



# Neural Network Design & Deployment

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# Contents

1	Pre	sentati	on 6
	1.1	Datab	ase handling
	1.2	Data 1	pre-processing
	1.3	Deep 1	network building
	1.4	Perfor	mances evaluation
	1.5	Hardw	vare exports
	1.6	Summ	ary 10
<b>2</b>	Abo	out N2	D2-IP 17
3	Per	formin	g simulations 11
•	3.1		ning the latest version of this manual
	3.2		num system requirements
	3.3		ning N2D2
	0.0	3.3.1	Prerequisites
		0.0.1	Red Hat Enterprise Linux (RHEL) 6
			Ubuntu
			Windows
		3.3.2	
		3.3.3	9
	9 4		1
	3.4		
	3.5		ne learning
	3.6		learned network
		3.6.1	Interpreting the results
			Recognition rate
			Confusion matrix
			Memory and computation requirements
			Kernels and weights distribution
		_	Output maps activity
	3.7	•	t a learned network
		3.7.1	C export
		3.7.2	CPP_OpenCL export         18
		3.7.3	CPP_TensorRT export
		3.7.4	CPP_cuDNN export
		3.7.5	C_HLS export
		3.7.6	Layer compatibility table
4	INI	file in	terface 22
-	4.1		K
	1.1	4.1.1	Properties
		4.1.2	Sections
		4.1.3	Case sensitivity
		4.1.4	Comments
		4.1.4 $4.1.5$	
		4.1.6	·
			1
	4.9	4.1.7	T. T
	4.2	_	ate inclusion syntax
		4.2.1	Variable substitution
		4.2.2	Control statements
			block

		if
		include
	4.3	Global parameters
	4.4	Databases
		4.4.1 MNIST
		4.4.2 GTSRB
		4.4.3 Directory
		Speech Commands Dataset
		4.4.4 Other built-in databases
		Actitracker_Database
		CIFAR10 Database
		CIFAR100_Database
		CKP_Database
		Caltech101_DIR_Database
		Caltech256_DIR_Database
		CaltechPedestrian_Database
		Cityscapes_Database
		Daimler_Database
		DOTA_Database
		<del>-</del>
		<del>-</del>
		GTSDB_DIR_Database
		ILSVRC2012_Database
		KITTI_Database
		KITTI_Road_Database
		KITTI_Object_Database
		LITISRouen_Database
		4.4.5 Dataset images slicing
	4.5	Stimuli data analysis
		4.5.1 Zero-mean and unity standard deviation normalization
		4.5.2 Substracting the mean image of the set
	4.6	Environment
		4.6.1 Built-in transformations
		AffineTransformation
		${\tt ApodizationTransformation} \; \ldots \; \ldots \; \ldots \; \ldots \; \ldots \; 36$
		ChannelExtractionTransformation
		${\tt ColorSpaceTransformation}  .  .  .  .  .  .  .  .  .  $
		DFTTransformation
2D2 IP only		${\tt DistortionTransformation}  .  .  .  .  .  .  .  .  .  $
2D2 IP only		EqualizeTransformation
2D2 IP only		ExpandLabelTransformation
		FilterTransformation
		FlipTransformation
2D2 IP only		${\tt GradientFilterTransformation} \; \ldots \; \ldots \; \ldots \; \ldots \; 39$
2D2 IP only		LabelSliceExtractionTransformation
		MagnitudePhaseTransformation
2D2 IP only		MorphologicalReconstructionTransformation
2D2 IP only		MorphologyTransformation
		NormalizeTransformation
		PadCropTransformation
2D2 IP only		RandomAffineTransformation
-		RangeAffineTransformation
		-

for  $\dots$ 

24

N2D2 IP only			RangeClippingTransformation
			RescaleTransformation
			ReshapeTransformation
N2D2 IP only			SliceExtractionTransformation
			ThresholdTransformation
			TrimTransformation
N2D2 IP only			WallisFilterTransformation
•	4.7	Netwo	rk layers
	1.,	4.7.1	Layer definition
		4.7.2	Weight fillers
		4.1.2	ConstantFiller
			HeFiller
			UniformFiller
		4 = 0	XavierFiller
		4.7.3	Weight solvers
			SGDSolver_Frame
			SGDSolver_Frame_CUDA
			AdamSolver_Frame
			AdamSolver_Frame_CUDA 49
		4.7.4	Activation functions
			Logistic
			LogisticWithLoss
			Rectifier 50
			Saturation
			Softplus
			Tanh
			TanhLeCun
		4.7.5	Anchor
		4.7.0	
		4 7 C	Outputs remapping
		4.7.6	Conv
			Configuration parameters (Frame models)
			Configuration parameters ( $Spike \mod els$ )
		4.7.7	Deconv
			Configuration parameters (Frame models)
		4.7.8	Pool
			Maxout example
			Configuration parameters (Spike models)
		4.7.9	Unpool
		4.7.10	ElemWise
			<b>Sum</b> operation
			AbsSum operation
			EuclideanSum operation
			Prod operation         61
			Max operation
			Examples
		4.7.11	1
		4.1.11	Configuration parameters (Frame models)
		1710	,
		4.1.12	Fc
			Configuration parameters (Frame models)
			Configuration parameters ( $Spike \mod ls$ )

		4.7.13	RDI
			Configuration parameters (Frame models)
		4.7.14	Softmax
		4.7.15	LRN
			Configuration parameters (Frame models)
		4.7.16	- ,
			Global layer parameters (Frame_CUDA models) 6
			Configuration parameters (Frame_CUDA models) 6
			Current restrictions
			Further development requirements
			Development guidance
		4.7.17	Dropout
			Configuration parameters (Frame models)
		4.7.18	Padding
			Resize
			Configuration parameters
		4.7.20	BatchNorm 6
			Configuration parameters (Frame models)
		4.7.21	Transformation
5	Tut	orials	7
	5.1	Learni	ng deep neural networks: tips and tricks
		5.1.1	Choose the learning solver
		5.1.2	Choose the learning hyper-parameters
		5.1.3	Convergence and normalization
	5.2	Buildi	ng a classifier neural network
	5.3	Buildi	ng a segmentation neural network
		5.3.1	Faces detection
		5.3.2	Gender recognition
		5.3.3	ROIs extraction
		5.3.4	Data visualization
	5.4	Transc	coding a learned network in spike-coding
		5.4.1	Render the network compatible with spike simulations
		5 4 2	Configure spike-coding parameters 8

N2D2 IP only

# 1 Presentation

The N2D2 platform is a comprehensive solution for fast and accurate Deep Neural Network (DNN) simulation and full and automated DNN-based applications building. The platform integrates database construction, data pre-processing, network building, benchmarking and hardware export to various targets. It is particularly useful for DNN design and exploration, allowing simple and fast prototyping of DNN with different topologies. It is possible to define and learn multiple network topology variations and compare the performances (in terms of recognition rate and computationnal cost) automatically. Export targets include CPU, DSP and GPU with OpenMP, OpenCL, Cuda, cuDNN and TensorRT programming models as well as custom hardware IP code generation with High-Level Synthesis for FPGA and dedicated configurable DNN accelerator IP<sup>1</sup>.

In the following, the first section describes the database handling capabilities of the tool, which can automatically generate learning, validation and testing data sets from any hand made database (for example from simple files directories). The second section briefly describes the data pre-processing capabilites built-in the tool, which does not require any external pre-processing step and can handle many data transformation, normalization and augmentation (for example using elastic distortion to improve the learning). The third section show an example of DNN building using a simple INI text configuration file. The fourth section show some examples of metrics obtained after the learning and testing to evaluate the performances of the learned DNN. Next, the fifth section introduces the DNN hardware export capabilities of the toolflow, which can automatically generate ready to use code for various targets such as embedded GPUs or full custom dedicated FPGA IP. Finally, we conclude by summarising the main features of the tool.

# 1.1 Database handling

The tool integrates everything needed to handle custom or hand made databases:

- Genericity: load image and sound, 1D, 2D or 3D data;
- Associate a label for each data point (useful for scene labeling for example) or a single label to each data file (one object/class per image for example), 1D or 2D labels;
- Advanced Region of Interest (ROI) handling:

Support arbitrary ROI shapes (circular, rectangular, polygonal or pixelwise defined); Convert ROIs to data point (pixelwise) labels;

Extract one or multiple ROIs from an initial dataset to create as many corresponding additional data to feed the DNN;

- Native support of file directory-based databases, where each sub-directory represents a different label. Most used image file formats are supported (JPEG, PNG, PGM...);
- Possibility to add custom datafile format in the tool without any change in the code base;
- Automatic random partitionning of the database into learning, validation and testing sets.

# 1.2 Data pre-processing

Data pre-processing, such as image rescaling, normalization, filtering... is directly integrated into the toolflow, with no need for external tool or pre-processing. Each pre-processing step is called a *transformation*.

The full sequence of transformations can be specified easily in a INI text configuration file. For example:

```
; First step: convert the image to grayscale
[env.Transformation-1]
Type=ChannelExtractionTransformation
CSChannel=Gray
```

<sup>&</sup>lt;sup>1</sup>Ongoing work

```
; Second step: rescale the image to a 29x29 size
[env.Transformation-2]
Type=RescaleTransformation
Width=29
Height=29
; Third step: apply histogram equalization to the image
[env.Transformation-3]
Type=EqualizeTransformation
; Fourth step (only during learning): apply random elastic distortions to the images to extent the
     learning set
[env.OnTheFlyTransformation]
Type=DistortionTransformation
ApplyTo=LearnOnly
ElasticGaussianSize=21
ElasticSigma=6.0
ElasticScaling=20.0
Scaling=15.0
Rotation=15.0
```

Example of pre-processing transformations built-in in the tool are:

- Image color space change and color channel extraction;
- Elastic distortion:
- Histogram equalization (including CLAHE);
- Convolutional filtering of the image with custom or pre-defined kernels (Gaussian, Gabor...);
- (Random) image flipping;
- (Random) extraction of fixed-size slices in a given label (for multi-label images)
- Normalization:
- Rescaling, padding/cropping, triming;
- Image data range clipping;
- (Random) extraction of fixed-size slices.

#### 1.3 Deep network building

The building of a deep network is straightforward and can be done withing the same INI configuration file. Several layer types are available: convolutional, pooling, fully connected, Radial-basis function (RBF) and softmax. The tool is highly modular and new layer types can be added without any change in the code base. Parameters of each layer type are modifiable, for example for the convolutional layer, one can specify the size of the convolution kernels, the stride, the number of kernels per input map and the learning parameters (learning rate, initial weights value...). For the learning, the data dynamic can be chosen between 16 bits (with NVIDIA® cuDNN<sup>2</sup>), 32 bit and 64 bit floating point numbers.

The following example, which will serve as the use case for the rest of this presentation, shows how to build a DNN with 5 layers: one convolution layer, followed by one MAX pooling layer, followed by two fully connected layers and a softmax output layer.

```
; Specify the input data format
[env]
SizeX=24
SizeY=24
BatchSize=12
; First layer: convolutional with 3x3 kernels
[conv1]
Input=env
Type=Conv
```

<sup>&</sup>lt;sup>2</sup>On future GPUs

```
KernelWidth=3
KernelHeight=3
NbOutputs=32
Stride=1
; Second layer: MAX pooling with pooling area 2x2
Input=conv1
Type=Pool
Pooling=Max
PoolWidth=2
PoolHeight=2
NbOutputs=32
Stride=2
Mapping.Size=1; one to one connection between convolution output maps and pooling input maps
; Third layer: fully connected layer with 60 neurons
[fc1]
Input=pool1
Type=Fc
NbOutputs=60
; Fourth layer: fully connected with 10 neurons
[fc2]
Input=fc1
Type=Fc
NbOutputs=10
; Final layer: softmax
[softmax]
Input=fc2
Type=Softmax
NbOutputs=10
WithLoss=1
[softmax.Target]
TargetValue=1.0
DefaultValue=0.0
```

The resulting DNN is shown in figure 1.

The learning is accelerated in GPU using the NVIDIA® cuDNN framework, integrated into the toolflow. Using GPU acceleration, learning times can be reduced typically by two orders of magnitude, enabling the learning of large databases within tens of minutes to a few hours instead of several days or weeks for non-GPU accelerated learning.

#### 1.4 Performances evaluation

The software automatically outputs all the information needed for the network applicative performances analysis, such as the recognition rate and the validation score during the learning; the confusion matrix during learning, validation and test; the memory and computation requirements of the network; the output maps activity for each layer, and so on, as shown in figure 2.

#### 1.5 Hardware exports

Once the learned DNN recognition rate performances are satisfying, an optimized version of the network can be automatically exported for various embedded targets. An automated network computation performances benchmarking can also be performed among different targets.

The following targets are currently supported by the toolflow:

• Plain C code (no dynamic memory allocation, no floating point processing);



Figure 1: Automatically generated and ready to learn DNN from the INI configuration file example.



Figure 2: Example of information automatically generated by the software during and after learning.

- C code accelerated with OpenMP;
- C code tailored for High-Level Synthesis (HLS) with Xilinx® Vivado® HLS; Direct synthesis to FPGA, with timing and utilization after routing;

Possibility to constrain the maximum number of clock cycles desired to compute the whole network;

FPGA utilization vs number of clock cycle trade-off analysis;

- OpenCL code optimized for either CPU/DSP or GPU;
- Cuda kernels, cuDNN and TensorRT code optimized for NVIDIA® GPUs.

Different automated optimizations are embedded in the exports:

- DNN weights and signal data precision reduction (down to 8 bit integers or less for custom FPGA IPs);
- Non-linear network activation functions approximations;
- Different weights discretization methods.

The exports are generated automatically and come with a Makefile and a working testbench, including the pre-processed testing dataset. Once generated, the testbench is ready to be compiled and executed on the target platform. The applicative performance (recognition rate) as well as the computing time per input data can then be directly mesured by the testbench.

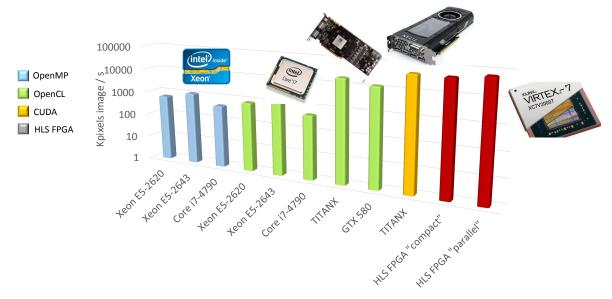


Figure 3: Example of network benchmarking on different hardware targets.

The figure 3 shows an example of benchmarking results of the previous DNN on different targets (in log scale). Compared to desktop CPUs, the number of input image pixels processed per second is more than one order of magnitude higher with GPUs and at least two orders of magnitude better with synthesized DNN on FPGA.

# 1.6 Summary

The N2D2 platform is today a complete and production ready neural network building tool, which does not require advanced knownledges in deep learning to be used. It is tailored for fast neural network applications generation and porting with minimum overhead in terms of database creation and management, data pre-processing, networks configuration and optimized code generation, which can save months of manual porting and verification effort to a single automated step in the tool.

# 2 About N2D2-IP

While N2D2 is our deep learning open-source core framework, some modules referred as "N2D2-IP" in the manual, are only available through custom license agreement with CEA LIST.

If you are interested in obtaining some of these modules, please contact our business developer for more information on available licensing options:

Sandrine VARENNE (Sandrine.VARENNE@cea.fr)

In addition to N2D2-IP modules, we can also provide our expertise to design specific solutions for integrating DNN in embedded hardware systems, where power, latency, form factor and/or cost are constrained. We can target CPU/DSP/GPU CoTS hardware as well as our own PNeuro (programmable) and DNeuro (dataflow) dedicated hardware accelerator IPs for DNN on FPGA or ASIC.

# 3 Performing simulations

#### 3.1 Obtaining the latest version of this manual

Before going further, please make sure you are reading the latest version of this manual. It is located in the manual sub-directory. To compile the manual in PDF, just run the following command:

cd manual && make

In order to compile the manual, you must have pdflatex and bibtex installed, as well as some common LaTeX packages.

- On Ubuntu, this can be done by installing the texlive and texlive-latex-extra software packages.
- On Windows, you can install the Miktex software, which includes everything needed and will install the required LaTeX packages on the fly.

#### 3.2 Minimum system requirements

• Supported processors:

ARM Cortex A15 (tested on Tegra K1)

ARM Cortex A53/A57 (tested on Tegra X1)

Pentium-compatible PC (Pentium III, Athlon or more-recent system recommended)

• Supported operating systems:

Windows  $\geq 7$  or Windows Server  $\geq 2012$ , 64 bits with Visual Studio  $\geq 2015.2$  (2015 Update 2)

GNU/Linux with GCC  $\geq 4.4$  (tested on RHEL  $\geq 6$ , Debian  $\geq 6$ , Ubuntu  $\geq 14.04$ )

- At least 256 MB of RAM (1 GB with GPU/CUDA) for MNIST dataset processing
- At least 150 MB available hard disk space + 350 MB for MNIST dataset processing

#### For CUDA acceleration:

- CUDA  $\geq 6.5$  and CuDNN  $\geq 1.0$
- NVIDIA GPU with CUDA compute capability ≥ 3 (starting from Kepler micro-architecture)
- $\bullet\,$  At least 512 MB GPU RAM for MNIST dataset processing

#### 3.3 Obtaining N2D2

#### 3.3.1 Prerequisites

Red Hat Enterprise Linux (RHEL) 6 Make sure you have the following packages installed:

- cmake
- gnuplot
- opencv
- opency-devel (may require the rhel-x86\_64-workstation-optional-6 repository channel)

Plus, to be able to use GPU acceleration:

• Install the CUDA repository package:

- Install cuDNN from the NVIDIA website: register to NVIDIA Developer and download the latest version of cuDNN. Simply copy the header and library files from the cuDNN archive to the corresponding directories in the CUDA installation path (by default: /usr/local/cuda/include and /usr/local/cuda/lib64, respectively).
- Make sure the CUDA library path (e.g. /usr/local/cuda/lib64) is added to the LD\_LIBRARY\_PATH environment variable.

**Ubuntu** Make sure you have the following packages installed, if they are available on your Ubuntu version:

- cmake
- gnuplot
- libopencv-dev
- libcv-dev
- libhighgui-dev

Plus, to be able to use GPU acceleration:

• Install the CUDA repository package matching your distribution. For example, for Ubuntu 14.04 64 bits:

```
wget http://developer.download.nvidia.com/compute/cuda/repos/ubuntu1404/x86_64/cuda-repo-ubuntu1404_7.5-18_amd64.deb
dpkg -i cuda-repo-ubuntu1404_7.5-18_amd64.deb
```

• Install the cuDNN repository package matching your distribution. For example, for Ubuntu 14.04 64 bits:

```
wget http://developer.download.nvidia.com/compute/machine-learning/repos/ubuntu1404/x86_64/nvidia-machine-learning-repo-ubuntu1404_4.0-2_amd64.deb
dpkg -i nvidia-machine-learning-repo-ubuntu1404_4.0-2_amd64.deb
```

Note that the cuDNN repository package is provided by NVIDIA for Ubuntu starting from version 14.04.

- Update the package lists: apt-get update
- Install the CUDA and cuDNN required packages:

```
apt-get install cuda-core-7-5 cuda-cudart-dev-7-5 cuda-cublas-dev-7-5 cuda-curand-dev-7-5 libcudnn5-dev
```

• Make sure there is a symlink to /usr/local/cuda:

```
ln -s /usr/local/cuda-7.5 /usr/local/cuda
```

• Make sure the CUDA library path (e.g. /usr/local/cuda/lib64) is added to the LD\_LIBRARY\_PATH environment variable.

