

supra -CON

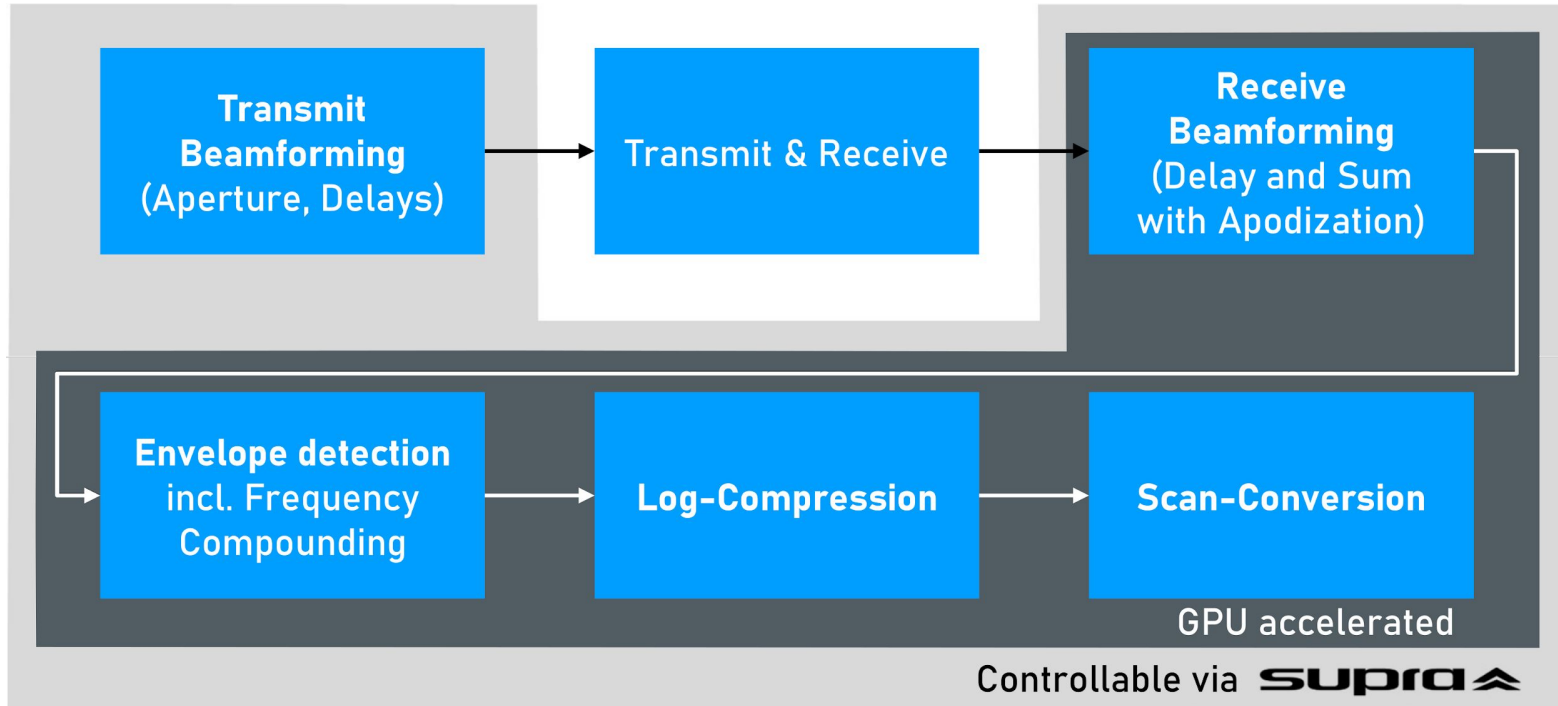
supra 

What is SUPRA?

