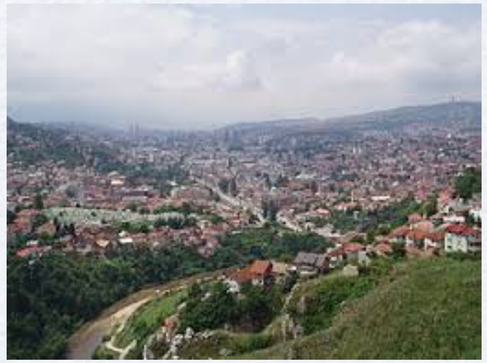
New Machine Learning Developments in ROOT/TMVA

L. Moneta (CERN EP-SFT)
on behalf of the ROOT/TMVA development team



ROOT Users Workshop

10-13 September 2018, Sarajevo

TMVA

- ROOT Machine Learning tools are provided in the package TMVA (Toolkit for MultiVariate Analysis)
- Provides a set of algorithms for standard HEP usage
- Used in LHC experiment production and in several analyses
  - several publications produced using TMVA
- Development done in collaboration with CERN experiments and HEP community
- HEP Software Foundation (HSF) community:
  - Machine Learning white paper
  - importance of providing internal machine learning software tools for HEP

Key Features of TMVA

- Facilitates HEP research, from detector to analysis
  - best suitable for HEP analysis with direct connection to ROOT I/O
  - written in C++
- Good performance (makes use of GPU and CPU parallelisation)
- Stability of interfaces
- Easy to use
- Long term support
- **Challenge in integrating new algorithms**
  - Machine learning world evolves very fast
- **Several features added recently** (e.g. deep learning)
- **Interfaces to integrate external tools** easily (from Python and R)

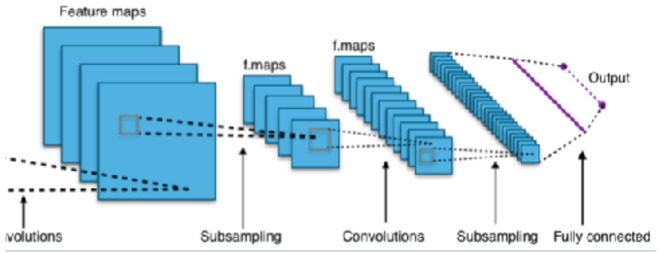
TMVA

Outline

- New machine learning methods for TMVA
  - Deep learning module
    - current status
    - performance tests with comparison to Keras/Tensorflow
- A look into the future
  - planned new developments
- Summary and conclusions

New Developments in TMVA

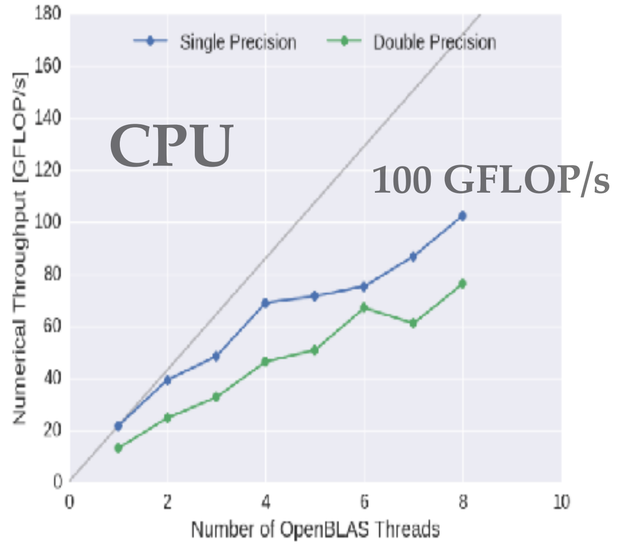
- New features available in latest ROOT version 6.14:
  - Extended Deep Learning Module with support for
    - Dense Layer
    - Convolutional Layer
    - Recurrent Layer

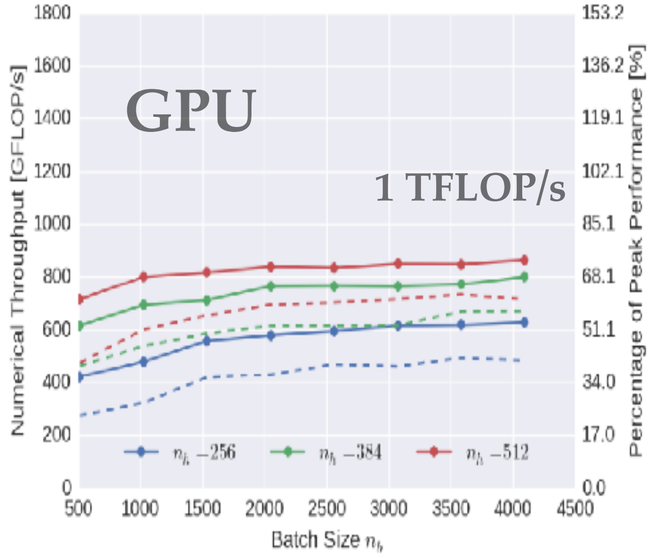


  - improved BDT performance using multi-thread parallelisation
  - improved Cross Validation
- And also available since few older versions:
  - interfaces to external tools (scikit-learn, Keras, R)

Deep Learning in TMVA

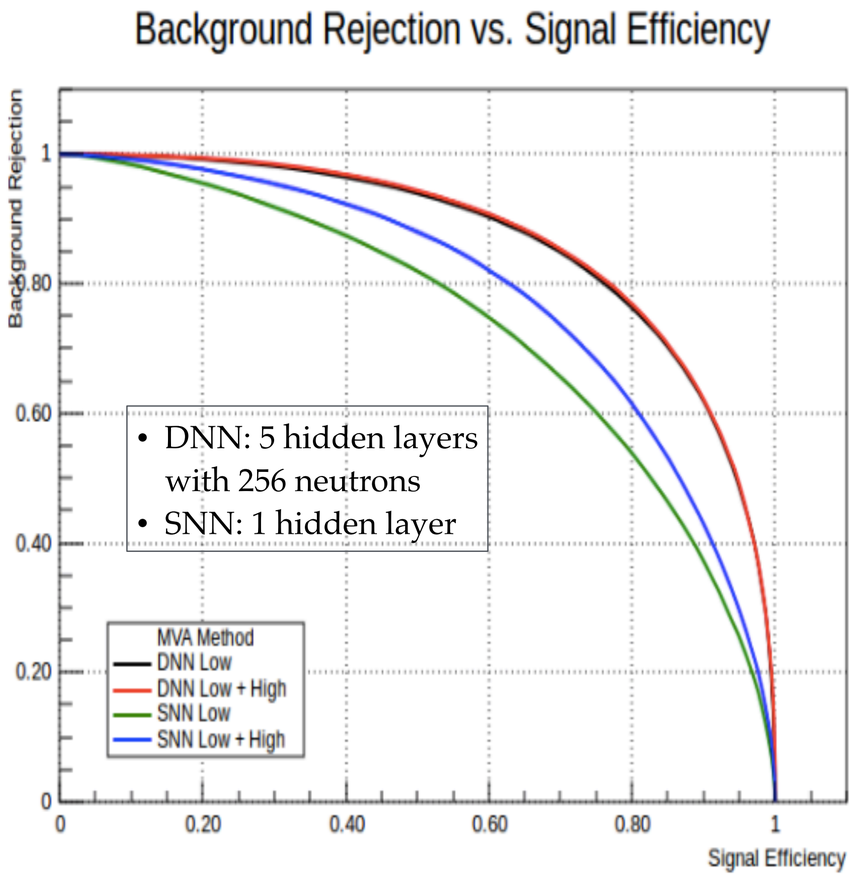
- Available for dense layers since ROOT 6.08
- extended the design in 6.14 to a new module supporting different layer types
  - parallel evaluation on CPU
    - implementation using OpenBlas and Intel TBB library
  - GPU support using CUDA
  - Excellent performance and high numerical throughput
    - see the presentation from S. Pfreundschuh