**Windows** On Windows 64 bits, Visual Studio  $\geq$  2015.2 (2015 Update 2) is required. Make sure you have the following software installed:

- CMake (http://www.cmake.org/): download and run the Windows installer.
- dirent.h C++ header (https://github.com/tronkko/dirent): to be put in the Visual Studio include path.
- Gnuplot (http://www.gnuplot.info/): the bin sub-directory in the install path needs to be added to the Windows PATH environment variable.
- OpenCV (http://opencv.org/): download the latest 2.x version for Windows and extract it to, for example, C:\OpenCV\. Make sure to define the environment variable OpenCV\_DIR to point to C:\OpenCV\opencv\build. Make sure to add the bin sub-directory (C:\OpenCV\opencv\build\x64 \vc12\bin) to the Windows PATH environment variable.

Plus, to be able to use GPU acceleration:

• Download and install CUDA toolkit 8.0 located at https://developer.nvidia.com/compute/cuda/8.0/prod/local\_installers/cuda\_8.0.44\_windows-exe:

```
rename cuda_8.0.44_windows-exe cuda_8.0.44_windows.exe cuda_8.0.44_windows.exe -s compiler_8.0 cublas_8.0 cublas_dev_8.0 cudart_8.0 curand_8.0 curand_dev_8.0
```

• Update the PATH environment variable:

set PATH=%ProgramFiles%\NVIDIA GPU Computing Toolkit\CUDA\v8.0\bin;%ProgramFiles%\NVIDIA GPU Computing Toolkit\CUDA\v8.0\libnvvp;%PATH%

• Download and install cuDNN 8.0 located at http://developer.download.nvidia.com/compute/redist/cudnn/v5.1/cudnn-8.0-windows7-x64-v5.1.zip (the following command assumes that you have 7-Zip installed):

```
7z x cudnn-8.0-windows7-x64-v5.1.zip
copy cuda\include\*.* ^
   "%ProgramFiles%\NVIDIA GPU Computing Toolkit\CUDA\v8.0\include\"
copy cuda\lib\x64\*.* ^
   "%ProgramFiles%\NVIDIA GPU Computing Toolkit\CUDA\v8.0\lib\x64\"
copy cuda\bin\*.* ^
   "%ProgramFiles%\NVIDIA GPU Computing Toolkit\CUDA\v8.0\bin\"
```

## 3.3.2 Getting the sources

Use the following command:

```
git clone git@github.com:CEA-LIST/N2D2.git
```

#### 3.3.3 Compilation

To compile the program:

```
mkdir build
cd build
cmake .. && make
```

On Windows, you may have to specify the generator, for example:

```
cmake .. -G"Visual Studio 14"
```

Then open the newly created N2D2 project in Visual Studio 2015. Select "Release" for the build target. Right click on ALL\_BUILD item and select "Build".

#### 3.4 Downloading training datasets

A python script located in the repository root directory allows you to select and automatically download some well-known datasets, like MNIST and GTSRB (the script requires Python 2.x with bindings for GTK 2 package):

```
./tools/install_stimuli_gui.py
```

By default, the datasets are downloaded in the path specified in the N2D2\_DATA environment variable, which is the root path used by the N2D2 tool to locate the databases. If the N2D2\_DATA variable is not set, the default value used is /local/\$USER/n2d2\_data/ (or /local/n2d2\_data/ if the USER environment variable is not set) on Linux and C:\n2d2\_data\ on Windows.

Please make sure you have write access to the N2D2\_DATA path, or if not set, in the default /local/\$USER/n2d2\_data/ path.

# 3.5 Run the learning

The following command will run the learning for 600,000 image presentations/steps and log the performances of the network every 10,000 steps:

```
./n2d2 "mnist24_16c4s2_24c5s2_150_10.ini" -learn 600000 -log 10000
```

Note: you may want to check the gradient computation using the -check option. Note that it can be extremely long and can occasionally fail if the required precision is too high.

#### 3.6 Test a learned network

After the learning is completed, this command evaluate the network performances on the test data set:

```
./n2d2 "mnist24_16c4s2_24c5s2_150_10.ini" -test
```

#### 3.6.1 Interpreting the results

**Recognition rate** The recognition rate and the validation score are reported during the learning in the *TargetScore\_\*/Success\_validation.png* file, as shown in figure 4.

Confusion matrix The software automatically outputs the confusion matrix during learning, validation and test, with an example shown in figure 5. Each row of the matrix contains the number of occurrences estimated by the network for each label, for all the data corresponding to a single actual, target label. Or equivalently, each column of the matrix contains the number of actual, target label occurrences, corresponding to the same estimated label. Idealy, the matrix should be diagonal, with no occurrence of an estimated label for a different actual label (network mistake).

The confusion matrix reports can be found in the simulation directory:

- TargetScore\_\*/ConfusionMatrix\_learning.png;
- TargetScore\_\*/ConfusionMatrix\_validation.png;
- TargetScore\_\*/ConfusionMatrix\_test.png.

Memory and computation requirements The software also report the memory and computation requirements of the network, as shown in figure 6. The corresponding report can be found in the *stats* sub-directory of the simulation.

**Kernels and weights distribution** The synaptic weights obtained during and after the learning can be analyzed, in terms of distribution (*weights* sub-directory of the simulation) or in terms of kernels (*kernels* sub-directory of the simulation), as shown in 7.

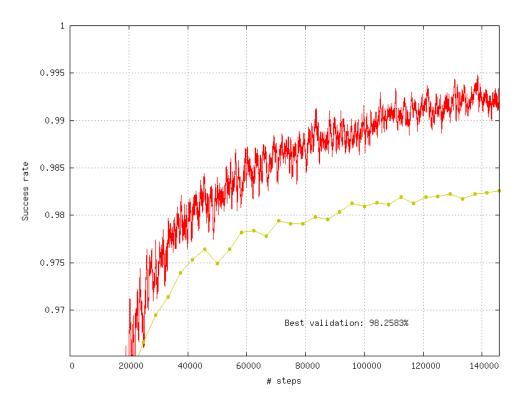


Figure 4: Recognition rate and validation score during learning.



Figure 5: Example of confusion matrix obtained after the learning.

**Output maps activity** The initial output maps activity for each layer can be visualized in the *outputs\_init* sub-directory of the simulation, as shown in figure 8.

# 3.7 Export a learned network

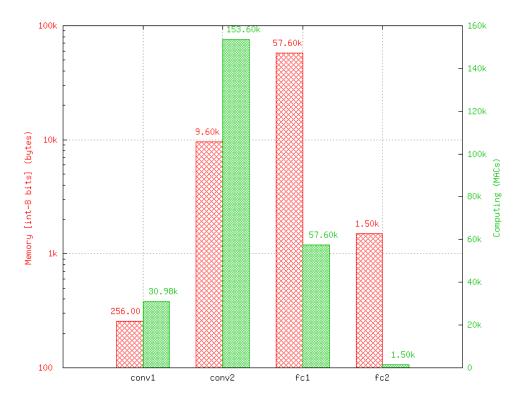


Figure 6: Example of memory and computation requirements of the network.

 $./n2d2 \ "mnist24\_16c4s2\_24c5s2\_150\_10.ini" \ -export \ CPP\_OpenCL$ 

# Export types:

- c C export using OpenMP;
- c\_hls C export tailored for HLS with Vivado HLS;
- CPP\_OpenCL C++ export using OpenCL;
- CPP\_Cuda C++ export using Cuda;
- CPP\_cuDNN C++ export using cuDNN;
- CPP\_TensorRT C++ export using tensorRT 2.1 API;
- SC\_Spike SystemC spike export.

Other program options related to the exports:

Option [default value]	Description
-nbbits [8]	Number of bits for the weights and signals. Must be 8, 16, 32
	or 64 for integer export, or -32, -64 for floating point export.
	The number of bits can be arbitrary for the c_hls export (for
	example, 6 bits). It must be -32 for the CPP_TensorRT export,
	the precision is directly set at runtime
-calib $[0]$	Number of stimuli used for the calibration. $0 = \text{no calibration}$
	(default), -1 = use the full test dataset for calibration
-calib-passes $\left[2\right]$	Number of KL passes for determining the layer output values
	distribution truncation threshold ( $0 = \text{use the max. value}$ ,
	no truncation)
-no-unsigned	If present, disable the use of unsigned data type in integer
	exports
-db-export [-1]	Max. number of stimuli to export $(0 = \text{no dataset export}, -1)$
	= unlimited)

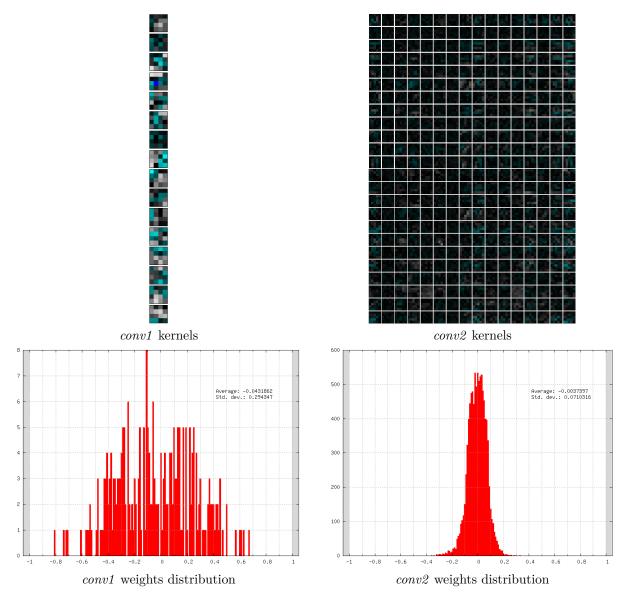


Figure 7: Example of kernels and weights distribution analysis for two convolutional layers.

# **3.7.1** C export

 $/T \setminus E /$ 

```
Test the exported network:

cd export_C_int8

make
./bin/n2d2_test

The result should look like:
...

1652.00/1762  (avg = 93.757094%)
1653.00/1763  (avg = 93.760635%)
1654.00/1764  (avg = 93.764172%)
Tested 1764 stimuli
Success rate = 93.764172%
Process time per stimulus = 187.548186 us (12 threads)

Confusion matrix:
```

1 /

0 /



Figure 8: Output maps activity example of the first convolutional layer of the network.

/	0 /	329	/	1 /	5	/	2	/
/	/	97.63%	/	0.30%	1.48%	/	0.59%	/
/	1 /	0	/	692	2	/	6	/
/	/	0.00%	/	98.86%	0.29%	/	0.86%	/
/	2 /	11	/	27 /	609	/	55	/
/	/	1.57%	/	3.85%	86.75%	/	7.83%	/
/	3 /	0	/	0 /	1	/	24	/
/	/	0.00%	/	0.00% /	4.00%	/	96.00%	/

T: Target E: Estimated

# 3.7.2 CPP\_OpenCL export

The OpenCL export can run the generated program in GPU or CPU architectures. Compilation features:

Preprocessor command [default value]	Description					
PROFILING [0]	Compile the binary with a synchronization be-					
	tween each layers and return the mean execution					
	time of each layer. This preprocessor option can					
	decrease performances.					
GENERATE_KBIN [0]	Generate the binary output of the OpenCL kernel					
	.cl file use. The binary is store in the /bin folder.					
LOAD_KBIN [0]	Indicate to the program to load an OpenCL ker-					
	nel as a binary from the /bin folder instead of a					
	.cl file.					
CUDA [0]	Use the CUDA OpenCL SDK locate at					
	/usr/local/cuda					
MALI [0]	Use the MALI OpenCL SDK locate at					
	$/usr/Mali_OpenCL_SDK_vXXX$					
INTEL [0]	Use the INTEL OpenCL SDK locate at					
	/opt/intel/opencl					
AMD [1]	Use the AMD OpenCL SDK locate at					
	/opt/AMDAPPSDK - XXX					

Program options related to the OpenCL export:

Option [default value]	Description
-cpu	If present, force to use a CPU architecture to run the program
-gpu	If present, force to use a GPU architecture to run the program
-batch [1]	Size of the batch to use
-stimulus [NULL]	Path to a specific input stimulus to test. For example: -
	stimulus $/stimulus/env0000.pgm$ command will test the file
	env0000.pgm of the stimulus folder.

Test the exported network:

cd export\_CPP\_OpenCL\_float32
make

# 3.7.3 CPP\_TensorRT export

The TensorRT API export can run the generated program in NVIDIA GPU architecture. It use CUDA, cuDNN and TensorRT API library. All the native TensorRT layers are supported. The export support from TensorRT 2.1 to TensorRT 5.0 versions.

Program options related to the TensorRT API export:

<sup>./</sup>bin/n2d2\_opencl\_test -gpu

Option [default value]	Description			
-batch [1]	Size of the batch to use			
-dev [0]	CUDA Device ID selection			
-stimulus [NULL]	Path to a specific input stimulus to test. For example: -			
	stimulus $/stimulus/env0000.pgm$ command will test the file			
	env0000.pgm of the stimulus folder.			
-prof	Activates the layer wise profiling mechanism. This option			
	can decrease execution time performance.			
-iter-build $\left[1 ight]$	Sets the number of minimization build iterations done by			
	the tensorRT builder to find the best layer tactics.			
-nbbits [-32]	Number of bits used for computation. Value -32 for Full			
	FP32 bits configuration, -16 for Half FP16 bits configuration			
	and 8 for INT8 bits configuration. When running INT8			
	mode for the first time, the TensorRT calibration process can			
	be very long. Once generated the generated calibration table			
	will be automatically reused. Supported compute mode in			
	function of the compute capability are provided here: https:			
	//docs.nvidia.com/cuda/cuda-c-programming-guide/			
	index.html#compute-capabilities.			

Test the exported network with layer wise profiling:

```
cd export_CPP_TensorRT_float32
make
./bin/n2d2_tensorRT_test -prof
```

The results of the layer wise profiling should look like:

```
(19\%)
 0.0219467 ms
  ******** POOL1: 0.00675573 ms
  (13\%)
(05\%)
  ******* POOL2: 0.00616047 ms
  (14\%)
  (19\%)
  ****** FC2: 0.0149013 ms
(13\%)
  (08\%)
Average profiled tensorRT process time per stimulus = 0.113932 ms
```

# 3.7.4 CPP\_cuDNN export

The cuDNN export can run the generated program in NVIDIA GPU architecture. It use CUDA and cuDNN library. Compilation features:

Preprocessor command [default value]	Description
PROFILING [0]	Compile the binary with a synchronization be-
	tween each layers and return the mean execution
	time of each layer. This preprocessor option can
	decrease performances.
ARCH32 [0]	Compile the binary with the 32-bits architecture
	compatibility.

Program options related to the cuDNN export:

Option [default value]	Description
-batch [1]	Size of the batch to use
-dev [0]	CUDA Device ID selection
-stimulus [NULL]	Path to a specific input stimulus to test. For example: -
	stimulus / stimulus / env0000.pgm command will test the file
	env0000.pgm of the stimulus folder.

Test the exported network:

cd export\_CPP\_cuDNN\_float32
make

./bin/n2d2\_cudnn\_test

# 3.7.5 C\_HLS export

Test the exported network:

cd export\_C\_HLS\_int8
make

./bin/n2d2\_test

Run the High-Level Synthesis (HLS) with Xilinx® Vivado® HLS:

vivado\_hls -f run\_hls.tcl

# 3.7.6 Layer compatibility table

Layer compatibility table in function of the export type:

Layer compatibility table		Export Type				
Layer compatibility table	С	HLS	OpenCL	TensorRT		
Convolution	Yes	Yes	Yes	Yes		
Pooling	Yes	Yes	Yes	Yes		
FullyConnected	Yes	Yes	Yes	Yes		
Softmax	Yes	No	Yes	Yes		
FractionalMaxPooling	Yes	No	Yes	No		
Deconvolution	No	No	No	Yes		
ElemWise	No	No	No	Yes		
Resize	Yes	No	No	Yes		
Padding	No	No	No	Yes		
LRN	No	No	No	Yes		
Anchor	No	No	No	Yes		
ObjectDetector	No	No	No	Yes		
ROIPooling	No	No	No	Yes		
RegionProposal	No	No	No	Yes		

Batchnormalization is not mentionned because batchnorm parameters are automatically fused with convolutions parameters with the command "-fuse".

# 4 INI file interface

The INI file interface is the primary way of using N2D2. It is a simple, lightweight and user-friendly format for specifying a complete DNN-based application, including dataset instanciation, data pre-processing, neural network layers instanciation and post-processing, with all its hyperparameters.

#### 4.1 Syntax

INI files are simple text files with a basic structure composed of sections, properties and values.

## 4.1.1 Properties

The basic element contained in an INI file is the property. Every property has a name and a value, delimited by an equals sign (=). The name appears to the left of the equals sign.

name=value

#### 4.1.2 Sections

Properties may be grouped into arbitrarily named sections. The section name appears on a line by itself, in square brackets ([ and ]). All properties after the section declaration are associated with that section. There is no explicit "end of section" delimiter; sections end at the next section declaration, or the end of the file. Sections may not be nested.

```
[section]
a=a
b=b
```

## 4.1.3 Case sensitivity

Section and property names are case sensitive.

#### 4.1.4 Comments

Semicolons (;) or number sign (#) at the beginning or in the middle of the line indicate a comment. Comments are ignored.

```
; comment text
a=a # comment text
a="a ; not a comment" ; comment text
```

#### 4.1.5 Quoted values

Values can be quoted, using double quotes. This allows for explicit declaration of whitespace, and/or for quoting of special characters (equals, semicolon, etc.).

## 4.1.6 Whitespace

Leading and trailing whitespace on a line are ignored.

#### 4.1.7 Escape characters

A backslash (\) followed immediately by EOL (end-of-line) causes the line break to be ignored.

# 4.2 Template inclusion syntax

Is is possible to recursively include templated INI files. For example, the main INI file can include a templated file like the following:

```
[inception@inception_model.ini.tpl]
INPUT=layer_x
SIZE=32
ARRAY=2 ; Must be the number of elements in the array
ARRAY[0].P1=Conv
ARRAY[0].P2=32
ARRAY[1].P1=Pool
ARRAY[1].P2=64
```

If the inception\_model.ini.tpl template file content is:

```
[{{SECTION_NAME}}_layer1]
Input={{INPUT}}
Type=Conv
NbOutputs={{SIZE}}

[{{SECTION_NAME}}_layer2]
Input={{SECTION_NAME}}_layer1
Type=Fc
NbOutputs={{SIZE}}

{% block ARRAY %}
[{{SECTION_NAME}}_array{{#}}]
Prop1=Config{{.P1}}
Prop2={{.P2}}
{% endblock %}
```

The resulting equivalent content for the main INI file will be:

```
[inception_layer1]
Input=layer_x
Type=Conv
NbOutputs=32

[inception_layer2]
Input=inception_layer1
Type=Fc
NbOutputs=32

[inception_array0]
Prop1=ConfigConv
Prop2=32

[inception_array1]
Prop1=ConfigPool
Prop2=64
```

The SECTION\_NAME template parameter is automatically generated from the name of the including section (before @).

#### 4.2.1 Variable substitution

{{VAR}} is replaced by the value of the VAR template parameter.

#### 4.2.2 Control statements

Control statements are between {% and %} delimiters.

block {%block ARRAY %} ... {%endblock %}

The # template parameter is automatically generated from the {%block ... %} template control statement and corresponds to the current item position, starting from 0.

for  ${\mbox{\for VAR in range([START, ]END])}}\$  ... {\mbox{\endfor }\mathcal{K}}

If START is not specified, the loop begins at 0 (first value of VAR). The last value of VAR is END-1.

if {%if VAR OP [VALUE] %} ... [{%else %}] ... {%endif %}
 OP may be ==, !=, exists or not\_exists.

include {%include FILENAME %}

# 4.3 Global parameters

Option [default value]	Description
DefaultModel [Transcode]	Default layers model. Can be Frame, Frame_CUDA, Transcode or
	Spike
DefaultDataType [Float32]	Default layers data type. Can be Float16, Float32 or Float64
SignalsDiscretization $\left[0 ight]$	Number of levels for signal discretization
FreeParametersDiscretization	Number of levels for weights discretization
[0]	

#### 4.4 Databases

The tool integrates pre-defined modules for several well-known database used in the deep learning community, such as MNIST, GTSRB, CIFAR10 and so on. That way, no extra step is necessary to be able to directly build a network and learn it on these database.