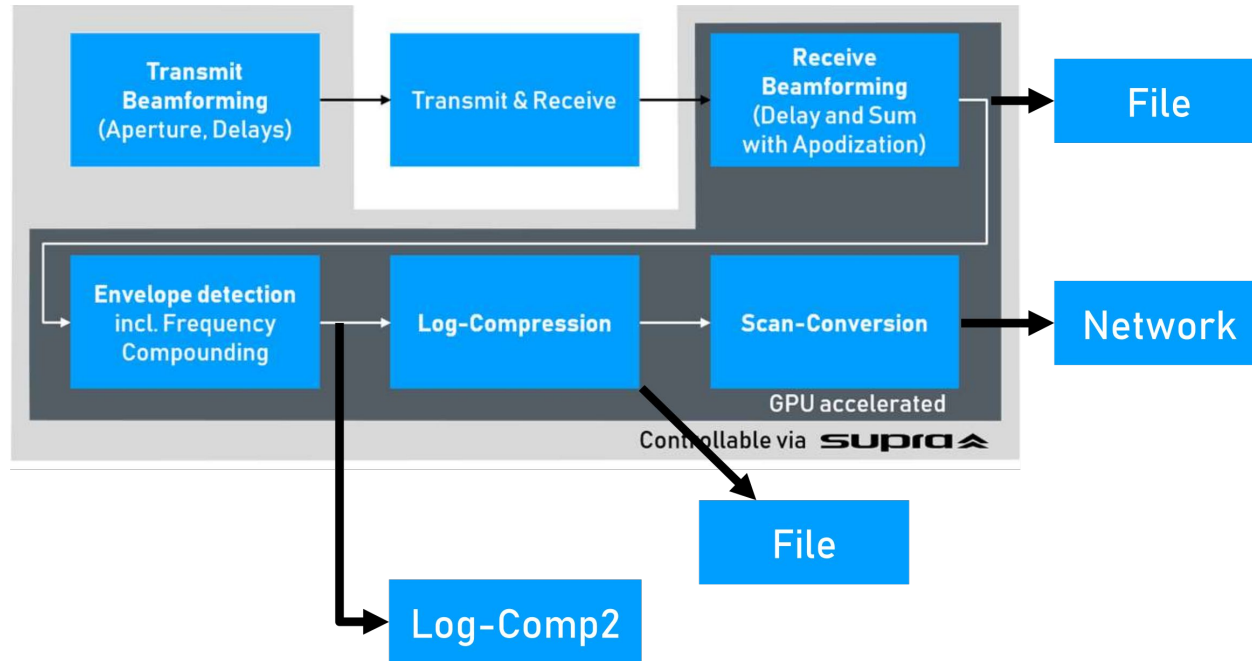
- Open-source pipeline for Ultrasound processing
- From Raw-data to final images
- Software-Defined for flexibility and accessibility
- CUDA-accelerated for real-time application
- Supports 2D and 3D imaging!

