

Call AI models in Gadgetron



Hui Xue

NHLBI, NIH
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National Heart, Lung,
and Blood Institute

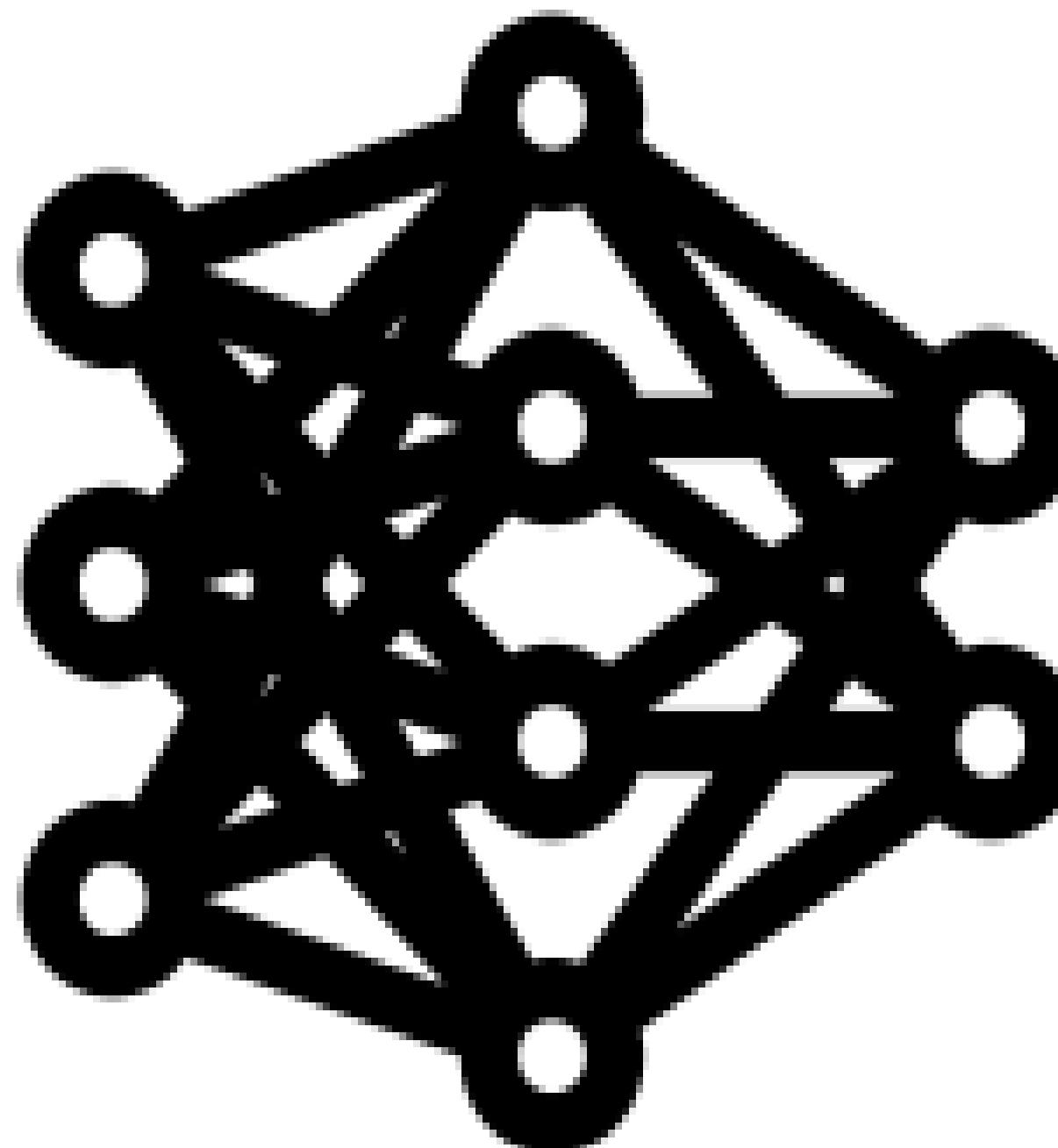
In this session



- **Motivation to bring in AI models**
- **Inline vs. offline**
- **Gadgetron InlineAI**
- **Example: Run Grappa as an AI model**
- **Example: Run cine landmark detection**
- **Demo with inline walk-through**
- **Final remarks**

Why we need to do something to run AI models

- Because vendor provided software lacks support



Trained NN

- ✓ Rich in computing power
- ✓ Flexible and updated software configuration
- ✓ Computing time matters less

Model
inference



Not so easy to
deploy NN

At this moment of 2020

- Limited computing power
- Often outdated software configuration
- Clinical relevant computing time

Inline vs Offline



←
data
Images,
report



Gadgetron
Computing engine

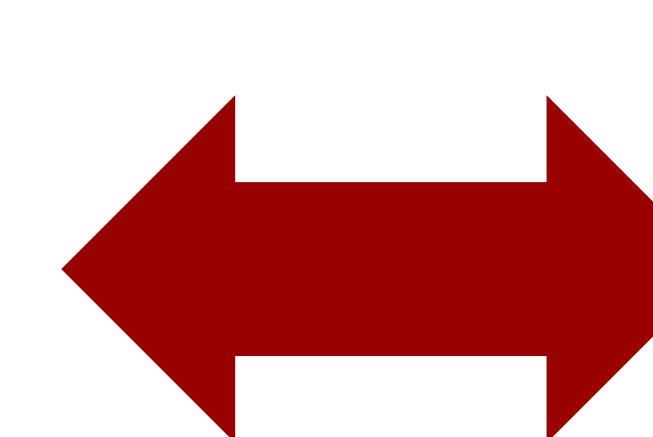
- Automated processing
- End-to-end from imaging to reporting
- Can be duplicated and on cloud



images



Software



Operator

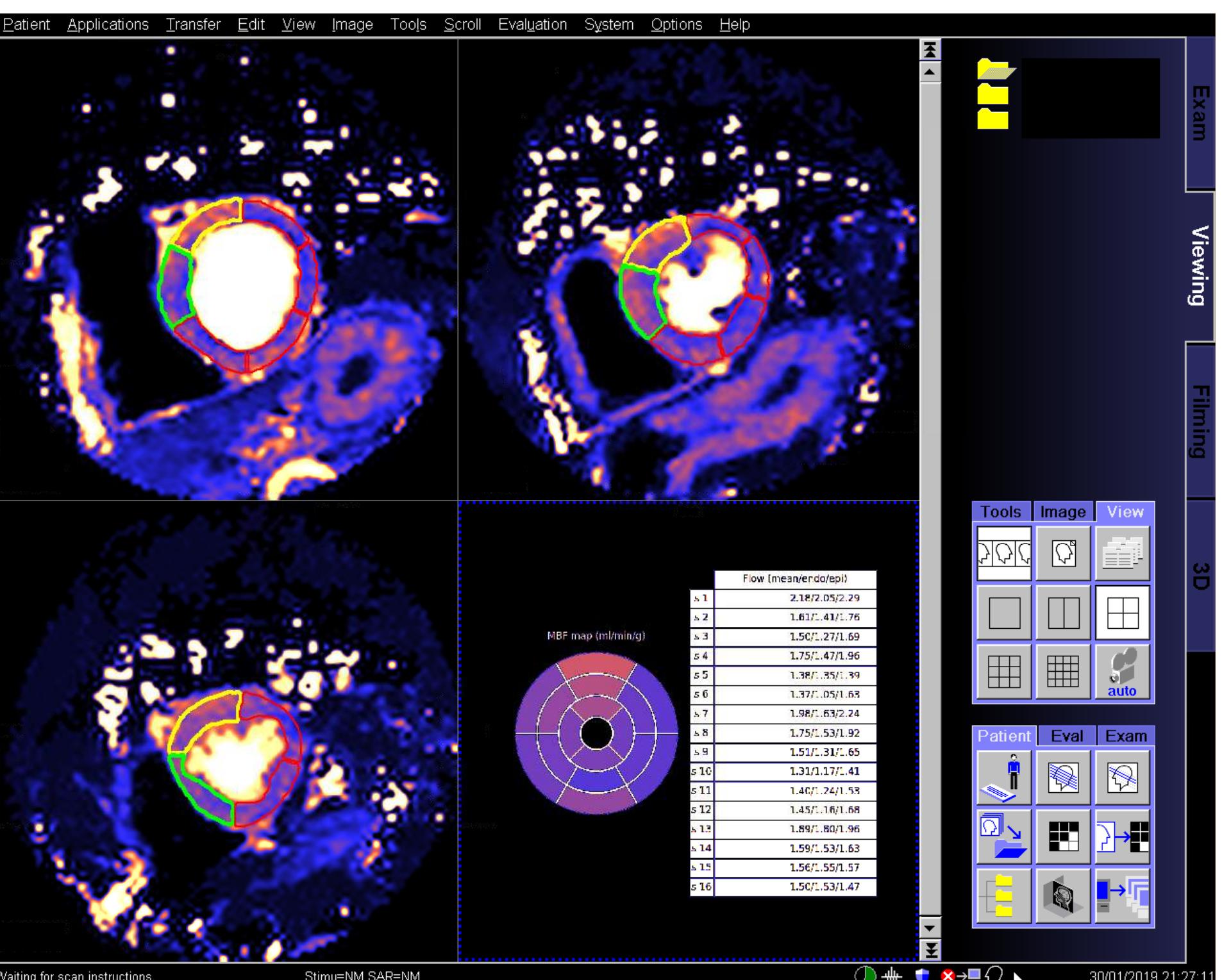
- Interactive processing
- Established workflow
- Expert time

