



Neural Network Design & Deployment

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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 IP¹.

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
```

¹Ongoing 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® cuDNN²), 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
```

²On future GPUs

```
KernelWidth=3
KernelHeight=3
NbChannels=32
Stride=1
; Second layer: MAX pooling with pooling area 2x2
Input=conv1
Type=Pool
Pooling=Max
PoolWidth=2
PoolHeight=2
NbChannels=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);

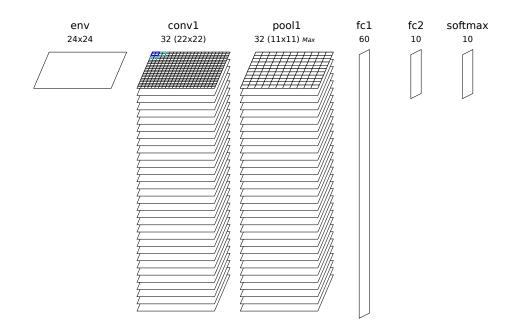


Figure 1: Automatically generated and ready to learn DNN from the INI configuration file example.

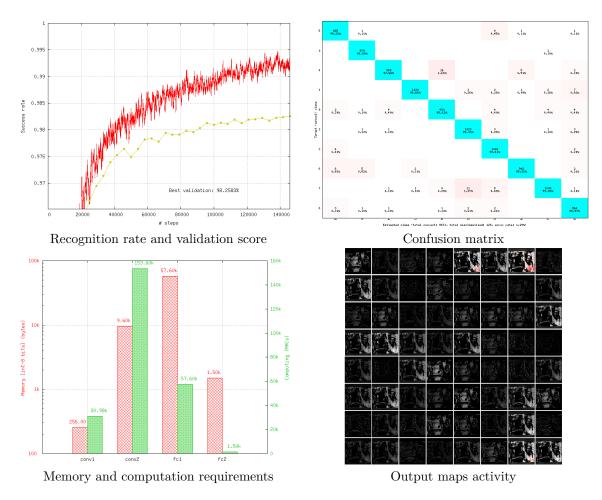


Figure 2: Example of information automatically generated by the software during and after learning.

- 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.

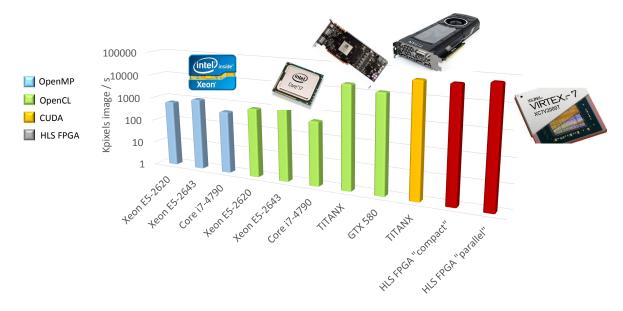


Figure 3: Example of network benchmarking on different hardware targets.

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 12 (2013) 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
- \bullet At least 150 MB available hard disk space + 350 MB for MNIST dataset processing

For CUDA acceleration:

- CUDA ≥ 6.5 and CuDNN ≥ 1.0
- NVIDIA GPU with CUDA compute capability ≥ 3 (starting from Kepler micro-architecture)
- 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
- opency
- 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:

```
rpm -Uhv http://developer.download.nvidia.com/compute/cuda/repos/rhel6/x86_64/cuda-repo-
    rhel6-7.5-18.x86_64.rpm
yum clean expire-cache
yum install cuda
```

- 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 ≥ 12 (2013) 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 12"
```

Then open the newly created N2D2 project in Visual Studio 12 (2013). 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;
- $\bullet \quad TargetScore_*/ConfusionMatrix_test.png.$

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.

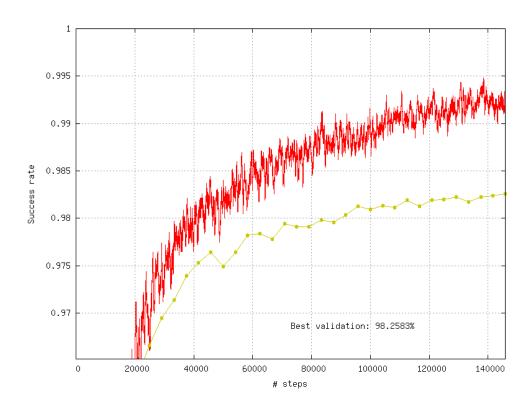


Figure 4: Recognition rate and validation score during learning.