<https://github.com/IFL-CAMP/supra>

General Ultrasound Processing & SUPRA



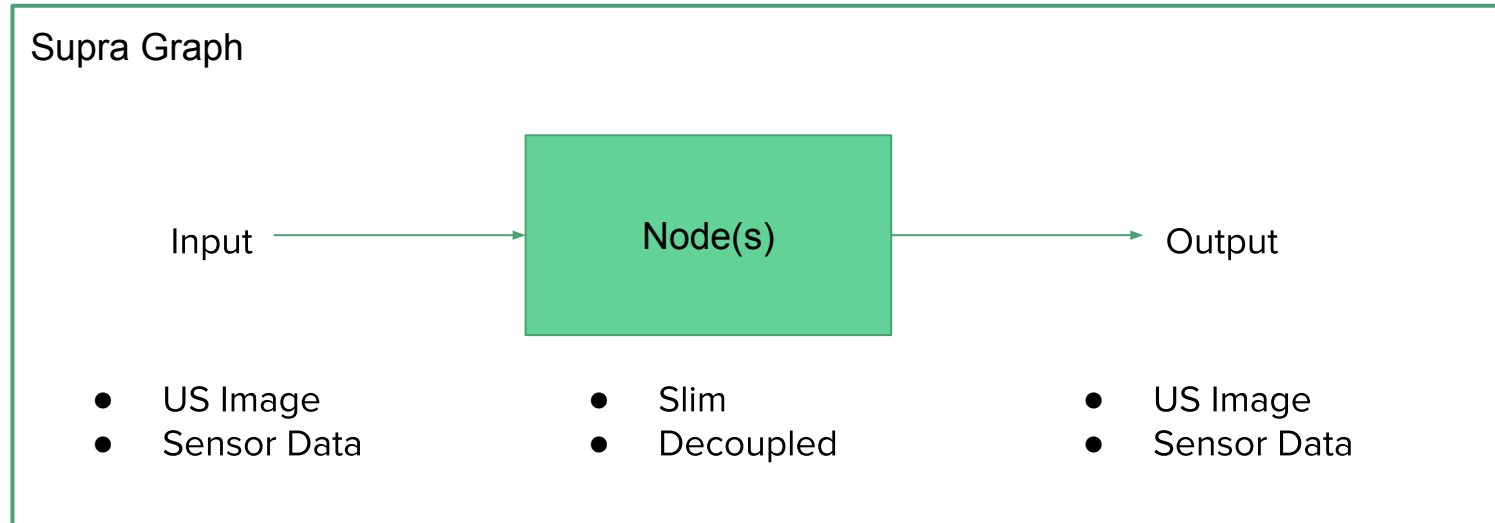
Flexibility in a Software-Defined Framework



SUPRA Design Goals

- Capability to map conventional ultrasound imaging pipeline in software
- Research platform for exploration of new methods
- Flexibility in use and extension
- Easy to use and extend
- Real-time use = High performance

Architecture - Compute Graph



Using SUPRA

What you can do with SUPRA?

- Data / Image acquisition
 - Compile and use it
- Realtime DeepLearning Inference



What you can do with SUPRA?

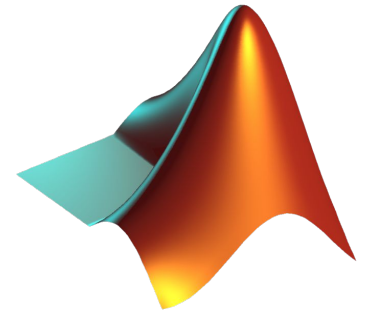
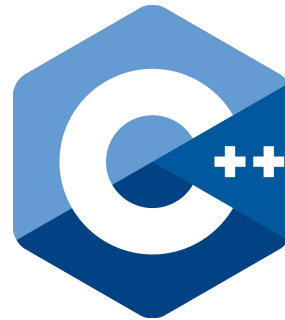
Current:

- Use WrappedPipeline from SupraLib



Upcoming:

- Matlab wrapper
- Python wrapper
- Other?



What you can do with SUPRA?

Add Features:

- New Reconstruction methods
- New Transducer Compatibility
- Advanced Ultrasound Filters
- Clinical/Research interfaces
- Further Tracking / Robotic Integrations
- AI integration

What you can do with SUPRA?

Development on framework itself

- Future architecture changes
- Addition and improvement of features
- Improved abstraction
- Additional GPU acceleration

Usage - Acquisition: Present Functionality

- 2D - Scanline imaging
 - Delay and Sum
 - Minimum Variance
- 3D Ultrasound
- Filters
 - Temporal
 - Bilateral
 - SRAD
- Pytorch integration
- Robotic tracking with ROS

<https://github.com/IFL-CAMP/supra/wiki#processing-nodes>

Usage - Acquisition: I/O (Cephasonics & Mock-files)

- Input: Open Device Interface
 - Supports Cephasonics *cuSDK*
 - Further support planned
- Input & Output: “Replay” Interface
 - config and mock file as input
- Output: ImFusion Stream
 - Contact ImFusion :)
- Output: MetalImage (MHD)
- Output: IGTL
- Output: ROS messages



