# New to MRI: simulator to the rescue

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# Laptop ready?





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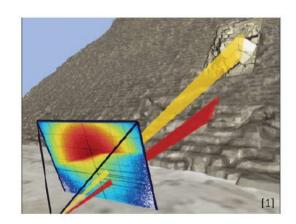




# **Background**

- Nanotechnology
- Muon tomography

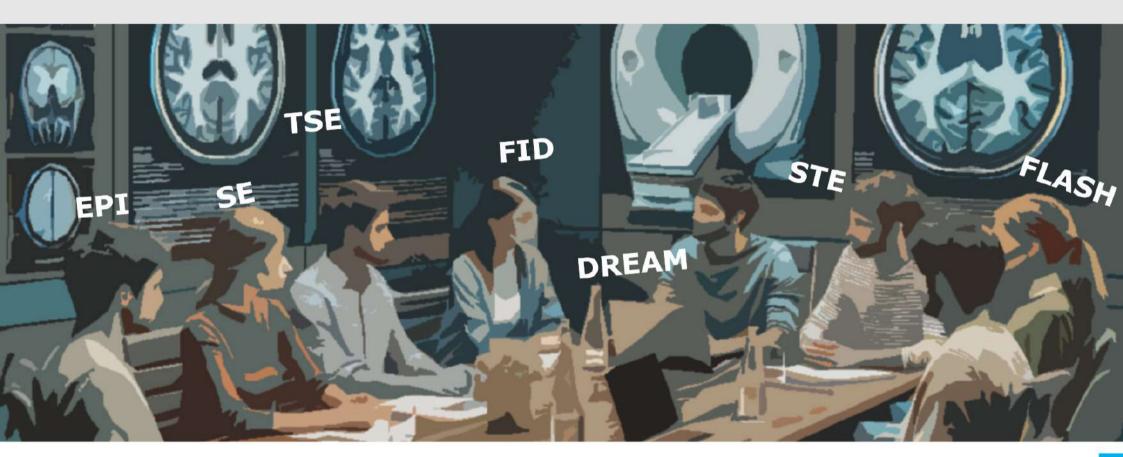








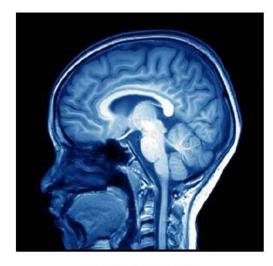
# **Hello MRI**







# **Hello MRI**



**ESCAPE SE** 



**DANTE** 



XÆ A-Xii

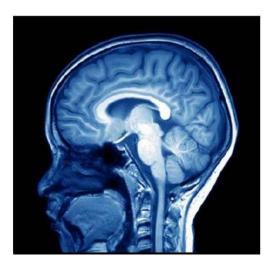




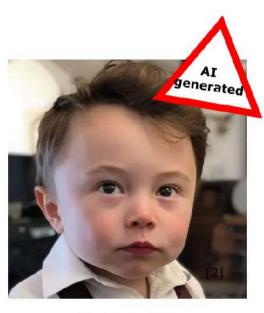
# **Hello MRI**



**ESCAPE SE** 



DANTE



XÆ A-Xii











Basic MRI physics (T1, T2\* relaxation)

**Basic sequences** 

More complicated sequences







MRzero, together with pypulseq, is super easy to use and you can try it yourself

- 03\_full\_acquisition\_simulations/notebooks/New\_to\_MRI\_Simulator\_to\_the\_Rescue/Talk2/
  - ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_Basic\_MRI\_Physics.ipynb
  - ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_ Basic\_MRI\_sequences.ipynb
  - ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_ More\_complicated\_sequences.ipynb
  - ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_Going\_to\_the\_scanner.ipynb
  - ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_HandsOn.ipynb





MRzero, together with pypulseq, is super easy to use and you can try it yourself

03\_full\_acquisition\_simulations/notebooks/New\_to\_MRI\_Simulator\_to\_the\_Rescue/Talk2/