Example of Inline AI: Cine Analysis and reporting



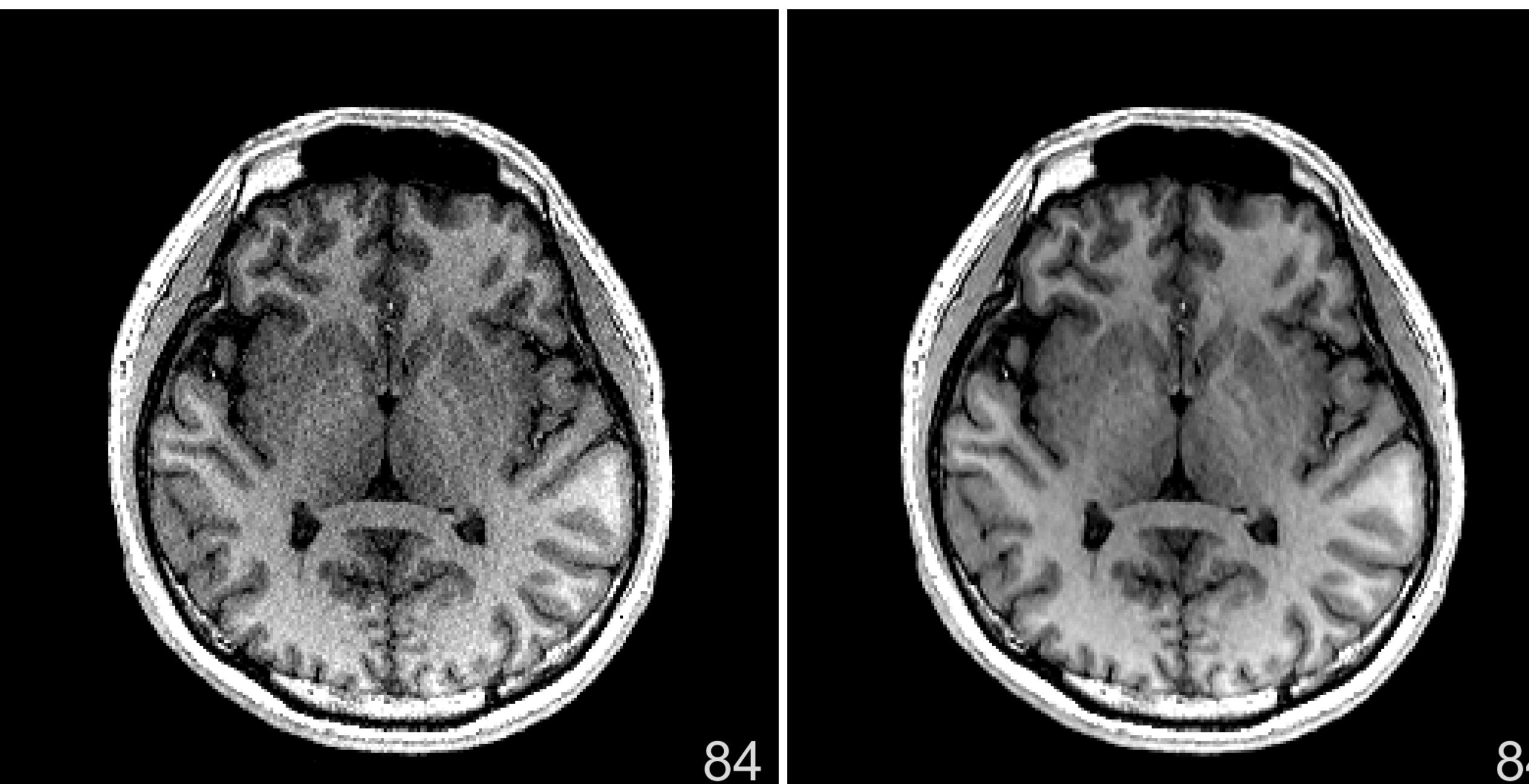
It is the time to do this

- Deep learning - new hope to promote Inline workflow

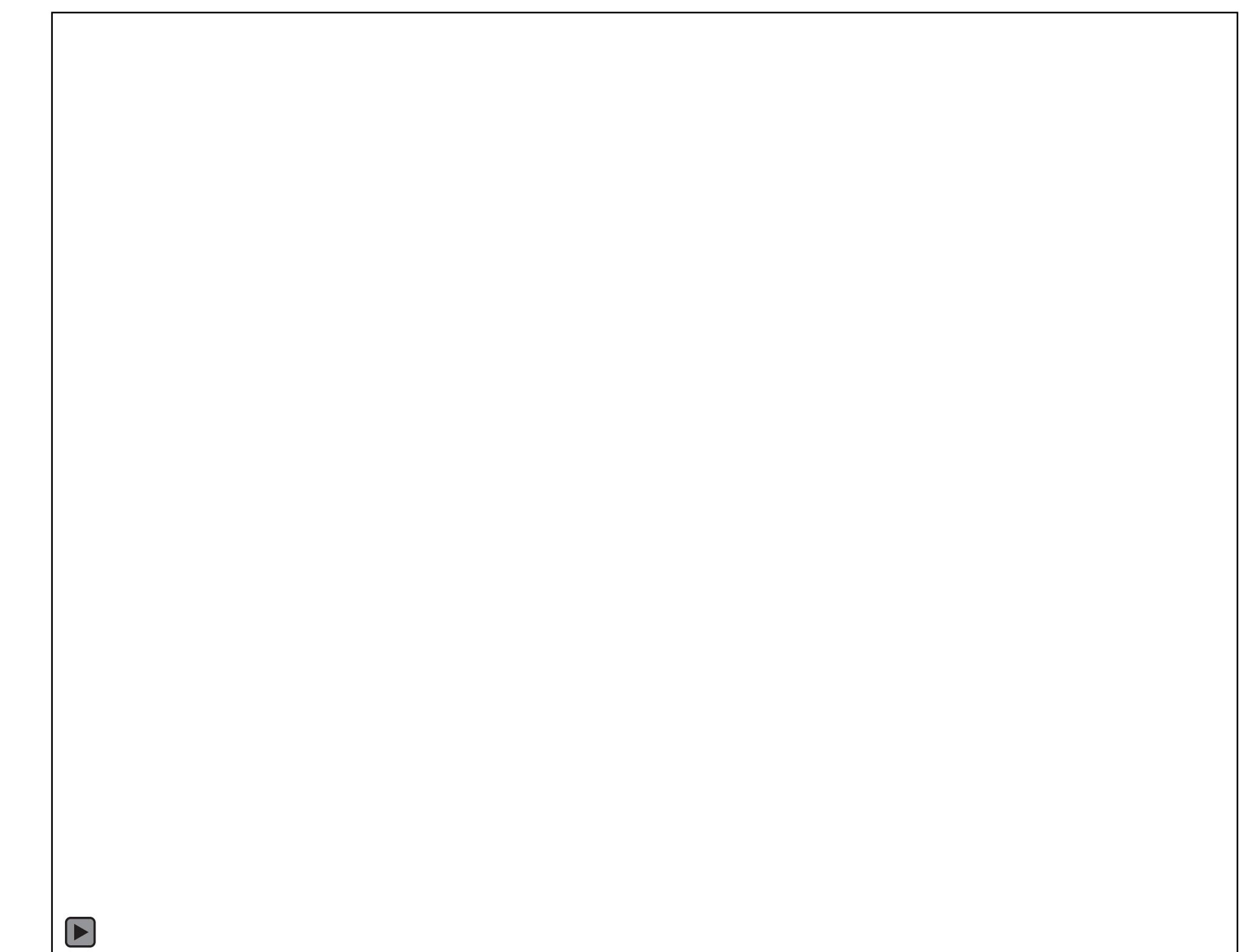


Cardiac perfusion

1. <https://arxiv.org/abs/1911.00625>
2. <https://onlinelibrary.wiley.com/doi/full/10.1002/mrm.28291>