<https://github.com/IFL-CAMP/supra/wiki#inputs>

Usage - Acquisition: Config File

- XML file
- specifies nodes
- specifies input and output
- Loaded at runtime

```
<?xml version="1.0" encoding="utf-8"?>
<supra_config>
  <devices>
    <inputs>
      <input type="UltrasoundInterfaceRawDataMock" id="US-Mock">
        <param name="mockDataFilename" type="string">data/linearProbe_IPCAI_128-2_0.raw</param>
        <param name="mockMetaDataFilename" type="string">data/linearProbe_IPCAI_128-2.mock</param>
      </input>
    </inputs>
    <outputs>
      <output type="MetaImageOutputDevice" id="MHD">
        <param name="createSequences" type="bool">1</param>
        <param name="filename" type="string">mhd_output</param>
      </output>
    </outputs>
  </nodes>
  <node type="BeamformingNode" id="BEAM">
    <param name="windowParameter" type="double">0.5</param>
    <param name="windowType" type="string">Hamming</param>
  </node>
  <node type="HilbertFirEnvelopeNode" id="DEMO" />
  <node type="FrequencyLimiterNode" id="LIMIT">
    <param name="maxFrequency" type="double">50</param>
  </node>
</supra_config>
```

Usage - Acquisition: Runtime Graph

- Interactive parameter changes
- Dynamic Graph construction
- Load/Save as Config Files

Interfaces

- GUI → Online processing
- Executor → Offline processing
- ROS → “Headless” online processing, control through ROS
- REST interface → Access & change parameters through web API
- WrappedPipeline → Use a SUPRA pipeline in other software

Using - Summary

- Nodes
- Configuration
 - Runtime
 - Config XMLs
- Interfaces

Adding Functionality

Important Data Structures

- SupraGraph
- AbstractNode
- Parameter system
- Containers
- RecordObject
 - USImage
 - USRawData
- USImageProperties
- RxBeamformerParameters

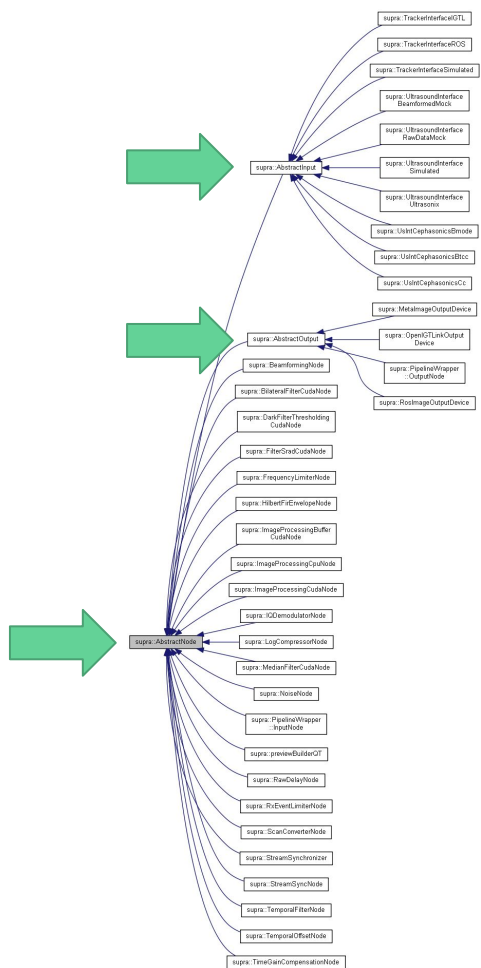
→ Doxygen documentation (can be activated in CMake!)