ESMRMB ET03 Mrzero to the rescue HandsOn.ipynb





MRzero, together with pypulseq, is super easy to use and you can try it yourself

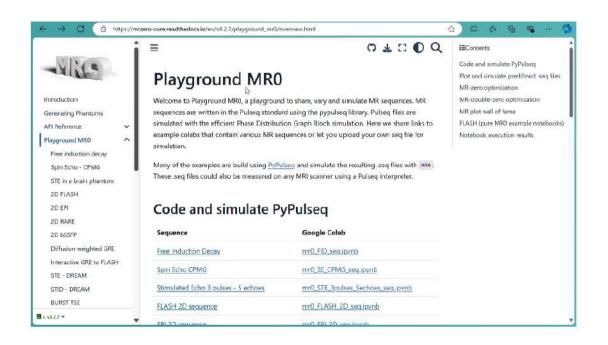
03\_full\_acquisition\_simulations/notebooks/New\_to\_MRI\_Simulator\_to\_the\_Rescue/Talk2/

Let's run it together at the end of the session



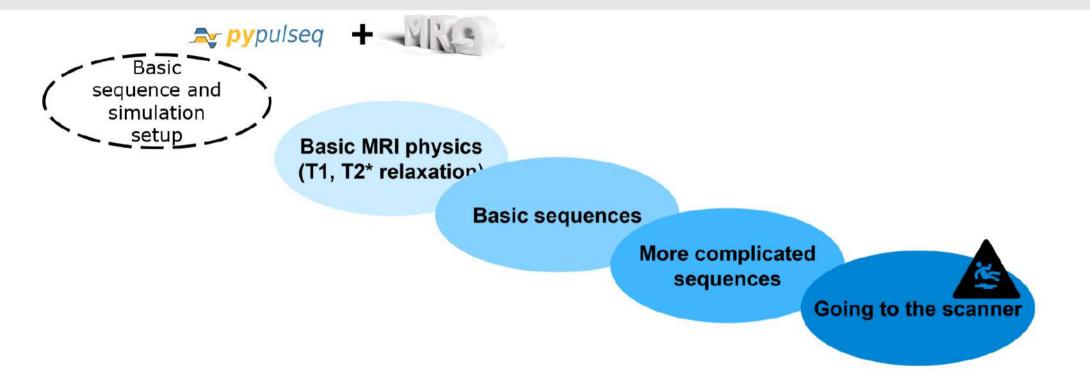


MRzero, together with pypulseq, is super easy to use and you can try it yourself More notebooks on the Playground-MR0 website:







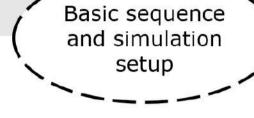






Install libraries

!pip install pypulseq==1.4.2
!pip install MRzeroCore







Prepare sequence and save as .seq file

```
# Build sequence: T2star relaxation
seqT2star = pp.Sequence()
seqT2star.add_block(rf1, gz)
seqT2star.add_block(gzr)
seqT2star.add_block(adc)
aux_check_timing(seqT2star)
seqT2star.write("FID.seq")
```

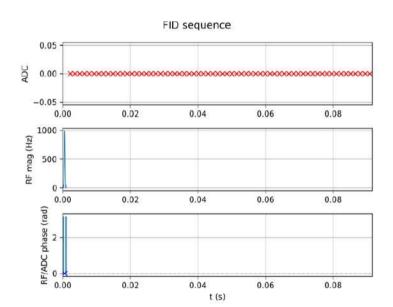


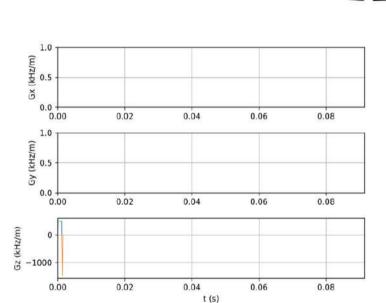






■ Prepare sequence and save as .seq file









Basic sequence and simulation

setup

Load the sequence as an MR-zero sequence object

```
seq0_T2star = mr0.Sequence.import_file('FID.seq')
```

