kel size: 1.7×1	.7×1.7mm	n Rel. Sl	NR: 1.00	: epse
Routine	Contrast	Resolution	Geometry	System	Physio	Diff S	equence	
	Slice gl	тоир	1 🔻 + -		F	oV read	240	mm
	S	lices	81 📜		Fo	∨ phase	100.0	%
	Dist. fa	nctor	0 📜 %		Slice ti	hickness	1.70	mm
F	Position Iso	center				TR	4110	ms
Orie	entation Tra	insversal				TE	88.40	ms
	Phase enc.	<i>dir.</i> A >> P						
				Multi-I	band acce	el. factor	3	•
Pha	se oversamp	ling	0 📜 %		Filter	None		
				Coil e	elements	HEA;HE		

TA: 8:01 PM: FIX	PAT: 2	Voxel size: 1.7×1	1.7×1.7mm Rel	. SNR: 0.71	: epse
Routine Contrast	Resolution Geo	metry System	Physio Diff	Sequence	
	Common iPA	T Filter Image	Filter Rawdata		
PAT mode	GRAPPA •				
Accel. factor	PE 2 -	Referen	nce scan mode	Single-shot	
Ref. lines	PE 24 ÷				

TA: 7:20 PM: FIX	Pat: (Off Vo	xel size: 1.7;	×1.7×1.7mm	ı Rel.	SNR: 1.00	: epse
Routine Contrast	Resolution	Geometry	System	Physio	Diff	Sequence	
	Common	ipat f	ilter Image	Filter Rav	vdata		
FoV re	ead 2	40 📜 mm					
FoV pha	ase 100	0.0 🚆 %					
Slice thickn	<i>ess</i> 1.	70 📜 mm					
Base resolut	tion 1	40 📜					
Phase resolut	tion 1	00 📜 %					
Phase partial Fou	rier 6/8						
						Interpolation	

Figure 5: GRAPPA, MB/SMA, pF

Diffusion Gradient Schemes: Diffusion Mode

If you are planning to do to Diffusion Tensor Imaging, you can choose 'MDDW' diffusion mode and select the desired number of diffusion gradient directions and a value for the b-value. For DTI, b-value of 1000 is common. If you need to acquire multi-shell diffusion data, a diffusion gradient scheme (diffusion vector set, DVS) must be imported by choosing 'Free' option. This .dvs file contains diffusion vectors for your experiment and your Physicist can help you get created one and set up the examcard.



TA: 8:01 PM: FIX	PAT: 2	Voxel size: 1.7	′×1.7×1.7mm Re	I. SNR: 0.71	: epse
Routine Contrast	Resolution Geon	netry System	Physio Diff	Sequence	
	Neuro	Body Com	posing		
Diffusion mode	Free •		Diff. we	ighted images	
Diff. directions	102		Trace we	ighted images	
Diffusion Scheme	Monopolar -			ADC maps	
Diff. weightings	2			FA maps	
b-Value	Averages			Mosaic	<u>v</u>
0 ÷ 3000 ÷				Tensor	
Noise	evel 40 ÷				

Diffusion vector set - Import/Export Use Import to load an external *.dvs file. Import Use Export to save the loaded vector set to a *.dvs file. Funct							
Export			Close				
Diffusion mode	Free	Diff. weighted images	<u>v</u>				
Diff. directions	102	Trace weighted images					
Diffusion Scheme	Monopolar	ADC maps					
Diff. weightings	2	FA maps					
b-Value	Averages	Mosaic	<u>v</u>				
0 3000		Tensor					

Coil Elements Selections

It's necessary and utmost important to make sure the receiver head coil is correctly plugged into the scanner before collecting data. We have three different head coils: 32CH, 64CH and 20CH. You can see if the head coil is correctly plugged on the screen of the scanner itself or check on the examcard tab shown below. Make sure both HEA and HEP parts are plugged in.

TA: 8:0′	1 PM: FIX	PAT: 2	2 Vox	el size: 1.7>	×1.7×1.7r	nm Rel.	. SNR: 0	.71	: epse
Routine	Contrast	Resolution	Geometry	System	Physio	Diff	Seque	nce	
	Coils Mis	cellaneous	Adjustments	Adjust V	'olume	pTx Vol	umes	Tx/Rx	
	HEA								
	HEP								
_									
	Body								

Figure 7: Head coil elements