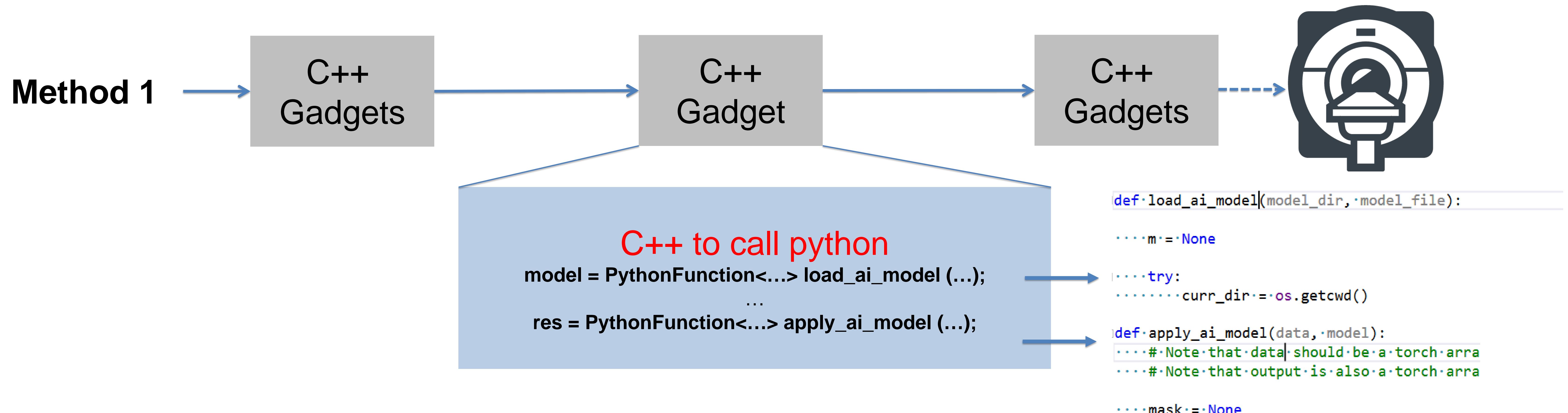


Denoising, 1mm²
MPRAGE



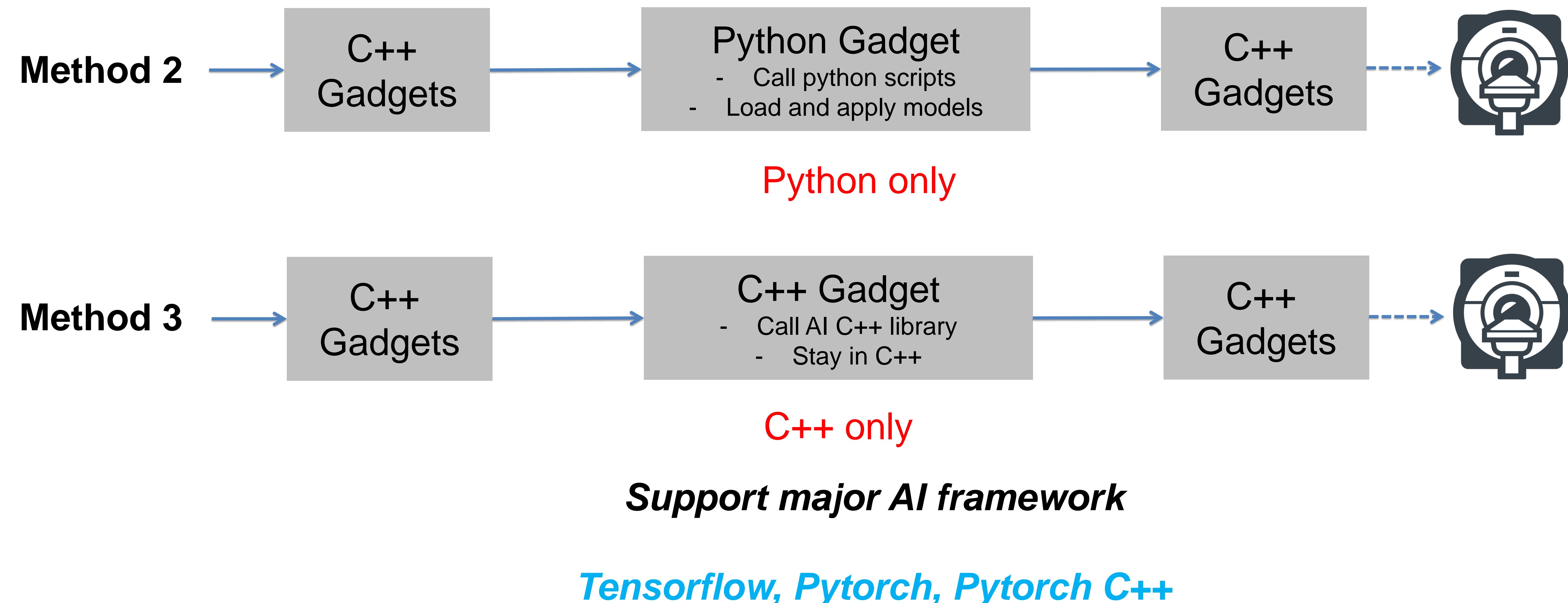
On scanner

Use Gadgetron to call AI models



Gadgetron Inline AI: Effective Model inference on MR scanner
<http://archive.ismrm.org/2019/4837.html>

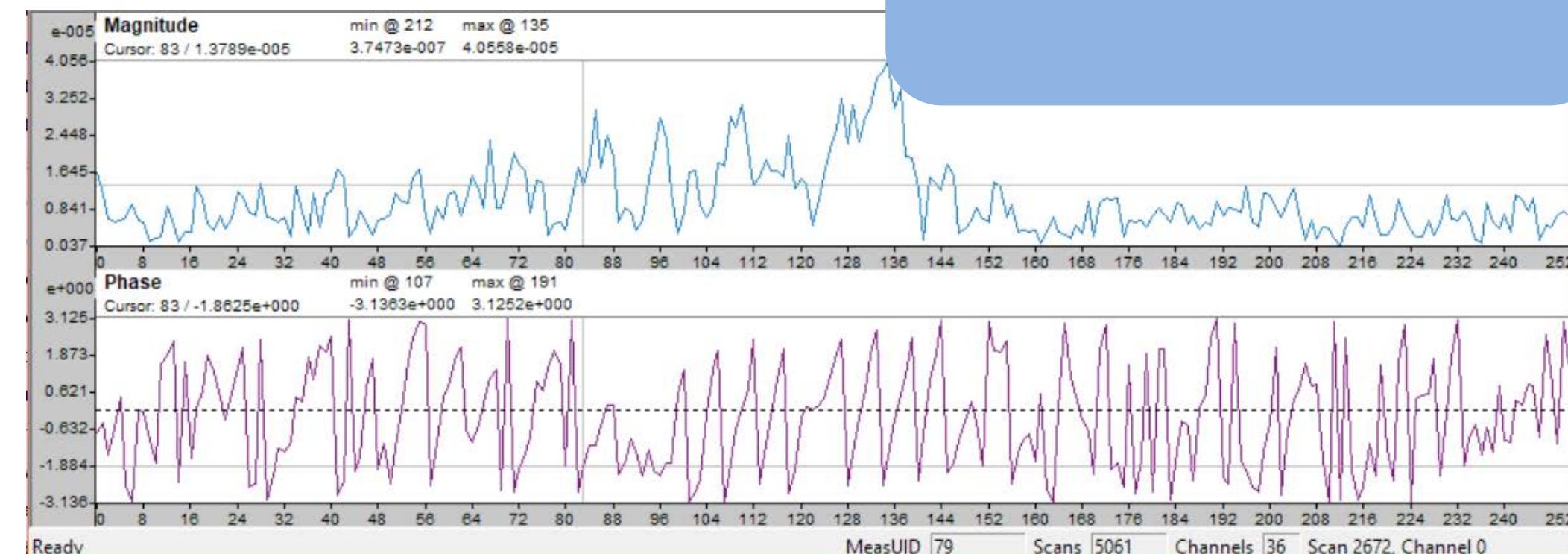
Use Gadgetron to call AI models



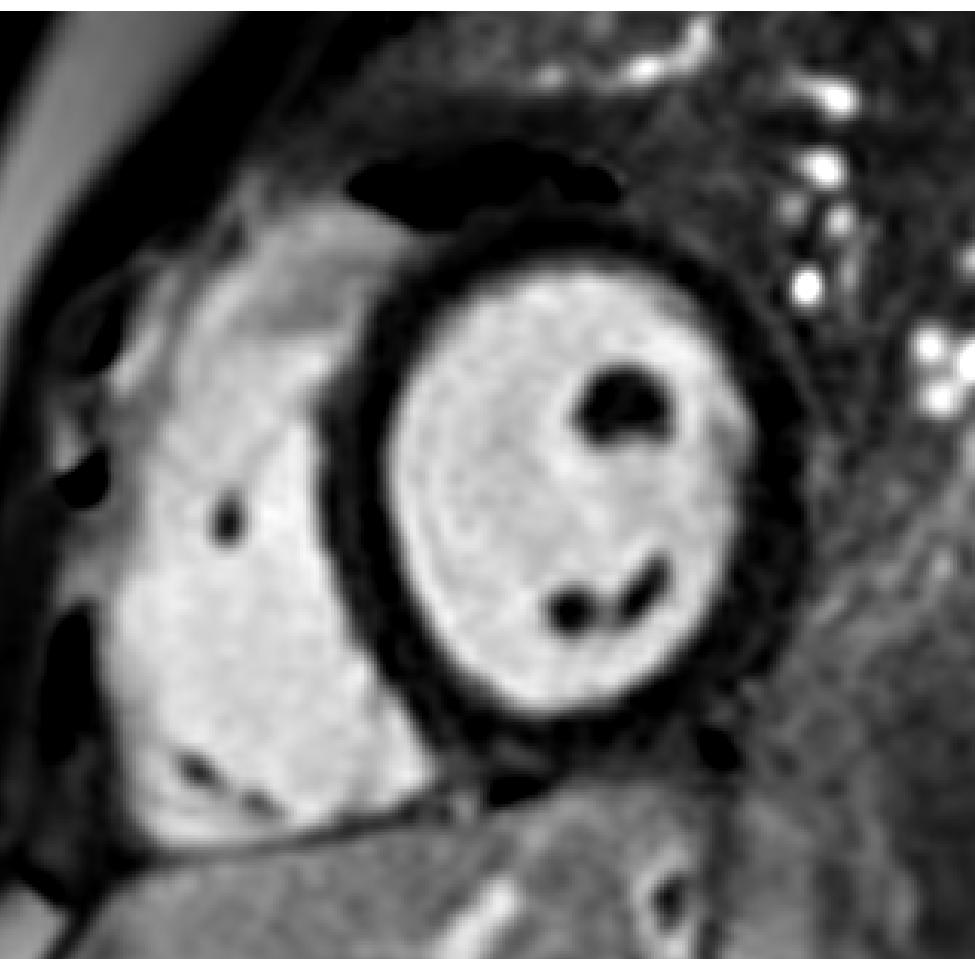
Gadgetron Inline AI: Effective Model inference on MR scanner
<http://archive.ismrm.org/2019/4837.html>

