



PyCBC: a device independent approach to CBC analysis

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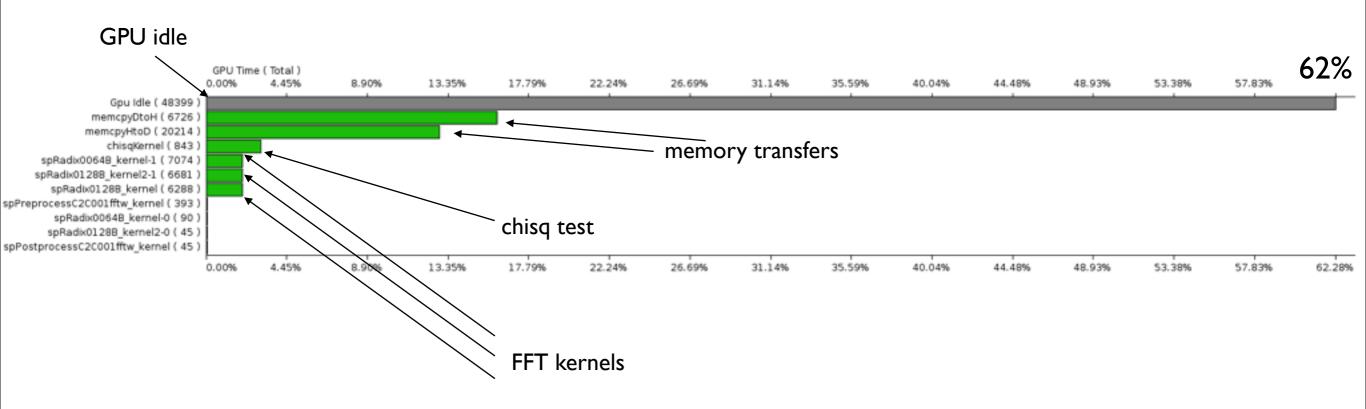
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Previous work

- Acceleration of lalapps_inspiral by GPUs obtained a speedup factor of 15 by performing all FFTs and the chi-squared test on a GPU. (CUDA) https://docid=40432
- Results of profiling: largest portions of time spent...
 - transferring memory to and from the GPU
 - running computations on the CPU (GPU is idle)
- Conclusion: a redesign focused on transparent GPU acceleration is needed

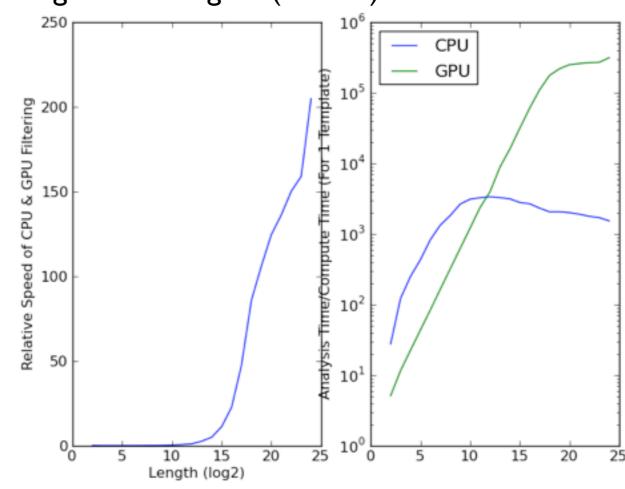






Preliminary Acceleration Investigations

- To optimize this, we need to implement more of the search engine (i.e., the whole hot loop) on the GPU.
 - The Budapest group has shown a possible speedup factor of 92 by porting the search engine completely to the GPU. (OpenCl) http://www.grid.kfki.hu/twiki/bin/view/RmkiVirgo/GPU_inspiral
 - The Syracuse group has shown even large gains (speedup factor of 200) are possible depending on the length of the signal. (CUDA)







Coding for the future

- Speed not the only issue
- We want something sustainable and that can bring us into the Adv. Detector Era
- Targeting offline compute intensive searches





Requirements of a novel software framework for CBC analysis

Design Requirements:

- flexible (easily reconfigurable from the top layer)
 - decouple algorithm from implementation
- transparent processing on different architectures
- transparent memory management
- better readability

• Code Requirements:

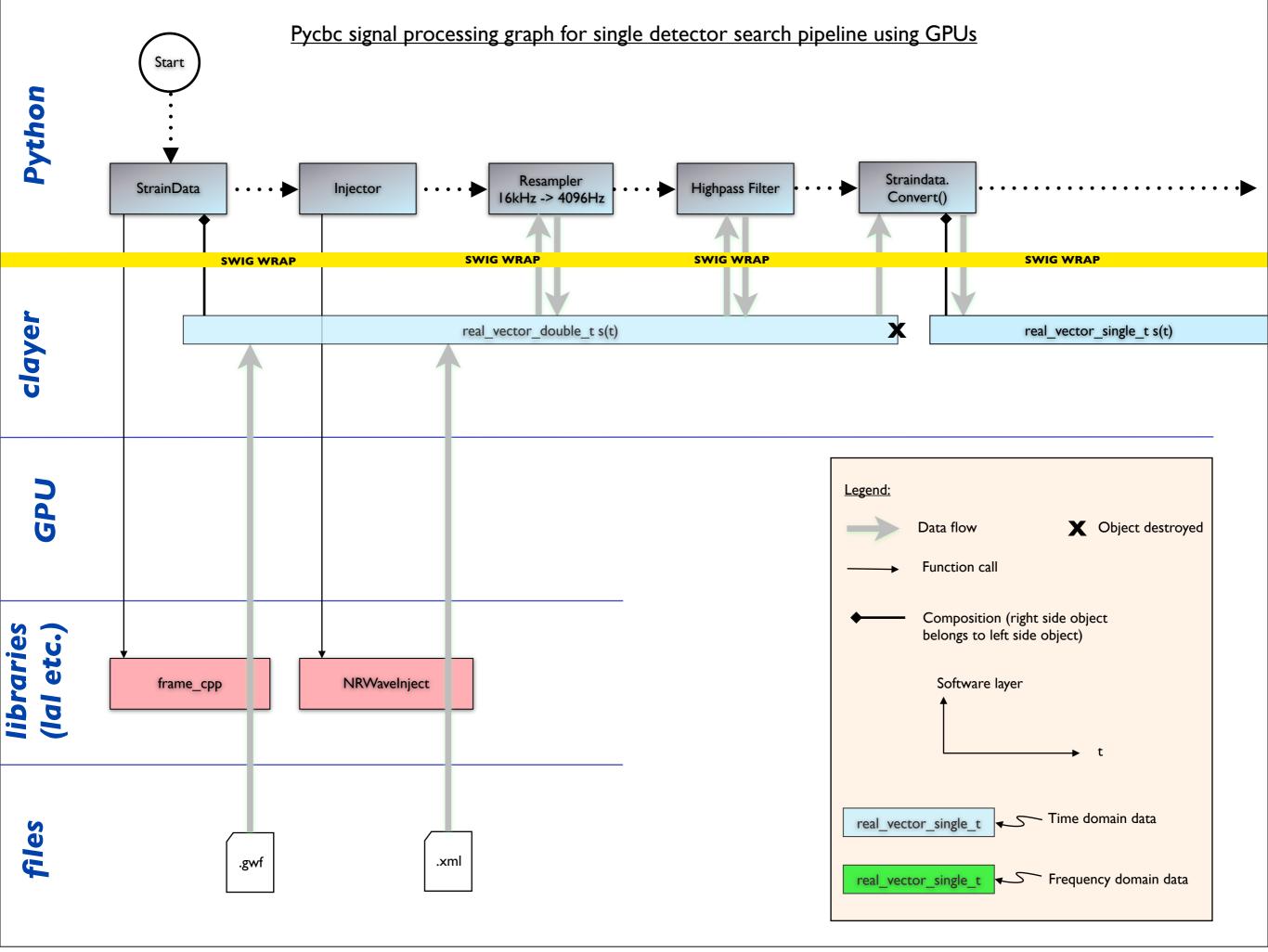
- modularity and simple API
 - simple processing python objects (object oriented design)
 - new features are encapsulated in new objects or extensions of existing objects
 - robustness from unit tests
- bindings to 3rd party libraries (e.g., lalsuite, fftw, numpy, etc.)
- auto documentation (epydoc)