SupraGraph

- The collection of Nodes and Connections
- Represents the “pipeline”
- Used by interfaces (frontends? GUI etc.) to access and modify nodes
- Usually only one
- Can read and write configuration XML
 - Nodes
 - Parameters
 - Connections

AbstractNode

- Base class for all nodes
(Special base classes AbstractInput and AbstractOutput predefine some behavior)
- Contains the parameter definitions and values
 - ValueRangeDictionary
 - ConfigurationDictionary
- Defines interface to access and change them



AbstractNode

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Need to overwrite functions:

`getNumInputs()`

`getNumOutputs()`

`getInput(size_t index)`

`getOutput(size_t index)`

Mostly generic

Node specific logic to
react to new
parameter values

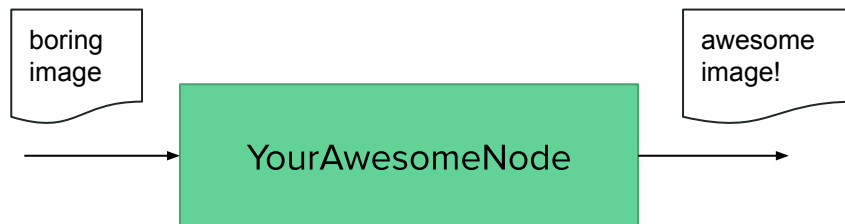
`configurationChanged()`

`configurationEntryChanged(
string configKey)`

How do Nodes do stuff?

How is YourAwesomeAlgorithm called?

Usual case: 1-Input, 1-Output



```
YourAwesomeNode::YourAwesomeNode (...)
```

```
{
```

```
    // Create the underlying tbb node for handling the message passing.  
    m_node = unique_ptr<NodeTypeQueueing>(new NodeTypeQueueing(graph, 1,
```

```
        [this](shared_ptr<RecordObject> inObj) -> shared_ptr<RecordObject> {  
            return YourAwesomeAlgorithm(inObj); }  
    );
```

```
});
```



Callback on arrival of input message!

Parameter System: ValueRangeDictionary

Every node has one!

Defines the parameters including

- Type
- Name (key)
- Display name
- Valid values
 - **Unrestricted**
 - Range
 - Discrete

```
template<typename ValueType>
void ValueRangeDictionary::set(
    const string & key,
    const ValueType & defaultValue,
    const string & displayName )
```

Example:

```
m_valueRangeDictionary.set<bool>(
    "additiveGaussCorrelated",
    false,
    "additive Gauss Correlated");
```


Parameter System: ValueRangeDictionary

Every node has one!

Defines the parameters including

- Type
- Name (key)
- Display name
- Valid values
 - Unrestricted
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```
template<typename ValueType>
void ValueRangeDictionary::set(
    const string & key,
    const ValueType & lowerBound,
    const ValueType & upperBound,
    const ValueType & defaultValue,
    const string & displayName)
```

Example:

```
m_valueRangeDictionary.set<double>(
    "speckleScale",
    0.0, 100.0, 25.0,
    "Speckle Scale");
```

Parameter System: ValueRangeDictionary

Every node has one!

Defines the parameters including

- Type
- Name (key)
- Display name
- Valid values
 - Unrestricted
 - Range
 - **Discrete**

```
template<typename ValueType>
void ValueRangeDictionary::set(
    const string & key,
    const vector<ValueType> & value,
    const ValueType & defaultValue,
    const string & displayName)
```

Example:

```
m_valueRangeDictionary.set<DataType>(
    "outputType",
    { TypeFloat, TypeUint8, TypeInt16 },
    TypeFloat,
    "Output type");
```

Parameter System: ConfigurationDictionary

Contains the actual parameter values

Used to get and set the params

Example:

```
m_speckleScale =  
    m_configurationDictionary.  
        get<double>("speckleScale");  
  
m_subArraySize =  
    m_configurationDictionary.  
        get<uint32_t>("subArraySize");
```

Container<T>

Thin wrapper around memory block

Handles allocation / deallocation

Follows RAII (Resource acquisition is initialization)

Keeps track:

- Location (Host / GPU)
- Associated CUDA stream
- Buffer size
- Data type

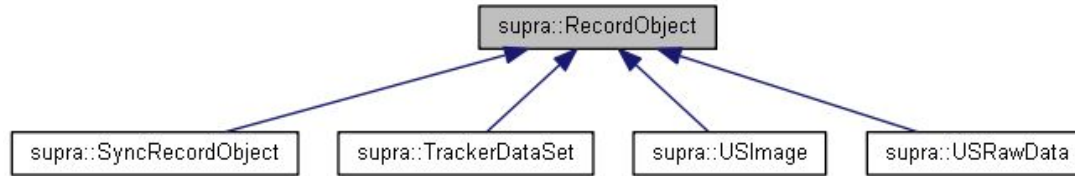
Hints:

- Use when you need GPU memory
- Also for moving data GPU ↔ Host

```
shared_ptr<Container<float> > data = ...;
if (!data->isGPU())
{
    data = make_shared<Container<float> >
        (LocationGpu, *data);
}
```

RecordObject

- The base class for data passing through the graph
- Contains timestamp



USImage

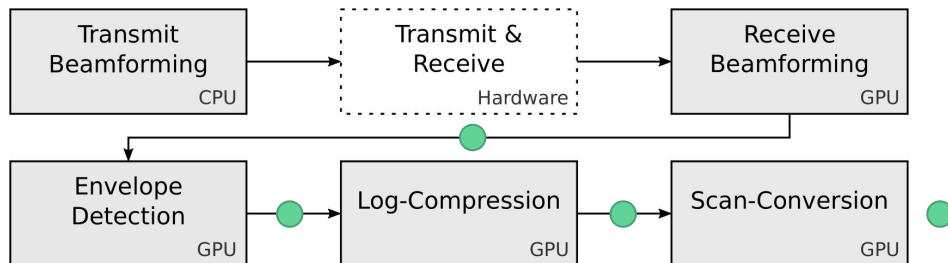
Describes an ultrasound image after beamforming

- Size
- USImageProperties
- Image data Container

Arbitrary element (“pixel”) datatypes

Possible states:

- Beamformed RF
- Envelope detected
- PreScan (Logcompressed scanline data)
- Scanconverted



USRawData

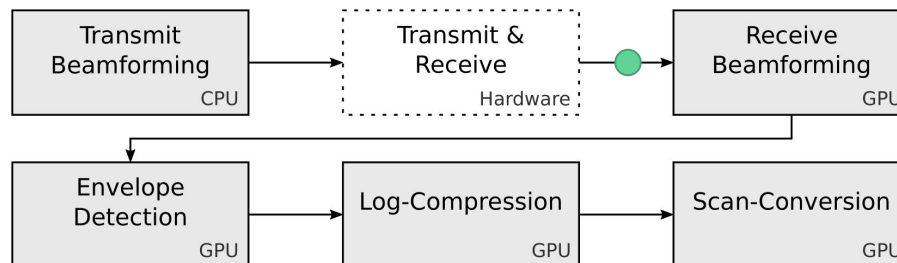
Describes an ultrasound image before beamforming

- Size i.e. #Scanlines, #elements, #temporal samples
- Sampling frequency
- USImageProperties
- RXBeamformerParameters
- Raw data Container

Arbitrary element (“Sample”) datatypes

Possible states:

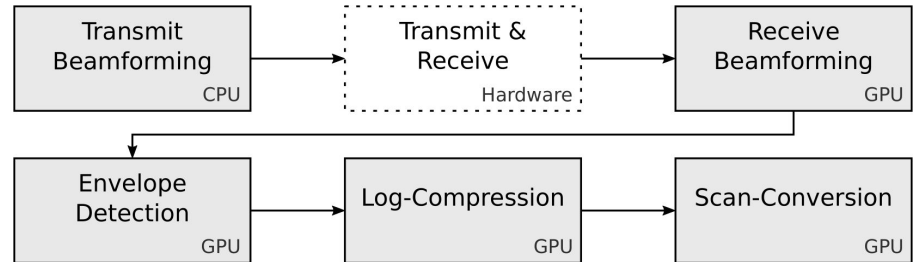
- Raw data / Channel data
- Delayed Raw



USImageProperties

Contains all the metadata to describe the meaning of an ultrasound image

- State (e.g. beamformed RF)
- Transducer type
- # Scanlines & geometry
- # Samples
- Acquisition depth



RXBeamformerParameters

The geometric, temporal and other low-level parameters to facilitate receive beamforming

- Location & direction of scanlines
- Transducer element locations
- Speed of sound
- Elements used for each transmit / receive event

Example Nodes

Perform same task

“Scale image intensities”

in slightly different ways

- ImageProcessing**Cpu**Node
- ImageProcessing**Cuda**Node
- ImageProcessing**BufferCuda**Node

Highly recommended!

Adding Nodes - Summary

- 1-Input, 1-Output is easy → Callback with your function
- Example nodes are good basis
- Use Containers!
- Make everything a parameter
- Use Buffer2/3 instead of manual index calculation

CUDA-basics in two slides (1)

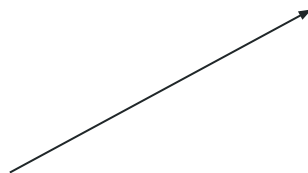
CPU 2D operation