Basic sequence and simulation setup

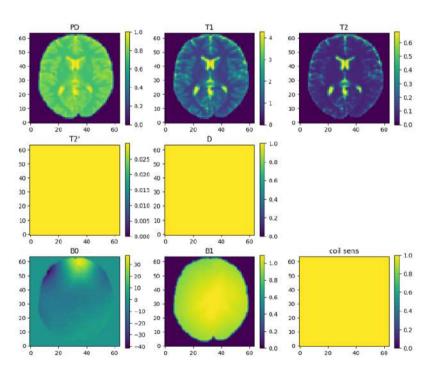


Load a digital phantom

```
sz = [Nread, Nphase]
phantom = mr0.VoxelGridPhantom.load_mat('numerical_brain_cropped.mat')
phantom = phantom.interpolate(sz[0], sz[1], 1)
obj_p = phantom.build()
phantom.plot()
```







Basic sequence and simulation setup





Basic MRI physics (T1, T2\* relaxation)

Simulate signal

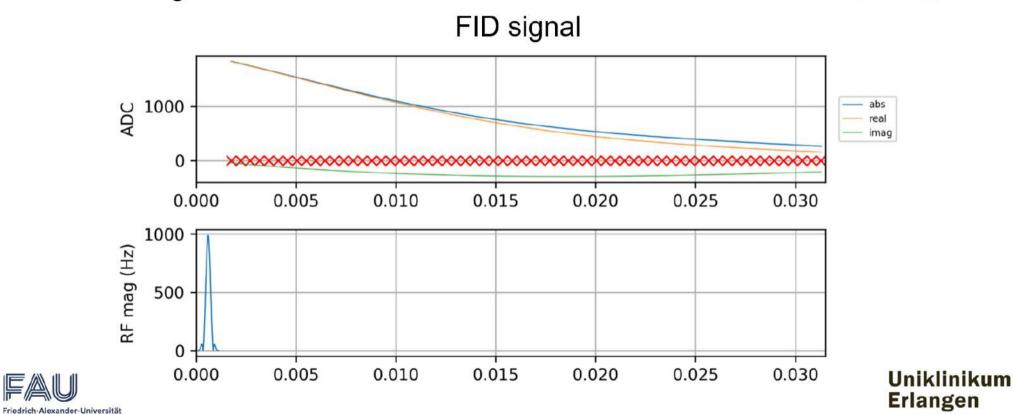
```
graph_T2star = mr0.compute_graph(seq0_T2star, obj_p, 200, 1e-3)
signal_T2star = mr0.execute_graph(graph_T2star, seq0_T2star, obj_p)
```





Basic MRI physics (T1, T2\* relaxation)

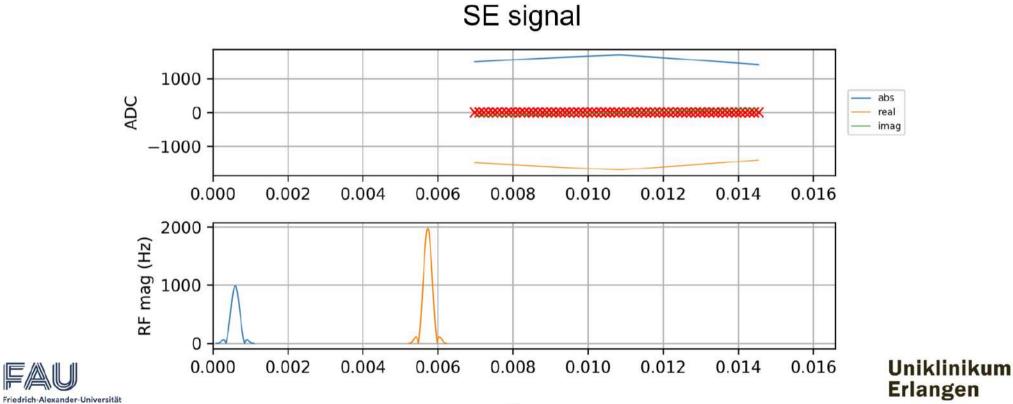
Simulate signal



13

Basic MRI physics (T1, T2\* relaxation)