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.

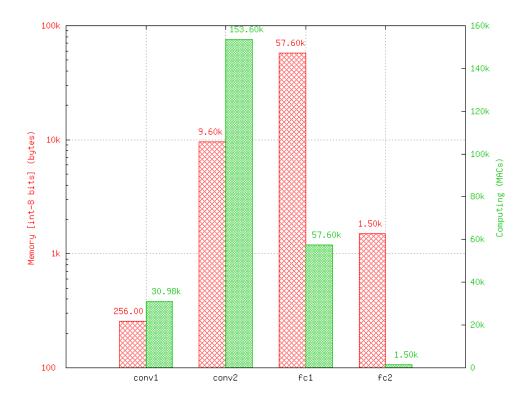


Figure 6: Example of memory and computation requirements of the network.

3.7 Export a learned 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
-uenv	If present, treat the input stimuli data as unsigned
-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)

N2D2 IP only

3.7.1 C export

Test the exported network:

cd export_C_int8
make

./bin/n2d2_test

The result should look like:

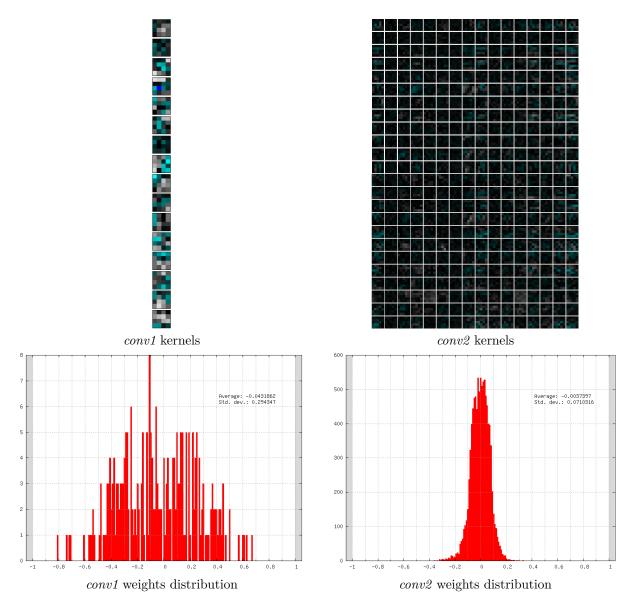


Figure 7: Example of kernels and weights distribution analysis for two convolutional layers.

```
\begin{array}{lll} \dots \\ 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) \end{array}
```

$Confusion\ matrix:$

/ T	\ E /	0	/	1	/	2	/	3 /
/	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% /



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

 $T{:}\ Target \ E{:}\ Estimated$

N2D2 IP only

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
./bin/n2d2_opencl_test -gpu
```

3.7.3 CPP_TensorRT export

The tensorRT 2.1 API export can run the generated program in NVIDIA GPU architecture. It use CUDA and tensorRT 2.1 API library. The currently supported layers by the tensorRT 2.1 export are: Convolutional, Pooling, Concatenation, Fully-Connected, Softmax and all activations type. Custom layers implementation through the plugin factory and generic 8-bits calibrations inference features are under development.

Program options related to the tensorRT 2.1 API 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.
-prof	Activates the layer wise profiling mechanism. This option
	can decrease execution time performance.
-iter-build $[1]$	Sets the number of minimization build iterations done by
	the tensorRT builder to find the best layer tactics.

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\%)
  ******* POOL2: 0.00616047 ms
(05\%)
  (14\%)
  (19\%)
  ****** FC2: 0.0149013 ms
(13\%)
  Average profiled tensor RT 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

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3.7.5 C_HLS export

Test the exported network:

cd export_C_HLS_int8
make

 $./{\tt bin/n2d2_test}$

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

vivado_hls -f run_hls.tcl

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
NbChannels={{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
NbChannels=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 {%for VAR in range([START,]END])%} ... {%endfor %}

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

include {%include FILENAME %}

4.3 Global parameters

Option [default value]	Description
DefaultModel [Transcode]	Default layers model. Can be Frame, Frame_CUDA, Transcode or
	Spike
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
$[\$N2D2_DATA/GTSRB]$	

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 $\left[0 ight]$	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 $\begin{bmatrix} P_1 & P_2 & P_3 \end{bmatrix}$
	(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 $\left[1 ight]$	If true, the stimuli are equi-partitioned for the learn/valida-
	tion/test sets, meaning that the same number of stimuli for
[0.0]	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,
_ [10]	regardless of their labels
Test $[1.0 ext{-Learn-Validation}]$	If PerLabelPartitioning is true, fraction of images used for the
	test; else, number of images used for the test, regardless of their labels
W-1: AP-t	List of space-separated valid stimulus file extensions (if left
ValidExtensions	empty, any file extension is considered a valid stimulus)
LoadMore [Name of an other section with the same options to load a
Loadhore []	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,
u u	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
```