Use Gadgetron to collect data for AI: rich data types for different applications

Reconstruction



MR Kspace data



Image

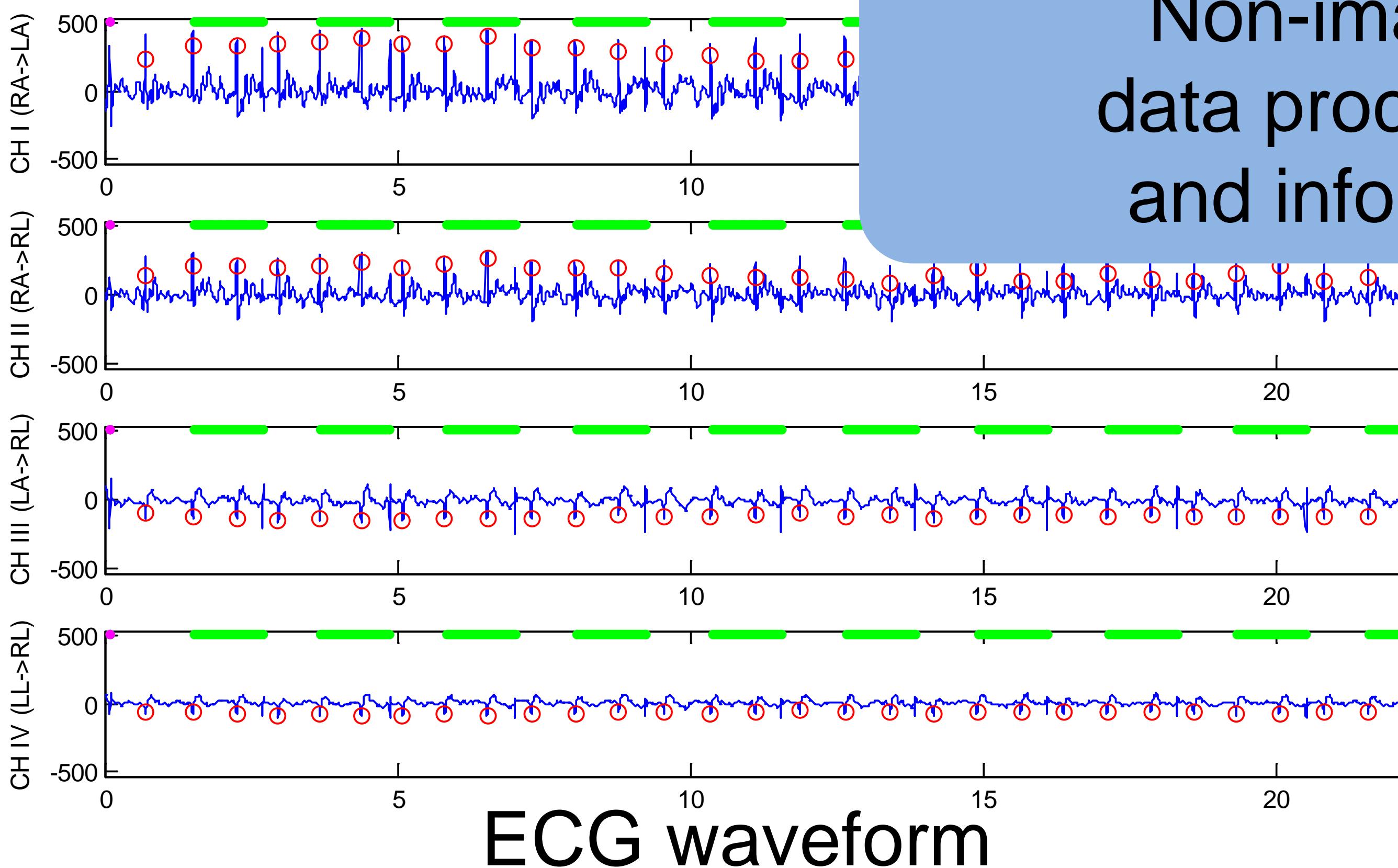
Image analysis

Diagnosis and feedback

ML meta data

```
</meta>
<meta>
  <name>E2</name>
  <value>0</value>
</meta>
<meta>
  <name>GADGETRON_DataRole</name>
  <value>Image</value>
  <value>MOCO</value>
  <value>AIE</value>
  <value>NORM</value>
  <value>PSIR</value>
</meta>
<meta>
  <name>GADGETRON_ImageComment</name>
  <value>GT</value>
  <value>x1000</value>
  <value>+4096</value>
  <value>MOCO</value>
  <value>AIE</value>
  <value>NORM</value>
  <value>PSIR</value>
</meta>
<meta>
  <name>GADGETRON_ImageNumber</name>
  <value>1</value>
</meta>
```