#### 4.4.1 MNIST

MNIST (LeCun et al., 1998) is already fractionned into a learning set and a testing set, with:

- 60,000 digits in the learning set;
- 10,000 digits in the testing set.

Example:

#### [database]

Type=MNIST\_IDX\_Database

Validation=0.2; Fraction of learning stimuli used for the validation [default: 0.0]

Option [default value]	Description
Validation $\left[0.0\right]$	Fraction of the learning set used for validation
DataPath	Path to the database
[\$N2D2_DATA/mnist]	

# 4.4.2 GTSRB

GTSRB (Stallkamp et al., 2012) is already fractionned into a learning set and a testing set, with:

- 39,209 digits in the learning set;
- 12,630 digits in the testing set.

Example:

#### [database]

Type=GTSRB\_DIR\_Database

Validation=0.2; Fraction of learning stimuli used for the validation [default: 0.0]

Option [default value]	Description
Validation $\left[0.0\right]$	Fraction of the learning set used for validation
DataPath	Path to the database
$[$ ext{N2D2\_DATA}/GTSRB]$	

#### 4.4.3 Directory

Hand made database stored in files directories are directly supported with the DIR\_Database module. For example, suppose your database is organized as following (in the path specified in the N2D2\_DATA environment variable):

GST/airplanes: 800 images
GST/car\_side: 123 images
GST/Faces: 435 images
GST/Motorbikes: 798 images

You can then instanciate this database as input of your neural network using the following parameters:

```
[database]
Type=DIR_Database
DataPath=${N2D2_DATA}/GST
Learn=0.4 ; 40% of images of the smallest category = 49 (0.4x123) images for each category will be used for learning
Validation=0.2 ; 20% of images of the smallest category = 25 (0.2x123) images for each category will be used for validation
; the remaining images will be used for testing
```

Each subdirectory will be treated as a different label, so there will be 4 different labels, named after the directory name.

The stimuli are equi-partitioned for the learning set and the validation set, meaning that the same number of stimuli for each category is used. If the learn fraction is 0.4 and the validation fraction is 0.2, as in the example above, the partitioning will be the following:

Label ID	Label name	Learn set	Validation set	Test set
0	airplanes	49	25	726
1	car_side	49	25	49
2	Faces	49	25	361
3	Motorbikes	49	25	724
	Total:	196	100	1860

# Mandatory option

Option [default value]	Description
DataPath	Path to the root stimuli directory
Learn	If PerLabelPartitioning is true, fraction of images used for
	the learning; else, number of images used for the learning,
	regardless of their labels
LoadInMemory $[0]$	Load the whole database into memory
Depth [1]	Number of sub-directory levels to include. Examples:

	Depth = 0: load stimuli only from the current directory
	(DataPath)  Depth = 1: load stimuli from DataPath and stimuli contained
	in the sub-directories of DataPath
	Depth < 0: load stimuli recursively from DataPath and all its
	sub-directories
LabelName []	Base stimuli label name
LabelDepth [1]	Number of sub-directory name levels used to form the stimuli
	labels. Examples:
	LabelDepth = -1: no label for all stimuli (label ID = -1)
	LabelDepth = 0: uses LabelName for all stimuli
	LabelDepth = 1: uses LabelName for stimuli in the current
	directory (DataPath) and LabelName/ $sub$ -directory name for
	stimuli in the sub-directories
PerLabelPartitioning $[1]$	If true, the stimuli are equi-partitioned for the learn/valida-
	tion/test sets, meaning that the same number of stimuli for
	each label is used
Validation $[0.0]$	If PerLabelPartitioning is true, fraction of images used for the
	validation; else, number of images used for the validation,
	regardless of their labels
Test $[1.0$ -Learn-Validation]	If PerLabelPartitioning is true, fraction of images used for the
	test; else, number of images used for the test, regardless of
	their labels
ValidExtensions []	List of space-separated valid stimulus file extensions (if left
	empty, any file extension is considered a valid stimulus)
LoadMore []	Name of an other section with the same options to load a
	different DataPath
ROIFile []	File containing the stimuli ROIs. If a ROI file is specified,
	LabelDepth should be set to -1
DefaultLabel []	Label name for pixels outside any ROI (default is no label,
[ ]	pixels are ignored)
ROIsMargin $[0]$	Number of pixels around ROIs that are ignored (and not
	considered as DefaultLabel pixels)

To load and partition more than one DataPath, one can use the LoadMore option:

```
[database]
Type=DIR_Database
DataPath=${N2D2_DATA}/GST
Learn=0.6
Validation=0.4
LoadMore=database.test

; Load stimuli from the "GST_Test" path in the test dataset
[database.test]
DataPath=${N2D2_DATA}/GST_Test
Learn=0.0
Test=1.0
; The LoadMore option is recursive:
; LoadMore=database.more

; [database.more]
; Load even more data here
```

**Speech Commands Dataset** Use with Speech Commands Data Set, released by the Google (Warden, 2018).

#### [database]

Type=DIR\_Database
DataPath=\${N2D2\_DATA}/speech\_commands\_v0.02
ValidExtensions=wav
IgnoreMasks=\*/\_background\_noise\_
Learn=0.6
Validation=0.2

# 4.4.4 Other built-in databases

Actitracker\_Database Actitracker database, released by the WISDM Lab (W. Lockhart et al., 2011).

Option [default value]	Description
Learn [0.6]	Fraction of data used for the learning
Validation $[0.2]$	Fraction of data used for the validation
UseUnlabeledForTest $\left[0 ight]$	If true, use the unlabeled dataset for the test
DataPath	Path to the database
[\$N2D2_DATA	
/WISDM_at_v2.0]	

# CIFAR10\_Database CIFAR10 database (Krizhevsky, 2009).

Option [default value]	Description
Validation $[0.0]$	Fraction of the learning set used for validation
DataPath	Path to the database
[\$N2D2_DATA/cifar-10-batches-	
bin	

# CIFAR100\_Database CIFAR100 database (Krizhevsky, 2009).

Option [default value]	Description
Validation $\left[0.0\right]$	Fraction of the learning set used for validation
UseCoarse $[0]$	If true, use the coarse labeling (10 labels instead of 100)
DataPath	Path to the database
$[\$N2D2\_DATA/cifar-100-binary]$	

CKP\_Database The Extended Cohn-Kanade (CK+) database for expression recognition (Lucey et al., 2010).

Option [default value]	Description
Learn	Fraction of images used for the learning
Validation $\left[0.0 ight]$	Fraction of images used for the validation
DataPath	Path to the database
[\$N2D2_DATA/cohn-kanade-	
images	

#### Caltech101\_DIR\_Database Caltech 101 database (Fei-Fei et al., 2004).

Option [default value]	Description
Learn	Fraction of images used for the learning
Validation $[0.0]$	Fraction of images used for the validation
IncClutter $[0]$	If true, includes the BACKGROUND_Google directory of
	the database
DataPath	Path to the database
[\$N2D2_DATA/	
101_ObjectCategories]	

### Caltech256\_DIR\_Database Caltech 256 database (Griffin et al., 2007).

Option [default value]	Description
Learn	Fraction of images used for the learning
Validation $\left[0.0 ight]$	Fraction of images used for the validation
IncClutter [0]	If true, includes the BACKGROUND_Google directory of
	the database
DataPath	Path to the database
[\$N2D2_DATA/	
256_ObjectCategories]	

#### CaltechPedestrian\_Database Caltech Pedestrian database (Dollár et al., 2009).

Note that the images and annotations must first be extracted from the seq video data located in the videos directory using the <code>dbExtract.m</code> Matlab tool provided in the "Matlab evaluation/labeling code" downloadable on the dataset website.

Assuming the following directory structure (in the path specified in the N2D2\_DATA environment variable):

- CaltechPedestrians/data-USA/videos/... (from the setxx.tar files)
- CaltechPedestrians/data-USA/annotations/... (from the setxx.tar files)
- CaltechPedestrians/tools/piotr\_toolbox/toolbox (from the Piotr's Matlab Toolbox archive)
- CaltechPedestrians/\*.m including dbExtract.m (from the Matlab evaluation/labeling code)

Use the following command in Matlab to generate the images and annotations:

```
cd([getenv('N2D2_DATA') '/CaltechPedestrians'])
addpath(genpath('tools/piotr_toolbox/toolbox')) % add the Piotr's Matlab Toolbox in the Matlab
    path
dbInfo('USA')
dbExtract()
```

Option [default value]	Description
Validation $[0.0]$	Fraction of the learning set used for validation
[1]	Use the same label for "person" and "people" bounding box
IncAmbiguous $\left[0\right]$	Include ambiguous bounding box labeled "person?" using the
	same label as "person"
DataPath	Path to the database images
[\$N2D2_DATA/	
CaltechPedestrians/data-	
USA/images]	
LabelPath	Path to the database annotations
[\$N2D2_DATA/	
CaltechPedestrians/data-	
USA/annotations]	

# Cityscapes\_Database Cityscapes database (Cordts et al., 2016).

Option [default value]	Description
IncTrainExtra $\left[0 ight]$	If true, includes the left 8-bit images - trainextra set (19,998
	images)
UseCoarse $[0]$	If true, only use coarse annotations (which are the only
	annotations available for the trainextra set)
$oxed{ t SingleInstanceLabels} egin{array}{c} 1 \end{bmatrix}$	If true, convert group labels to single instance labels (for
	example, cargroup becomes car)
DataPath	Path to the database images
[\$N2D2_DATA/	
Cityscapes/leftImg8bit] or	
[\$CITYSCAPES_DATASET] if defined	
LabelPath []	Path to the database annotations (deduced from DataPath if
	left empty)

# Daimler\_Database Daimler Monocular Pedestrian Detection Benchmark (Daimler Pedestrian).

Option [default value]	Description
Learn [1.0]	Fraction of images used for the learning
Validation $\left[0.0 ight]$	Fraction of images used for the validation
Test $[0.0]$	Fraction of images used for the test
Fully [0]	When activate it use the test dataset to learn. Use only on
	fully-cnn mode

# DOTA\_Database DOTA database (Xia et al., 2017).

Option [default value]	Description
Learn	Fraction of images used for the learning
DataPath	Path to the database
$[$N2D2\_DATA/DOTA]$	
LabelPath	Path to the database labels list file

# FDDB\_Database Face Detection Data Set and Benchmark (FDDB) (Jain and Learned-Miller, 2010).

Option [default value]	Description
Learn	Fraction of images used for the learning
Validation $[0.0]$	Fraction of images used for the validation
DataPath	Path to the images (decompressed originalPics.tar.gz)
$[\$ ext{N2D2\_DATA}/ ext{FDDB}]$	
LabelPath	Path to the annotations (decompressed FDDB-folds.tgz)
$[\$ ext{N2D2\_DATA}/ ext{FDDB}]$	

# GTSDB\_DIR\_Database GTSDB database (Houben et al., 2013).

Option [default value]	Description
Learn	Fraction of images used for the learning
Validation $\left[0.0 ight]$	Fraction of images used for the validation
DataPath	Path to the database
[\$N2D2_DATA/FullIJCNN2013]	

#### ILSVRC2012\_Database ILSVRC2012 database (Russakovsky et al., 2015).

Option [default value]	Description
Learn	Fraction of images used for the learning
DataPath	Path to the database
[\$N2D2_DATA/ILSVRC2012]	
LabelPath	Path to the database labels list file
[\$N2D2_DATA	
/ILSVRC2012/synsets.txt]	

KITTI\_Database The KITTI Database provide ROI which can be use for autonomous driving and environment perception. The database provide 8 labeled different classes. Utilization of the KITTI Database is under licensing conditions and request an email registration. To install it you have to follow this link: <a href="http://www.cvlibs.net/datasets/kitti/eval\_tracking.php">http://www.cvlibs.net/datasets/kitti/eval\_tracking.php</a> and download the left color images (15 GB) and the trainling labels of tracking data set (9 MB). Extract the downloaded archives in your \$N2D2\_DATA/KITTI folder.

Option [default value]	Description
Learn $[0.8]$	Fraction of images used for the learning
Validation $[0.2]$	Fraction of images used for the validation

KITTI\_Road\_Database The KITTI Road Database provide ROI which can be used to road segmentation. The dataset provide 1 labeled class (road) on 289 training images. The 290 test images are not labeled. Utilization of the KITTI Road Database is under licensing conditions and request an email registration. To install it you have to follow this link: http://www.cvlibs.net/datasets/kitti/eval\_road.php and download the "base kit" of (0.5 GB) with left color images, calibration and training labels. Extract the downloaded archive in your \$N2D2\_DATA/KITTI folder.

Option [default value]	Description
Learn [0.8]	Fraction of images used for the learning
Validation $[0.2]$	Fraction of images used for the validation

KITTI\_Object\_Database The KITTI Object Database provide ROI which can be use for autonomous driving and environment perception. The database provide 8 labeled different classes on 7481 training images. The 7518 test images are not labeled. The whole database provide 80256 labeled objects. Utilization of the KITTI Object Database is under licensing conditions and request an email registration. To install it you have to follow this link: http://www.cvlibs.net/datasets/kitti/eval\_object.php?obj\_benchmark and download the "lef color images" (12 GB) and the training labels of object data set (5 MB). Extract the downloaded archives in your \$N2D2\_DATA/KITTI\_Object folder.

Option [default value]	Description
Learn [0.8]	Fraction of images used for the learning
Validation $[0.2]$	Fraction of images used for the validation

# LITISRouen\_Database LITIS Rouen audio scene dataset (Rakotomamonjy and Gasso, 2014).

Option [default value]	Description
Learn $[0.4]$	Fraction of images used for the learning
Validation $\left[0.4\right]$	Fraction of images used for the validation
DataPath	Path to the database
$[\$N2D2\_DATA/data\_rouen]$	

#### 4.4.5 Dataset images slicing

It is possible to automatically slice images from a dataset, with a given slice size and stride, using the .slicing attribute. This effectively increases the number of stimuli in the set.

```
[database.slicing]
ApplyTo=NoLearn
Width=2048
Height=1024
StrideX=2048
StrideY=1024
RandomShuffle=1 ; 1 is the default value
```

The RandomShuffle option, enabled by default, randomly shuffle the dataset after slicing. If disabled, the slices are added in order at the end of the dataset.

# 4.5 Stimuli data analysis

You can enable stimuli data reporting with the following section (the name of the section must start with env.StimuliData):

```
[env.StimuliData-raw]
ApplyTo=LearnOnly
LogSizeRange=1
LogValueRange=1
```

The stimuli data reported for the full MNIST learning set will look like:

```
env. StimuliData-raw\ data:
Number\ of\ stimuli:\ 60000
Data\ width\ range:\ [28\ ,\ 28]
Data\ height\ range:\ [28\ ,\ 28]
Data\ channels\ range:\ [1\ ,\ 1]
Value\ range:\ [0\ ,\ 255]
Value\ mean:\ 33.3184
Value\ std.\ dev.:\ 78.5675
```

## 4.5.1 Zero-mean and unity standard deviation normalization

It it possible to normalize the whole database to have zero mean and unity standard deviation on the learning set using a RangeAffineTransformation transformation:

```
; Stimuli normalization based on learning set global mean and std.dev.

[env.Transformation-normalize]

Type=RangeAffineTransformation

FirstOperator=Minus

FirstValue=[env.StimuliData-raw]_GlobalValue.mean

SecondOperator=Divides

SecondValue=[env.StimuliData-raw]_GlobalValue.stdDev
```

The variables \_GlobalValue.mean and \_GlobalValue.stdDev are automatically generated in the [env. StimuliData-raw] block. Thanks to this facility, unknown and arbitrary database can be analysed and normalized in one single step without requiring any external data manipulation.

After normalization, the stimuli data reported is:

```
env. StimuliData-normalized data:

Number of stimuli: 60000

Data width\ range: [28,\ 28]

Data height range: [28,\ 28]

Data channels range: [1,\ 1]

Value range: [-0.424074,\ 2.82154]

Value mean: 2.64796e-07

Value std.\ dev.:\ 1
```

Where we can check that the global mean is close to 0 and the standard deviation is 1 on the whole dataset. The result of the transformation on the first images of the set can be checked in the generated *frames* folder, as shown in figure 9.

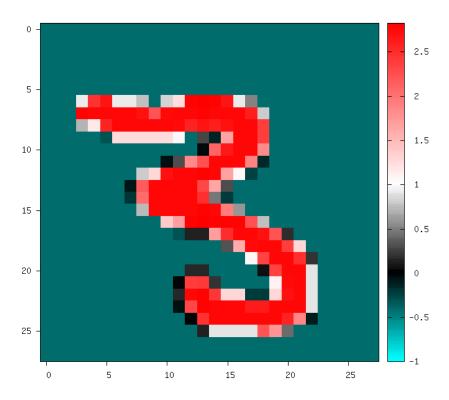


Figure 9: Image of the set after normalization.

# 4.5.2 Substracting the mean image of the set

Using the StimuliData object followed with an AffineTransformation, it is also possible to use the mean image of the dataset to normalize the data:

```
[env.StimuliData-meanData]
ApplyTo=LearnOnly
MeanData=1 ; Provides the _MeanData parameter used in the transformation

[env.Transformation]
Type=AffineTransformation
FirstOperator=Minus
FirstValue=[env.StimuliData-meanData]_MeanData
```

The resulting global mean image can be visualized in env.StimuliData-meanData/meanData.bin.png an is shown in figure 10.

After this transformation, the reported stimuli data becomes:

```
env. StimuliData-processed \ data: \\ Number \ of \ stimuli: 60000 \\ Data \ width \ range: [28, 28] \\ Data \ height \ range: [28, 28] \\ Data \ channels \ range: [1, 1] \\ Value \ range: [-139.554, 254.979] \\ Value \ mean: -3.45583e-08 \\ Value \ std. \ dev: 66.1288 \\ \\
```

The result of the transformation on the first images of the set can be checked in the generated frames folder, as shown in figure 11.

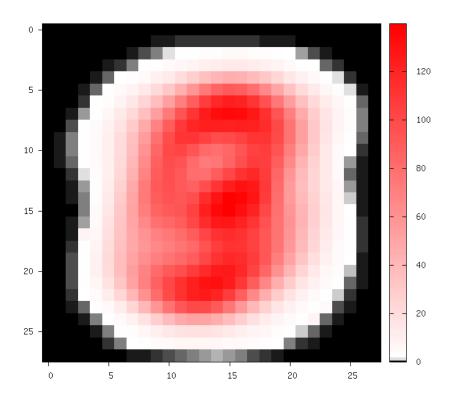


Figure 10: Global mean image generated by StimuliData with the MeanData parameter enabled.

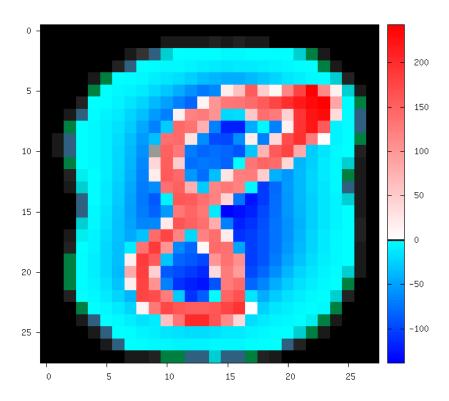


Figure 11: Image of the set after the AffineTransformation substracting the global mean image (keep in mind that the original image value range is [0, 255]).