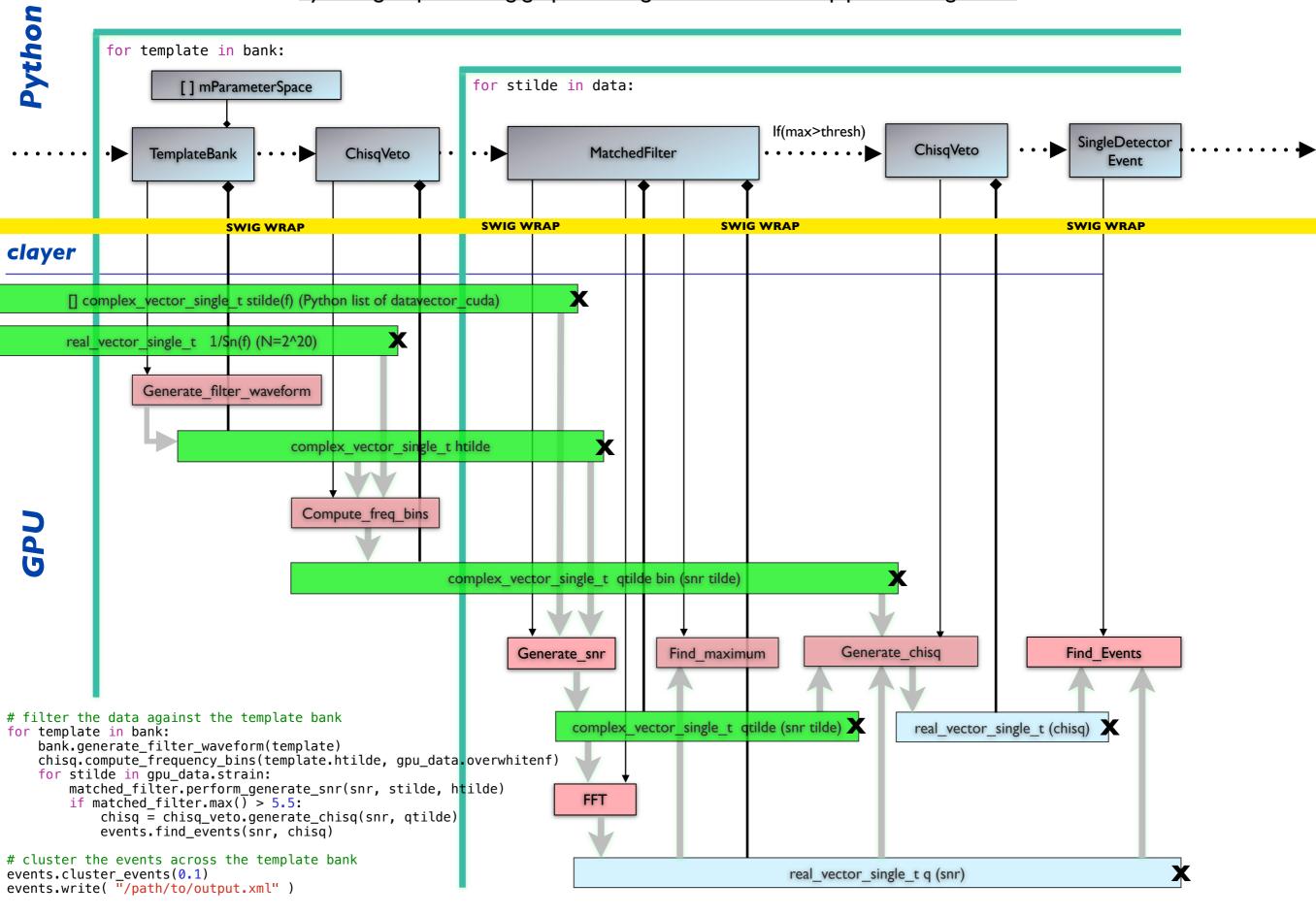
Inspiral processing graphs

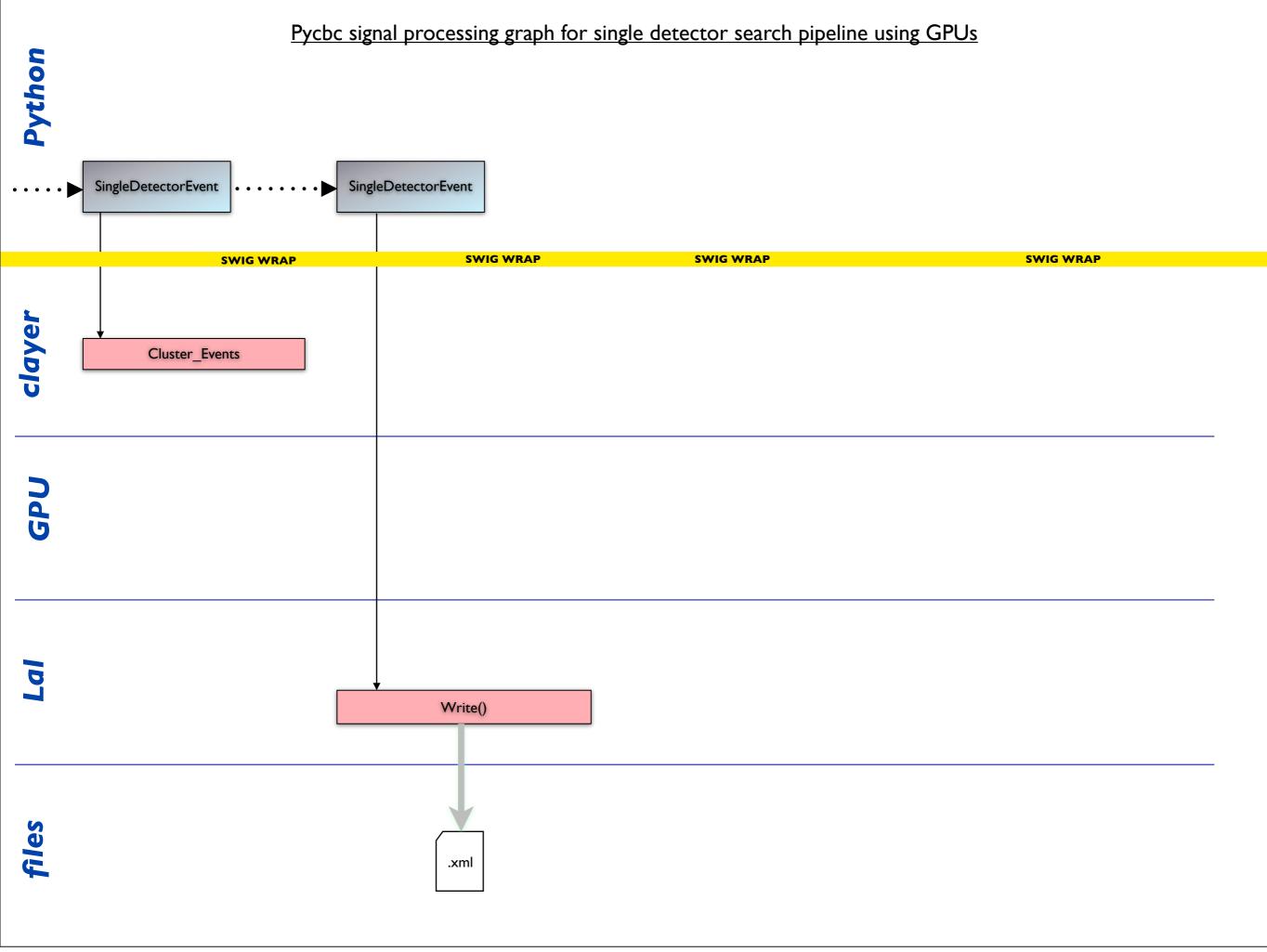
- Planned how this code will look based on lalapps_inspiral
 - in particular where different processes will be run
- Created processing graphs
 - Shows how functions and kernels are called and where they live (layers)
 - Shows how data flows
 - Shows memory objects
 - Where they live
 - Who is the owner
 - Which kind of data do they hold
 - How long they live