```
for (size_t y = 0; y < height; y++) {  
    for (size_t x = 0; x < width; x++) {  
        WorkType inPixel =  
            pInputImage[x + y*width];  
        WorkType value = inPixel * m_factor;  
        pOutputImage[x + y*width] = value;  
    }  
}
```

CUDA 2D operation

```
dim3 blockSize(32, 4, 1);  
dim3 gridSize(  
    (size.x + blockSize.x - 1) / blockSize.x,  
    (size.y + blockSize.y - 1) / blockSize.y);  
processKernel <<<gridSize, blockSize, 0,  
inImageData->getStream() >>> (  
    inImageData->get(),  
    size,  
    factor,  
    outImageData->get());  
cudaSafeCall(cudaPeekAtLastError());
```

Kernel call



CUDA-basics in two slides (2)

“Equivalent” loops for kernel

implicit in kernel call, **order unknown!**

```
for (blockIndex.x = 0; blockIndex.x < gridSize.x; blockIndex.x++) {
  for (blockIndex.y = 0; blockIndex.y < gridSize.y; blockIndex.y++) {
    for (threadIndex.x = 0; threadIndex.x < blockDim.x; threadIndex.x++) {
      for (threadIndex.y = 0; threadIndex.y < blockDim.y; threadIndex.y++) {
        size_t x = blockDim.x*blockIdx.x + threadIdx.x;
        size_t y = blockDim.y*blockIdx.y + threadIdx.y;
        if (x < width && y < height)
        {
          WorkType inPixel = pInputImage[x + y*width];
          WorkType value = inPixel * m_factor;
          pOutputImage[x + y*width] = value;
        }
      }
    }
  }
}
```

Code in
kernel
function

CUDA tips

- Start from example CUDA node!
- Ensure all containers are in GPU memory

```
data->isGPU()
```
- Avoid GPU synchronization where possible
 - Except for debugging
- Add error checking after every kernel call