# 4.6 Environment

The environment simply specify the input data format of the network (width, height and batch size). Example:

# [env] SizeX=24 SizeY=24 BatchSize=12 ; [default: 1]

Option [default value]	Description
SizeX	Environment width
SizeY	Environment height
NbChannels $[1]$	Number of channels (applicable only if there is no env.
	ChannelTransformation[])
BatchSize [1]	Batch size
CompositeStimuli $\left[0\right]$	If true, use pixel-wise stimuli labels
CachePath []	Stimuli cache path (no cache if left empty)
StimulusType [SingleBurst]	Method for converting stimuli into spike trains. Can be any
	of SingleBurst, Periodic, JitteredPeriodic Or Poissonian
DiscardedLateStimuli $\left[1.0\right]$	The pixels in the pre-processed stimuli with a value above
	this limit never generate spiking events
PeriodMeanMin $[50\ \mathtt{TimeMs}]$	Mean minimum period $\overline{T_{min}}$ , used for periodic temporal cod-
	ings, corresponding to pixels in the pre-processed stimuli with
	a value of 0 (which are supposed to be the most significant
	pixels)
PeriodMeanMax $[12 \; \mathtt{TimeS}]$	Mean maximum period $\overline{T_{max}}$ , used for periodic temporal
	codings, corresponding to pixels in the pre-processed stimuli
	with a value of 1 (which are supposed to be the least signifi-
	cant pixels). This maximum period may be never reached if
	DiscardedLateStimuli is lower than 1.0
PeriodRelStdDev $\left[0.1\right]$	Relative standard deviation, used for periodic temporal cod-
	ings, applied to the spiking period of a pixel
PeriodMin $[11 \text{ TimeMs}]$	Absolute minimum period, or spiking interval, used for peri-
	odic temporal codings, for any pixel

#### 4.6.1 Built-in transformations

There are 6 possible categories of transformations:

- env.Transformation[...] Transformations applied to the input images before channels creation;
- env.OnTheFlyTransformation[...] On-the-fly transformations applied to the input images before channels creation;
- env.ChannelTransformation[...] Create or add transformation for a specific channel;
- env.ChannelOnTheFlyTransformation[...] Create or add on-the-fly transformation for a specific channel;
- env.ChannelsTransformation[...] Transformations applied to all the channels of the input images:
- env.ChannelsOnTheFlyTransformation[...] On-the-fly transformations applied to all the channels of the input images.

# Example:

# [env.Transformation] Type=PadCropTransformation Width=24 Height=24

Several transformations can applied successively. In this case, to be able to apply multiple transformations of the same category, a different suffix ([...]) must be added to each transformation.

The transformations will be processed in the order of appearance in the INI file regardless of their suffix.

Common set of parameters for any kind of transformation:

Option [default value]	Description
ApplyTo [All]	Apply the transformation only to the specified stimuli sets.
	Can be:
	LearnOnly: learning set only
	ValidationOnly: validation set only
	TestOnly: testing set only
	NoLearn: validation and testing sets only
	NoValidation: learning and testing sets only
	NoTest: learning and validation sets only
	All: all sets (default)

# Example:

```
[env.Transformation-1]
Type=ChannelExtractionTransformation
CSChannel=Gray
[env.Transformation-2]
Type=RescaleTransformation
Width=29
Height=29
[env.Transformation-3]
Type=EqualizeTransformation
[env.OnTheFlyTransformation]
Type=DistortionTransformation
ApplyTo=LearnOnly ; Apply this transformation for the Learning set only
ElasticGaussianSize=21
ElasticSigma=6.0
ElasticScaling=20.0
Scaling=15.0
Rotation=15.0
```

List of available transformations:

**AffineTransformation** Apply an element-wise affine transformation to the image with matrixes of the same size.

Option [default value]	Description
FirstOperator	First element-wise operator, can be Plus, Minus, Multiplies,
	Divides
FirstValue	First matrix file name
SecondOperator [Plus]	Second element-wise operator, can be Plus, Minus, Multiplies,
	Divides
SecondValue []	Second matrix file name

The final operation is the following, with A the image matrix,  $B_{1st}$ ,  $B_{2nd}$  the matrixes to add/substract/multiply/divide and  $\odot$  the element-wise operator:

$$f(A) = \left(A \stackrel{\odot}{op_{1st}} B_{1st}\right) \stackrel{\odot}{op_{2nd}} B_{2nd}$$

# ApodizationTransformation Apply an apodization window to each data row.

Option [default value]	Description
Size	Window total size (must match the number of data columns)
WindowName [Rectangular]	Window name. Possible values are:
	Rectangular: Rectangular
	Hann: Hann
	Hamming: Hamming
	Cosine: Cosine
	Gaussian: Gaussian
	Blackman: Blackman
	Kaiser: Kaiser

# Gaussian window Gaussian window.

Option [default value]	Description
$WindowName. {\tt Sigma} \ [0.4]$	Sigma

# Blackman window Blackman window.

Option [default value]	Description
	Alpha

# Kaiser window Kaiser window.

Option [default value]	Description
WindowName.Beta $[5.0]$	Beta

# ChannelExtractionTransformation Extract an image channel.

Option	Description
CSChannel	Blue: blue channel in the BGR colorspace, or first channel of
	any colorspace
	Green: green channel in the BGR colorspace, or second chan-
	nel of any colorspace
	Red: red channel in the BGR colorspace, or third channel of
	any colorspace
	Hue: hue channel in the HSV colorspace
	Saturation: saturation channel in the HSV colorspace
	Value: value channel in the HSV colorspace
	Gray: gray conversion
	Y: Y channel in the YCbCr colorspace
	Съ: Cb channel in the YCbCr colorspace
	Cr: Cr channel in the YCbCr colorspace

 ${\tt ColorSpaceTransformation} \quad {\tt Change \ the \ current \ image \ colorspace}.$ 

Option	Description
ColorSpace	BGR: convert any gray, BGR or BGRA image to BGR
	RGB: convert any gray, BGR or BGRA image to RGB
	HSV: convert BGR image to HSV
	HLS: convert BGR image to HLS
	YCrCb: convert BGR image to YCrCb
	CIELab: convert BGR image to CIELab
	CIELuv: convert BGR image to CIELuv
	RGB_to_BGR: convert RGB image to BGR
	RGB_to_HSV: convert RGB image to HSV
	RGB_to_HLS: convert RGB image to HLS
	RGB_to_YCrCb: convert RGB image to YCrCb
	RGB_to_CIELab: convert RGB image to CIELab
	RGB_to_CIELuv: convert RGB image to CIELuv
	HSV_to_BGR: convert HSV image to BGR
	HSV_to_RGB: convert HSV image to RGB
	HLS_to_BGR: convert HLS image to BGR
	HLS_to_RGB: convert HLS image to RGB
	YCrCb_to_BGR: convert YCrCb image to BGR
	YCrCb_to_RGB: convert YCrCb image to RGB
	CIELab_to_BGR: convert CIELab image to BGR
	CIELab_to_RGB: convert CIELab image to RGB
	CIELuv_to_BGR: convert CIELuv image to BGR
	CIELuv_to_RGB: convert CIELuv image to RGB

Note that the default colorspace in N2D2 is BGR, the same as in OpenCV.

**DFTTransformation** Apply a DFT to the data. The input data must be single channel, the resulting data is two channels, the first for the real part and the second for the imaginary part.

Option [default value]	Description
TwoDimensional $[1]$	If true, compute a 2D image DFT. Otherwise, compute the
	1D DFT of each data row

Note that this transformation can add zero-padding if required by the underlying FFT implementation.

N2D2 IP only DistortionTransformation Apply elastic distortion to the image. This transformation is generally used on-the-fly (so that a different distortion is performed for each image), and for the learning only.

Option [default value]	Description
ElasticGaussianSize $\left[15\right]$	Size of the gaussian for elastic distortion (in pixels)
ElasticSigma $\left[6.0\right]$	Sigma of the gaussian for elastic distortion
ElasticScaling $\left[0.0\right]$	Scaling of the gaussian for elastic distortion
Scaling $[0.0]$	Maximum random scaling amplitude $(+/-, in percentage)$
Rotation $[0.0]$	Maximum random rotation amplitude (+/-, in °)

N2D2 IP only EqualizeTransformation Image histogram equalization.

Option [default value]	Description
Method [Standard]	Standard: standard histogram equalization
	CLAHE: contrast limited adaptive histogram equalization
CLAHE_ClipLimit [40.0]	Threshold for contrast limiting (for CLAHE only)
CLAHE_GridSize [8]	Size of grid for histogram equalization (for Clahe only). Input
	image will be divided into equally sized rectangular tiles. This
	parameter defines the number of tiles in row and column.

### N2D2 IP only ExpandLabelTransformation Expand single image label (1x1 pixel) to full frame label.

# FilterTransformation Apply a convolution filter to the image.

Option [default value]	Description
Kernel	Convolution kernel. Possible values are:
	*: custom kernel
	Gaussian: Gaussian kernel
	Log: Laplacian Of Gaussian kernel
	Dog: Difference Of Gaussian kernel
	Gabor: Gabor kernel

### \* kernel Custom kernel.

Option	Description
Kernel.SizeX $\left[0 ight]$	Width of the kernel (numer of columns)
Kernel.SizeY $\left[0 ight]$	Height of the kernel (number of rows)
Kernel.Mat	List of row-major ordered coefficients of the kernel

If both Kernel.SizeX and Kernel.SizeY are 0, the kernel is assumed to be square.

### Gaussian kernel Gaussian kernel.

Option [default value]	Description
Kernel.SizeX	Width of the kernel (numer of columns)
Kernel.SizeY	Height of the kernel (number of rows)
Kernel.Positive $\left[1 ight]$	If true, the center of the kernel is positive
Kernel.Sigma $[\sqrt{2.0}]$	Sigma of the kernel

# LoG kernel Laplacian Of Gaussian kernel.

Option [default value]	Description
Kernel.SizeX	Width of the kernel (numer of columns)
Kernel.SizeY	Height of the kernel (number of rows)
Kernel.Positive $\left[1 ight]$	If true, the center of the kernel is positive
Kernel.Sigma $[\sqrt{2.0}]$	Sigma of the kernel

# DoG kernel Difference Of Gaussian kernel kernel.

Option [default value]	Description
Kernel.SizeX	Width of the kernel (numer of columns)
Kernel.SizeY	Height of the kernel (number of rows)
Kernel.Positive $\left[1 ight]$	If true, the center of the kernel is positive
Kernel.Sigma1 $[2.0]$	Sigma1 of the kernel
Kernel.Sigma2 $[1.0]$	Sigma2 of the kernel

### Gabor kernel Gabor kernel.

Option [default value]	Description
Kernel.SizeX	Width of the kernel (numer of columns)
Kernel.SizeY	Height of the kernel (number of rows)
Kernel.Theta	Theta of the kernel
Kernel.Sigma $[\sqrt{2.0}]$	Sigma of the kernel
Kernel.Lambda $\left[10.0\right]$	Lambda of the kernel
Kernel.Psi $[\pi/2.0]$	Psi of the kernel
Kernel.Gamma $[0.5]$	Gamma of the kernel

# FlipTransformation Image flip transformation.

Option [default value]	Description
HorizontalFlip $[0]$	If true, flip the image horizontally
$\texttt{VerticalFlip} \ [0]$	If true, flip the image vertically
RandomHorizontalFlip $\left[0 ight]$	If true, randomly flip the image horizontally
RandomVerticalFlip $\left[0 ight]$	If true, randomly flip the image vertically

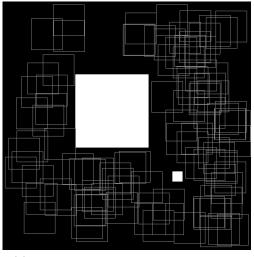
# ${\tt N2D2\ IP}\ \mathit{only}\ \ \textbf{GradientFilterTransformation}\ \ \mathrm{Compute}\ \mathrm{image}\ \mathrm{gradient}.$

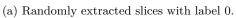
Option [default value]	Description
Scale $[1.0]$	Scale to apply to the computed gradient
Delta $[0.0]$	Bias to add to the computed gradient
GradientFilter [Sobel]	Filter type to use for computing the gradient. Possible
	options are: Sobel, Scharr and Laplacian
KernelSize [3]	Size of the filter kernel (has no effect when using the Scharr
	filter, which kernel size is always 3x3)
ApplyToLabels $\left[0 ight]$	If true, use the computed gradient to filter the image label and
	ignore pixel areas where the gradient is below the Threshold.
	In this case, only the labels are modified, not the image
InvThreshold $\left[0 ight]$	If true, ignored label pixels will be the ones with a low
	gradient (low contrasted areas)
Threshold $\left[0.5\right]$	Threshold applied on the image gradient
Label []	List of labels to filter (space-separated)
[1.0]	Rescale the image by this factor before applying the gradient
	and the threshold, then scale it back to filter the labels

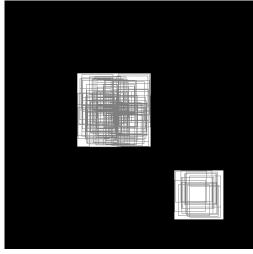
N2D2 IP only LabelSliceExtractionTransformation Extract a slice from an image belonging to a given label.

Option [default value]	Description
Width	Width of the slice to extract
Height	Height of the slice to extract
Label [-1]	Slice should belong to this label ID. If -1, the label ID is
	random
$\texttt{RandomRotation} \; [0]$	If true, extract randomly rotated slices
RandomRotationRange $[0.0\ 360.0]$	Range of the random rotations, in degrees, counterclockwise
	(if RandomRotation is enabled)
SlicesMargin $[0]$	Positive or negative, indicates the margin around objects
	that can be extracted in the slice
$\texttt{KeepComposite} \; [0]$	If false, the 2D label image is reduced to a single value
	corresponding to the extracted object label (useful for patches
	classification tasks). Note that if SlicesMargin is $> 0$ , the 2D
	label image may contain other labels before reduction. For
	pixel-wise segmentation tasks, set KeepComposite to true.

This transformation is useful to learn sparse object occurrences in a lot of background. If the dataset is very unbalanced towards background, this transformation will ensure that the learning is done on a more balanced set of every labels, regardless of their actual pixel-wise ratio.







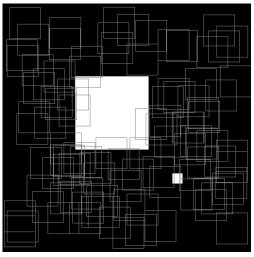
(b) Randomly extracted slices with label 1.

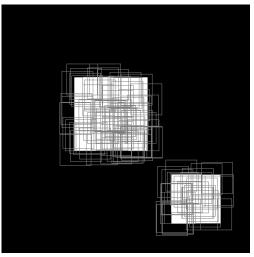
Figure 12: Illustration of the working behavior of LabelSliceExtractionTransformation with SlicesMargin = 0.

When SlicesMargin is 0, only slices that fully include a given label are extracted, as shown in figure 12. The behavior with SlicesMargin < 0 is illustrated in figure 13. Note that setting a negative SlicesMargin larger in absolute value than Width/2 or Height/2 will lead in some (random) cases in incorrect slice labels in respect to the majority pixel label in the slice.

MagnitudePhaseTransformation Compute the magnitude and phase of a complex two channels input data, with the first channel x being the real part and the second channel y the imaginary part. The resulting data is two channels, the first one with the magnitude and the second one with the phase.

Option [default value]	Description
LogScale $[0]$	If true, compute the magnitude in log scale





- (a) Randomly extracted slices including label 0.
- (b) Randomly extracted slices including label 1.

Figure 13: Illustration of the working behavior of LabelSliceExtractionTransformation with SlicesMargin = -32.

The magnitude is:

$$M_{i,j} = \sqrt{x_{i,j}^2 + x_{i,j}^2}$$

If LogScale = 1, compute  $M'_{i,j} = log(1 + M_{i,j})$ . The phase is:

$$\theta_{i,j} = atan2(y_{i,j}, x_{i,j})$$

N2D2 IP only MorphologicalReconstructionTransformation Apply a morphological reconstruction transformation to the image. This transformation is also useful for post-processing.

Option [default value]	Description
Operation	Morphological operation to apply. Can be:
	ReconstructionByErosion: reconstruction by erosion operation
	ReconstructionByDilation: reconstruction by dilation opera-
	tion
	OpeningByReconstruction: opening by reconstruction operation
	ClosingByReconstruction: closing by reconstruction operation
Size	Size of the structuring element
${\tt ApplyToLabels} \; [0]$	If true, apply the transformation to the labels instead of the
	image
Shape [Rectangular]	Shape of the structuring element used for morphology opera-
	tions. Can be Rectangular, Elliptic or Cross.
NbIterations $[1]$	Number of times erosion and dilation are applied for opening
	and closing reconstructions

N2D2 IP only MorphologyTransformation Apply a morphology transformation to the image. This transformation is also useful for post-processing.

Option [default value]	Description
Operation	Morphological operation to apply. Can be:
	Erode: erode operation $(=erode(src))$
	Dilate: dilate operation $(= dilate(src))$
	Opening: opening operation $(open(src) = dilate(erode(src)))$
	Closing: closing operation $(close(src) = erode(dilate(src)))$
	Gradient: morphological gradient (= $dilate(src) - erode(src)$ )
	TopHat: top hat $(= src - open(src))$
	BlackHat: black hat $(=close(src) - src)$
Size	Size of the structuring element
${\tt ApplyToLabels} \; [0]$	If true, apply the transformation to the labels instead of the
	image
Shape [Rectangular]	Shape of the structuring element used for morphology opera-
	tions. Can be Rectangular, Elliptic or Cross.
NbIterations $[1]$	Number of times erosion and dilation are applied

# $\label{thm:normalize} \textbf{Normalize the image}.$

Option [default value]	Description
Norm [MinMax]	Norm type, can be:
	L1: L1 normalization
	L2: L2 normalization
	Linf: Linf normalization
	MinMax: min-max normalization
NormValue $\left[1.0\right]$	Norm value (for L1, L2 and Linf)
	Such that $  data  _{L_p} = NormValue$
NormMin $[0.0]$	Min value (for MinMax only)
	Such that $min(data) = NormMin$
NormMax $[1.0]$	Max value (for MinMax only)
	Such that $max(data) = NormMax$
PerChannel $[0]$	If true, normalize each channel individually

# ${\tt PadCropTransformation} \quad {\rm Pad/crop} \ {\rm the} \ {\rm image} \ {\rm to} \ {\rm a} \ {\rm specified} \ {\rm size}.$

Option [default value]	Description
Width	Width of the padded/cropped image
Height	Height of the padded/cropped image
PaddingBackground [MeanColor]	Background color used when padding. Possible values:
	MeanColor: pad with the mean color of the image
	BlackColor: pad with black

 ${\tt N2D2\ IP\ only}$  RandomAffineTransformation Apply a global random affine transformation to the values of the image.