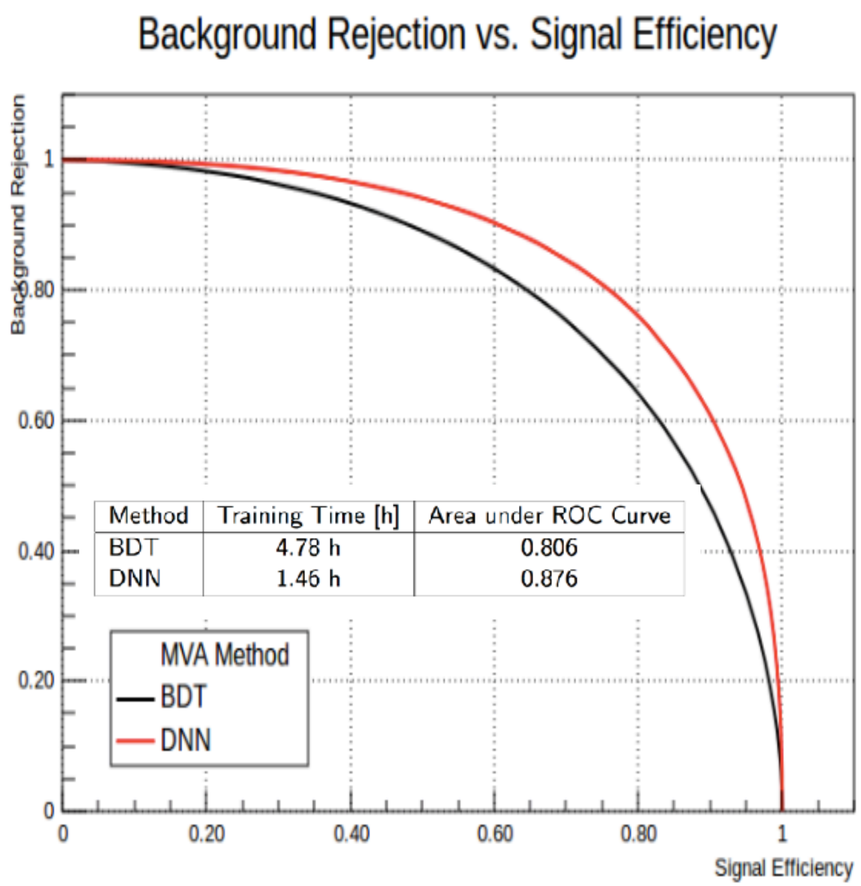


Deep Learning Performance

DNN vs Standard ANN



DNN vs BDT



- Using Higgs public dataset (from UCI) with 11M events
- Significant improvements compared to shallow networks and BDT

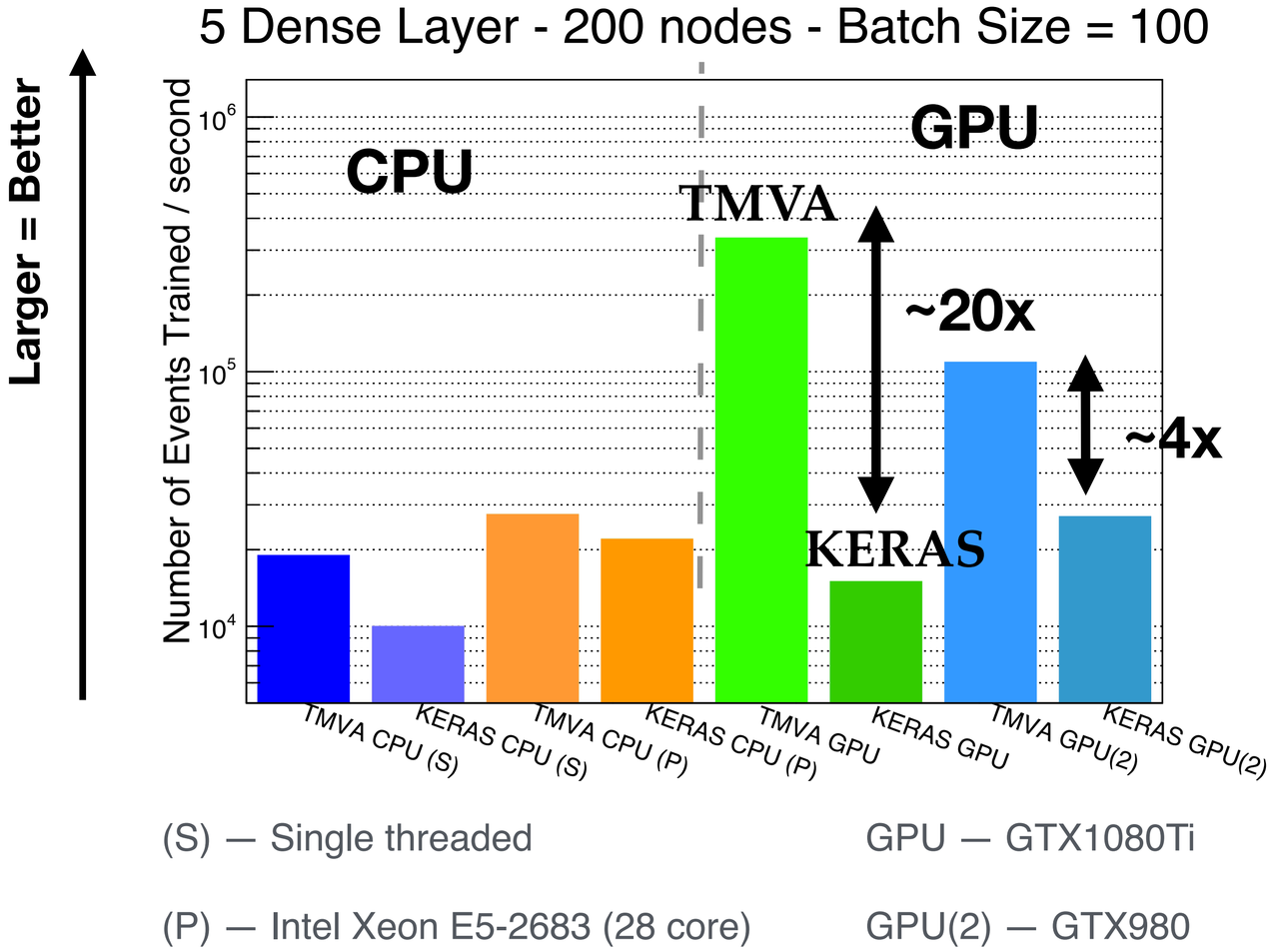
DNN Training Performance

Training time — Dense networks

- Higgs UCI dataset with 11M Events
- TMVA vs. Keras/Tensorflow
- "Out-of-the-box" performance

Excellent TMVA performance !

- How does it scale ?
  e.g. increasing batch size ?



(S) — Single threaded

(P) — Intel Xeon E5-2683 (28 core)