Non-imaging
data processing
and info fusion



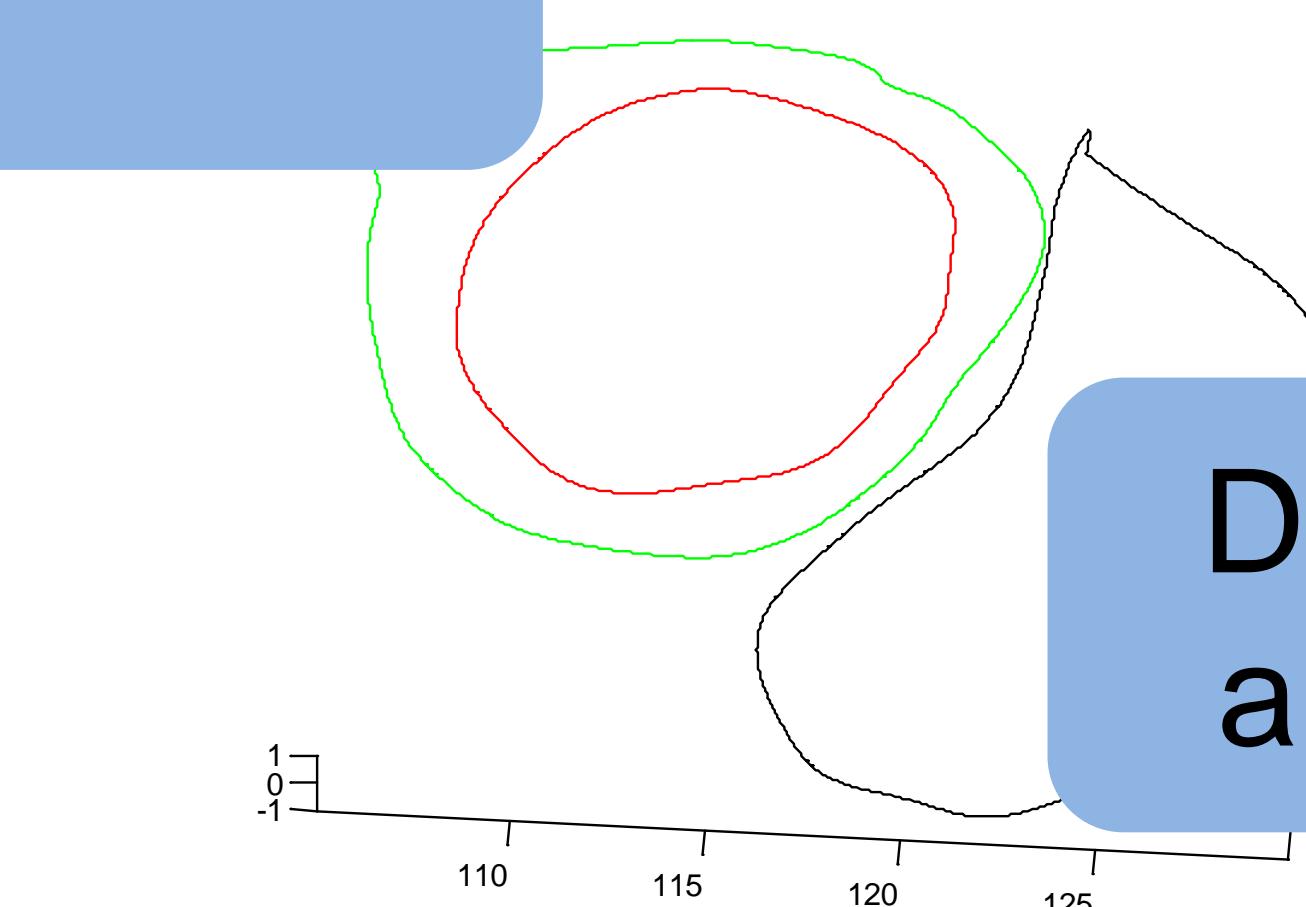
ECG waveform

- Support all steps of MR imaging

- New data types can be added

- Designed to support AI R&D

Data labelling
and collection



Contours and landmarks

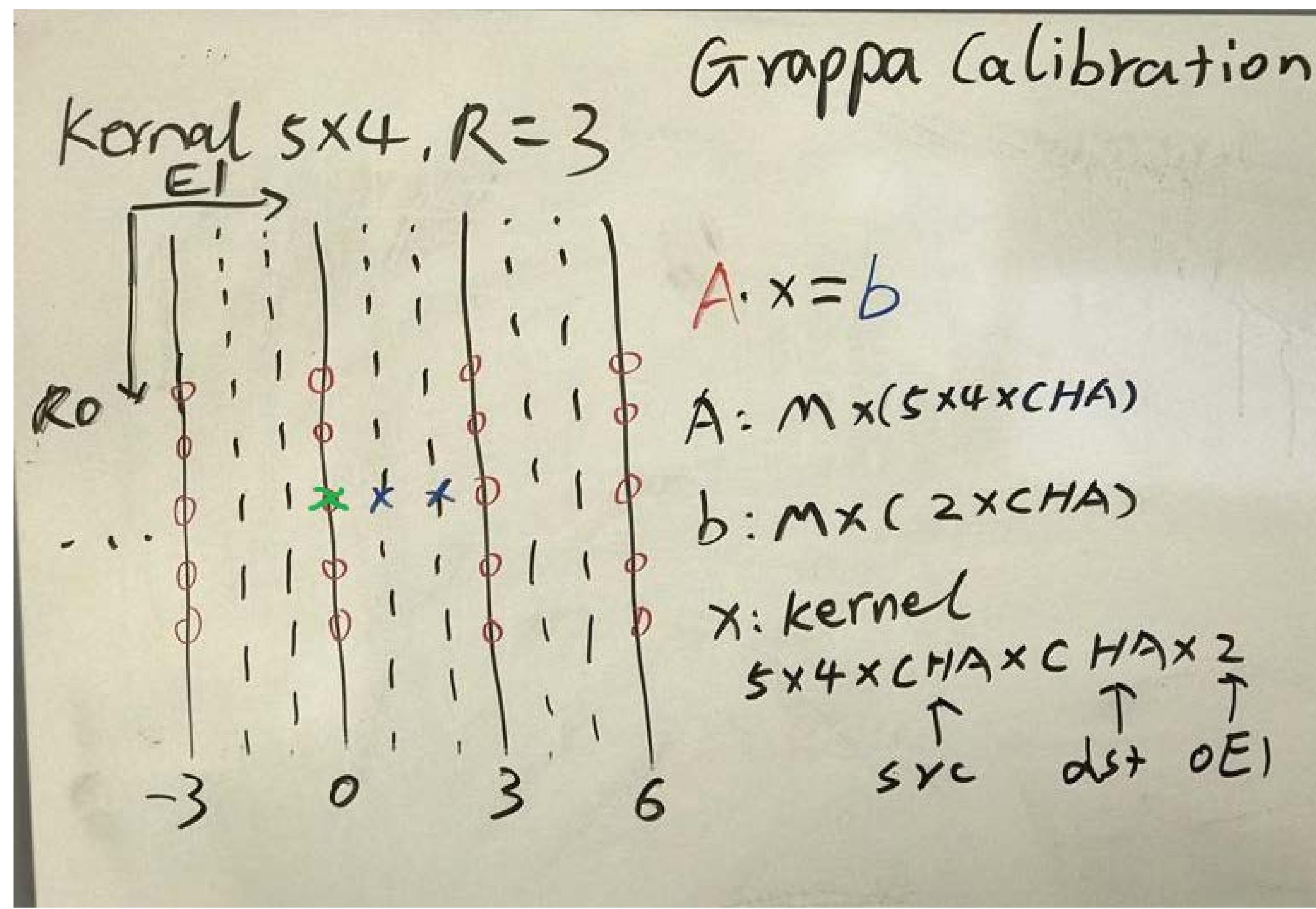


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Example: Run Grappa in Pytorch

Let's look at Grappa

Recall Grappa is to estimate kspace interpolation kernel to fill in unacquired phase encoding line:

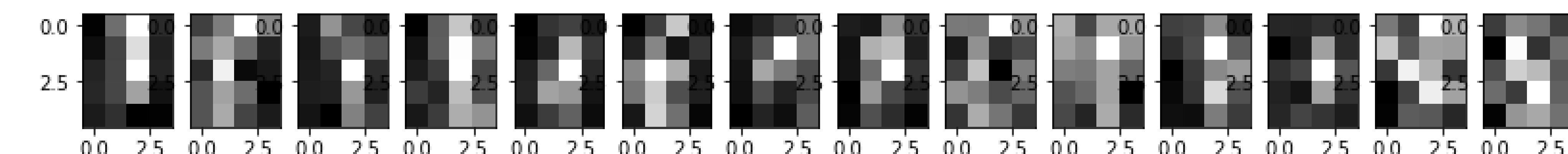


Perform Grappa reconstruction by calling the Gadgetron python wrapper for Grappa:

Kernel size $5 \times 4 \times \text{SrcCHA} \times \text{DstCHA}$

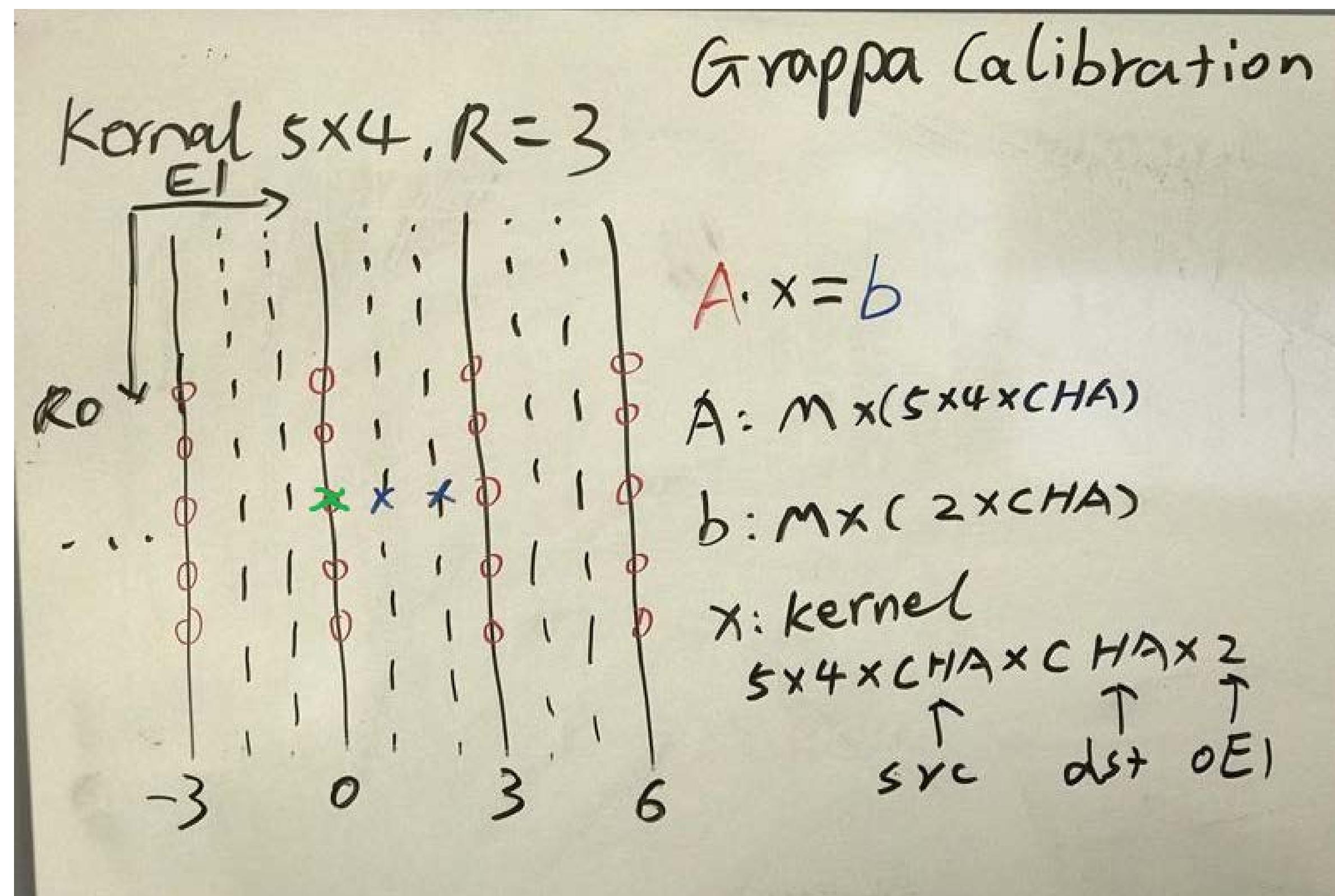
Calibration is solved by :