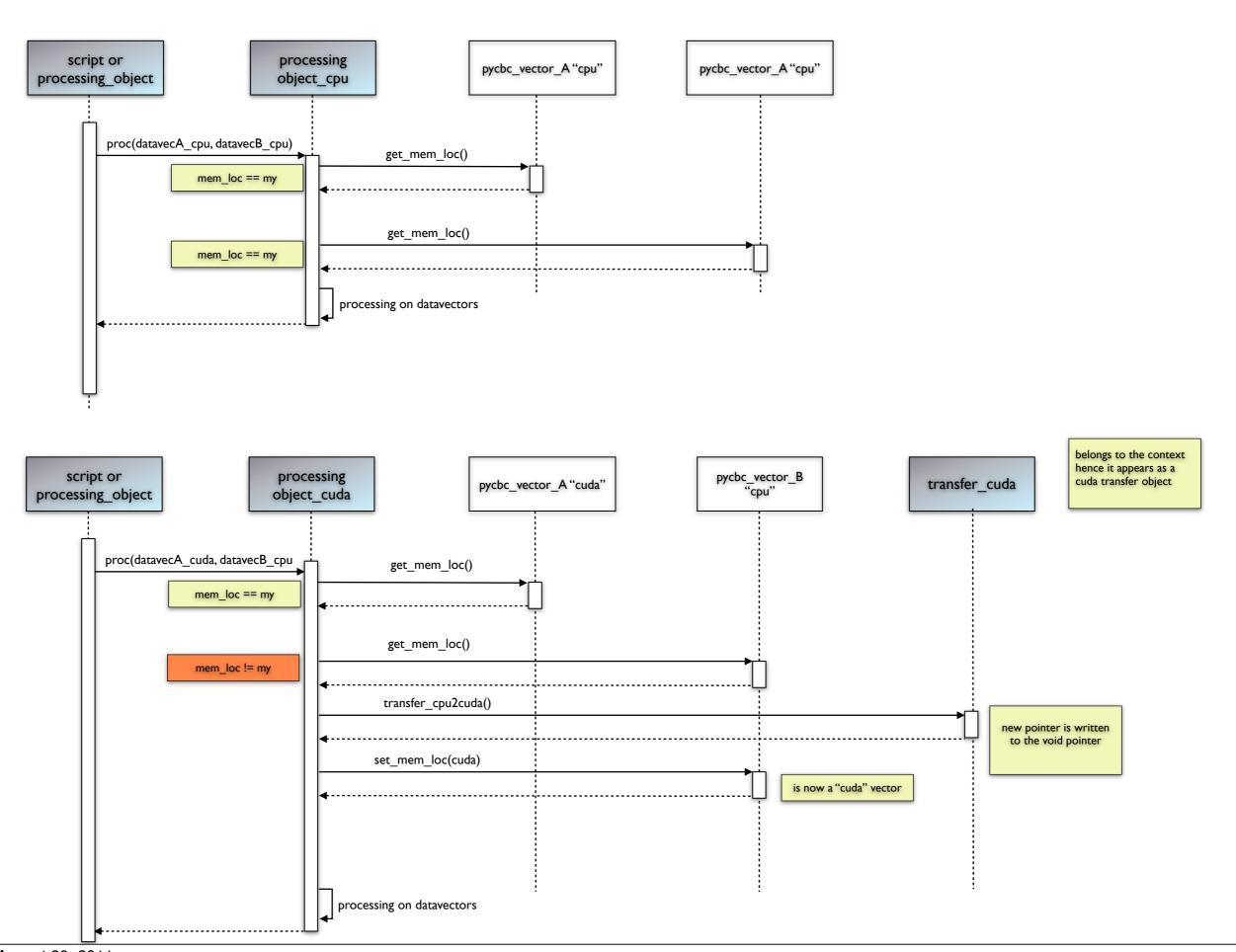
Pycbc signal processing graph for single detector search pipeline using GPUs [typ. 10³] ParameterSpace Overwhiten render() TemplateBank. Overwhitening Straindata. Straindata *= Straindata. **TemplateBank** PreConditionData() render() Segmenting() Filter Overw.-Filt **SWIG WRAP SWIG WRAP SWIG WRAP** X _vector_single_t s(t) [typ. 15 segments freq data from typ 256s timedomain data -> (N=2^20)] complex_vector_single_t_stilde(f) (Python list of datavector_cpu) X FFTs of overlapping input data real_vector_single_t 1/Sn(f) alias inverse PSD alias overwhitening filter N=2^20 render render median - inverse _mul() IFFT - truncate - FFT (inplace) [] complex_vector_single_t stilde(f) real_vector_single_t 1/Sn(f) **GLUE XMLread** files .xml