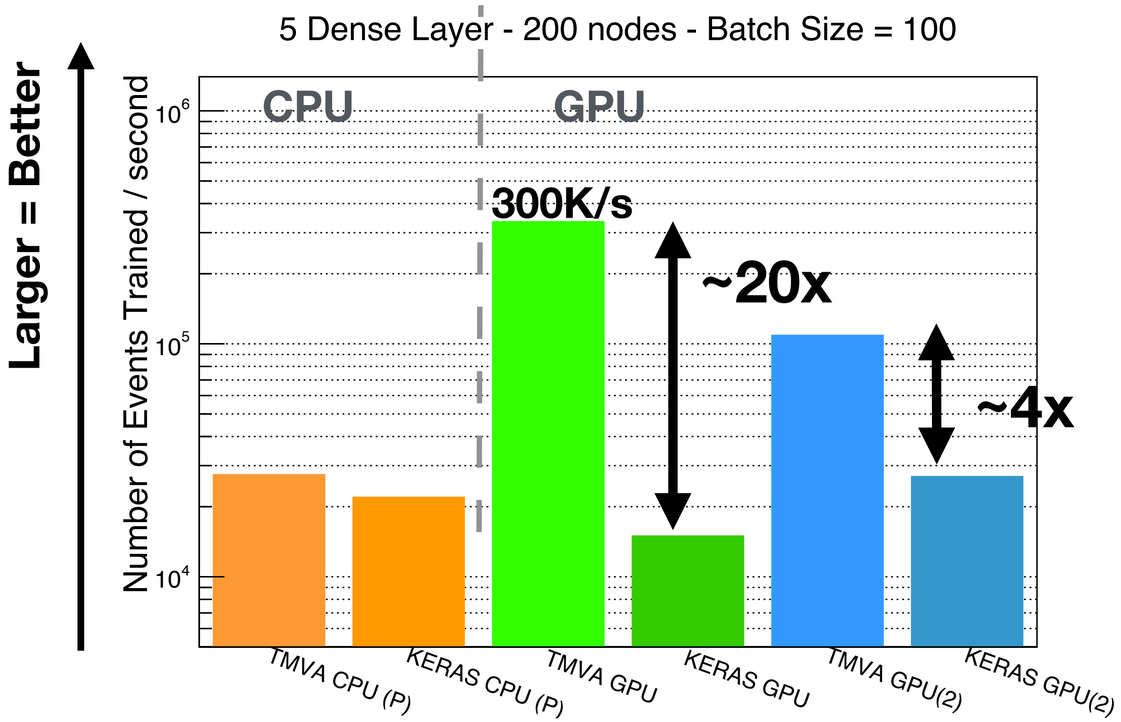
GPU — GTX1080Ti

GPU(2) — GTX980

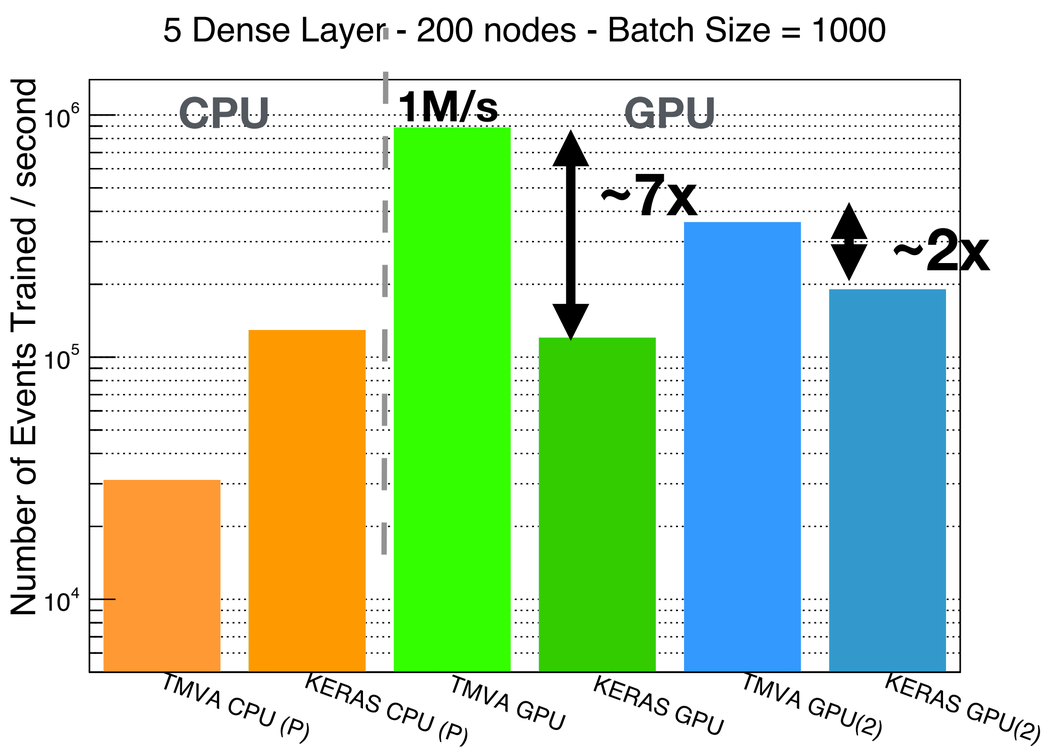
DNN Training Performance

TMVA vs. Keras/Tensorflow on CPU and GPU using a typical HEP dataset

Batch size 100



Batch size 1000



- Key difference is GPU utilisation
  - Tensorflow optimised for large operations

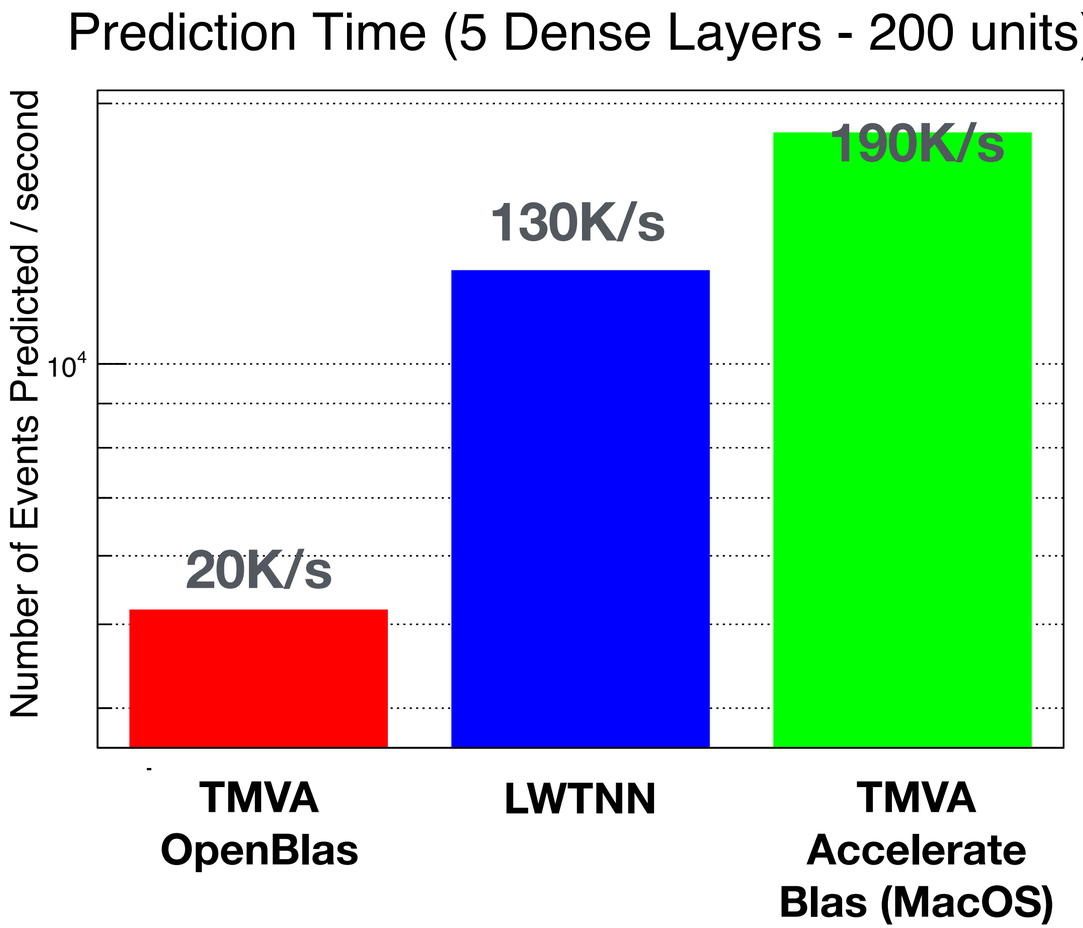
CPU— Intel Xeon E5-2683 (28 core)

GPU — GTX1080Ti

GPU(2) — GTX980

Evaluation Performance

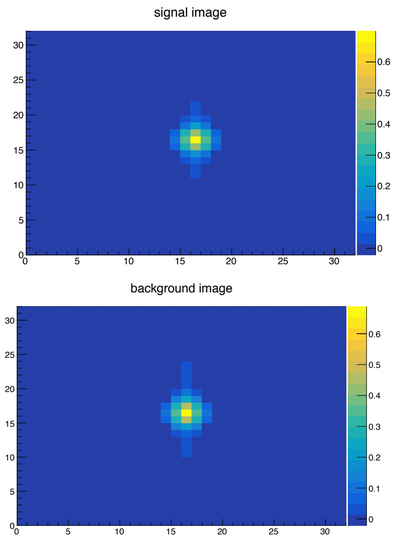
- **Single event evaluation time** for 5 layer network
  - For time critical applications — e.g. on-line reconstruction
  - Fast! 1.5 times speedup over specialised libraries like LWTNN when using optimised Blas library exploiting vectorisation
- For batched evaluation, same story as training