```
cudaSafeCall(cudaPeekAtLastError());
```
- Don't worry about optimization at the beginning

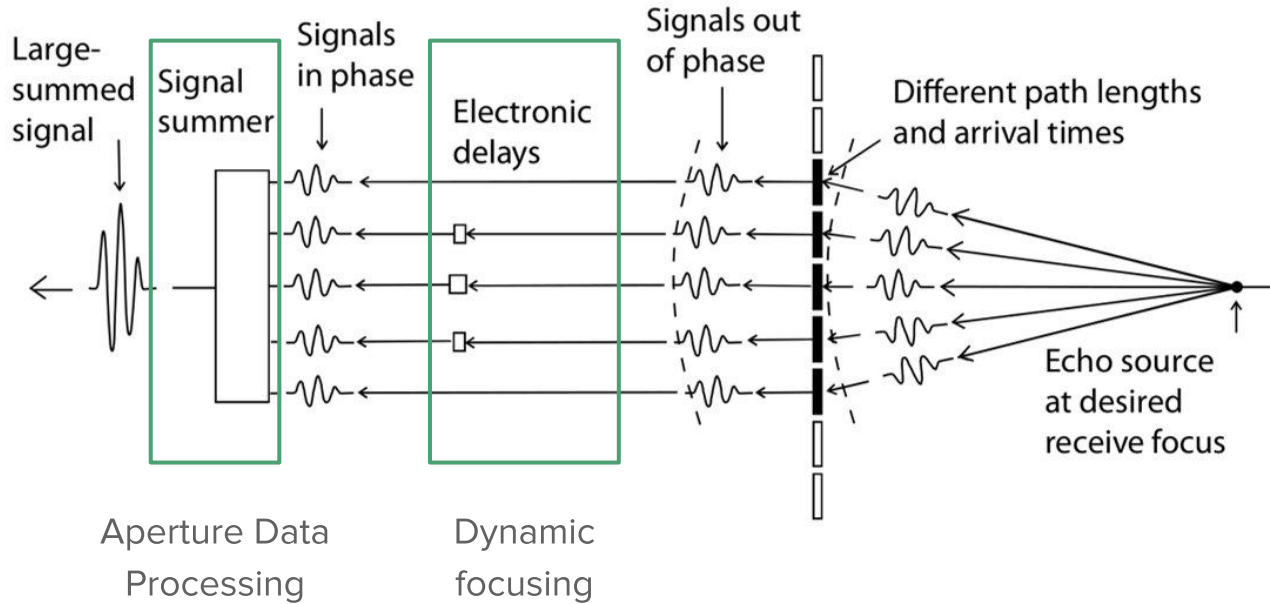
Beamforming in SUPRA

Beamforming in two slides (1)



Hoskins, P. R., Martin, K. R., & Thrush, A. R. (2019). *Diagnostic Ultrasound: physics and equipment* (3rd ed.). page 8: CRC Press.

Beamforming in two slides (2)



From Hoskins, P. R., Martin, K. R., & Thrush, A. R. (2019). *Diagnostic Ultrasound: physics and equipment* (3rd ed.). page 49: CRC Press.

Beamforming: what I left out

- Transmit beamforming (TX)
- Acoustic properties of tissue
- Acoustic interactions
- and more...

See “Diagnostic Ultrasound: Physics and Equipment (3rd ed.)” for more

Beamforming in SUPRA

- Focusing and Aperture data processing combined
 - Delay-and-Sum
 - Coherence Factor weighting
- Focusing and Aperture data processing separate
 - Minimum Variance
 - DeepFormer

The Future

Architecture / Future Development goals

Architectural

Functionality

Compatibility

Architecture / Future Development goals

Architectural

- Option for grouping Parameters
 - apply several parameters at the same time while ensuring valid parameter state at all times
- Transducer-specification in files instead of code

Architecture / Future Development goals

Functionality

- Plane wave / Diverging wave (2D & 3D)
- Ultrasound File Format (UFF) input and output (HDF5 based)
- Pre-beamformed processing
- Increase modularity in fused Rx beamformer

Architecture / Future Development goals

Compatibility

- Wider variety of compatible devices
- “Clinical” interfaces via REST-API

Roadmap: Proposal for v0.1

- Current state
- Extended documentation
 - Improve Doxygen “low-level”
 - High-level documentation (“Getting started, using...”)
- Code file structure rework



Roadmap: Proposal for v0.2

- Complete REST-API
- MATLAB Interface
- Mock files replaced with HDF5
- Support for Cephasonics curvi-linear probe



Architecture / Future Development goals

What do you think is missing?

