

ICE and Gadgetron Reconstruction Based on the Pulseq Framework

Qingping Chen

Division of Medical Physics, Dept. Of Radiology, University Medical Center Freiburg, Germany

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Outline

- Workflow for data acquisition and image reconstruction
- How to enable ICE/Gadgetron (based on s21_GRE2D_GRAPPA: https://github.com/pulseq/ISMRM-Virtual-Meeting--November-15-17-2023/tree/main/tutorials/day2_imageReconstruction)
 - ADC labeling & label debugging
 - sequence definitions
 - interface setting
- Example reconstructions
 - MPRAGE with GRAPPA = 2
 - Multi-slice EPI with navigator and ramp sampling



Workflow

- Sequence design in the Pulseq software:
 - ADC labeling
 - sequence definitions (seq.setDefinition(...))
- Extended Pulseq interpreter sequence
 - Load ADC labels and sequence definitions
 - Link to ICE and Gadgetron^{1,2}
- ICE is integrated to Siemens, while Gadgetron is vendor-independent by employing ISMRMRD³ data format (Siemens, GE, Philip, Bruker, etc).

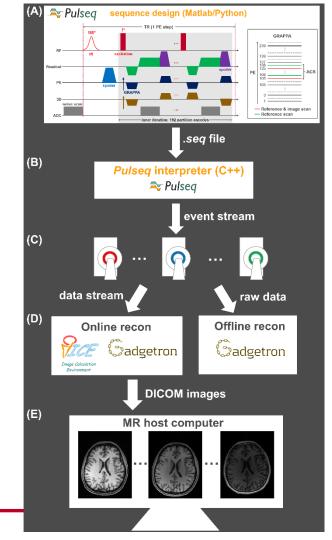
Gadgetron: <a href="https://github.com/gadgetron/gadgetr

ISMRMRD: https://github.com/ismrmrd/ge_to_ismrmrd

IceGadgetron: https://github.com/NHLBI-

MR/ISMRM2019_demo/tree/master/Siemens_Scanner_Setup

1. Hansen, et al., MRM, 2013. 2. Xue, et al., MRM, 2015. 3. Inati, et al. MRM, 2017.



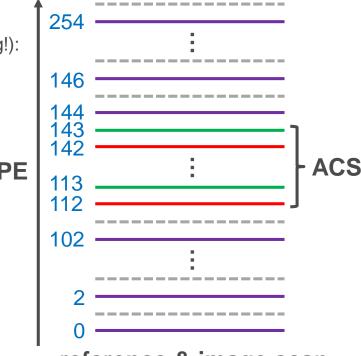
ADC labeling

- Phase-encoded index labeling (caution: c-style numbering!):
 - Index setting

```
mr.makeLabel( 'SET', 'LIN', 0 )
```

- Index increment
 - mr.makeLabel('INC', 'LIN', 2)
- ACS flag labeling:
 - Reference & image scan
 - mr.makeLabel('SET', 'REF', true)
 - mr.makeLabel('SET', 'IMA', true)
 - Reference scan
 - mr.makeLabel('SET', 'REF', true)
 - > mr.makeLabel('SET', 'IMA', false)
 - Image scan
 - mr.makeLabel('SET', 'REF', false)
 - > mr.makeLabel('SET','IMA', false)

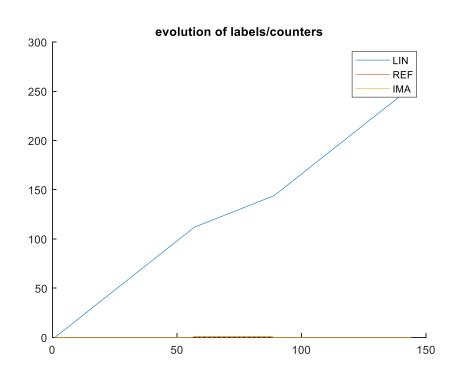
GRAPPA



- reference & image scan
- reference scan
- image scan

ADC debugging

```
adc_lbl =
seq.evalLabels('evolution','adc');
figure ;
hold on ;
plot(adc_lbl.LIN) ;
plot(adc_lbl.REF) ;
plot(adc_lbl.IMA) ;
```