Pycbc signal processing graph for single detector search pipeline using GPUs

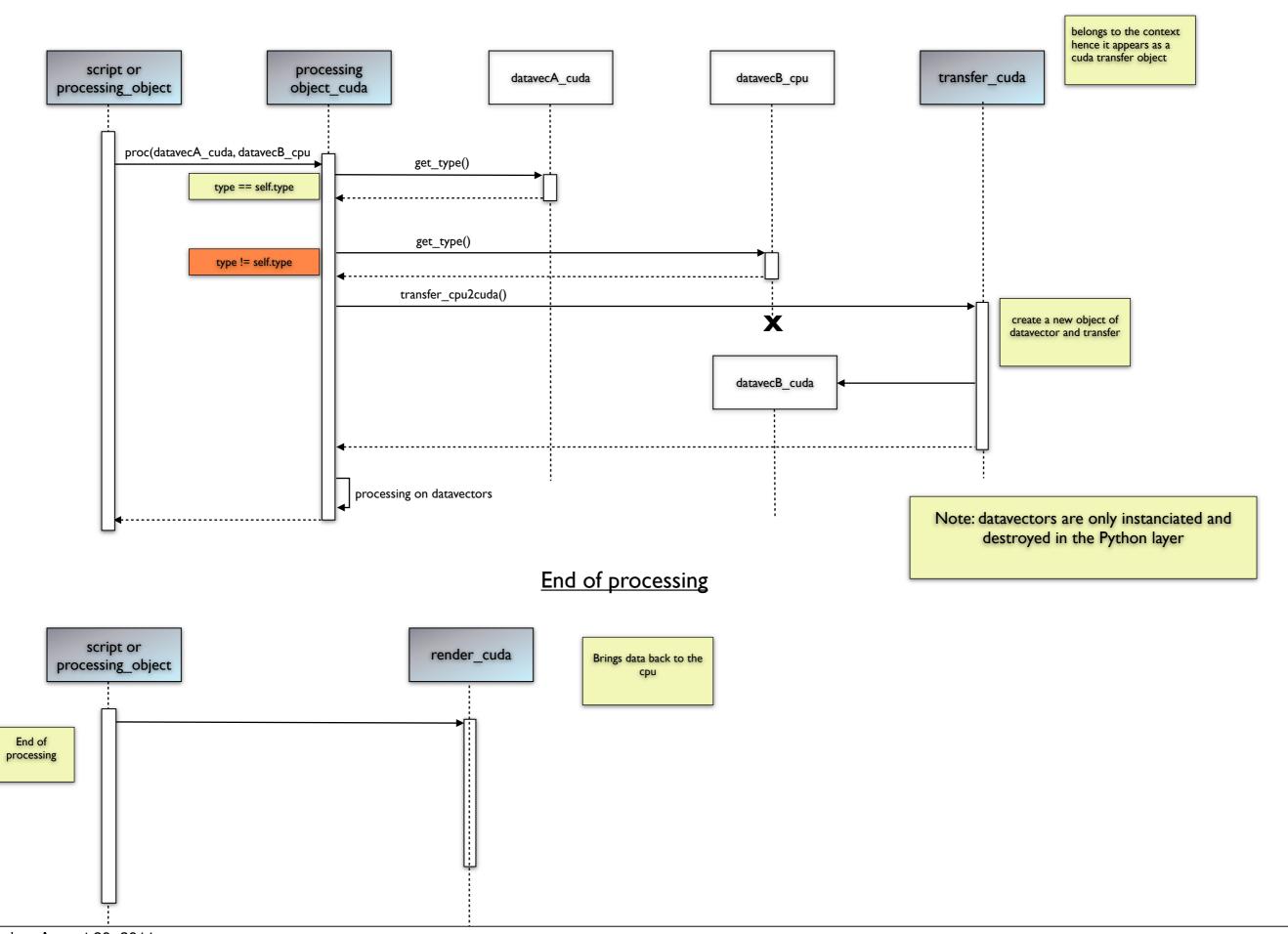