Option [default value]	Description
GainRange [1.0 1.0]	Random gain $(\alpha)$ range (identical for all channels)
$\texttt{GainRange[*]} \ [1.0 \ 1.0]$	Random gain $(\alpha)$ range for channel *. Mutually exclusive
	with GainRange. If any specified, a different random gain will
	always be sampled for each channel. Default gain is 1.0 (no
	gain) for missing channels  The gain control the contract of the image
BiasRange [0.0 0.0]	The gain control the <i>contrast</i> of the image Random bias $(\beta)$ range (identical for all channels)
BiasRange [*] $[0.0 \ 0.0]$	Random bias $(\beta)$ range (identical for all channels)
	with BiasRange. If any specified, a different random bias will
	always be sampled for each channel. Default bias is 0.0 (no
	bias) for missing channels
	The bias control the <i>brightness</i> of the image
$\texttt{GammaRange} \; [1.0 \; 1.0]$	Random gamma $(\gamma)$ range (identical for all channels)
$\texttt{GammaRange[*]} \ [1.0 \ 1.0]$	Random gamma $(\gamma)$ range for channel *. Mutually exclusive
	with GammaRange. If any specified, a different random gamma
	will always be sampled for each channel. Default gamma is
	1.0 (no change) for missing channels
GainVarProb [1.0]	The gamma control more or less the <i>exposure</i> of the image
GainvarProb [1.0]	Probability to have a gain variation for each channel. If only one value is specified, the same probability applies to all
	the channels. In this case, the same gain variation will be
	sampled for all the channels only if a single range if specified
	for all the channels using GainRange. If more than one value
	is specified, a different random gain will always be sampled
	for each channel, even if the probabilities and ranges are
	identical
BiasVarProb $\left[1.0\right]$	Probability to have a bias variation for each channel. If only
	one value is specified, the same probability applies to all
	the channels. In this case, the same bias variation will be
	sampled for all the channels only if a single range if specified for all the channels using BiasRange. If more than one value
	is specified, a different random bias will always be sampled
	for each channel, even if the probabilities and ranges are
	identical
GammaVarProb [1.0]	Probability to have a gamma variation for each channel. If
	only one value is specified, the same probability applies to
	all the channels. In this case, the same gamma variation
	will be sampled for all the channels only if a single range if
	specified for all the channels using GammaRange. If more than
	one value is specified, a different random gamma will always
	be sampled for each channel, even if the probabilities and
DisjointGamma $[0]$	ranges are identical If true, gamma variation and gain/bias variation are mutu-
DISJOINGAMMA [O]	ally exclusive. The probability to have a random gamma
	variation is therefore GammaVarProb and the probability to have
	a gain/bias variation is 1-GammaVarProb.
ChannelsMask	If not empty, specifies on which channels the transformation
	is applied. For example, to apply the transformation only to
	the first and third channel, set ChannelsMask to 1 0 1

The equation of the transformation is:

$$S = \begin{cases} \text{numeric\_limits} < T > :: \text{max}() & \text{if is\_integer} < T > \\ 1.0 & \text{otherwise} \end{cases}$$

$$v(i,j) = \text{cv::saturate\_cast} < \text{T} > \left(\alpha \left(\frac{v(i,j)}{S}\right)^{\gamma} S + \beta.S\right)$$

RangeAffineTransformation Apply an affine transformation to the values of the image.

Option [default value]	Description
FirstOperator	First operator, can be Plus, Minus, Multiplies, Divides
FirstValue	First value
SecondOperator [Plus]	Second operator, can be Plus, Minus, Multiplies, Divides
SecondValue $\left[0.0\right]$	Second value

The final operation is the following:

$$f(x) = (x \underset{op_{1st}}{\circ} val_{1st}) \underset{op_{2nd}}{\circ} val_{2nd}$$

RangeClippingTransformation Clip the value range of the image.

Option [default value]	Description
RangeMin $[min(data)]$	Image values below RangeMin are clipped to 0
RangeMax $[max(data)]$	Image values above RangeMax are clipped to 1 (or the maximum
	integer value of the data type)

RescaleTransformation Rescale the image to a specified size.

Option [default value]	Description
Width	Width of the rescaled image
Height	Height of the rescaled image
KeepAspectRatio $\left[0 ight]$	If true, keeps the aspect ratio of the image
ResizeToFit $\left[1 ight]$	If true, resize along the longest dimension when
	KeepAspectRatio is true

ReshapeTransformation Reshape the data to a specified size.

Option [default value]	Description
NbRows	New number of rows
NbCols $[0]$	New number of cols $(0 = \text{no check})$
NbChannels $[0]$	New number of channels $(0 = \text{no change})$

2D2 IP only SliceExtractionTransformation Extract a slice from an image.

Option [default value]	Description
Width	Width of the slice to extract
Height	Height of the slice to extract
OffsetX $[0]$	X offset of the slice to extract
OffsetY $[0]$	Y offset of the slice to extract
${\tt RandomOffsetX} \; [0]$	If true, the X offset is chosen randomly
RandomOffsetY $\left[0\right]$	If true, the Y offset is chosen randomly
$\texttt{RandomRotation} \; [0]$	If true, extract randomly rotated slices
RandomRotationRange $[0.0\ 360.0]$	Range of the random rotations, in degrees, counterclockwise
	(if RandomRotation is enabled)
RandomScaling $\left[0\right]$	If true, extract randomly scaled slices
RandomScalingRange $\left[0.8\ 1.2\right]$	Range of the random scaling (if RandomRotation is enabled)
AllowPadding $\left[0\right]$	If true, zero-padding is allowed if the image is smaller than
	the slice to extract

**ThresholdTransformation** Apply a thresholding transformation to the image. This transformation is also useful for post-processing.

Option [default value]	Description
Threshold	Threshold value
OtsuMethod $\left[0 ight]$	Use Otsu's method to determine the optimal threshold (if
	true, the Threshold value is ignored)
Operation [Binary]	Thresholding operation to apply. Can be:
	Binary
	BinaryInverted
	Truncate
	ToZero
	ToZeroInverted
MaxValue $[1.0]$	Max. value to use with Binary and BinaryInverted operations

### TrimTransformation Trim the image.

Option [default value]	Description
NbLevels	Number of levels for the color discretization of the image
Method [Discretize]	Possible values are:
	Reduce: discretization using K-means
	Discretize: simple discretization

### N2D2 IP only WallisFilterTransformation Apply Wallis filter to the image.

Option [default value]	Description
Size	Size of the filter
Mean $\left[0.0 ight]$	Target mean value
StdDev $[1.0]$	Target standard deviation
PerChannel $[0]$	If true, apply Wallis filter to each channel individually (this
	parameter is meaningful only if Size is 0)

# 4.7 Network layers

# 4.7.1 Layer definition

Common set of parameters for any kind of layer.

Option [default value]	Description
Input	Name of the section(s) for the input layer(s). Comma sepa-
	rated
Туре	Type of the layer. Can be any of the type described below
Model [DefaultModel]	Layer model to use
DataType [DefaultDataType]	Layer data type to use. Please note that some layers may
	not support every data type.
ConfigSection []	Name of the configuration section for layer

To specify that the back-propagated error must be computed at the output of a given layer (generally the last layer, or output layer), one must add a target section named *LayerName*. Target:

```
...
[LayerName.Target]
TargetValue=1.0 ; default: 1.0
DefaultValue=0.0 ; default: -1.0
```

### 4.7.2 Weight fillers

Fillers to initialize weights and biases in the different type of layer. Usage example:

```
[conv1]
...
WeightsFiller=NormalFiller
WeightsFiller.Mean=0.0
WeightsFiller.StdDev=0.05
...
```

The initial weights distribution for each layer can be checked in the weights\_init folder, with an example shown in figure 14.

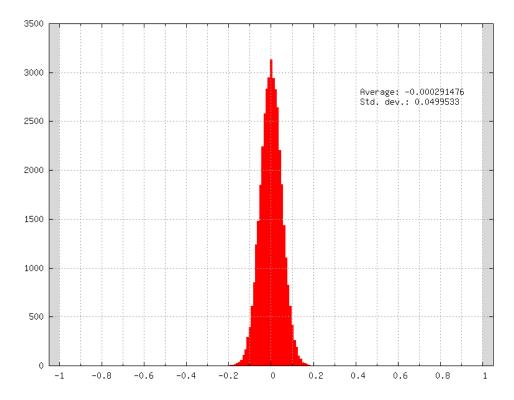


Figure 14: Initial weights distribution of a layer using a normal distribution (NormalFiller) with a 0 mean and a 0.05 standard deviation.

#### ConstantFiller Fill with a constant value.

Option	Description
FillerName. Value	Value for the filling

HeFiller Fill with an normal distribution with normalized variance taking into account the rectifier nonlinearity (He et al., 2015). This filler is sometimes referred as MSRA filler.

Option [default value]	Description
FillerName.VarianceNorm	Normalization, can be FamIn, Average or FamOut
[FanIn]	
FillerName.Scaling $[1.0]$	Scaling factor

Use a normal distribution with standard deviation  $\sqrt{\frac{2.0}{n}}$ .

- n = fan-in with FanIn, resulting in  $Var(W) = \frac{2}{fan\text{-}in}$
- $n=\frac{(fan\hbox{-}in\hbox{+}fan\hbox{-}out)}{2}$  with Average, resulting in  $Var(W)=\frac{4}{fan\hbox{-}in\hbox{+}fan\hbox{-}out}$
- n=fan-out with FanOut, resulting in  $Var(W)=\frac{2}{fan\text{-}out}$

#### NormalFiller Fill with a normal distribution.

Option [default value]	Description
FillerName.Mean $[0.0]$	Mean value of the distribution
$FillerName. { t StdDev} \ [1.0]$	Standard deviation of the distribution

### UniformFiller Fill with an uniform distribution.

Option [default value]	Description
FillerName.Min $[0.0]$	Min. value
FillerName.Max $[1.0]$	Max. value

**XavierFiller** Fill with an uniform distribution with normalized variance (Glorot and Bengio, 2010).

Option [default value]	Description
FillerName. VarianceNorm	Normalization, can be FamIn, Average or FamOut
[FanIn]	
FillerName.Distribution	Distribution, can be Uniform or Normal
[Uniform]	
FillerName.Scaling $[1.0]$	Scaling factor

Use an uniform distribution with interval [-scale, scale], with  $scale = \sqrt{\frac{3.0}{n}}$ .

- n = fan-in with FanIn, resulting in  $Var(W) = \frac{1}{fan\text{-}in}$
- $n = \frac{(fan-in+fan-out)}{2}$  with Average, resulting in  $Var(W) = \frac{2}{fan-in+fan-out}$
- n=fan-out with FanOut, resulting in  $Var(W)=\frac{1}{fan\text{-}out}$

#### 4.7.3 Weight solvers

SGDSolver\_Frame SGD Solver for Frame models.

Option [default value]	Description
SolverName.LearningRate	Learning rate
[0.01]	
SolverName . Momentum $[0.0]$	Momentum
$SolverName$ . Decay $\left[0.0 ight]$	Decay
SolverName.	Learning rate decay policy. Can be any of None, StepDecay,
LearningRatePolicy [None]	ExponentialDecay, InvTDecay, PolyDecay
SolverName.	Learning rate step size (in number of stimuli)
LearningRateStepSize $\left[1\right]$	
SolverName . LearningRateDecay	Learning rate decay
[0.1]	
SolverName. Clamping $[0]$	If true, clamp the weights and bias between -1 and 1
SolverName.Power $[0.0]$	Polynomial learning rule power parameter
$SolverName. { t MaxIterations}$	Polynomial learning rule maximum number of iterations
[0.0]	

The learning rate decay policies are the following:

- StepDecay: every SolverName.LearningRateStepSize stimuli, the learning rate is reduced by a factor SolverName.LearningRateDecay;
- ExponentialDecay: the learning rate is  $\alpha = \alpha_0 \exp(-kt)$ , with  $\alpha_0$  the initial learning rate SolverName.LearningRate, k the rate decay SolverName.LearningRateDecay and t the step number (one step every SolverName.LearningRateStepSize stimuli);
- InvTDecay: the learning rate is  $\alpha = \alpha_0/(1+kt)$ , with  $\alpha_0$  the initial learning rate SolverName. LearningRate, k the rate decay SolverName.LearningRateDecay and t the step number (one step every SolverName.LearningRateStepSize stimuli).
- InvDecay: the learning rate is  $\alpha = \alpha_0 * (1 + kt)^{-n}$ , with  $\alpha_0$  the initial learning rate Solver-Name.LearningRate, k the rate decay SolverName.LearningRateDecay, t the current iteration and n the power parameter SolverName.Power
- PolyDecay: the learning rate is  $\alpha = \alpha_0 * (1 \frac{k}{t})^n$ , with  $\alpha_0$  the initial learning rate Solver-Name. LearningRate, k the current iteration, t the maximum number of iteration SolverName. MaxIterations and n the power parameter SolverName. Power

SGDSolver\_Frame\_CUDA SGD Solver for Frame\_CUDA models.

Option [default value]	Description
SolverName . Learning Rate	Learning rate
[0.01]	
$SolverName$ . Momentum $\left[0.0 ight]$	Momentum
$SolverName$ . Decay $\left[0.0 ight]$	Decay
SolverName .	Learning rate decay policy. Can be any of None, StepDecay,
LearningRatePolicy [None]	ExponentialDecay, InvTDecay
SolverName .	Learning rate step size (in number of stimuli)
LearningRateStepSize $\left[1 ight]$	
SolverName . Learning Rate Decay	Learning rate decay
[0.1]	
$SolverName. {\tt Clamping} \ [0]$	If true, clamp the weights and bias between -1 and 1

The learning rate decay policies are identical to the ones in the SGDSolver\\_Frame solver.

AdamSolver\_Frame Adam Solver for Frame models (Kingma and Ba, 2014).

Option [default value]	Description
SolverName.LearningRate	Learning rate (stepsize)
[0.001]	
SolverName.Beta1 $[0.9]$	Exponential decay rate of these moving average of the first
	moment
SolverName.Beta2 $[0.999]$	Exponential decay rate of these moving average of the second
	moment
SolverName.Epsilon [1.0e-8]	Epsilon

AdamSolver\_Frame\_CUDA Adam Solver for Frame\_CUDA models (Kingma and Ba, 2014).

Option [default value]	Description	
SolverName.LearningRate	Learning rate (stepsize)	
[0.001]		
$SolverName. {\tt Beta1} \ [0.9]$	Exponential decay rate of these moving average of the first	
	moment	
SolverName.Beta2 $[0.999]$	Exponential decay rate of these moving average of the second	
	moment	
SolverName.Epsilon $[1.0e-8]$	Epsilon	

#### 4.7.4 Activation functions

Activation function to be used at the output of layers. Usage example:

```
[conv1]
...
ActivationFunction=Rectifier
ActivationFunction.LeakSlope=0.01
ActivationFunction.Clipping=20
...
```

Logistic activation function.

Logistic With Loss activation function.

### Rectifier or ReLU activation function.

Option [default value]	Description
	Leak slope for negative inputs
ActivationFunction.Clipping	Clipping value for positive outputs
[0.0]	

Saturation Saturation activation function.

Softplus Softplus activation function.

Tanh Tanh activation function.

Computes  $y = tanh(\alpha x)$ .

Option [default value]	Description
ActivationFunction.Alpha $\left[1.0\right]$	$\alpha$ parameter

**TanhLeCun** Tanh activation function with an  $\alpha$  parameter of  $1.7159 \times (2.0/3.0)$ .

### 4.7.5 Anchor

Anchor layer for Faster R-CNN or Single Shot Detector.

Option [default value]	Description	
Input	This layer takes one or two inputs. The total number of	
	input channels must be ScoresCls $+4$ , with ScoresCls being	
	equal to 1 or 2.	
Anchor[*]	Anchors definition. For each anchor, there must be two	
	space-separated values: the root area and the aspect ratio.	
ScoresCls	Number of classes per anchor. Must be 1 (if the scores input	
	uses logistic regression) or 2 (if the scores input is a two-class	
	softmax layer)	
FeatureMapWidth	Reference width use to scale anchors coordinate.	
[StimuliProvider.Width]		
FeatureMapHeight	Reference height use to scale anchors coordinate.	
[StimuliProvider.Height]		

# Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
PositiveIoU $\left[0.7 ight]$	all Frame	Assign a positive label for anchors whose IoU overlap
		is higher than PositiveIoU with any ground-truth box
NegativeIoU $\left[0.3 ight]$	all Frame	Assign a negative label for non-positive anchors whose
		IoU overlap is lower than NegativeIoU for all ground-
		truth boxes
LossLambda $\left[10.0\right]$	all Frame	Balancing parameter $\lambda$
LossPositiveSample $\left[128\right]$	all Frame	Number of random positive samples for the loss com-
		putation

LossNegativeSample $\left[128\right]$	all Frame	Number of random negative samples for the loss com-
		putation

### Usage example:

```
; RPN network: cls layer
[scores]
Input=...
Type=Conv
KernelWidth=1
KernelHeight=1
; 18 channels for 9 anchors
NbOutputs=18
[scores.softmax]
Input=scores
Type=Softmax
NbOutputs=[scores]NbOutputs
WithLoss=1
; RPN network: coordinates layer
[coordinates]
Input=...
Type=Conv
KernelWidth=1
KernelHeight=1
; 36 channels for 4 coordinates x 9 anchors
NbOutputs=36
; RPN network: anchors
[anchors]
Input=scores.softmax,coordinates
Type=Anchor
ScoresCls=2; using a two-class softmax for the scores
Anchor[0]=32 1.0
Anchor[1]=48 1.0
Anchor[2]=64 1.0
Anchor[3]=80 1.0
Anchor[4]=96 1.0
Anchor[5]=112 1.0
Anchor[6]=128 1.0
Anchor[7]=144 1.0
Anchor[8]=160 1.0
ConfigSection=anchors.config
[anchors.config]
PositiveIoU=0.7
NegativeIoU=0.3
LossLambda=1.0
```

Outputs remapping Outputs remapping allows to convert *scores* and *coordinates* output feature maps layout from another ordering that the one used in the N2D2 Anchor layer, during weights import/export.

For example, lets consider that the imported weights corresponds to the following output feature maps ordering:

```
0 anchor[0].y
1 anchor[0].x
```

```
2 anchor[0].h
3 anchor[0].w
4 anchor[1].y
5 anchor[1].x
6 anchor[1].h
7 anchor[1].w
8 anchor[2].y
9 anchor[2].x
10 anchor[2].h
11 anchor[2].w
   The output feature maps ordering required by the Anchor layer is:
0 anchor[0].x
1 anchor[1].x
2 anchor[2].x
3 anchor[0].y
4 anchor[1].y
5 anchor[2].y
```

The feature maps ordering can be changed during weights import/export:

```
; RPN network: coordinates layer
[coordinates]
Input=...
Type=Conv
KernelWidth=1
KernelHeight=1
; 36 channels for 4 coordinates x 9 anchors
NbOutputs=36
...
ConfigSection=coordinates.config
[coordinates.config]
WeightsExportFormat=HWCO; Weights format used by TensorFlow
OutputsRemap=1:4,0:4,3:4,2:4
```

#### 4.7.6 Conv

6 anchor[0].w 7 anchor[1].w 8 anchor[2].w 9 anchor[0].h 10 anchor[1].h 11 anchor[2].h

Convolutional layer.