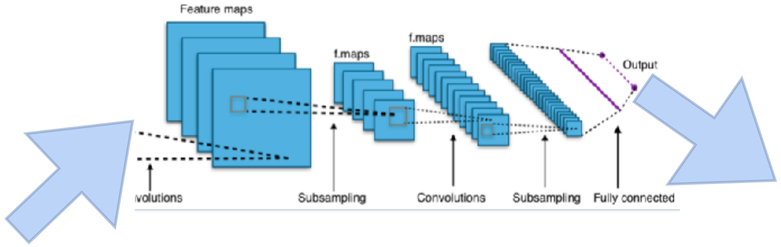


Convolutional Neural Network

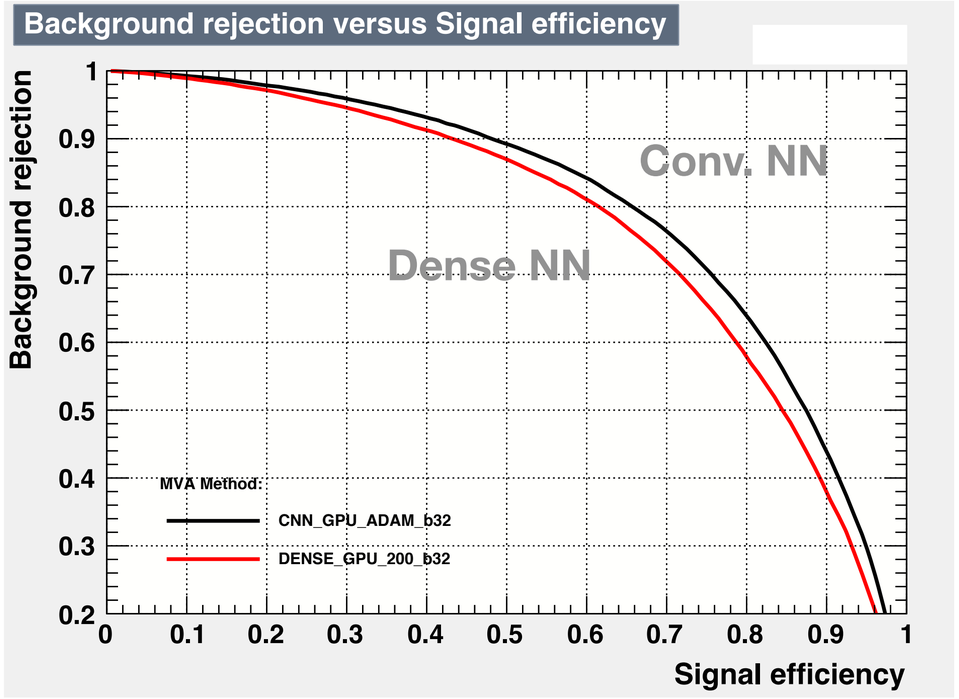
- Available in latest ROOT version (6.14)
- Supporting CPU parallelization, GPU is now also available
  - parallelisation and code optimisation is essential
- Image dataset from simulated particle showers from an electromagnetic calorimeter
  - distinguish electron from photon showers

input 32x32 images





Convolutional + Pooling + Dense layers

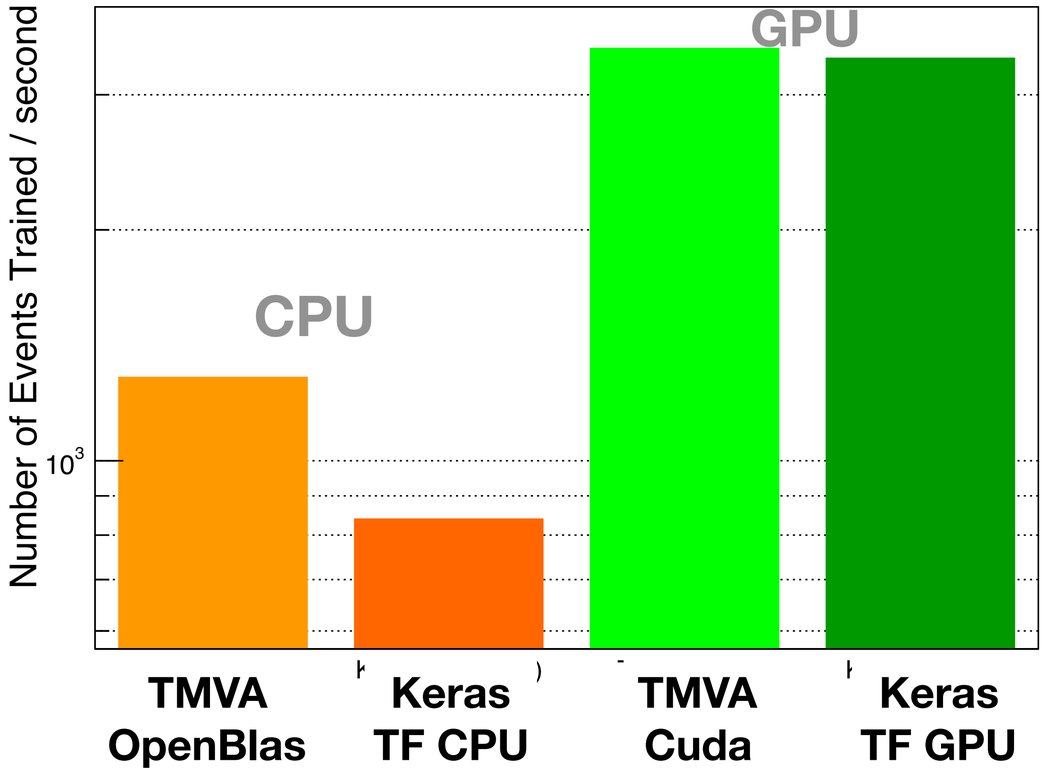


CNN Training Performance

CNN performance for TMVA CPU and GPU

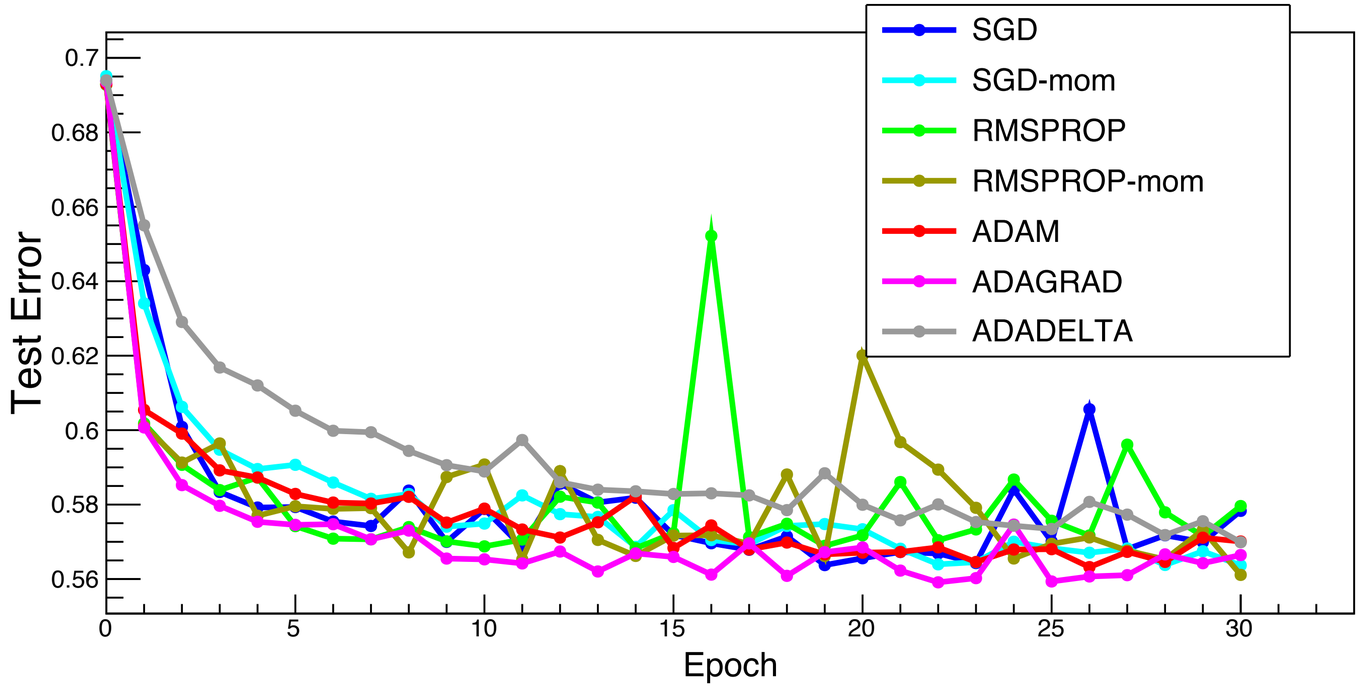
- Simulated particle showers from electromagnetic calorimeter image dataset
- TMVA GPU is now available in ROOT master
- again excellent TMVA performance for typical HEP networks !
- Code run already at same speed as Keras/Tensorflow on small/medium batch sizes
  - further optimisations are possible