Device memory flow to and from cpu



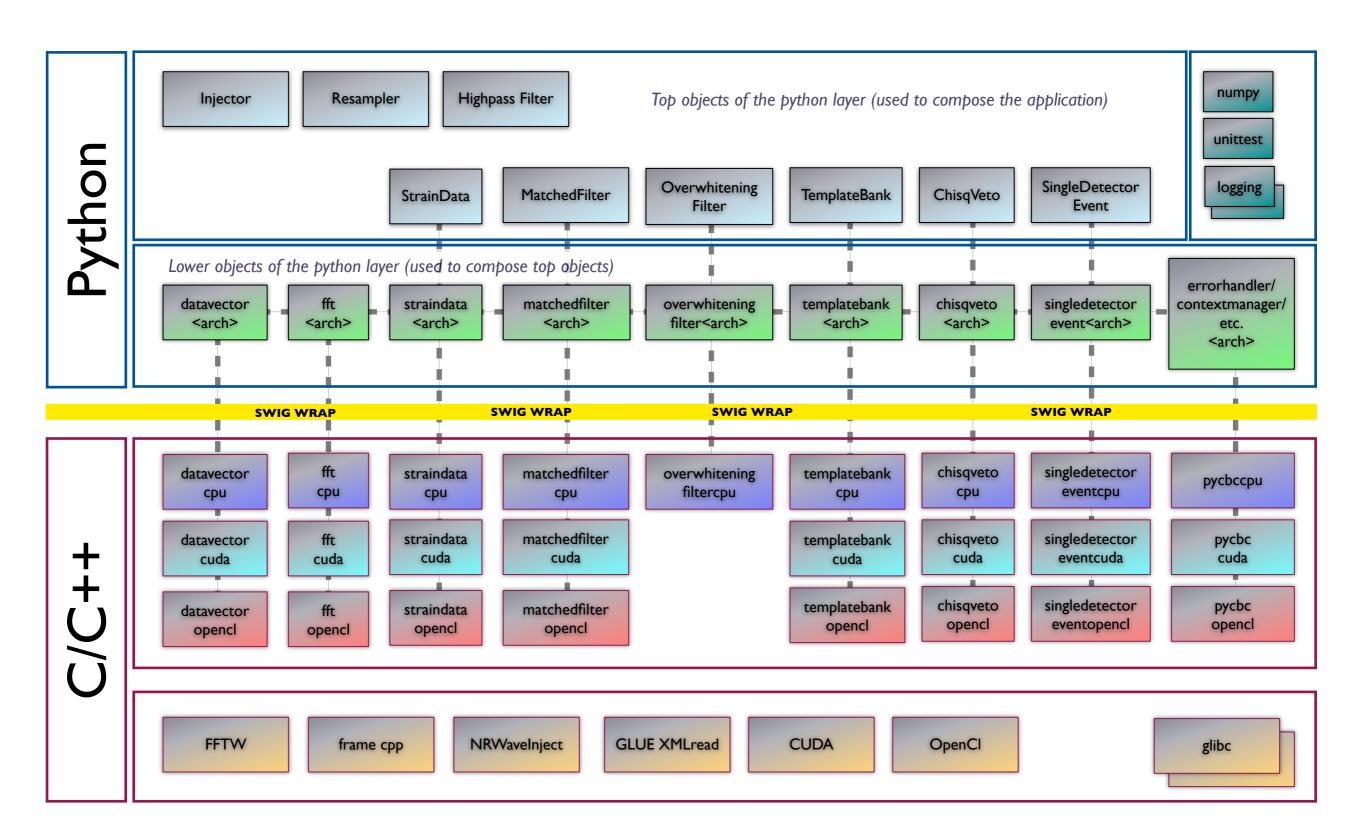
Device memory flow to and from cpu object orientated