Simulate signal



**Basic sequences** 

Simulate

```
signal_EPI = mr0.execute_graph(graph_EPI, seq0_EPI, obj_p)
```

Reconstruct image from signal

```
# kspace reordering
kspace_EPI = torch.reshape((signal_EPI), (Nphase, Nread)).clone().t()
kspace_EPI[:,0::2] = torch.flip(kspace_EPI[:,0::2],[0] )[:,:]

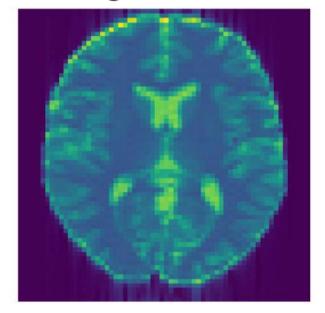
# FFT
spectrum_EPI = torch.fft.fftshift(kspace_EPI)
image_EPI = torch.fft.fft2(spectrum_EPI)
image_EPI = torch.fft.ifftshift(image_EPI)
```





**Basic sequences** 

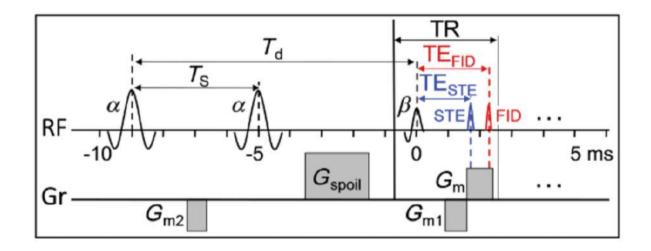
# Magnitude EPI







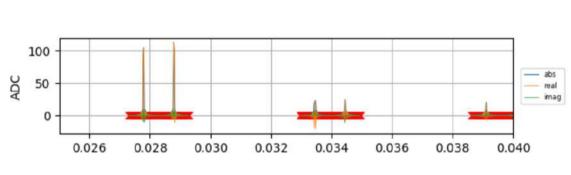
More complicated sequences

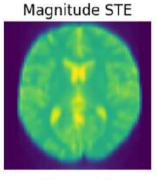


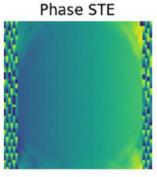




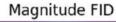
■ DREAM – B1, B0, TxRx mapping sequence

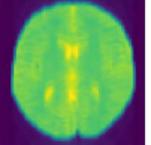




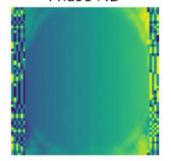


# More complicated sequences





Phase FID

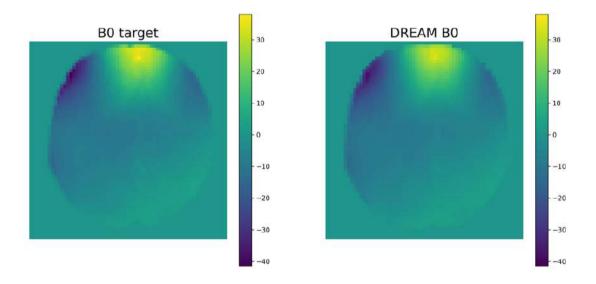






More complicated sequences

■ DREAM – B1, B0, TxRx mapping sequence

















Problem number #1

#### Time to code:

03\_full\_acquisition\_simulations/notebooks/New\_to\_MRI\_Simulator\_t o\_the\_Rescue/Talk2/

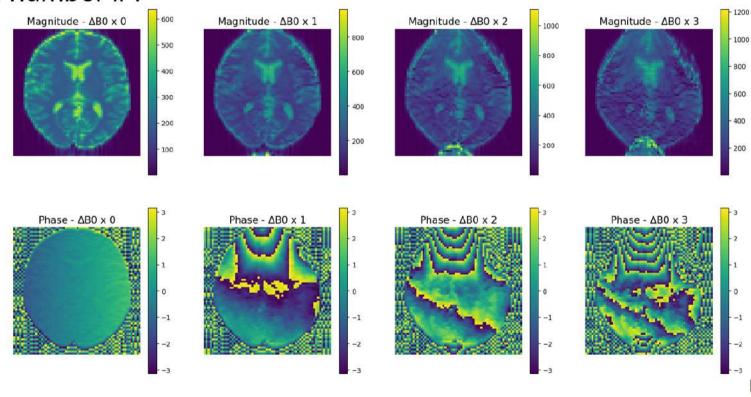
ESMRMB\_ET03\_Mrzero\_to\_the\_rescue\_HandsOn.ipynb