4.4.4 Other built-in databases

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 $[0.0]$	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 $[0.0]$	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 $[0.0]$	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/	
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 dbExtract.m 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
igspace SingleLabel [1]	Use the same label for "person" and "people" bounding box
IncAmbiguous $[0]$	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]	

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

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)
$[\$N2D2_DATA/FDDB]$	
LabelPath	Path to the annotations (decompressed FDDB-folds.tgz)
[\$N2D2_DATA/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]	

KITTY_Database KITTY Database.

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

KITTY_Road_Database KITTY Road Database. The KITTY Road Database provide ROI which can be used to road segmentation.

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

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.

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
```

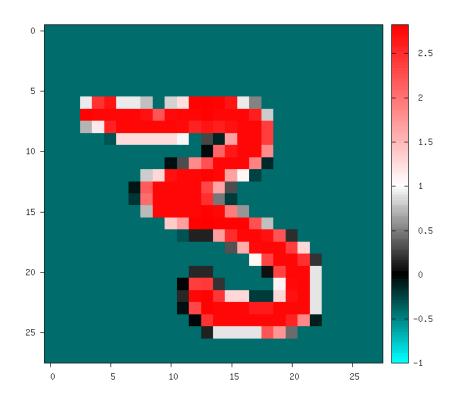


Figure 9: Image of the set after normalization.

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

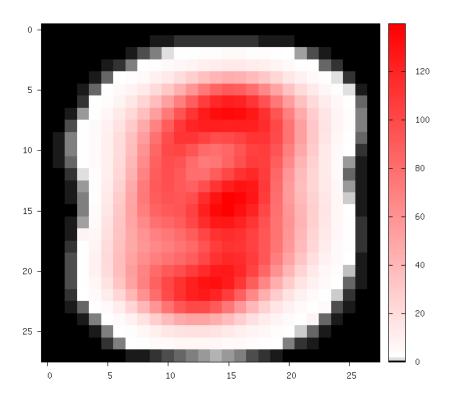


Figure 10: Global mean image generated by StimuliData with the MeanData parameter enabled.

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.

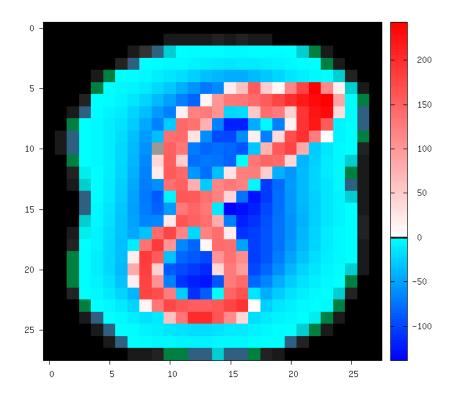


Figure 11: Image of the set after the AffineTransformation substracting the global mean image (keep in mind that the original image value range is [0, 255]).

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 $[0]$	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)
$\texttt{PeriodMeanMax} \; [12 \; \texttt{TimeS}]$	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 $[11\ \mathtt{TimeMs}]$	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

I	
	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.Sigma $[0.4]$	Sigma

Blackman window Blackman window.

Option [default value]	Description
WindowName.Alpha $[0.16]$	Alpha

Kaiser window Kaiser window.

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

$\textbf{ChannelExtractionTransformation} \quad \text{Extract an image channel}.$

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
	сь: 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: if the image is in grayscale, convert it in BGR
	HSV
	HLS
	YCrCb
	CIELab
	CIELuv

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 $\left[1\right]$	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 $[15]$	Size of the gaussian for elastic distortion (in pixels)
ElasticSigma $\left[6.0\right]$	Sigma of the gaussian for elastic distortion
ElasticScaling $[0.0]$	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 $[0]$	Width of the kernel (numer of columns)
Kernel.SizeY $[0]$	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 $\left[2.0\right]$	Sigma1 of the kernel
Kernel.Sigma2 $\left[1.0\right]$	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 $\left[0.5\right]$	Gamma of the kernel

FlipTransformation Image flip transformation.

Option [default value]	Description
HorizontalFlip $\left[0 ight]$	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

N2D2 IP only GradientFilterTransformation Compute image 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 $\left[0\right]$	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)
GradientScale $\left[1.0\right]$	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

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

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
ApplyToLabels $\left[0 ight]$	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