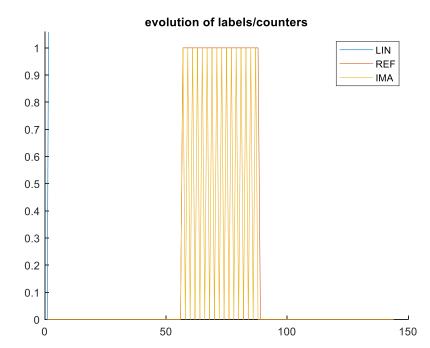


LIN: PE index



ADC debugging

```
adc_lbl =
seq.evalLabels('evolution','adc');
figure ;
hold on ;
plot(adc_lbl.LIN) ;
plot(adc_lbl.REF) ;
plot(adc_lbl.IMA) ;
```



REF: ACS reference scan flag IMA: ACS reference&image scan flag

Sequence definitions

2D multi-slice mode

```
seq.setDefinition('SlicePositions', 0);
seq.setDefinition('SliceThickness', sliceThickness);
seq.setDefinition('SliceGap', sliceGap);
```

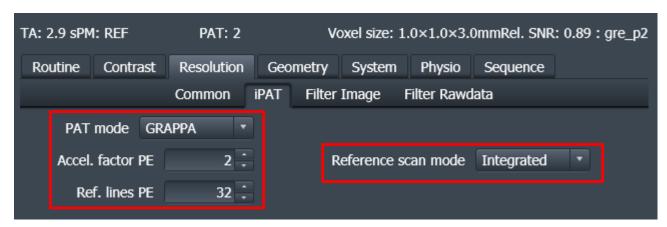
Additional information required by GRAPPA

```
seq.setDefinition('kSpaceCenterLine', centerLineIdx-1);
seq.setDefinition('PhaseResolution', phaseResolution);
```



Interface setting

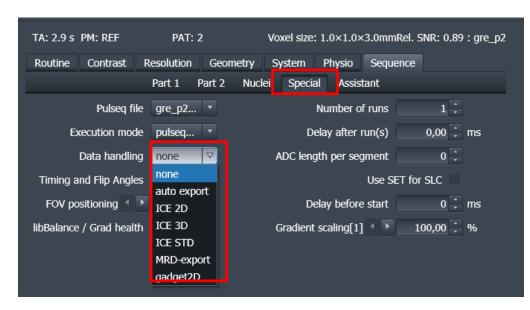
- Turn on iPAT setting: PAT mode
- Set Accel. factor PE, Ref. lines PE, and Reference scan mode
- Caution: GRAPPA mode must be turned on for ICE/Gadgetron recon. Do not forget to turn it off after data acquisition.





Interface setting

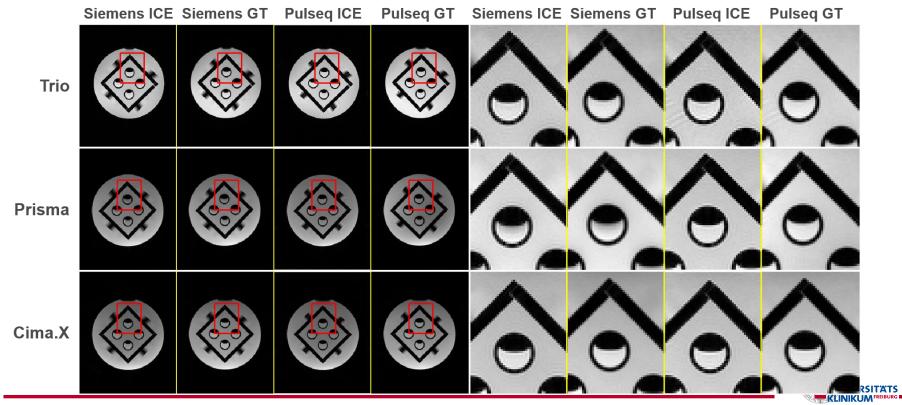
- Select "Data handling"
 - none: no data handling
 - auto export: NIH data catcher
 - ICE 2D: 2D recon (VB, VE, etc)
 - ICE 3D: 3D recon (VB, VE, etc)
 - ICE STD: 2D&3D recon (XA, etc)
 - MRD-export: ISMRMRD¹ data export
 - gadget2D: gadgetron online recon