#### ■ Problem number #1





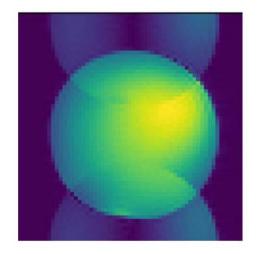
Uniklinikum Erlangen









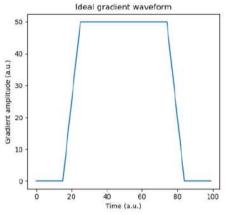


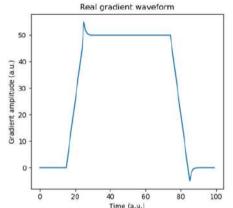








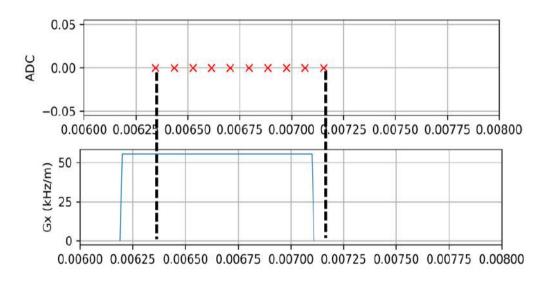


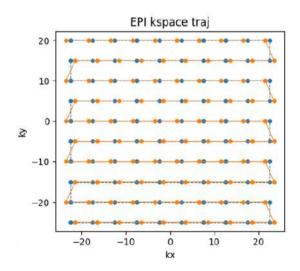






# Going to the scanner

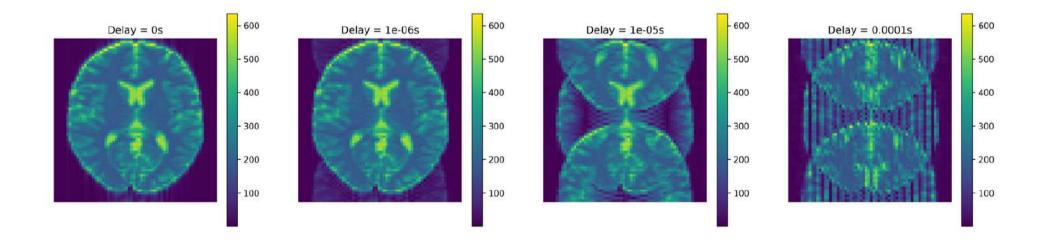








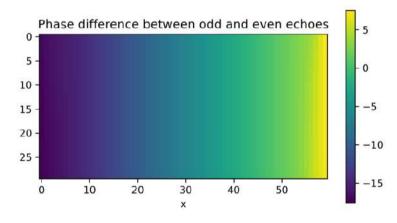












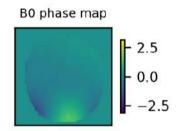
Time delay = linear phase factor

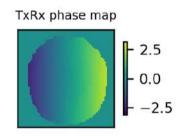


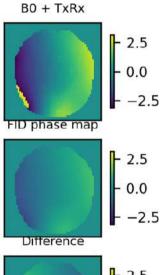


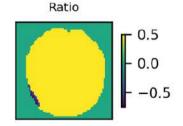


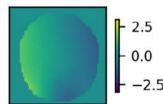
#### ■ Problem number #3









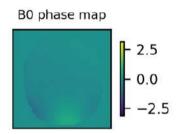


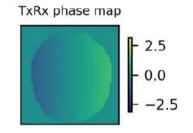


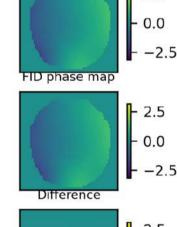
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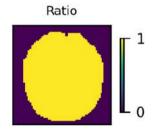
#### ■ Problem number #3