4 Convolutional layers, Batch size = 32



New Deep Learning Optimizers

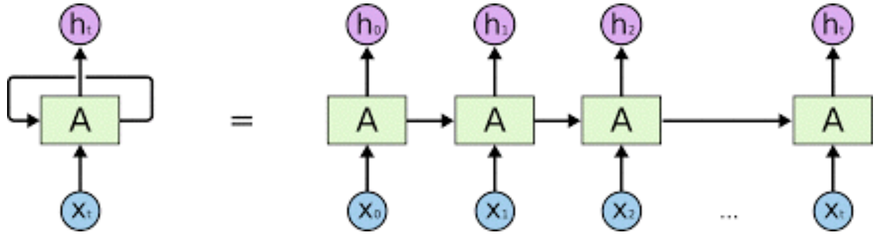
- Integrated in TMVA master new deep learning optimisers
  - In addition to SGD (Stochastic Gradient Descent) added
    - support acceleration using momentum
    - ADAM (new default)
    - ADADelta
    - ADAGrad
    - RMSProp



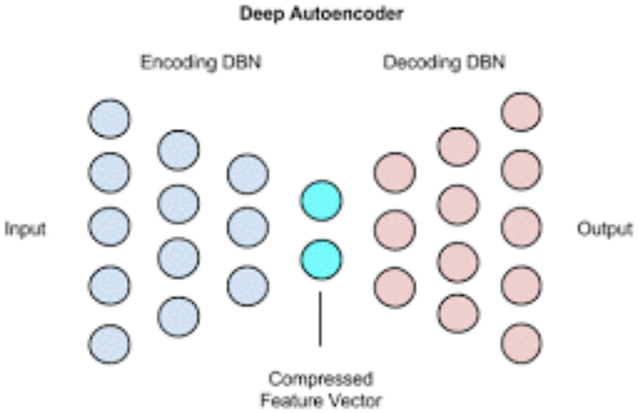
With these new optimisers need less epochs (iterations) to converge !

Other Deep Learning Developments

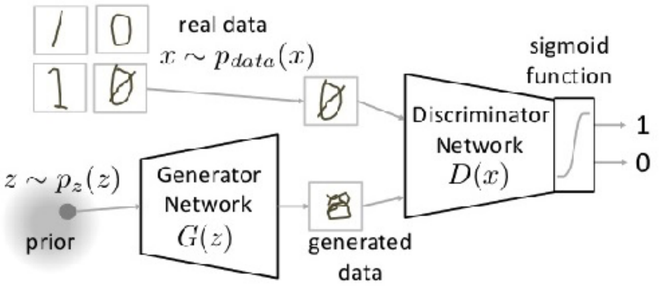
- **Recurrent Neural Network**
  - useful for time-dependent data
  - first version available in 6.14
  - extending to support LSTM layer



- **Deep Auto Encoder**
  - dimensionality reduction (pre-processing tool)
  - unsupervised tool (e.g. for anomaly detection)
  - usable also for generating models,
    Variational Auto Encoder (VAE)



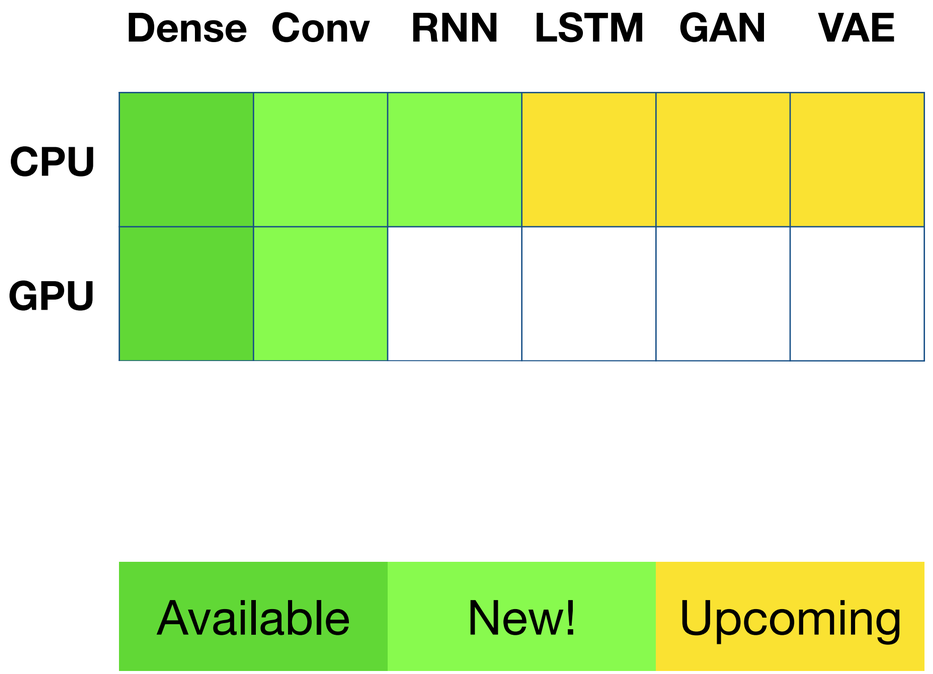
- **Generative Adversarial Network (GAN)**
  - model generation tool (fast simulation)



GSOC projects during this summer for these new developments

Summary Deep Learning

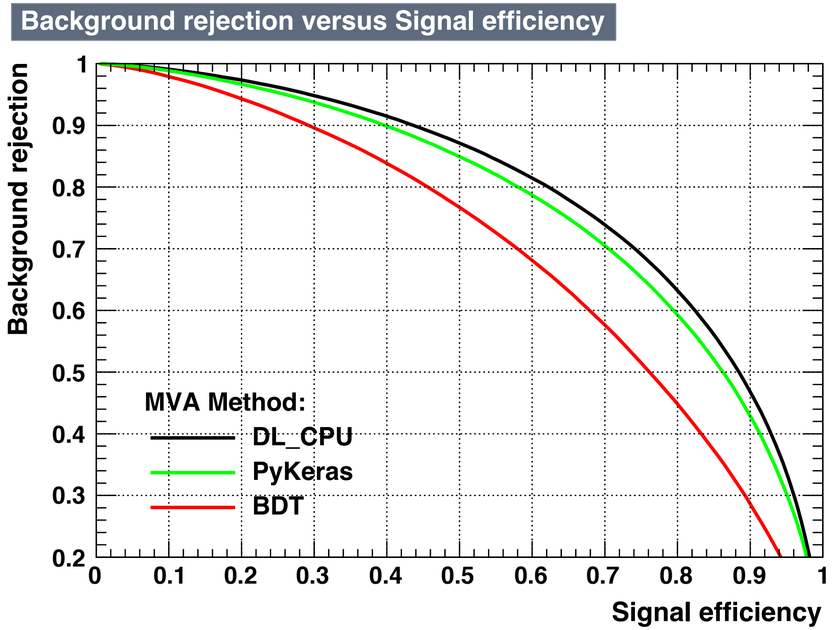
- Recent additions
  - Convolutional and recurrent layers
  - new optimisers complementing SGD
- Development ongoing!
  - LSTM (and also GRU) layers
  - GAN and VAE for event generation