- 1) Assmeble data matrix A and RHS b
- 2) Solve $Ax=b$ for kernel x



Example: Run Grappa in Pytorch

- Let's look at Grappa



Grappa recon involves applying kernel to acquired data

for every missing kspace point in data:

1. Extract the $5 \times 4 \times \text{CHA}$ acquired data A
2. Multiply dataA with ker : $\text{res} = \text{np.dot}(\text{dataA}, \text{ker})$ here ker is $(5 \times 4 \times \text{CHA}, 2 \times \text{CHA})$ res is $(1, 2 \times \text{CHA})$
3. Fill res to the data array for missing data

Example: Run Grappa as AI model

Wrap grappa to call in Python

```
namespace Gadgetron
{
    class EXPORTMRICORE grappa2D
    {
        public:
            typedef std::complex<float> T;
            grappa2D();
            ~grappa2D();
            int initialize(size_t accelFactor, size_t kR0, size_t kNE1, bool fitItself, double thres);
            int calib(hoNDArray<T> acsSrc, hoNDArray<T> acsDst);
            // kspace [R0, E1, srcCHA], perform recon using ker_
            hoNDArray<T> recon(hoNDArray<T> kspace, bool periodic_boundary_condition);
            // if kspace is already converted to data matrix (dataA and dataAInd), perform recon using ker_
            hoNDArray<T> recon_data_matrix(hoNDArray<T> dataA, hoNDArray<unsigned short> dataAInd, size_t R0, size_t E1, bool periodic_boundary_condition);
            // if recon == dataA*ker_, fill back recon points to result kspace
            hoNDArray<T> recon_fill_back_kspace(hoNDArray<T> recon, hoNDArray<unsigned short> dataAInd, size_t R0, size_t E1);

            hoNDArray<T> get_A()
            {
                return A_;
            }
    };
}
```

toolboxes\mri_core\mri_core_grappa_python.h

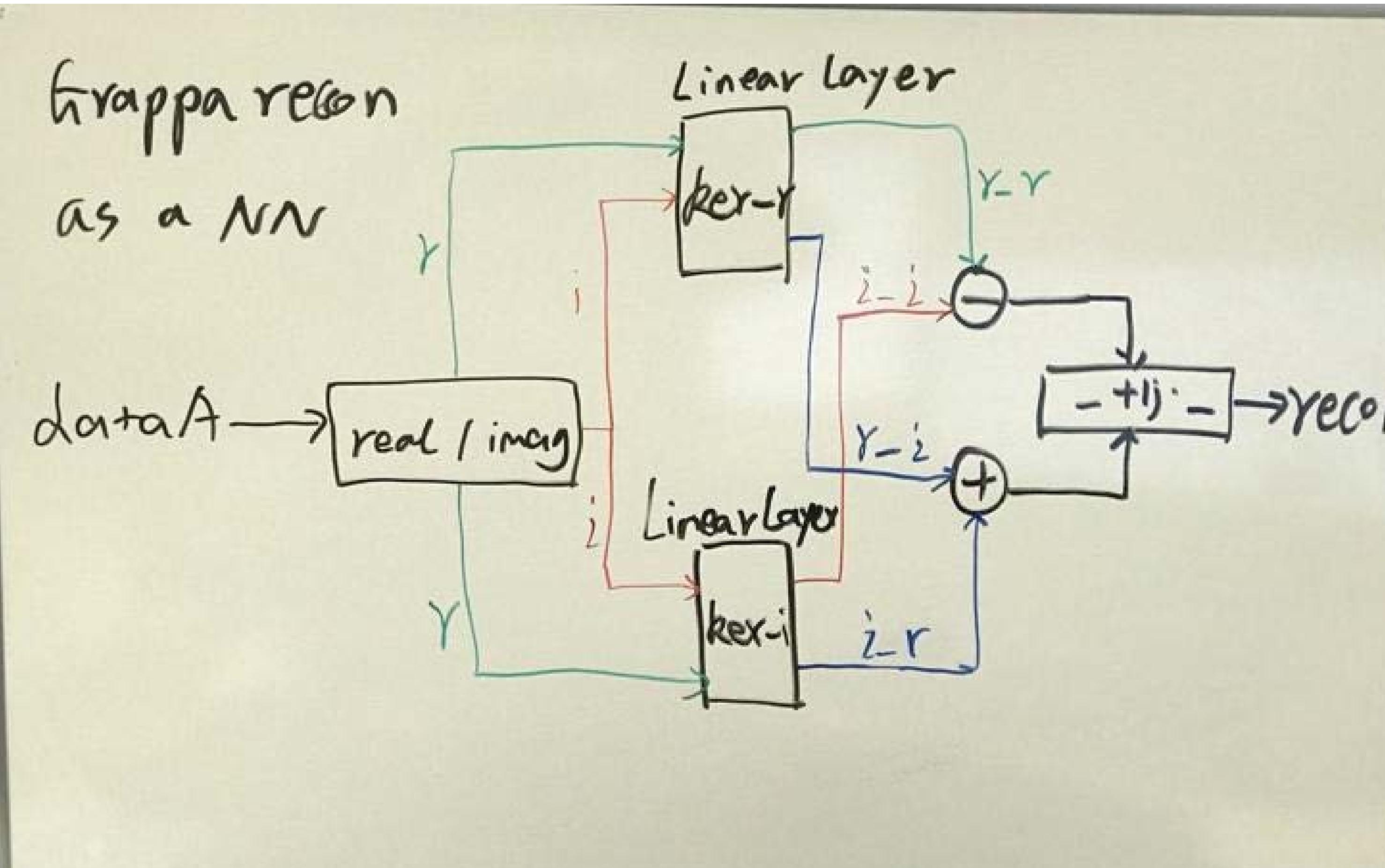
```
BOOST_PYTHON_MODULE(gadgetron_toolbox_mri_core_python)
{
    // for test purpose
    def("hello", hello);

    class_<Gadgetron::grappa2D>("grappa2D")
        .def("initialize", &Gadgetron::grappa2D::initialize)
        .def("calib", &Gadgetron::grappa2D::calib)
        .def("recon", &Gadgetron::grappa2D::recon)
        .def("recon_data_matrix", &Gadgetron::grappa2D::recon_data_matrix)
        .def("recon_fill_back_kspace", &Gadgetron::grappa2D::recon_fill_back_kspace)
        .def("get_A", &Gadgetron::grappa2D::get_A)
        .def("get_B", &Gadgetron::grappa2D::get_B)
        .def("get_ker", &Gadgetron::grappa2D::get_ker)
        .def("get_data_A", &Gadgetron::grappa2D::get_data_A)
        .def("get_data_A_index", &Gadgetron::grappa2D::get_data_A_index)
        .def("help", &Gadgetron::grappa2D::help)
        .def("status", &Gadgetron::grappa2D::status)
    ;
}
```

toolboxes\mri_core\mri_core_python.h

Example: Run Grappa as AI model

■ Implement Grappa recon as NN



```
class GrappaAI(nn.Module):
    def __init__(self, ·Din, ·Dout, ·params):
        .....
        super(GrappaAI, ·self). __init__()

        self.verbose = ·params[ 'verbose' ]
        self.with_bias = ·False
        self.grappa_weights_r = ·params[ 'grappa_weights_r' ]
        self.grappa_weights_i = ·params[ 'grappa_weights_i' ]
        .....
        #·for·real·and·imag
        self.input_layer_r = ·torch.nn.Linear(int(Din/2), ·int(Dout/2), ·bias=self.with_bias)
        self.input_layer_i = ·torch.nn.Linear(int(Din/2), ·int(Dout/2), ·bias=self.with_bias)

        if ·self.verbose:
            print("···GrappaAI···input·size·(%d), ·output·size·(%d), ·with·bias·%s" ·%·(Din, ·Dout, ·self.with_bias))

            print("--->·Set·grappa·weights")
            self.input_layer_r.weight.data = ·self.grappa_weights_r
            self.input_layer_i.weight.data = ·self.grappa_weights_i
        .....
    def forward(self, ·x):

        #·split·x·to·real·and·img
        M, ·N = ·x.shape

        N_h = ·int(N/2)

        x_r = ·x[:, ·0:N_h]
        x_i = ·x[:, ·N_h:N]

        r_r = ·self.input_layer_r(x_r)
        i_i = ·self.input_layer_i(x_i)

        r_i = ·self.input_layer_r(x_i)
        i_r = ·self.input_layer_i(x_r)

        res_r = ·r_r - ·i_i
        res_i = ·r_i + ·i_r

        out = ·torch.cat([res_r, ·res_i], ·dim=1)
```