Option [default value]	Description		
KernelWidth	Width of the kernels		
KernelHeight	Height of the kernels		
KernelDepth []	Depth of the kernels (implies 3D kernels)		
	OR		
KernelSize []	Kernels size (implies 2D square kernels)		
OR			
KernelDims []	List of space-separated dimensions for N-D kernels		
NbOutputs	Outputs Number of output channels		
SubSampleX [1]	SubSampleX [1] X-axis subsampling factor of the output feature maps		
SubSampleY [1]	ubSampleY [1] Y-axis subsampling factor of the output feature maps		
SubSampleZ []	SampleZ [] Z-axis subsampling factor of the output feature maps		
OR			

SubSample [1]	Subsampling factor of the output feature maps		
	OR		
SubSampleDims []	List of space-separated subsampling dimensions for N-D kernels		
StrideX [1]	X-axis stride of the kernels		
StrideY [1]	Y-axis stride of the kernels		
StrideZ []	Z-axis stride of the kernels		
U	OR		
Stride [1]	Stride of the kernels		
	OR		
StrideDims []	List of space-separated stride dimensions for N-D kernels		
PaddingX [0]	X-axis input padding		
PaddingY [0]	Y-axis input padding		
PaddingZ []	Z-axis input padding		
[0]	OR		
Padding [0]	Input padding		
	OR		
PaddingDims	List of space-separated padding dimensions for N-D kernels		
DilationX [1]	X-axis dilation of the kernels		
DilationY [1]	Y-axis dilation of the kernels		
DilationZ []	Z-axis dilation of the kernels		
	OR		
Dilation $\left[1 ight]$	Dilation of the kernels		
	OR		
DilationDims []	List of space-separated dilation dimensions for N-D kernels		
ActivationFunction [Tanh]	Activation function. Can be any of Logistic, LogisticWithLoss,		
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh		
WeightsFiller	Weights initial values filler		
[NormalFiller(0.0, 0.05)]			
BiasFiller	Biases initial values filler		
[NormalFiller(0.0, 0.05)]			
Mapping.NbGroups	Mapping: number of groups		
mapping.modroups []	(mutually exclusive with all other Mapping.* options)		
Mapping.ChannelsPerGroup	Mapping: number of channels per group		
rapping. Chammers erdroup	(mutually exclusive with all other Mapping.* options)		
Manusium Giary [1]	`		
Mapping.SizeX [1]	Mapping canvas pattern default width  Mapping canvas pattern default height		
Mapping.SizeY [1]	Mapping canvas pattern default height Mapping canvas pattern default size		
Mapping.Size [1]	Mapping canvas pattern default size		
w [1]	(mutually exclusive with Mapping.SizeX and Mapping.SizeY)		
Mapping.StrideX [1]	Mapping canvas default X-axis step		
Mapping.StrideY [1]	Mapping canvas default Y-axis step		
Mapping.Stride $\left[1 ight]$	Mapping canvas default step		
	(mutually exclusive with Mapping.StrideX and Mapping.StrideY)		
Mapping.OffsetX $[0]$	Mapping canvas default X-axis offset		
Mapping.OffsetY $[0]$	Mapping canvas default Y-axis offset		
Mapping.Offset $\left[0 ight]$	Mapping canvas default offset		
	(mutually exclusive with Mapping.OffsetX and Mapping.OffsetY)		
Mapping.NbIterations $\left[0 ight]$	Mapping canvas pattern default number of iterations (0		
	means no limit)		
t Mapping(in).SizeX [1]	Mapping canvas pattern default width for input layer in		
Mapping(in).SizeY [1]	Mapping canvas pattern default height for input layer in		

Mapping(in).Size $[1]$	Mapping canvas pattern default size for input layer in	
	(mutually exclusive with Mapping(in).SizeX and	
	Mapping(in).SizeY)	
Mapping(in).StrideX $\left[1 ight]$	Mapping canvas default X-axis step for input layer in	
Mapping(in).StrideY $[1]$	Mapping canvas default Y-axis step for input layer in	
Mapping(in).Stride $[1]$	Mapping canvas default step for input layer in	
	(mutually exclusive with Mapping(in).StrideX and	
	Mapping(in).StrideY)	
Mapping(in).OffsetX $\left[0 ight]$	Mapping canvas default X-axis offset for input layer in	
Mapping(in).OffsetY $\left[0 ight]$	Mapping canvas default Y-axis offset for input layer in	
Mapping(in).Offset $\left[0 ight]$	Mapping canvas default offset for input layer in	
	(mutually exclusive with Mapping(in).OffsetX and	
	Mapping(in).OffsetY)	
Mapping(in).NbIterations $\left[0 ight]$	Mapping canvas pattern default number of iterations for	
	input layer in (0 means no limit)	
WeightsSharing []	Share the weights with an other layer	
BiasesSharing []	Share the biases with an other layer	

# Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
NoBias $[0]$	all Frame	If true, don't use bias
Solvers.*	all Frame	Any solver parameters
WeightsSolver.*	all Frame	Weights solver parameters, take precedence over the
		Solvers.* parameters
BiasSolver.*	all Frame	Bias solver parameters, take precedence over the
		Solvers.* parameters
WeightsExportFormat	all Frame	Weights import/export format. Can be ochw or ochw,
[OCHW]		with 0 the output feature map, c the input feature map
		(channel), H the kernel row and W the kernel column, in
		the order of the outermost dimension (in the leftmost
		position) to the innermost dimension (in the rightmost
		position)
$\texttt{WeightsExportFlip} \; [0]$	all Frame	If true, import/export flipped kernels

# Configuration parameters (Spike models)

Experimental option (implementation may be wrong or susceptible to change)

Option [default value]	Model(s)	Description
IncomingDelay [1 TimePs	all Spike	Synaptic incoming delay $w_{delay}$
$;100 \; \mathtt{TimeFs}]$		
Threshold $\left[1.0\right]$	Spike, Spike_RRAM	Threshold of the neuron $I_{thres}$
BipolarThreshold $\left[1 ight]$	Spike, Spike_RRAM	If true, the threshold is also applied to the absolute
		value of negative values (generating negative spikes)
Leak $\left[0.0 ight]$	Spike, Spike_RRAM	Neural leak time constant $\tau_{leak}$ (if 0, no leak)
Refractory $\left[0.0\right]$	Spike, Spike_RRAM	Neural refractory period $T_{refrac}$

WeightsRelInit $[0.0;0.05]$	Spike	Relative initial synaptic weight $w_{init}$
WeightsMinMean $[1;0.1]$	Spike_RRAM	Mean minimum synaptic weight $w_{min}$
WeightsMaxMean $[100;10.0]$	Spike_RRAM	Mean maximum synaptic weight $w_{max}$
WeightsMinVarSlope $[0.0]$	Spike_RRAM	OXRAM specific parameter
WeightsMinVarOrigin $[0.0]$	Spike_RRAM	OXRAM specific parameter
WeightsMaxVarSlope $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
WeightsMaxVarOrigin $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
WeightsSetProba $\left[1.0 ight]$	Spike_RRAM	Intrinsic SET switching probability $P_{SET}$ (upon receiv-
		ing a SET programming pulse). Assuming uniform
		statistical distribution (not well supported by experi-
		ments on RRAM)
WeightsResetProba $\left[1.0\right]$	Spike_RRAM	Intrinsic RESET switching probability $P_{RESET}$ (upon
		receiving a RESET programming pulse). Assuming
		uniform statistical distribution (not well supported by
		experiments on RRAM)
SynapticRedundancy $\left[1 ight]$	Spike_RRAM	Synaptic redundancy (number of RRAM device per
		synapse)
BipolarWeights $\left[0 ight]$	Spike_RRAM	Bipolar weights
BipolarIntegration $\left[0 ight]$	Spike_RRAM	Bipolar integration
LtpProba $\left[0.2\right]$	Spike_RRAM	Extrinsic STDP LTP probability (cumulative with in-
		trinsic SET switching probability $P_{SET}$ )
LtdProba $\left[0.1\right]$	Spike_RRAM	Extrinsic STDP LTD probability (cumulative with
		intrinsic RESET switching probability $P_{RESET}$ )
StdpLtp $[1000 \text{ TimePs}]$	Spike_RRAM	STDP LTP time window $T_{LTP}$
InhibitRefractory $[0]$	Spike_RRAM	Neural lateral inhibition period $T_{inhibit}$
TimePs]		
EnableStdp [1]	Spike_RRAM	If false, STDP is disabled (no synaptic weight change)
RefractoryIntegration $[1]$	Spike_RRAM	If true, reset the integration to 0 during the refractory
		period
${\tt DigitalIntegration} \; [0]$	Spike_RRAM	If false, the analog value of the devices is integrated,
		instead of their binary value

# 4.7.7 Deconv

 ${\bf De convolution layer.}$ 

Option [default value]	Description	
KernelWidth	Width of the kernels	
KernelHeight	Height of the kernels	
KernelDepth []	Depth of the kernels (implies 3D kernels)	
	OR	
KernelSize []	Kernels size (implies 2D square kernels)	
OR		
KernelDims []	List of space-separated dimensions for N-D kernels	
NbOutputs	Number of output channels	
StrideX [1]	X-axis stride of the kernels	
StrideY [1]	Y-axis stride of the kernels	
StrideZ []	Z-axis stride of the kernels	
	OR	
Stride [1]	Stride of the kernels	

	OP	
	List of space-separated stride dimensions for N-D kernels	
PaddingX [0]	X-axis input padding	
PaddingY [0]	Y-axis input padding	
Padding		
PaddingZ []	Z-axis input padding	
Padding [0]	OR   Input padding	
radding [O]	OR	
PaddingDims	List of space-separated padding dimensions for N-D kernels	
DilationX [1]	X-axis dilation of the kernels	
DilationY [1]	Y-axis dilation of the kernels	
Dilation [1]   Dilation [7]	Z-axis dilation of the kernels	
DITACTORZ []	OR	
Dilation [1]	Dilation of the kernels	
Direction [1]	OR	
DilationDims	List of space-separated dilation dimensions for N-D kernels	
ActivationFunction [Tanh]	Activation function. Can be any of Logistic, LogisticWithLoss,	
Accivation miction [idin]	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
WeightsFiller	Weights initial values filler	
NormalFiller(0.0, 0.05)	Weights initial values liner	
BiasFiller	Biases initial values filler	
[NormalFiller(0.0, 0.05)]	Diases initial values inier	
Mapping.NbGroups	Mapping: number of groups	
mapping.wbdroups	(mutually exclusive with all other Mapping.* options)	
   Mapping.ChannelsPerGroup	Mapping: number of channels per group	
mapping.chammersreigroup	(mutually exclusive with all other Mapping.* options)	
$\mid$ Mapping.SizeX $[1]$	Mapping canvas pattern default width	
Mapping.SizeY [1]	Mapping canvas pattern default height	
Mapping.Size [1]	Mapping canvas pattern default size	
mapping.bize [1]	(mutually exclusive with Mapping.SizeX and Mapping.SizeY)	
Mapping.StrideX $[1]$	Mapping canvas default X-axis step	
Mapping.StrideY [1]	Mapping canvas default Y-axis step	
Mapping.Stride [1]	Mapping canvas default step	
mapping.bulide [1]	(mutually exclusive with Mapping.StrideX and Mapping.StrideY)	
Mapping.OffsetX $[0]$	Mapping canvas default X-axis offset	
Mapping.OffsetY [0]	Mapping canvas default Y-axis offset	
Mapping.Offset [0]	Mapping canvas default 1-axis offset  Mapping canvas default offset	
mapping.oiiset [O]	(mutually exclusive with Mapping.OffsetX and Mapping.OffsetY)	
Mapping.NbIterations $[0]$	Mapping canvas pattern default number of iterations (0	
	means no limit)	
Mapping(in).SizeX $[1]$	Mapping canvas pattern default width for input layer in	
Mapping(in).SizeY [1]	Mapping canvas pattern default height for input layer in	
Mapping(in).Size [1]	Mapping canvas pattern default size for input layer in	
uabbing(in).pize [1]	(mutually exclusive with Mapping(in).SizeX and	
	(Inditidally exclusive with mapping(in).Sizek and Mapping(in).SizeY)	
Mapping(in).StrideX $[1]$	Mapping canvas default X-axis step for input layer in	
Mapping(in).StrideX [1]	Mapping canvas default Y-axis step for input layer in	
Mapping(in).Stride[1]	Mapping canvas default 1-axis step for input layer in	
	(mutually exclusive with Mapping(in). StrideX and	
	(Mapping(in).Stridex and Mapping(in).Stridex and	
Monning(in) Officety [O]	Mapping canvas default X-axis offset for input layer in	
Mapping(in).OffsetX $\left[0 ight]$	mapping canvas default A-axis offset for input layer in	

Mapping(in).OffsetY $\left[0 ight]$	Mapping canvas default Y-axis offset for input layer in	
Mapping(in).Offset $\left[0 ight]$	Mapping canvas default offset for input layer in	
	(mutually exclusive with Mapping(in).Offset% and	
	Mapping(in).OffsetY)	
Mapping(in).NbIterations $\left[0 ight]$	Mapping canvas pattern default number of iterations for	
	input layer in (0 means no limit)	
WeightsSharing []	Share the weights with an other layer	
BiasesSharing []	Share the biases with an other layer	

### Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
NoBias $[0]$	all Frame	If true, don't use bias
BackPropagate $[1]$	all Frame	If true, enable backpropagation
Solvers.*	all Frame	Any solver parameters
WeightsSolver.*	all Frame	Weights solver parameters, take precedence over the
		Solvers.* parameters
BiasSolver.*	all Frame	Bias solver parameters, take precedence over the
		Solvers.* parameters
WeightsExportFormat	all Frame	Weights import/export format. Can be ochw or ochw,
[OCHW]		with 0 the output feature map, c the input feature map
		(channel), H the kernel row and W the kernel column, in
		the order of the outermost dimension (in the leftmost
		position) to the innermost dimension (in the rightmost
		position)
WeightsExportFlip $\left[0 ight]$	all Frame	If true, import/export flipped kernels

#### 4.7.8 Pool

Pooling layer.

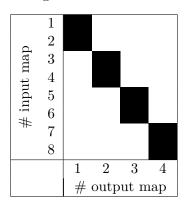
There are two CUDA models for this cell:

- Frame\_CUDA, which uses CuDNN as back-end and only supports one-to-one input to output map connection;
- Frame\_EXT\_CUDA, which uses custom CUDA kernels and allows arbitrary connections between input and output maps (and can therefore be used to implement Maxout or both Maxout and Pooling simultaneously).

**Maxout example** In the following INI section, one implements a Maxout between each consecutive pair of 8 input maps:

[maxout\_layer]
Input=...
Type=Pool
Model=Frame\_EXT\_CUDA
PoolWidth=1
PoolHeight=1
NbOutputs=4
Pooling=Max
Mapping.SizeY=2
Mapping.StrideY=2

The layer connectivity is the following:



O-4:[1-f1, 1 ]	D:	
Option [default value]	Description	
Pooling	Type of pooling (Max or Average)	
PoolWidth	Width of the pooling area	
PoolHeight	Height of the pooling area	
PoolDepth []	Depth of the pooling area (implies 3D pooling area)	
	OR	
PoolSize []	Pooling area size (implies 2D square pooling area)	
n	OR	
PoolDims	List of space-separated dimensions for N-D pooling area	
NbOutputs	Number of output channels	
StrideX [1]	X-axis stride of the pooling area	
StrideY [1]	Y-axis stride of the pooling area	
StrideZ []	Z-axis stride of the pooling area	
	OR	
Stride $[1]$	Stride of the pooling area	
	OR	
StrideDims []	List of space-separated stride dimensions for N-D pooling	
	area	
PaddingX [0]	X-axis input padding	
PaddingY [0]	Y-axis input padding	
PaddingZ []	Z-axis input padding	
	OR	
Padding $[0]$	Input padding	
OR		
PaddingDims []	List of space-separated padding dimensions for N-D pooling	
	area	
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,	
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
Mapping.NbGroups []	Mapping: number of groups	
	(mutually exclusive with all other Mapping.* options)	
Mapping.ChannelsPerGroup	Mapping: number of channels per group	
	(mutually exclusive with all other Mapping.* options)	
Mapping.SizeX [1]	Mapping canvas pattern default width	
Mapping.SizeY [1]	Mapping canvas pattern default height	
Mapping.Size [1]	Mapping canvas pattern default size	
	(mutually exclusive with Mapping.SizeX and Mapping.SizeY)	
Mapping.StrideX [1]	Mapping canvas default X-axis step	
Mapping.StrideY [1]	Mapping canvas default Y-axis step	
110,[+]	TT O same as a same a same a same	

$\mid$ Mapping.Stride $[1]$	Mapping canvas default step
	(mutually exclusive with Mapping.StrideX and Mapping.StrideY)
Mapping.OffsetX $[0]$	Mapping canvas default X-axis offset
Mapping.OffsetY $[0]$	Mapping canvas default Y-axis offset
Mapping.Offset $[0]$	Mapping canvas default offset
	(mutually exclusive with Mapping.OffsetX and Mapping.OffsetY)
Mapping.NbIterations $[0]$	Mapping canvas pattern default number of iterations (0 means no limit)
Mapping(in).SizeX $[1]$	Mapping canvas pattern default width for input layer in
Mapping(in).SizeY $[1]$	Mapping canvas pattern default height for input layer in
Mapping(in).Size $\left[1 ight]$	Mapping canvas pattern default size for input layer in
	(mutually exclusive with Mapping(in).SizeX and
	Mapping(in).SizeY)
Mapping(in).StrideX $\left[1 ight]$	Mapping canvas default X-axis step for input layer in
Mapping(in).StrideY $\left[1 ight]$	Mapping canvas default Y-axis step for input layer in
Mapping(in).Stride $\left[1 ight]$	Mapping canvas default step for input layer in
	(mutually exclusive with Mapping(in).StrideX and
	Mapping(in).StrideY)
Mapping(in).OffsetX $\left[0 ight]$	Mapping canvas default X-axis offset for input layer in
Mapping(in).OffsetY $\left[0 ight]$	Mapping canvas default Y-axis offset for input layer in
Mapping(in).Offset $\left[0 ight]$	Mapping canvas default offset for input layer in
	(mutually exclusive with Mapping(in).Offset% and
	Mapping(in).OffsetY)
Mapping(in).NbIterations $\left[0 ight]$	Mapping canvas pattern default number of iterations for
	input layer in (0 means no limit)

# Configuration parameters (Spike models)

Option [default value]	Model(s)	Description
IncomingDelay [1 TimePs	all Spike	Synaptic incoming delay $w_{delay}$
$;100 \; \mathtt{TimeFs}]$		
value		

# **4.7.9** Unpool

Unpooling layer.

Option [default value]	Description	
Pooling	Type of pooling (Max or Average)	
PoolWidth Width of the pooling area		
PoolHeight	Height of the pooling area	
PoolDepth []	Depth of the pooling area (implies 3D pooling area)	
OR		
PoolSize []	Pooling area size (implies 2D square pooling area)	
	OR	
PoolDims []	List of space-separated dimensions for N-D pooling area	
NbOutputs	Number of output channels	

ArgMax	Name of the associated pool layer for the argmax (the pool	
J	layer input and the unpool layer output dimension must	
	match)	
StrideX [1]	X-axis stride of the pooling area	
StrideY [1]	Y-axis stride of the pooling area	
StrideZ []	Z-axis stride of the pooling area	
	OR	
Stride [1]	Stride of the pooling area	
	OR	
StrideDims []	List of space-separated stride dimensions for N-D pooling	
	area	
PaddingX $[0]$	X-axis input padding	
PaddingY $[0]$	Y-axis input padding	
PaddingZ []	Z-axis input padding	
	OR	
Padding $[0]$	Input padding	
	OR	
PaddingDims []	List of space-separated padding dimensions for N-D pooling	
	area	
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,	
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
Mapping.NbGroups []	Mapping: number of groups	
_	(mutually exclusive with all other Mapping.* options)	
Mapping.ChannelsPerGroup []	Mapping: number of channels per group	
	(mutually exclusive with all other Mapping.* options)	
Mapping.SizeX $[1]$	Mapping canvas pattern default width	
Mapping.SizeY [1]	Mapping canvas pattern default height	
Mapping.Size $[1]$	Mapping canvas pattern default size	
	(mutually exclusive with Mapping.SizeX and Mapping.SizeY)	
Mapping.StrideX [1]	Mapping canvas default X-axis step	
Mapping.StrideY [1]	Mapping canvas default Y-axis step	
Mapping.Stride $[1]$	Mapping canvas default step	
[o]	(mutually exclusive with Mapping.StrideY and Mapping.StrideY)	
Mapping.OffsetX $[0]$	Mapping canvas default X-axis offset	
Mapping.OffsetY [0]	Mapping canvas default Y-axis offset	
Mapping.Offset $[0]$	Mapping canvas default offset	
w · w · [0]	(mutually exclusive with Mapping.OffsetX and Mapping.OffsetY)	
Mapping.NbIterations $[0]$	Mapping canvas pattern default number of iterations (0	
Monning(in) Gi-v [1]	means no limit)  Mapping capyag pattern default width for input layer in	
Mapping(in).SizeX [1] Mapping(in).SizeY [1]	Mapping canvas pattern default width for input layer in Mapping canvas pattern default height for input layer in	
Mapping(in).Size [1]	Mapping canvas pattern default size for input layer in (mutually exclusive with Mapping(in).SizeX and	
	(Inditially exclusive with Mapping(in).Sizex and Mapping(in).SizeY)	
Mapping(in).StrideX [1]	Mapping canvas default X-axis step for input layer in	
Mapping(in).StrideY [1]	Mapping canvas default Y-axis step for input layer in	
Mapping(in).Strider[1]	Mapping canvas default 1-axis step for input layer in	
	(mutually exclusive with Mapping(in).StrideX and	
	Mapping(in).StrideY)	
Mapping(in).OffsetX $[0]$	Mapping canvas default X-axis offset for input layer in	
Mapping(in).OffsetY [0]	Mapping canvas default Y-axis offset for input layer in	
rapping(in).uiiseti [U]	mapping canvas default 1-axis onset for input layer in	

${\tt Mapping(in).Offset} \ \big[ 0 \big]$	Mapping canvas default offset for input layer in (mutually exclusive with Mapping(in).OffsetX and
	Mapping(in).OffsetY)
Mapping(in).NbIterations $\left[0 ight]$	Mapping canvas pattern default number of iterations for
	input layer in (0 means no limit)

#### 4.7.10 ElemWise

Element-wise operation layer.