TMVA Interfaces

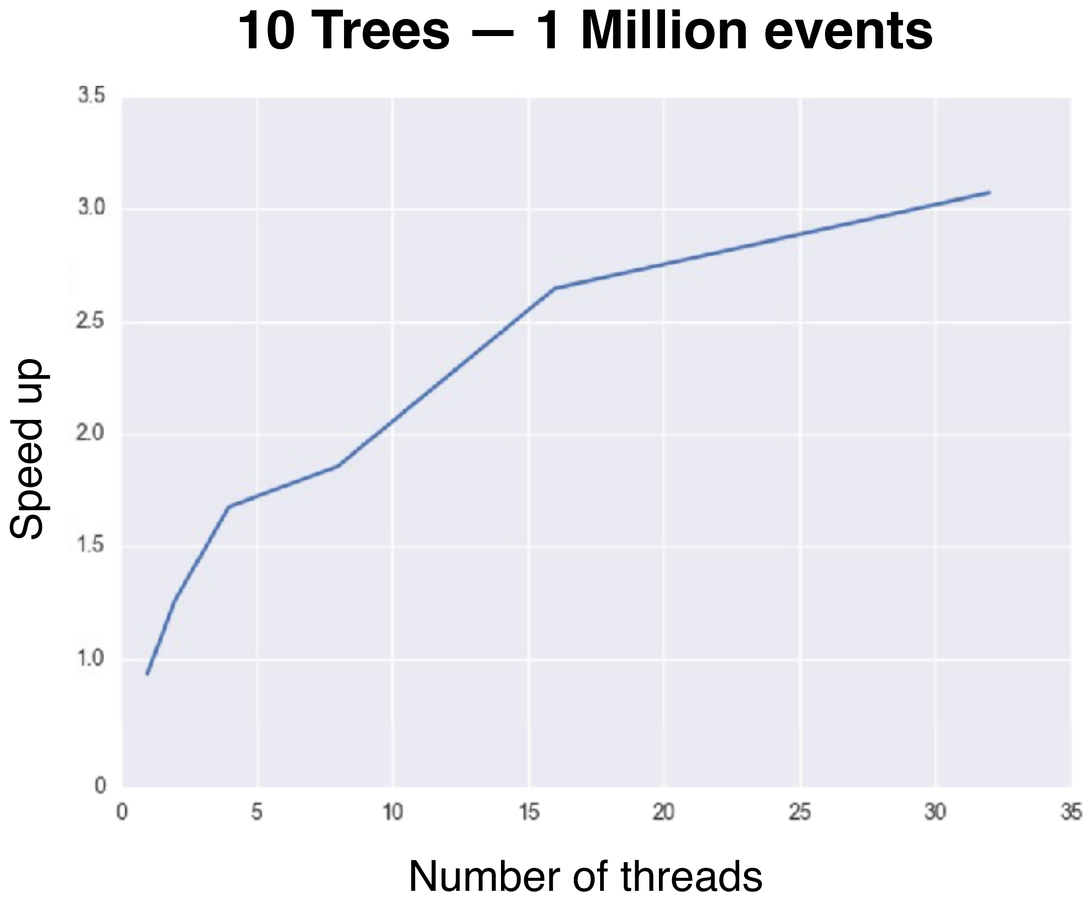
External tools are available as additional methods in TMVA and they can be trained and evaluated as any other internal ones.

- **RMVA**: **Interface to ML methods in R**
  - c50, xgboost, RSNNS, e1071
- **PYMVA**: **Interface to Python ML**
  - **scikit-learn**
    - with RandomForest,
      Gradiend Tree Boost, Ada Boost
  - **Keras** (Theano + Tensorflow)
    - support model definition in Python and then training and evaluation in TMVA
  - working on direct mapping from ROOT tree to Numpy arrays
    - see Stefan's presentation



Boosted Decision Tree

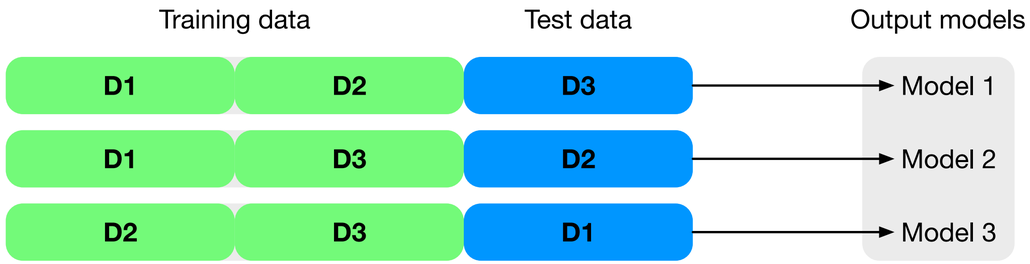
- Boosting is serial ⟶ Can't construct all trees in parallel
- Training time speed up ~1.6x with 4 threads approaching ~3x asymptotically
- To use, just add
  ROOT::EnableImplicitMT()
  to your code

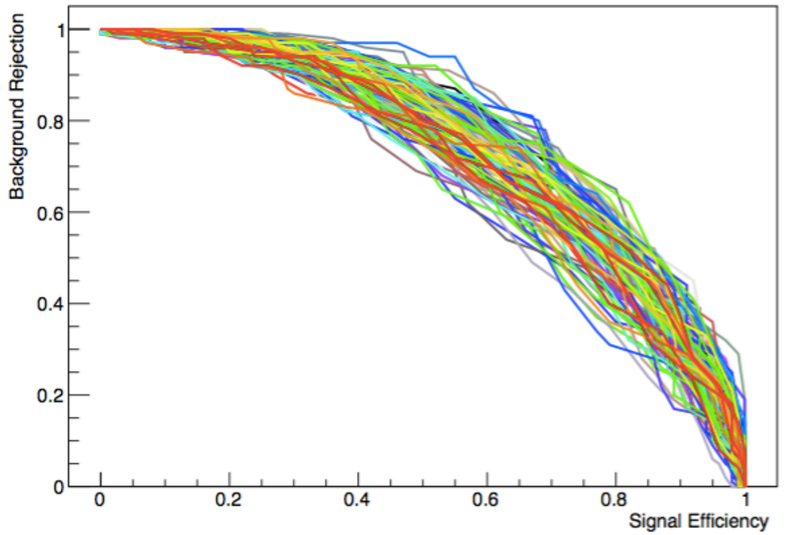


Original slide by Andrew Carnes

Cross Validation in TMVA

- TMVA supports k-fold cross-validation





  - Integration with TMVA analysis tools (e.g. GUI)
  - support for "CV in application"
- Hyper-parameter tuning
  - find optimised parameters (BDT-SVM)
- Parallel execution of folds in CV
  - using multi-processes execution in on a single node
  - foreseen to provide parallelisation in a cluster using Spark or MPI
- See Kim's presentation

Future Developments

- Our aim is to provide to the users community efficient physics workflows
  - tools for efficient
    - data loading (using new RDataFrame)
    - integration with external ML tools
    - training of commonly used architectures
    - deployment and inference of trained models
- TMVA efficiently connects input data to ML algorithms
  - we are defining new user interfaces (see Stefan's presentation)

Summary

Machine learning methods

- Dense, Convolutional and Recurrent networks in TMVA
- Excellent training + evaluation time performance
- Training in parallel Boosted Decision Trees

Workflow improvements

- Cross validation analysis and parallelisation

Future:

- Efficient physics workflows connecting input data to algorithms
  - integration with new RDataFrame and mapping to Numpy
  - fast deployment and inference of trained models

Conclusions

- Very active development happening in TMVA
  - several new features released recently
  - and even more expected in a near future
    - thanks to many student contributions (e.g. from Google Summer of Code)
- Strong competition, but hopefully still good reasons for continuing using TMVA !
- Feedback from users essential
  - best way to contribute is with Pull Request in GitHub https://github.com/root-project/root
  - ROOT Forum for user support with a category dedicated to TMVA https://root.cern/forum
  - JIRA for reporting ROOT bugs: https://sft.its.cern.ch/jira
  - or just contact us (TMVA developers) directly for any questions or issues