Pycbc Layers and packages



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Inspiral analysis in a single slide...

```
import sys
import random
# preliminary hard coded path to packages
sys.path.append('/Users/kawies/dev/src/pycbc')
from pycbc.pycbc import CPUDeviceContext as ProcessingTargetContext
from pycbc.straindata.straindata_cpu import StrainDataCpu as StrainData
from pycbc.templatebank.templatebank_cpu import TemplateBankCpu as TemplateBank
from pycbc.matchedfilter.matchedfilter_cpu import MatchedFilterCpu as
MatchedFilter
from pycbc.chisqveto.chisqveto_cpu import ChisqVetoCpu as ChisqVeto
from pycbc.singledetectorevent.singledetectorevent cpu import
SingleDetectorEventCpu as SingleDetectorEvent
from pycbc.datavector.datavectorcpu import real_vector_single_t as
SnrResultTimeSeries
import logging
logging.basicConfig(level=logging.DEBUG,
                    format='%(name)s %(asctime)s %(levelname)s %(message)s',
                    filename='pycbc_min_pipeline.log',
                    filemode='w')
logger= logging.getLogger('pycbc.main script')
start message = 'Starting pycbc single detector minimal pipeline ...'
logger.debug(start message)
print start_message
# setup straindata
search_time = 128 # typ design spec: 2048
sample freq = 256 # typ design spec: 4096
length =
              search_time * sample_freq
segments = 15
gps start time= 871147532
gps end time= gps start time + search time
interferometer = "H1"
# setup chisa veto
chisq_bins = 16
# setup event finder
snr threshold = 5.5
with ProcessingTargetContext(1) as context:
   strain_data = StrainData(gps_start_time, gps_end_time,
                        segments, sample_freq,
                        interferometer, context)
   # initialize straindata w/ noise
    for i in range(length):
        tmp = random.uniform(-1,1)
        strain data.time series[i] = tmp
```

```
# convert straindata to single precision
   strain data.convert to single preci()
   # segmenting straindata and transform into frequency domain
   strain data.perform fft segments()
   # transfer straindata to appropriate memory space on target device
   strain_data.render()
   # create 5 templates (testing the iterator of TemplateBank)
   bank = TemplateBank(5, strain_data.segments_length,
                          strain_data.segments_delta_x, context)
    logger.debug("instantiated TemplateBank w/ waveform length: {0}"
                 .format(bank.waveform_length))
   # create matched filter (only generate snr() has to be implemented
   # for the minimal pipeline)
   matched_filter = MatchedFilter(strain_data.segments_length, context)
    logger.debug("instantiated MatchedFilter w/ length: {0}".format
    (matched_filter.length))
   # instantiate result vectors
   snr = SnrResultTimeSeries(strain_data.segments_length,
                             strain_data.segments_delta_x, context)
    logger.debug("instantiated SnrResultTimeSeries as {0}".format(repr(snr)))
   # instantiate chisq veto
   chisq veto = ChisqVeto(strain data.segments length, chisq bins, context)
    logger.debug("instantiated ChisqVeto as {0}".format(repr(chisq_veto)))
   # instantiate event finder
   events = SingleDetectorEvent(snr_threshold, context)
    logger.debug("instantiated SingleDetectorEvent as {0}".format(repr(events)))
   # filter the data against the template bank
   for template in bank:
       htilde = bank.perform generate waveform(template)
       for stilde in strain_data:
           matched filter.perform generate snr(stilde, htilde, snr)
           if matched filter.max() > snr threshold:
               chisq = chisq veto.generate chisg(snr, gtilde)
               events.find events(snr, chisq)
   # cluster the events across the template bank
   events.cluster events(0.1)
   events.write("/path/to/output.xml")
   # leaving the ProccessingTargetContext NOW (destroy the device context)
   end_message = '... end of pycbc single detector minimal pipeline.'
logger.debug(end message)
print end message
```





Current Status

- Codes in progress
 - prototype OpenCL implementation (Budapest)
 - prototype CUDA implementation (Syracuse)
 - prototype framework (AEI)
 - working on integrating different implementations
- Started work as a GPU project
- Currently in a stage where we are rewriting lalapps_inspiral and want to open it up to the group
- Starting to have regular telecons (Tuesdays 2 hours before the CBC telecon)
- For discussion, please mail cbc+pycbc@gravity.phys.uwm.edu
- Can view repository at https://sugwg-git.phy.syr.edu/dokuwiki/doku.php?
 id=pycbc:home