gadgets/python/models/grappa_ai.py

Example: Run Grappa as AI model

■ Integrate GrappaAI into Generic chain

Generic_Cartesian_Grappa_AI.xml

```
#pragma once

#include "GenericReconCartesianGrappaGadget.h"
#include "python_toolbox.h"

namespace Gadgetron {

class EXPORTGADGETSMRICORE GenericReconCartesianGrappaAIGadget : public GenericReconCartesianGrappaGadget
{
public:
    GADGET_DECLARE(GenericReconCartesianGrappaAIGadget);

    typedef GenericReconCartesianGrappaGadget::BaseClass;
    typedef typename BaseClass::ReconObjType ReconObjType;
    typedef std::complex<float> T;

    GenericReconCartesianGrappaAIGadget();
    ~GenericReconCartesianGrappaAIGadget();

protected:
    // ...
    // gadget functions
    // ...
    /// [Nort·Sor1·SLC]
    std::vector<std::vector<hoNDArray<T>>> kernels_;
    std::vector<std::vector<boost::python::object>> models_;
    /// [R0·E1·E2·1·N·S·SLC]
    std::vector<IsmmrdImageArray> recon_res_grappa_ai_;

    // gadgetron home
    std::string gt_home_;

    // default interface function
    virtual int process_config(ACE_Message_Block* mb);
    virtual int process(Gadgetron::GadgetContainerMessage<IsmmrdReconData*>* m1);

    // calibration, if only one dst channel is prescribed, the GrappaOne is used
    virtual void perform_calib(IsmmrdReconBit& recon_bit, ReconObjType& recon_obj, size_t encoding);

    // unwrapping or coil combination
    virtual void perform_unwrapping(IsmmrdReconBit& recon_bit, ReconObjType& recon_obj, size_t encoding);

    virtual int close(unsigned long flags);
};

}
```

- Extend the generic recon
- Implement a new Recon gadget
- Derived from CartesianGrappa gadget to reuse components
- Call Python GrappaAI module

Demo and code walk through



Grappa AI recon

a) Grappa python binding

toolboxes\mri_core\mri_core_grappa_python.h

toolboxes\mri_core\mri_core_python.h

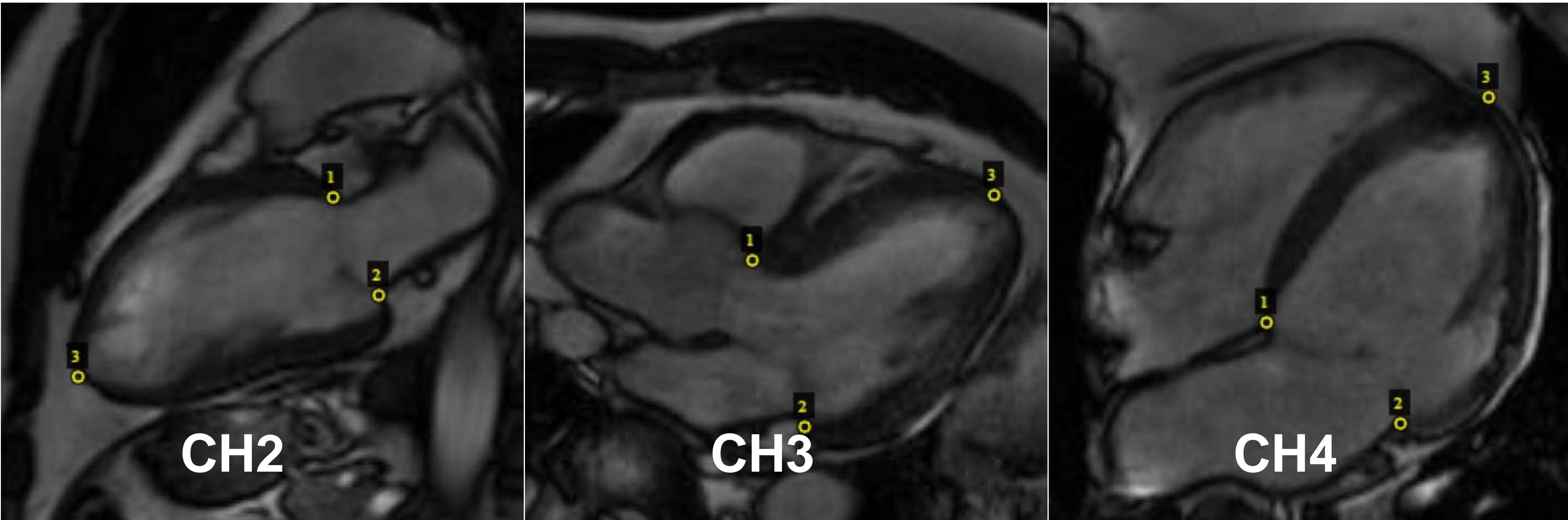
b) Gappa python module

gadgets/python/models/grappa_ai.py

- Compare C++ Grappa with this AI Grappa

Example: AI for image analysis

- Call pre-trained AI model, report results on line



Detect landmarks in cardiac images

Example: AI for image analysis

■ A Res-Unet CNN

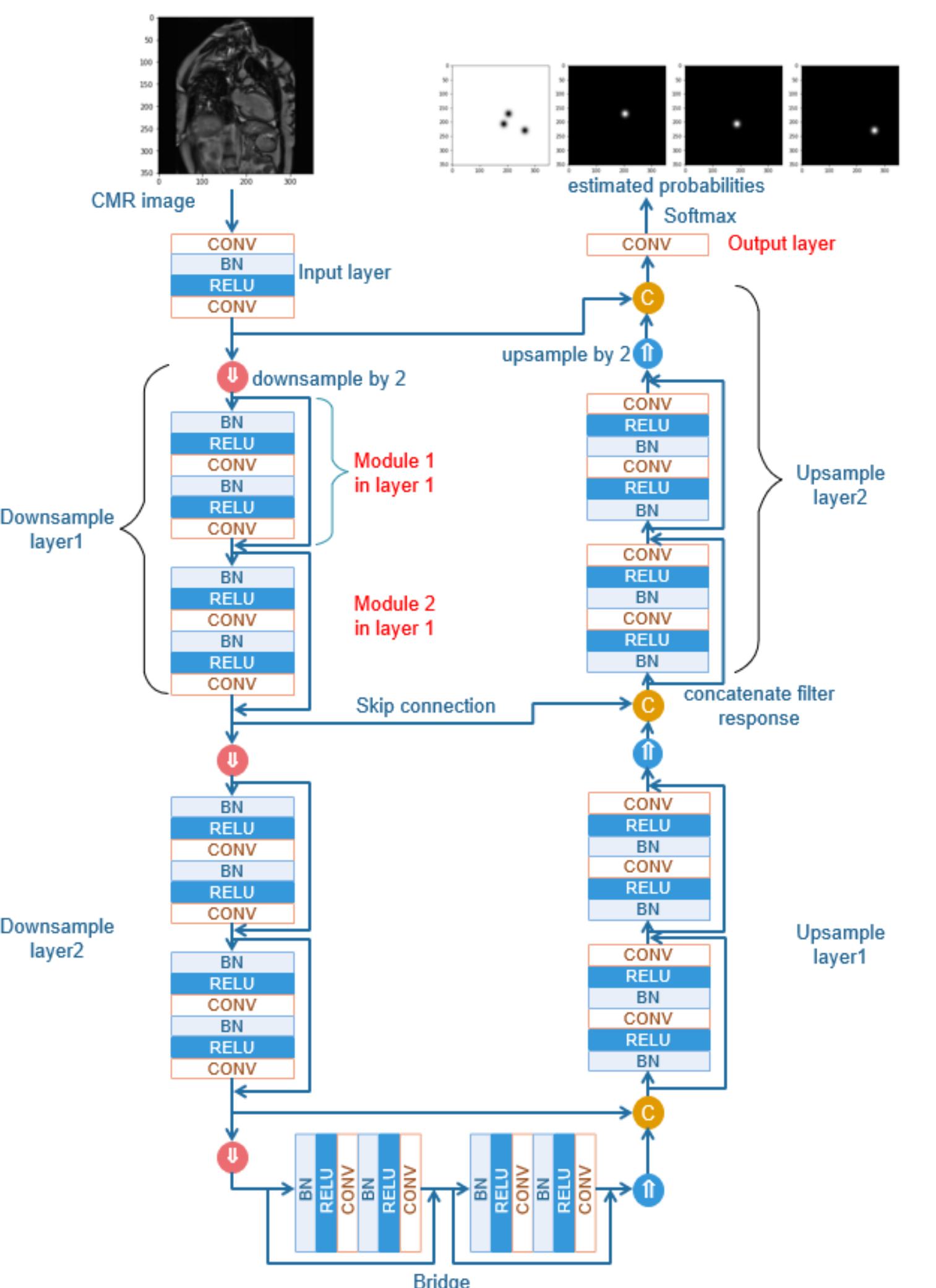


Figure 4. The backbone CNN network developed for landmark detection has a U-net structure. More layers can be inserted to both downsampling and upsampling branches. More modules can be inserted into each layer. The output layer can be a convolution for semantic segmentation or fully connected layer for regression task.