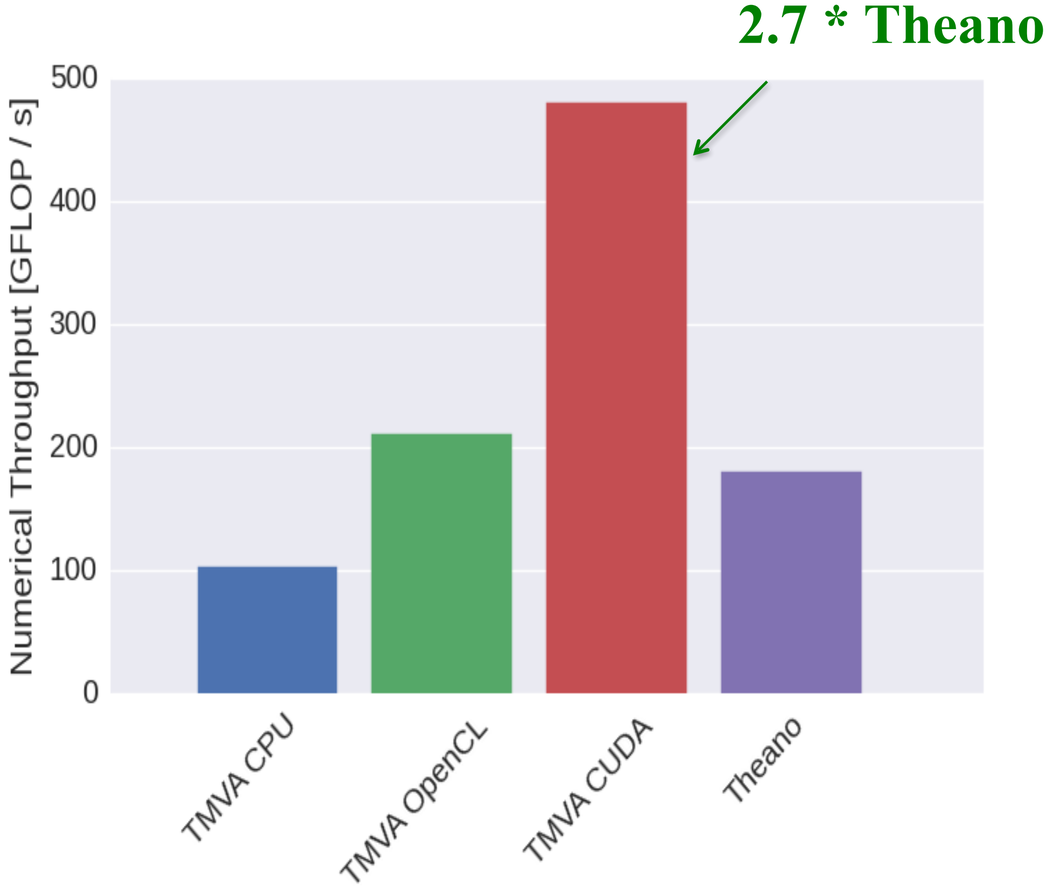
TMVA Contributors

Lorenzo Moneta	Algorithm development, Integration and support
Sergei Gleyzer	Analyzer Tools, Algorithm Development
Omar Zapata Mesa	PyMVA, RMVA, Modularity, Parallelization and Integration
Kim Albertsson	Multi-class for BDT, cross validation/evaluation and support
Stefan Wunsch	Keras Interface, integration, improved data handling
Peter Speckmeyer	Deep Learning CPU
Simon Pfreundschuh (GSOC 2016)	Deep Learning CPU and GPU
Vladimir Ilievski (GSOC 2017)	New Deep Learning module, Convolutional layers
Saurav Shekkar (GSOC 2017)	New Deep Learning module and Recurrent layers
Manos Stergiadis (GSOC 2018)	GPU support for CNN
Ravi Selvam (GSOC 2018)	New optimisers for deep learning
Adrian Bevan, Tom Stevenson	SVMs, Cross-Validation, Hyperparameter Tuning
Attila Bagoly (GSOC 2016)	Jupyter Integration, Visualization, Output
Paul Seyfert	Performance optimization
Andrew Carnes	Regression, Loss Functions, BDT Parallelization
Anurshee Rankawat, Siddhartha Rao, Harsit Prasad	GSOC 2008 projects: GAN, VAE and LSTM

And with continued invaluable contributions from Andreas Hoecker, Helge Voss, Eckhard v.Thorne, Jörg Stelzer

Backup Slides

Deep Learning Performance



Network:

- 20 input nodes,
- 5 hidden layers with 256 nodes each,
- *tanh* activation functions,
- squared error loss
- batch size = 1024
- Single precision

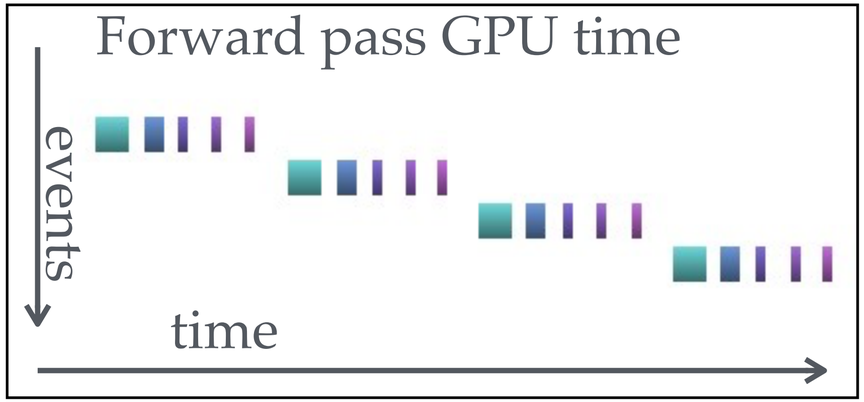
Training Data:

- Random data from a linear mapping $\mathbb{R}^n \rightarrow \mathbb{R}$

Excellent throughput compared to Theano on same GPU

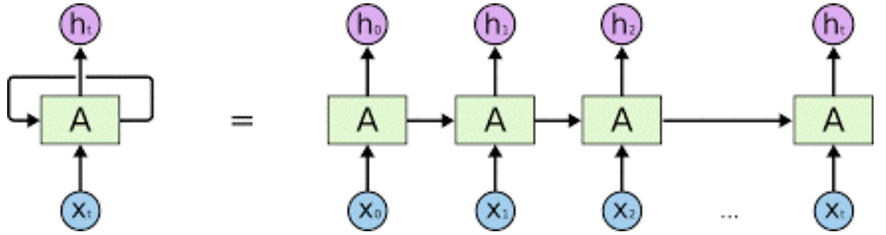
CNN Developments

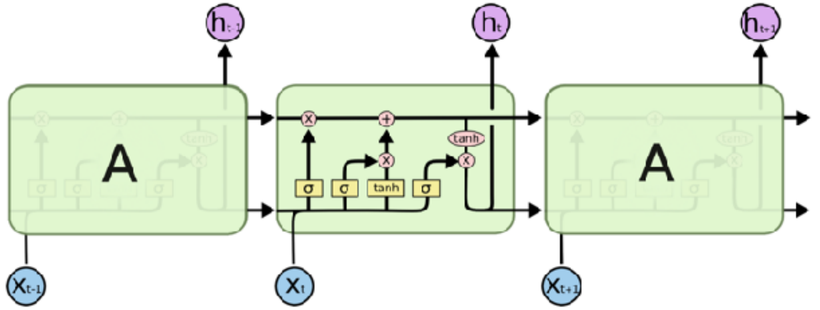
- GSOC student developed GPU implementation
  - excellent performances already obtained
  - further optimisation are possible using parallelisation within batches
    - require modifying used data structure for GPU
      - using a tensor with (batch size x depth x image size)
      - perform loop for events in a batch in the GPU
  - Further planned developments
    - support for 1D and 3D convolutions
    - implement transpose (inverse) convolution (for generative models)

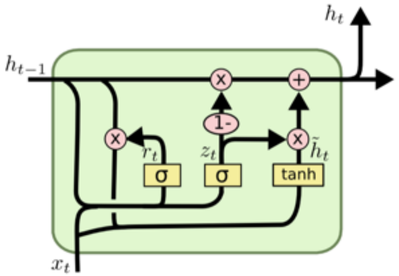


Recurrent Network

- Added in 6.14 first implementation of a recurrent layer
  - GSOC 2017 project
- RNN are very useful for time depend data
  - several applications in HEP (e.g. flavour tagging)
- 2018 GSOC project for developing a LSTM layer
  - LSTM (Long Short Term Memory) can cope with long term dependencies in the sequence
- Work is not completed, but plan to complete and integrate first version before end of the year
- Once LSTM layer is available also GRU (Gated Recurrent Unit) can be implemented