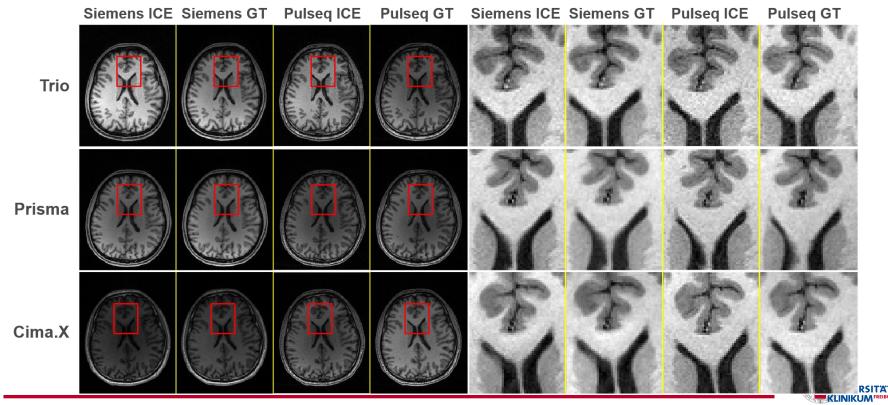
- Caution: IceGadgetron² needs to be installed in the Siemens scanner for online Gadgetron reconstruction.
 - 1. Inati, et al. MRM, 2017.; 2. Xue, et al., MRM, 2015.



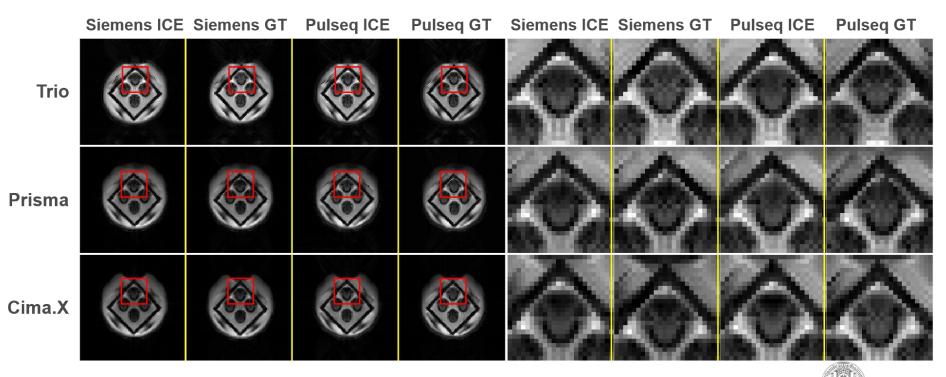
MPRAGE with GRAPPA = 2



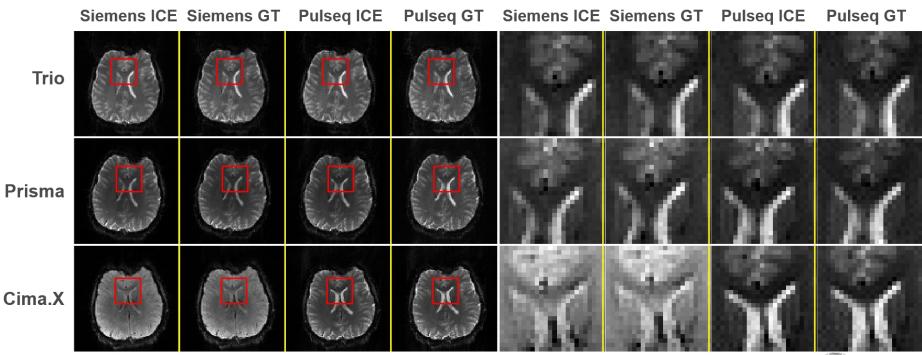
MPRAGE with GRAPPA = 2



Multi-slice EPI with navigator and ramp sampling



Multi-slice EPI with navigator and ramp sampling





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