Trained on 11,426 images from 1,905 patients

25 CONV layers

Pytorch 1.5.0

Example: AI for image analysis



CMR_RTCine_LAX_AI.xml

gadgets/cmr/CmrRealTimeLAXCineAIAnalysisGadget.h/cpp

```
    // load model
    PythonFunction<boost::python::object>::load_model_cmr_landmark_detection("gadgetron_cmr_landmark_detection", "load_model_cmr_landmark_detection");
    boost::python::object model = load_model_cmr_landmark_detection(this->gt_home_, this->lax_landmark_detection_model.value());
    bp::incref(model.ptr());

    // apply model
    {
        boost::python::object* pModel = &model;
        PythonFunction<hoNDArray<float>, hoNDArray<float>>::perform_cmr_landmark_detection("gadgetron_cmr_landmark_detection", "perform_cmr_landmark_detection");
        std::tie(pts, probs) = perform_cmr_landmark_detection(lax_images, *pModel, 1.0, 8, 0.1, this->oper_R0.value(), this->oper_E1.value());
    }
```

gadgets/cmr/config/gadgetron_cmr_landmark_detection.py

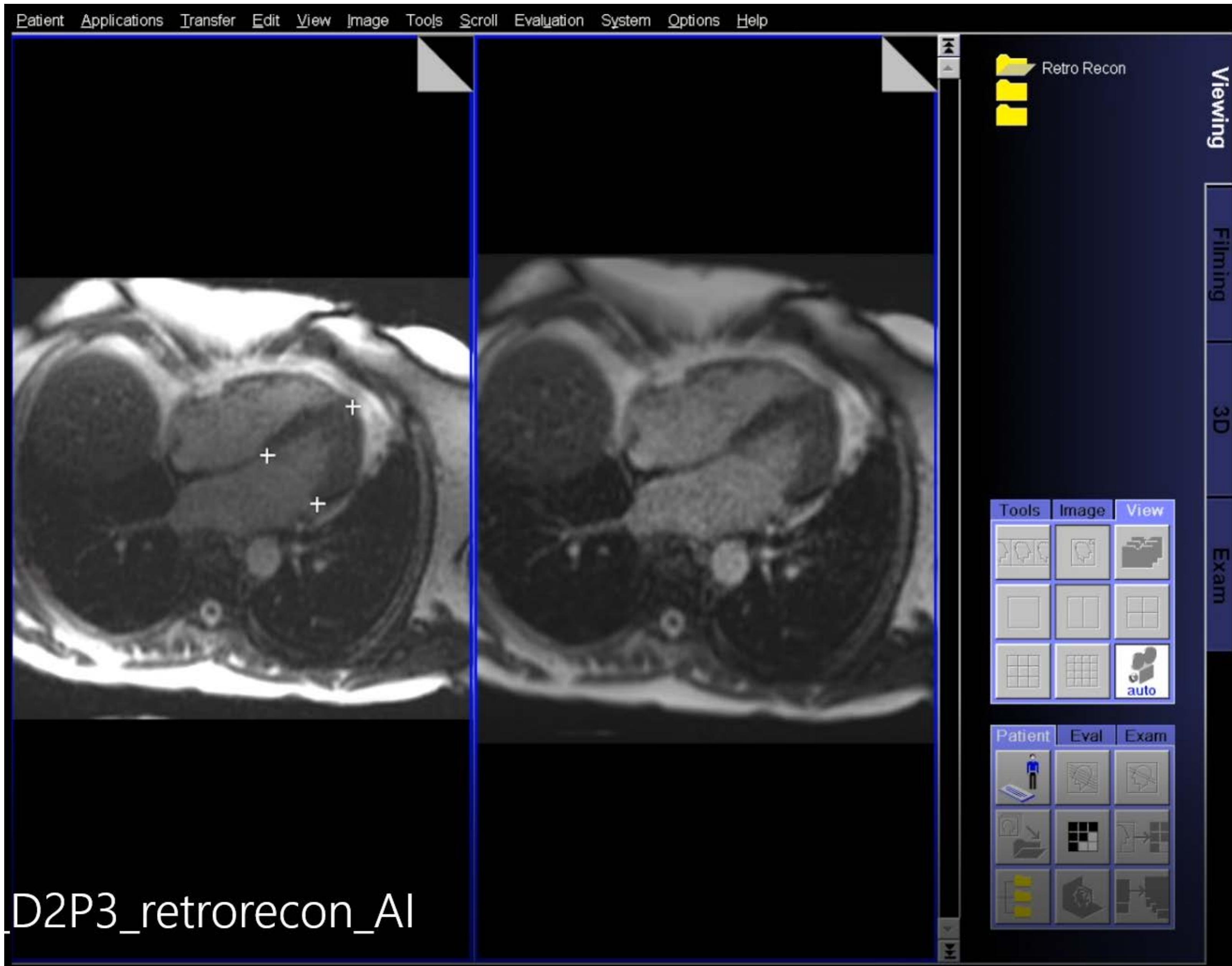
```
def load_model_cmr_landmark_detection(model_dir, model_file):
    m = None
    try:
        print("Load cmr landmark model: %s" % os.path.join(model_dir, model_file), file=sys.stderr)
        t0 = time.time()
        m = torch.jit.load(os.path.join(model_dir, model_file))
        t1 = time.time()
        print("Model loading took %f seconds" % (t1 - t0), file=sys.stderr)
        sys.stderr.flush()
    except Exception as e:
        print("Error happened in load_model_cmr_landmark_detection for %s" % model_file, file=sys.stderr)
        print(e)

    return m
```



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Example: AI for image analysis



Demo: AI for CMR landmark detection

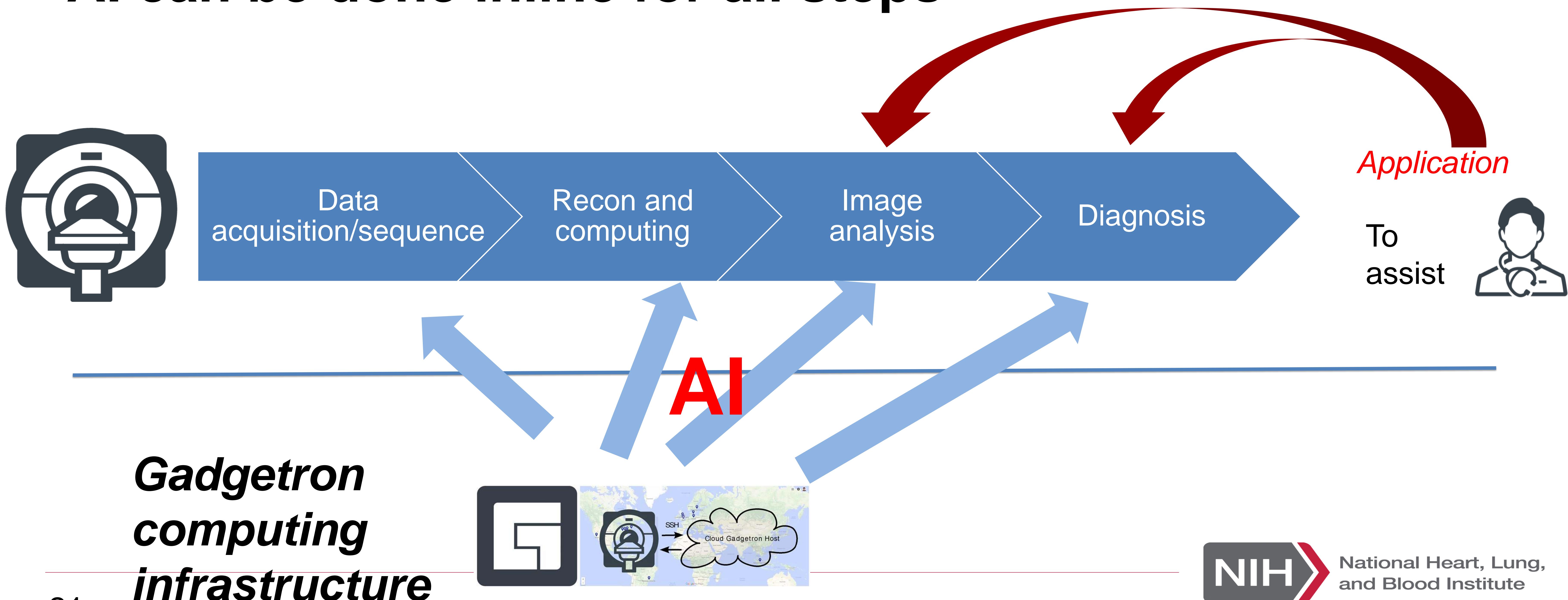
- Code walk through
 - C++ side
 - gadgets/cmr/CmrRealTimeLAXCineAIAnalysisGadget.h/cpp
 - Python side
 - gadgets/cmr/config/gadgetron_cmr_landmark_detection.py

To run this demo:

```
gadgetron_ismrmrd_client -f 2ch_RT_cine.h5 -c CMR_RTCine_LAX_AI.xml -a localhost -p 9002 -F hdr -G  
CMR_RTCine_LAX_AI.xml -o ref_20200626_084537.h5
```

Final remarks: exciting time ahead

- Imaging, computing, analysis and diagnosis
- AI can be done inline for all steps





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