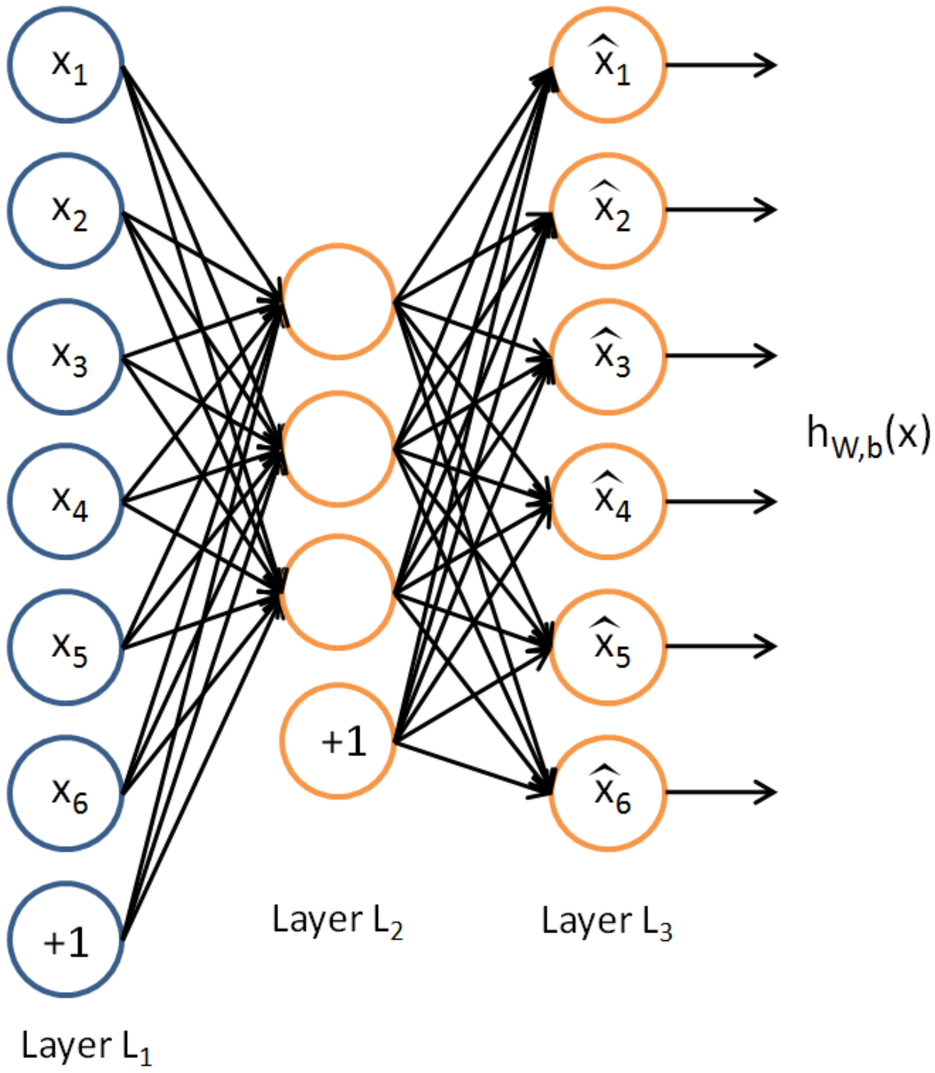






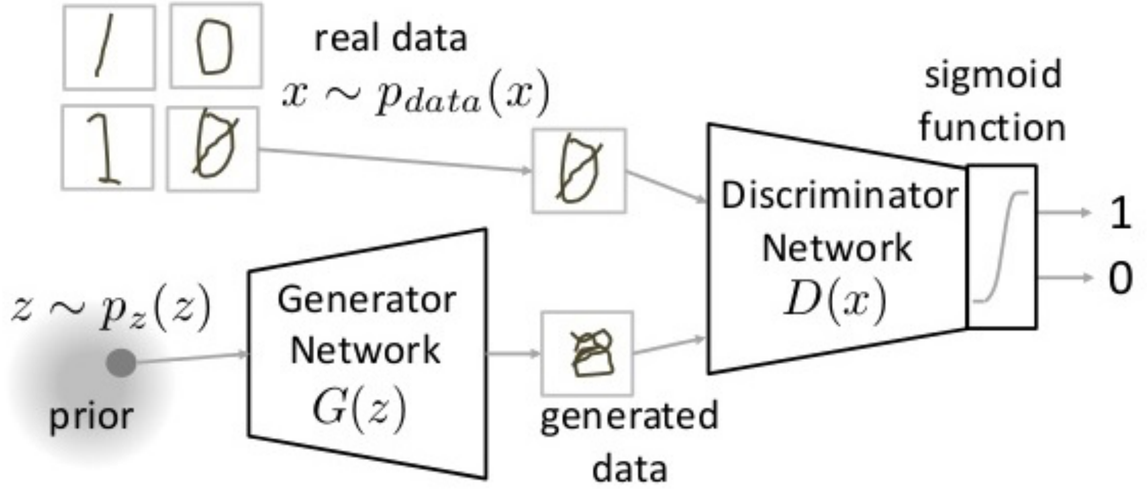
Deep Auto-encoder

- A neural network trained to learn the input data
  - Unsupervised machine learning methods
  - Useful for dimensionality reduction or anomaly detection
  - Can be used also as a generator
    - Variational Auto-encoder
- GSOC project on developing auto-encoders
  - implemented Kullback-Leiber divergence
  - MethodAE class for building auto-encoders
- Plan to integrate it in TMVA



Generative Adversarial Model

- GSOC project on developing a class for creating and training GAN based on the current TMVA DL library
- MethodGAN class
  - plan to complete and integrate in the ROOT master in the next months



Training of GAN can be difficult, min-max game optimisation

$$\min_G \max_D V(D,G) = \mathbb{E}_{\boldsymbol{x}\sim p_{\text{data}}(\boldsymbol{x})}[\log D(\boldsymbol{x})] + \mathbb{E}_{\boldsymbol{z}\sim p_{\boldsymbol{z}}(\boldsymbol{z})}[\log(1 - D(G(\boldsymbol{z})))].$$

Example PyMVA with Keras

Define model for Keras

Define the Keras model in Python

```
In [5]: # Define model
model = Sequential()
model.add(Dense(32, init='glorot_normal', activation='relu',
        input_dim=numVariables))
model.add(Dropout(0.5))
model.add(Dense(32, init='glorot_normal', activation='relu'))
model.add(Dropout(0.5))
model.add(Dense(2, init='glorot_uniform', activation='softmax'))

# Set loss and optimizer
model.compile(loss='categorical_crossentropy', optimizer=Adam(),
        metrics=['categorical_accuracy',])

# Store model to file
model.save('model.h5')

# Print summary of model
model.summary()
```

Book the method as any others of TMVA

Book methods

Just run the cells that contain the classifiers you want to try.

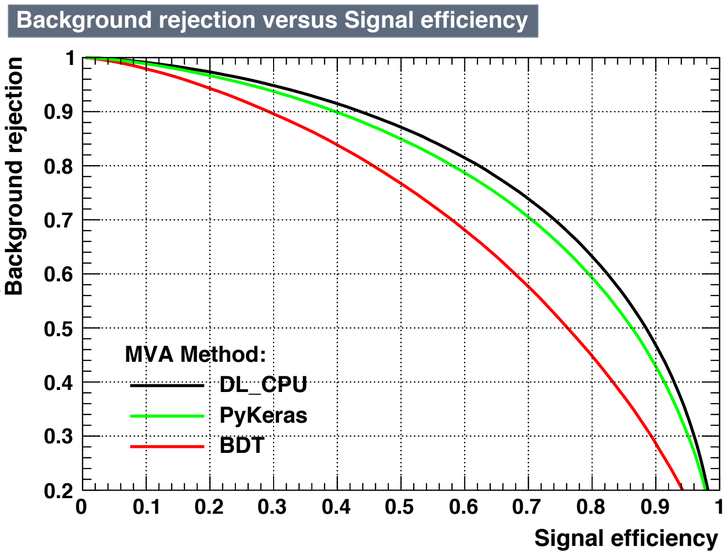
```
In [6]: # Keras interface with previously defined model
factory.BookMethod(dataloader, ROOT.TMVA.Types.kPyKeras, 'PyKeras',
        'H:!V:VarTransform=G:FilenameModel=model.h5:'+\
        'NumEpochs=10:BatchSize=32:'+\
        'TriesEarlyStopping=3')

Out[6]: <ROOT.TMVA::MethodPyKeras object ("PyKeras") at 0x77e48b0>
```

PyMVA with Keras

Train,Test and Evaluate inside TMVA (using TMVA::Factory)

Run training, testing and evaluation

```
In [8]: factory.TrainAllMethods()

Factory                  : Train all methods

In [9]: factory.TestAllMethods()

Factory                  : Test all methods
Factory                  : Test method: PyKeras
:

In [10]: factory.EvaluateAllMethods()

Factory                  : Evaluate all methods
Factory                  : Evaluate classifier: PyKeras
:
```



Examine result with TMVA GUI