Option [default value]	Description
NbOutputs	Number of output neurons
Operation	Type of operation (Sum, AbsSum, EuclideanSum, Prod, or Max)
Weights $\left[1.0 ight]$	Weights for the Sum, AbsSum, and EuclideanSum operation, in
	the same order as the inputs
Shifts $[0.0]$	Shifts for the Sum and EuclideanSum operation, in the same
	order as the inputs
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh

Given N input tensors  $T_i$ , performs the following operation:

Sum operation  $T_{out} = \sum_{1}^{N} (w_i T_i + s_i)$ 

AbsSum operation  $T_{out} = \sum_{1}^{N} (w_i | T_i |)$ 

EuclideanSum operation  $T_{out} = \sqrt{\sum_{1}^{N} \left(w_i T_i + s_i\right)^2}$ 

Prod operation  $T_{out} = \prod_{1}^{N}(T_i)$ 

Max operation  $T_{out} = MAX_1^N(T_i)$ 

**Examples** Sum of two inputs  $(T_{out} = T_1 + T_2)$ :

[elemwise\_sum]
Input=layer1,layer2
Type=ElemWise
NbOutputs=[layer1]NbOutputs
Operation=Sum

Weighted sum of two inputs, by a factor 0.5 for layer1 and 1.0 for layer2 ( $T_{out} = 0.5 \times T_1 + 1.0 \times T_2$ ):

[elemwise\_weighted\_sum]
Input=layer1,layer2
Type=ElemWise
NbOutputs=[layer1]NbOutputs
Operation=Sum
Weights=0.5 1.0

Single input scaling by a factor 0.5 and shifted by 0.1 ( $T_{out} = 0.5 \times T_1 + 0.1$ ):

```
[elemwise_scale]
Input=layer1
Type=ElemWise
NbOutputs=[layer1]NbOutputs
Operation=Sum
Weights=0.5
Shifts=0.1
```

Absolute value of an input  $(T_{out} = |T_1|)$ :

```
[elemwise_abs]
Input=layer1
Type=ElemWise
NbOutputs=[layer1]NbOutputs
Operation=Abs
```

### 4.7.11 FMP

Fractional max pooling layer (Graham, 2014).

Option [default value]	Description
NbOutputs	Number of output channels
ScalingRatio	Scaling ratio. The output size is $round\left(\frac{\text{input size}}{\text{scaling ratio}}\right)$ .
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh

# Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
Overlapping $[1]$	all Frame	If true, use overlapping regions, else use disjoint regions
PseudoRandom $[1]$	all Frame	If true, use pseudorandom sequences, else use random
		sequences

### 4.7.12 Fc

Fully connected layer.

Option [default value]	Description
NbOutputs	Number of output neurons
WeightsFiller	Weights initial values filler
[NormalFiller(0.0, 0.05)]	
BiasFiller	Biases initial values filler
[NormalFiller(0.0, 0.05)]	
ActivationFunction [Tanh]	Activation function. Can be any of Logistic, LogisticWithLoss,
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh

### Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
NoBias $[0]$	all Frame	If true, don't use bias
BackPropagate $[1]$	all Frame	If true, enable backpropagation
Solvers.*	all Frame	Any solver parameters
WeightsSolver.*	all Frame	Weights solver parameters, take precedence over the
		Solvers.* parameters
BiasSolver.*	all Frame	Bias solver parameters, take precedence over the
		Solvers.* parameters
DropConnect $[1.0]$	Frame	If below 1.0, fraction of synapses that are disabled with
		drop connect

# Configuration parameters (Spike models)

Option [default value]	Model(s)	Description
IncomingDelay [1 TimePs	all Spike	Synaptic incoming delay $w_{delay}$
$;100 \; \mathtt{TimeFs}]$		, and the second
Threshold $\left[1.0\right]$	Spike, Spike_RRAM	Threshold of the neuron $I_{thres}$
BipolarThreshold $\left[1 ight]$	Spike, Spike_RRAM	If true, the threshold is also applied to the absolute
		value of negative values (generating negative spikes)
Leak $\left[0.0\right]$	Spike, Spike_RRAM	Neural leak time constant $\tau_{leak}$ (if 0, no leak)
Refractory $\left[0.0\right]$	Spike, Spike_RRAM	Neural refractory period $T_{refrac}$
TerminateDelta $\left[0 ight]$	Spike, Spike_RRAM	Terminate delta
WeightsRelInit $[0.0;0.05]$	Spike	Relative initial synaptic weight $w_{init}$
WeightsMinMean $[1;0.1]$	Spike_RRAM	Mean minimum synaptic weight $w_{min}$
WeightsMaxMean $[100;10.0]$	Spike_RRAM	Mean maximum synaptic weight $w_{max}$
WeightsMinVarSlope $\left[0.0\right]$	Spike_RRAM	OXRAM specific parameter
WeightsMinVarOrigin $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
WeightsMaxVarSlope $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
WeightsMaxVarOrigin $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
WeightsSetProba $\left[1.0\right]$	Spike_RRAM	Intrinsic SET switching probability $P_{SET}$ (upon receiv-
WeightsResetProba $\left[1.0\right]$	Spike_RRAM	ing a SET programming pulse). Assuming uniform statistical distribution (not well supported by experiments on RRAM) Intrinsic RESET switching probability $P_{RESET}$ (upon receiving a RESET programming pulse). Assuming uniform statistical distribution (not well supported by experiments on RRAM)
${\tt SynapticRedundancy} \ [1]$	Spike_RRAM	Synaptic redundancy (number of RRAM device per synapse)
BipolarWeights $\left[0 ight]$	Spike_RRAM	Bipolar weights
BipolarIntegration $[0]$	Spike_RRAM	Bipolar integration
LtpProba $\left[0.2\right]$	Spike_RRAM	Extrinsic STDP LTP probability (cumulative with in-
		trinsic SET switching probability $P_{SET}$ )
LtdProba $\left[0.1\right]$	Spike_RRAM	Extrinsic STDP LTD probability (cumulative with
		intrinsic RESET switching probability $P_{RESET}$ )
${\tt StdpLtp} \; [1000 \; {\tt TimePs}]$	Spike_RRAM	STDP LTP time window $T_{LTP}$
${\tt InhibitRefractory} \qquad [0$	Spike_RRAM	Neural lateral inhibition period $T_{inhibit}$
TimePs]		
${\tt EnableStdp} \ [1]$	Spike_RRAM	If false, STDP is disabled (no synaptic weight change)

RefractoryIntegration $[1]$	Spike_RRAM	If true, reset the integration to 0 during the refractory
DigitalIntegration $igl[0igr]$	Spike_RRAM	period  If false, the analog value of the devices is integrated, instead of their binary value

### ${\tt N2D2\ IP}\ only$

### 4.7.13 Rbf

Radial basis function fully connected layer.

Option [default value]	Description
NbOutputs	Number of output neurons
CentersFiller	Centers initial values filler
[NormalFiller(0.5, 0.05)]	
ScalingFiller	Scaling initial values filler
[NormalFiller(10.0, 0.05)]	

# Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
Solvers.*	all Frame	Any solver parameters
CentersSolver.*	all Frame	Centers solver parameters, take precedence over the
		Solvers.* parameters
${\tt ScalingSolver.}^{m{*}}$	all Frame	Scaling solver parameters, take precedence over the
		Solvers.* parameters
RbfApprox [None]	Frame	Approximation for the Gaussian function, can be any
		of: None, Rectangular Or SemiLinear

### 4.7.14 Softmax

Softmax layer.

Option [default value]	Description	
NbOutputs	Number of output neurons	
WithLoss $\left[0 ight]$	Softmax followed with a multinomial logistic layer	
GroupSize $[0]$	Softmax is applied on groups of outputs. The group size	
	must be a divisor of NbOutputs parameter.	

The softmax function performs the following operation, with  $a_{x,y}^i$  and  $b_{x,y}^i$  the input and the output respectively at position (x,y) on channel i:

$$b_{x,y}^{i} = \frac{\exp(a_{x,y}^{i})}{\sum\limits_{j=0}^{N} \exp(a_{x,y}^{j})}$$

and

$$da_{x,y}^i = \sum_{j=0}^N \left(\delta_{ij} - a_{x,y}^i\right) a_{x,y}^j db_{x,y}^j$$

When the WithLoss option is enabled, compute the gradient directly in respect of the cross-entropy loss:

$$L_{x,y} = \sum_{j=0}^{N} t_{x,y}^{j} \log(b_{x,y}^{j})$$

In this case, the gradient output becomes:

$$\mathrm{d}a_{x,y}^i = \mathrm{d}b_{x,y}^i$$

with

$$\mathrm{d}b_{x,y}^i = t_{x,y}^i - b_{x,y}^i$$

### 4.7.15 LRN

Local Response Normalization (LRN) layer.

Option [default value]	Description
NbOutputs	Number of output neurons

The response-normalized activity  $b_{x,y}^i$  is given by the expression:

$$b_{x,y}^{i} = \frac{a_{x,y}^{i}}{\left(k + \alpha \sum_{j=max(0,i-n/2)}^{min(N-1,i+n/2)} \left(a_{x,y}^{j}\right)^{2}\right)^{\beta}}$$

### Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
N [5]	all Frame	Normalization window width in elements
Alpha $[1.0\mathrm{e} ext{-}4]$	all Frame	Value of the alpha variance scaling parameter in the
		normalization formula
Beta $[0.75]$	all Frame	Value of the beta power parameter in the normalization
		formula
к [2.0]	all Frame	Value of the k parameter in normalization formula

### 4.7.16 LSTM

Long Short Term Memory Layer (Hochreiter and Schmidhuber, 1997).

Global layer parameters (Frame\_CUDA models)

Option [default value]	Description
SeqLength	Maximum sequence length that the LSTM can take as an
	input.
BatchSize	Number of sequences used for a single weights actualisation
	process: size of the batch.
InputDim	Dimension of every element composing a sequence.
HiddenSize	Dimension of the LSTM inner state and output.
SingleBackpropFeeding $\left[1 ight]$	If disabled return the full output sequence.
Bidirectional $\left[0 ight]$	If enabled, build a bidirectional structure.
AllGatesWeightsFiller	All Gates weights initial values filler.
AllGatesBiasFiller	All Gates bias initial values filler.
WeightsInputGateFiller	Input gate previous layer and recurrent weights initial values
	filler. Take precedence over AllGatesWeightsFiller parameter.
WeightsForgetGateFiller	Forget gate previous layer and recurrent weights initial values
	filler. Take precedence over AllGatesWeightsFiller parameter.
WeightsCellGateFiller	Cell gate (or new memory) previous layer and recurrent
	weights initial values filler. Take precedence over All-
	GatesWeightsFiller parameter.
WeightsOutputGateFiller	Output gate previous layer and recurrent weights initial
	values filler. Take precedence over AllGatesWeightsFiller
	parameter.
BiasInputGateFiller	Input gate previous layer and recurrent bias initial values
	filler. Take precedence over AllGatesBiasFiller parameter.
BiasRecurrentForgetGateFiller	Forget gate recurrent bias initial values filler. Take prece-
	dence over AllGatesBiasFiller parameter. Often set to 1.0 to
	show better convergence performance.
BiasPreviousLayerForgetGateFill	Forget gate previous layer bias initial values filler. Take
	precedence over AllGatesBiasFiller parameter.
BiasCellGateFiller	Cell gate (or new memory) previous layer and recurrent bias
	initial values filler. Take precedence over AllGatesBiasFiller
	parameter.
BiasOutputGateFiller	Output gate previous layer and recurrent bias initial values
	filler. Take precedence over AllGatesBiasFiller parameter.
HxFiller	Recurrent previous state initialisation. Often set to 0.0
CxFiller	Recurrent previous LSTM inner state initialisation. Often
	set to 0.0

# Configuration parameters (Frame\_CUDA models)

Option [default value]	Model(s)	Description
Solvers.*	all Frame	Any solver parameters
Dropout $[0.0]$	all Frame	The probability with which the value from input would
		be dropped.
InputMode []	all Frame	If enabled, drop the matrix multiplication of the input
		data.
Algo $[0]$	all Frame	Allow to choose different cuDNN implementation. Can
		be 0 : STANDARD, 1 : STATIC, 2 : DYNAMIC. Case
		1 and 2 aren't supported yet.

#### Current restrictions :

- Only Frame\_Cuda version is supported yet.
- The implementation only support input sequences with a fixed length associated with a single label.
- CuDNN structures requires the input data to be ordered as [1, InputDim, BatchSize, SeqLength]. Depending on the use case (like sequential-MNIST), the input data would need to be shuffled between the stimuli provisder and the RNN in order to process batches of data. No shuffling layer is yet operational. In that case, set batch to one for first experiments.

#### Further development requirements

When it comes to RNN, two main factors needs to be considered to build proper interfaces:

- 1. Whether the input data has a variable or a fixed length over the data base, that is to say whether the input data will have a variable or fixed Sequence length. Of course the main strength of a RNN is to process variable length data.
- 2. Labelling granularity of the input data, that is to say whether every elements of a sequence is labelled or the sequence itself has only one label.

For instance, let's consider sentences as sequences of words in which every word would be part of a vocabulary. Sentences could have a variable length and every element/word would have a label. In that case, every relevant element of the output sequence from the recurrent structure is turned into a prediction throught a fully connected layer with a linear activation fonction and a softmax. On the opposite, using sequential-MNIST database, the sequence length would be the same regarding every image and there is only one label for an image. In that case, the last element of the output sequence is the most relevant one to be turned into a prediction as it carries the information of the entire input sequence.

To provide flexibility according to these factors, the first implementation choice is to set a maximum sequence length emphSeqLength as an hyperparameter that the User provide. Variable length senquences can be processed by padding the remaining steps of the input sequence. Then two cases occur as the labeling granularity is scaled at each element of the sequence or scaled at the sequence itself:

- 1. The sequence itself has only one label: The model has a fixed size with one fully connected mapped to the relevant element of the output sequence according to the input sequence.
- 2. Every elements of a sequence is labelled:

The model has a fixed size with one big fully connected (or Tmax fully connected) mapped to the relevant elements of the output sequence according to the input sequence. The remaining elements need to be masked so it doesn't influence longer sequences.

#### Development guidance

• Replace the inner local variables of LSTMCell\_Frame\_Cuda with a generic layer of shuffling (on device) to enable the process of data batch.

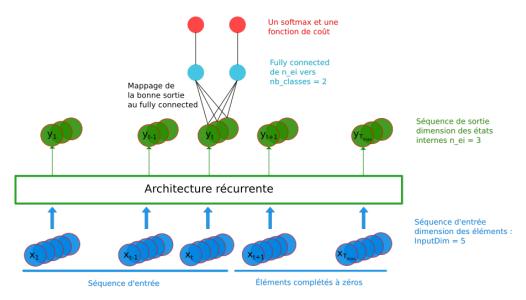


Figure 15: RNN model: variable sequence length and labeling scaled at the sequence

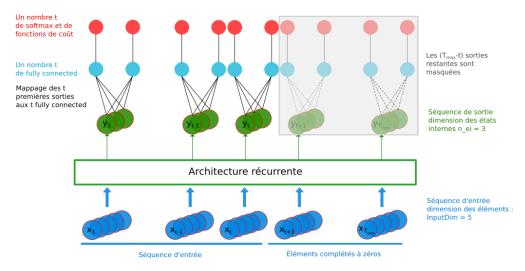


Figure 16: RNN model : variable sequence length and labeling scaled at each element of the sequence

- Develop some kind of label embedding within the layer to better articulate the labeling granularity of the input data.
- Adapt structures to support the STATIC and DYNAMIC algorithm of cuDNN functions.

### 4.7.17 Dropout

Dropout layer (Srivastava et al., 2012).

Option [default value]	Description
NbOutputs	Number of output neurons

### Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
Dropout $[0.5]$	all Frame	The probability with which the value from input would
		be dropped

68/83

# Configuration parameters

Option [default value]	Model(s)	Description
AlignCorners $[True]$	all Frame	Corner alignement mode if BilinearTF is used as inter-
		polation mode

# 4.7.20 BatchNorm

Batch Normalization layer (Ioffe and Szegedy, 2015).

Option [default value]	Description	
NbOutputs	Number of output neurons	
ActivationFunction [Tanh]	Activation function. Can be any of Logistic, LogisticWithLoss,	
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
ScalesSharing []	Share the scales with an other layer	
BiasesSharing []	Share the biases with an other layer	
MeansSharing []	Share the means with an other layer	
VariancesSharing []	Share the variances with an other layer	

# Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
Solvers.*	all Frame	Any solver parameters
ScaleSolver.*	all Frame	Scale solver parameters, take precedence over the
		Solvers.* parameters
BiasSolver.*	all Frame	Bias solver parameters, take precedence over the
		Solvers.* parameters
Epsilon $[0.0]$	all Frame	Epsilon value used in the batch normalization formula.
		If 0.0, automatically choose the minimum possible
		value.
MovingAverageMomentum	all Frame	MovingAverageMomentum: used for the moving av-
[0.1]		erage of batch-wise means and standard deviations
		during training. The closer to 1.0, the more it will
		depend on the last batch.

### 4.7.21 Transformation

Transformation layer, which can apply any transformation described in 4.6.1. Useful for fully CNN post-processing for example.

Option [default value]	Description	
NbOutputs	Number of outputs	
Transformation	Name of the transformation to apply	

The Transformation options must be placed in the same section. Usage example for fully CNNs:

```
[post.Transformation-thres]
Input=...; for example, network's logistic of softmax output layer
NbOutputs=1
Type=Transformation
Transformation=ThresholdTransformation
Operation=ToZero
Threshold=0.75

[post.Transformation-morpho]
Input=post.Transformation-thres
NbOutputs=1
Type=Transformation
Transformation=MorphologyTransformation
Operation=Opening
Size=3
```

### 5 Tutorials

### 5.1 Learning deep neural networks: tips and tricks

#### 5.1.1 Choose the learning solver

Generally, you should use the SGD solver with a momentum (typical value for the momentum: 0.9). It generalizes better, often significantly better, than adaptive methods like Adam (Wilson et al., 2017).

Adaptive solvers, like Adam, may be used for fast exploration and prototyping, thanks to their fast convergence.

#### 5.1.2 Choose the learning hyper-parameters

You can use the <code>-find-lr</code> option available in the <code>n2d2</code> executable to automatically find the best learning rate for a given neural network.

Usage example:

./n2d2 model.ini -find-lr 10000

This command starts from a very low learning rate (1.0e-6) and increase it exponentially to reach the maximum value (10.0) after 10000 steps, as shown in figure 17. The loss change during this phase is then plotted in function of the learning rate, as shown in figure 18.

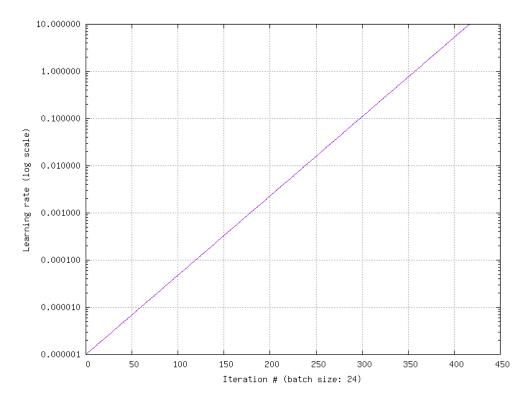


Figure 17: Exponential increase of the learning rate over the specified number of iterations, equals to the number of steps divided by the batch size (here: 24).