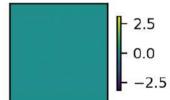






B0 + TxRx





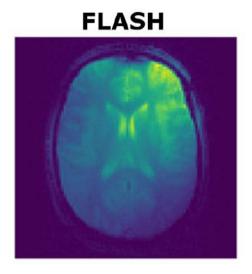


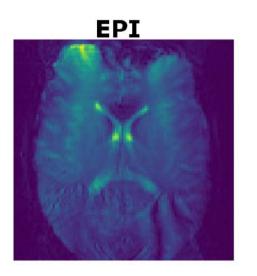




■ Problem number #4

Just a few weeks ago...looking to have the same FOV for a FLASH and EPI

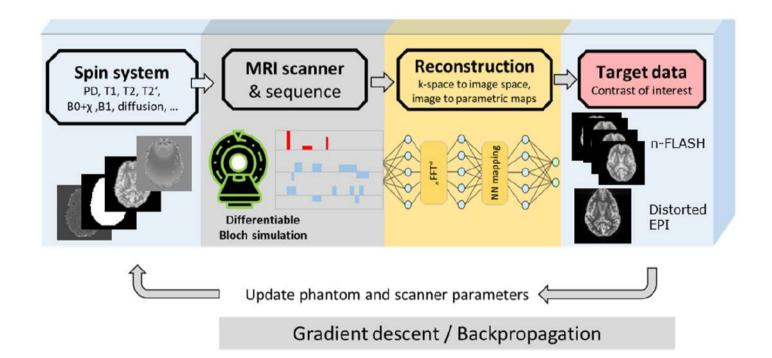








#### You can do this and much more







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- Determine the best readout for transient state
- Test the effect of more dummy pulses and variale flip angles
- Optimize sequences
- Jointly reconstruct different readout images
- Perform parameter quantification
- Optimize kT-points for pTx





#### You can do this and much more

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Visit the UKER posters to learn more

0011, 0112, 0113, 0421, 0422, 0423, 0456, 0483, 0484





#### Conclusion

- MR-zero is a great simulation tool for introduction to MRI.
- Integrated with pypulseq, it allows easy and fast sequence simulation.
- Its full diferentiability allows end-to-end optimization.







#### Conclusion

- There are several opportunities to get started throughout the year:
  - MRTwin course at FAU
  - ISMRM 2023 pulseq course
  - Hands-on MR Physics with Pulseq, Freiburg 2024







#### References

- [1] Morishima, K., Kuno, M., Nishio, A., et al. (2017). Discovery of a big void in Khufu's Pyramid by observation of cosmic-ray muons. Nature, 552, 386–390. <a href="https://doi.org/10.1038/nature24647">https://doi.org/10.1038/nature24647</a>
- [2] Farhat, A. (n.d.). Al-generated image of Elon Musk as a baby. Twitter. Retrieved from <a href="https://twitter.com/alifarhat79">https://twitter.com/alifarhat79</a>
- [3] Endres, J., Weinmüller, S., Dang, H. N., & Zaiss, M. (2024). Phase distribution graphs for fast, differentiable, and spatially encoded Bloch simulations of arbitrary MRI sequences. Magnetic Resonance in Medicine. https://doi.org/10.1002/mrm.30055
- [4] Nehrke, K., Börnert, P. (2012). DREAM—a novel approach for robust, ultrafast, multislice B1 mapping. <a href="https://doi.org/10.1002/mrm.24158">https://doi.org/10.1002/mrm.24158</a>
- [5] Baum, T. <a href="https://mrzero-core.readthedocs.io/en/v0.2.7/playground-mr0/mr0">https://mrzero-core.readthedocs.io/en/v0.2.7/playground-mr0/mr0</a> DREAM STE seq.html#dream-ste-seq