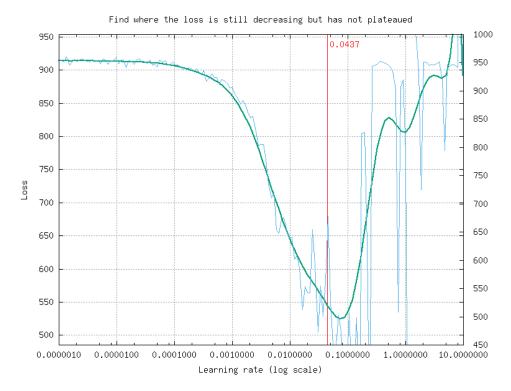


Figure 18: Loss change as a function of the learning rate.

Note that in N2D2, the learning rate is automatically normalized by the global batch size  $(N \times \texttt{IterationSize})$  for the SGDSolver. A simple linear scaling rule is used, as recommanded in (Goyal et al., 2017). The effective learning rate  $\alpha_{\text{eff}}$  applied for parameters update is therefore:

$$\alpha_{\rm eff} = \frac{\alpha}{N \times {\tt IterationSize}} \ {\rm with} \ \alpha = {\tt LearningRate}$$

Typical values for the SGDSolver are:

Solvers.LearningRate=0.01 Solvers.Decay=0.0001 Solvers.Momentum=0.9

### 5.1.3 Convergence and normalization

Deep networks (> 30 layers) and especially residual networks usually don't converge without normalization. Indeed, batch normalization is almost always used. *ZeroInit* is a method that can be used to overcome this issue without normalization (Zhang et al., 2019).

### 5.2 Building a classifier neural network

For this tutorial, we will use the classical MNIST handwritten digit dataset. A driver module already exists for this dataset, named MNIST\_IDX\_Database.

To instantiate it, just add the following lines in a new INI file:

```
[database]
Type=MNIST_IDX_Database
Validation=0.2; Use 20% of the dataset for validation
```

In order to create a neural network, we first need to define its input, which is declared with a [sp] section (sp for StimuliProvider). In this section, we configure the size of the input and the batch size:

```
[sp]
SizeX=32
SizeY=32
BatchSize=128
```

We can also add pre-processing transformations to the *StimuliProvider*, knowing that the final data size after transformations must match the size declared in the [sp] section. Here, we must rescale the MNIST 28x28 images to match the 32x32 network input size.

```
[sp.Transformation_1]
Type=RescaleTransformation
Width=[sp]SizeX
Height=[sp]SizeY
```

Next, we declare the neural network layers. In this example, we reproduced the well-known LeNet network. The first layer is a 5x5 convolutional layer, with 6 channels. Since there is only one input channel, there will be only 6 convolution kernels in this layer.

```
[conv1]
Input=sp
Type=Conv
KernelWidth=5
KernelHeight=5
NbOutputs=6
```

The next layer is a 2x2 MAX pooling layer, with a stride of 2 (non-overlapping MAX pooling).

```
[pool1]
Input=conv1
Type=Pool
PoolWidth=2
PoolHeight=2
NbOutputs=[conv1]NbOutputs
Stride=2
Pooling=Max
Mapping.Size=1; One to one connection between input and output channels
```

The next layer is a 5x5 convolutional layer with 16 channels.

```
[conv2]
Input=pool1
Type=Conv
KernelWidth=5
KernelHeight=5
NbOutputs=16
```

Note that in LeNet, the [conv2] layer is not fully connected to the pooling layer. In N2D2, a custom mapping can be defined for each input connection. The connection of n-th output map to the inputs is defined by the n-th column of the matrix below, where the rows correspond to the inputs.

```
Mapping(pool1)=\
1 0 0 0 1 1 1 0 0 1 1 1 1 0 1 1 \
1 1 0 0 0 1 1 1 0 0 1 1 1 1 0 1 \
1 1 0 0 0 1 1 1 0 0 1 0 1 1 1 \
1 1 1 0 0 0 1 1 1 0 0 1 0 1 1 1 \
0 1 1 1 0 0 1 1 1 1 0 0 1 0 1 1 \
0 0 1 1 1 0 0 1 1 1 1 0 1 0 1 \
0 0 0 1 1 1 0 0 1 1 1 1 0 1 1 1
```

Another MAX pooling and convolution layer follow:

```
[pool2]
Input=conv2
Type=Pool
PoolWidth=2
PoolHeight=2
```

```
NbOutputs=[conv2]NbOutputs
Stride=2
Pooling=Max
Mapping.Size=1

[conv3]
Input=pool2
Type=Conv
KernelWidth=5
KernelHeight=5
NbOutputs=120
```

The network is composed of two fully-connected layers of 84 and 10 neurons respectively:

```
[fc1]
Input=conv3
Type=Fc
NbOutputs=84

[fc2]
Input=fc1
Type=Fc
NbOutputs=10
```

Finally, we use a softmax layer to obtain output classification probabilities and compute the loss function.

```
[softmax]
Input=fc2
Type=Softmax
NbOutputs=[fc2]NbOutputs
WithLoss=1
```

In order to tell N2D2 to compute the error and the classification score on this softmax layer, one must attach a N2D2 *Target* to this layer, with a section with the same name suffixed with .Target:

```
[softmax.Target]
```

By default, the activation function for the convolution and the fully-connected layers is the hyperbolic tangent. Because the [fc2] layer is fed to a softmax, it should not have any activation function. We can specify it by adding the following line in the [fc2] section:

```
[fc2]
...
ActivationFunction=Linear
```

In order to improve further the networks performances, several things can be done:

• Use ReLU activation functions. In order to do so, just add the following in the [conv1], [conv2], [conv3] and [fc1] layer sections:

```
ActivationFunction=Rectifier
```

For the ReLU activation function to be effective, the weights must be initialized carefully, in order to avoid dead units that would be stuck in the  $]-\infty,0]$  output range before the ReLU function. In N2D2, one can use a custom WeightsFiller for the weights initialization. For the ReLU activation function, a popular and efficient filler is the so-called XavierFiller (see the 4.7.2 section for more information):

```
WeightsFiller=XavierFiller
```

• Use dropout layers. Dropout is highly effective to improve the network generalization capacity. Here is an example of a dropout layer inserted between the [fc1] and [fc2] layers:

```
[fc1]
...
```

```
[fc1.drop]
Input=fc1
Type=Dropout
NbOutputs=[fc1]NbOutputs

[fc2]
Input=fc1.drop ; Replaces "Input=fc1"
...
```

• Tune the learning parameters. You may want to tune the learning rate and other learning parameters depending on the learning problem at hand. In order to do so, you can add a configuration section that can be common (or not) to all the layers. Here is an example of configuration section:

```
[conv1]
...
ConfigSection=common.config

[...]
...
[common.config]
NoBias=1
WeightsSolver.LearningRate=0.05
WeightsSolver.Decay=0.0005
Solvers.LearningRatePolicy=StepDecay
Solvers.LearningRateStepSize=[sp]_EpochSize
Solvers.LearningRateDecay=0.993
Solvers.Clamping=-1.0:1.0
```

For more details on the configuration parameters for the Solver, see section 4.7.3.

• Add input distortion. See for example the DistortionTransformation (section 4.6.1).

The complete INI model corresponding to this tutorial can be found in *models/LeNet.ini*. In order to use CUDA/GPU accelerated learning, the default layer model should be switched to Frame\_CUDA. You can enable this model by adding the following line at the top of the INI file (before the first section):

DefaultModel=Frame\_CUDA

### 5.3 Building a segmentation neural network

In this tutorial, we will learn how to do image segmentation with N2D2. As an example, we will implement a face detection and gender recognition neural network, using the IMDB-WIKI dataset. First, we need to instanciate the IMDB-WIKI dataset built-in N2D2 driver:

```
[database]
Type=IMDBWIKI_Database
WikiSet=1 ; Use the WIKI part of the dataset
IMDBSet=0 ; Don't use the IMDB part (less accurate annotation)
Learn=0.90
Validation=0.05
DefaultLabel=background ; Label for pixels outside any ROI (default is no label, pixels are ignored)
```

We must specify a default label for the background, because we want to learn to differenciate faces from the background (and not simply ignore the background for the learning).

The network input is then declared:

```
[sp]
SizeX=480
```

```
SizeY=360
BatchSize=48
CompositeStimuli=1
```

In order to work with segmented data, i.e. data with bounding box annotations or pixel-wise annotations (as opposed to a single label per data), one must enable the CompositeStimuli option in the [sp] section.

We can then perform various operations on the data before feeding it to the network, like for example converting the 3-channels RGB input images to single-channel gray images:

```
[sp.Transformation-1]
Type=ChannelExtractionTransformation
CSChannel=Gray
```

We must only rescale the images to match the networks input size. This can be done using a RescaleTransformation, followed by a PadCropTransformation if one want to keep the images aspect ratio.

```
[sp.Transformation-2]
Type=RescaleTransformation
Width=[sp]SizeX
Height=[sp]SizeY
KeepAspectRatio=1 ; Keep images aspect ratio

; Required to ensure all the images are the same size
[sp.Transformation-3]
Type=PadCropTransformation
Width=[sp]SizeX
Height=[sp]SizeY
```

A common additional operation to extend the learning set is to apply random horizontal mirror to images. This can be achieved with the following FlipTransformation:

```
[sp.OnTheFlyTransformation-4]
Type=FlipTransformation
RandomHorizontalFlip=1
ApplyTo=LearnOnly; Apply this transformation only on the learning set
```

Note that this is an *on-the-fly* transformation, meaning it cannot be cached and is re-executed every time even for the same stimuli. We also apply this transformation only on the learning set, with the ApplyTo option.

Next, the neural network can be described:

```
[conv1.1]
Input=sp
Type=Conv
...

[pool1]
...
[...]
...

[fc2]
Input=drop1
Type=Conv
...

[drop2]
Input=fc2
Type=Dropout
NbOutputs=[fc2] NbOutputs
```

A full network description can be found in the *IMDBWIKI.ini* file in the *models* directory of N2D2. It is a fully-CNN network.

Here we will focus on the output layers required to detect the faces and classify their gender. We start from the [drop2] layer, which has 128 channels of size 60x45.

#### 5.3.1 Faces detection

We want to first add an output stage for the faces detection. It is a 1x1 convolutional layer with a single 60x45 output map. For each output pixel, this layer outputs the probability that the pixel belongs to a face.

```
[fc3.face]
Input=drop2
Type=Conv
KernelWidth=1
KernelHeight=1
NbOutputs=1
Stride=1
ActivationFunction=LogisticWithLoss
WeightsFiller=XavierFiller
ConfigSection=common.config; Same solver options that the other layers
```

In order to do so, the activation function of this layer must be of type LogisticWithLoss.

We must also tell N2D2 to compute the error and the classification score on this softmax layer, by attaching a N2D2 Target to this layer, with a section with the same name suffixed with .Target:

```
[fc3.face.Target]
LabelsMapping=${N2D2_MODELS}/IMDBWIKI_target_face.dat
; Visualization parameters
NoDisplayLabel=0
LabelsHueOffset=90
```

In this *Target*, we must specify how the dataset annotations are mapped to the layer's output. This can be done in a separate file using the LabelsMapping parameter. Here, since the output layer has a single output per pixel, the target value can only be 0 or 1. A target value of -1 means that this output is ignored (no error back-propagated). Since the only annotations in the IMDB-WIKI dataset are faces, the mapping described in the *IMDBWIKI\_target\_face.dat* file is easy:

```
# background
background 0
# padding (*) is ignored (-1)
* -1
# not background = face
default 1
```

#### 5.3.2 Gender recognition

We can also add a second output stage for gender recognition. Like before, it would be a 1x1 convolutional layer with a single 60x45 output map. But here, for each output pixel, this layer would output the probability that the pixel represents a female face.

```
[fc3.gender]
Input=drop2
Type=Conv
KernelWidth=1
KernelHeight=1
NbOutputs=1
Stride=1
ActivationFunction=LogisticWithLoss
WeightsFiller=XavierFiller
```

#### ConfigSection=common.config

The output layer is therefore identical to the face's output layer, but the target mapping is different. For the target mapping, the idea is simply to ignore all pixels not belonging to a face and affect the target 0 to male pixels and the target 1 to female pixels.

```
[fc3.gender.Target]
LabelsMapping=${N2D2_MODELS}/IMDBWIKI_target_gender.dat
; Only display gender probability for pixels detected as face pixels
MaskLabelTarget=fc3.face.Target
MaskedLabel=1
```

The content of the IMDBWIKI target gender.dat file would therefore look like:

```
# background
# ?-* (unknown gender)
# padding
default -1
# male gender
M-? 0 # unknown age
M-0 0
M-1 0
M-20
M-98 0
M-99 0
# female gender
F-? 1 # unknown age
F-0 1
F-1 1
F-2 1
F-98 1
F-99 1
```

#### 5.3.3 ROIs extraction

The next step would be to extract detected face ROIs and assign for each ROI the most probable gender. To this end, we can first set a detection threshold, in terms of probability, to select face pixels. In the following, the threshold is fixed to 75% face probability:

```
[post.Transformation-thres]
Input=fc3.face
Type=Transformation
NbOutputs=1
Transformation=ThresholdTransformation
Operation=ToZero
Threshold=0.75
```

We can then assign a target of type TargetROIs to this layer that will automatically create the bounding box using a segmentation algorithm.

```
[post.Transformation-thres.Target-face]
Type=TargetROIs
MinOverlap=0.33; Min. overlap fraction to match the ROI to an annotation
FilterMinWidth=5; Min. ROI width
FilterMinHeight=5; Min. ROI height
FilterMinAspectRatio=0.5; Min. ROI aspect ratio
FilterMaxAspectRatio=1.5; Max. ROI aspect ratio
LabelsMapping=${N2D2_MODELS}/IMDBWIKI_target_face.dat
```

In order to assign a gender to the extracted ROIs, the above target must be modified to:

```
[post.Transformation-thres.Target-gender]
Type=TargetR0Is
R0IsLabelTarget=fc3.gender.Target
Min0verlap=0.33
FilterMinWidth=5
FilterMinHeight=5
FilterMinAspectRatio=0.5
FilterMaxAspectRatio=1.5
LabelsMapping=${N2D2_MODELS}/IMDBWIKI_target_gender.dat
```

Here, we use the fc3.gender.Target target to determine the most probable gender of the ROI.

#### 5.3.4 Data visualization

For each *Target* in the network, a corresponding folder is created in the simulation directory, which contains learning, validation and test confusion matrixes. The output estimation of the network for each stimulus is also generated automatically for the test dataset and can be visualized with the ./test.py helper tool. An example is shown in figure 19.

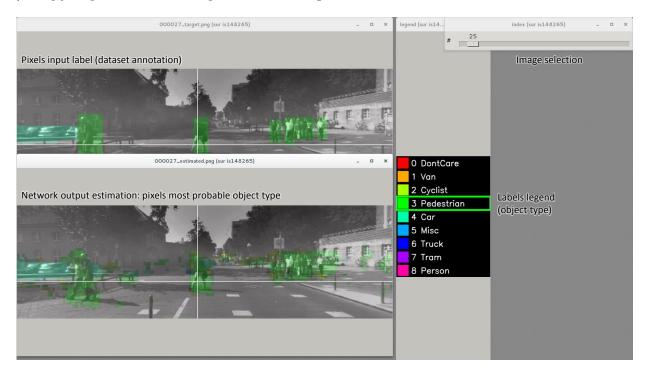


Figure 19: Example of the target visualization helper tool.

#### 5.4 Transcoding a learned network in spike-coding

N2D2 embeds an event-based simulator (historically known as 'Xnet') and allows to transcode a whole DNN in a spike-coding version and evaluate the resulting spiking neural network performances. In this tutorial, we will transcode the LeNet network described in section 5.2.

#### 5.4.1 Render the network compatible with spike simulations

The first step is to specify that we want to use a transcode model (allowing both formal and spike simulation of the same network), by changing the DefaultModel to:

#### DefaultModel=Transcode\_CUDA

In order to perform spike simulations, the input of the network must be of type *Environment*, which is a derived class of *StimuliProvider* that adds spike coding support. In the INI model file, it

is therefore necessary to replace the [sp] section by an [env] section and replace all references of sp to env.

Note that these changes have at this point no impact at all on the formal coding simulations. The beginning of the INI file should be:

```
DefaultModel=Transcode_CUDA
; Database
[database]
Type=MNIST_IDX_Database
Validation=0.2 ; Use 20% of the dataset for validation

; Environment
[env]
SizeX=32
SizeY=32
BatchSize=128

[env.Transformation_1]
Type=RescaleTransformation
Width=[env]SizeX
Height=[env]SizeY
[conv1]
Input=env
...
```

The dropout layer has no equivalence in spike-coding inference and must be removed:

```
...

[fc1.drop]

Input=fc1

Type=Dropout

NbOutputs=[fc1]NbOutputs

[fc2]

Input=fc1.drop

...
```

The softmax layer has no equivalence in spike-coding inference and must be removed as well. The *Target* must therefore be attached to [fc2]:

```
[softmax]
Input=fc2
Type=Softmax
NbOutputs=[fc2]NbOutputs
WithLoss=1
[softmax.Target]
[fc2.Target]
```

The network is now compatible with spike-coding simulations. However, we did not specify at this point how to translate the input stimuli data into spikes, nor the spiking neuron parameters (threshold value, leak time constant...).

#### 5.4.2 Configure spike-coding parameters

The first step is to configure how the input stimuli data must be coded into spikes. To this end, we must attach a configuration section to the *Environment*. Here, we specify a periodic coding with random initial jitter with a minimum period of 10 ns and a maximum period of 100 us:

```
[env]
...
ConfigSection=env.config

[env.config]
; Spike-based computing
StimulusType=JitteredPeriodic
PeriodMin=1,000,000 ; unit = fs
PeriodMeanMin=10,000,000 ; unit = fs
PeriodMeanMax=100,000,000,000 ; unit = fs
PeriodRelStdDev=0.0
```

The next step is to specify the neurons parameters, that will be common to all layers and can therefore be specified in the [common.config] section. In N2D2, the base spike-coding layers use a Leaky Integrate-and-Fire (LIF) neuron model. By default, the leak time constant is zero, resulting to simple Integrate-and-Fire (IF) neurons.

Here we simply specify that the neurons threshold must be the unity, that the threshold is only positive and that there is no incoming synaptic delay:

```
[common.config]
...
; Spike-based computing
Threshold=1.0
BipolarThreshold=0
IncomingDelay=0
```

Finally, we can limit the number of spikes required for the computation of each stimulus by adding a decision delta threshold at the output layer:

```
[fc2]
...
ConfigSection=common.config,fc2.config

[fc2.Target]

[fc2.config]
; Spike-based computing
TerminateDelta=4
BipolarThreshold=1
```

The complete INI model corresponding to this tutorial can be found in *models/LeNet\_Spike.ini*. Here is a summary of the steps required to reproduce the whole experiment:

```
./n2d2 "$N2D2_MODELS/LeNet.ini" -learn 6000000 -log 100000
./n2d2 "$N2D2_MODELS/LeNet_Spike.ini" -test
```

The final recognition rate reported at the end of the spike inference should be almost identical to the formal coding network (around 99% for the LeNet network).

Various statistics are available at the end of the spike-coding simulation in the  $stats\_spike$  folder and the  $stats\_spike.log$  file. Looking in the  $stats\_spike.log$  file, one can read the following line towards the end of the file:

```
Read events per virtual synapse per pattern (average): 0.654124
```

This line reports the average number of accumulation operations per synapse per input stimulus in the network. If this number if below 1.0, it means that the spiking version of the network is more efficient than its formal counterpart in terms of total number of operations!

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