$\label{thm:normalize} \textbf{Normalize Transformation} \quad \text{Normalize the image}.$

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

PadCropTransformation Pad/crop the image to a specified 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

N2D2 IP only RandomAffineTransformation Apply a global random affine transformation to the values of the image.

Option [default value]	Description
GainVar	Random gain is in range ±GainVar
BiasVar $\left[0.0 ight]$	Random bias is in range ±BiasVar

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:

$$f(x) = (x \stackrel{o}{op_{1st}} val_{1st}) \stackrel{o}{op_{2nd}} val_{2nd}$$

Ponly 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)

 $\label{lem:RescaleTransformation} \ \ Rescale \ the \ image \ to \ a \ specified \ size.$

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 $[1]$	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 $(0 = \text{no change})$

${\tt N2D2\ IP}\ {\it only}\ \ {\tt SliceExtractionTransformation}\ \ {\tt Extract}\ a\ slice\ from\ an\ image.$

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
$\texttt{RandomOffsetX} \; [0]$	If true, the X offset is chosen randomly
RandomOffsetY $\left[0\right]$	If true, the Y offset is chosen randomly
AllowPadding $\left[0 ight]$	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
$\texttt{MaxValue} \; [1.0]$	Max. value to use with Binary and BinaryInverted operations

$\label{thm:thm:mage} \textbf{TrimTransformation} \quad \text{Trim the image}.$

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 $[0.0]$	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
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 12.

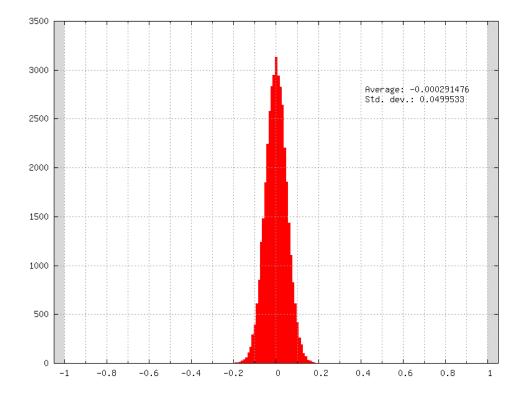


Figure 12: Initial weights distribution of a layer using a normal distribution (NormalFiller) with a 0 mean and a 0.05 standard deviation.

ConstantFiller Fill with a constant value.

Option	Description
FillerName. Value	Value for the filling

NormalFiller Fill with a normal distribution.

Option [default value]	Description
FillerName.Mean $[0.0]$	Mean value of the distribution
FillerName . 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]	

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

- n=fan-in with FamIm, resulting in $Var(W)=\frac{1}{fan\text{-}in}$
- $n = \frac{(fan\text{-}in + fan\text{-}out)}{2}$ with Average, resulting in $Var(W) = \frac{2}{fan\text{-}in + fan\text{-}out}$
- 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 $[1]$	
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.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 α_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 α_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 α_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 α_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.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
SolverName.	Learning rate step size (in number of stimuli)
LearningRateStepSize $\left[1 ight]$	
SolverName . LearningRateDecay	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.

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 Logistic activation function.

Logistic With Loss activation function.

Rectifier or ReLU activation function.

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

Saturation Saturation activation function.

Softplus Softplus activation function.

Tanh Tanh activation function.

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

Option [default value]	Description
ActivationFunction.Alpha $\left[1.0 ight]$	α parameter

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

4.7.5 Anchor

Anchor layer for Faster R-CNN.

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)	

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 $\left[0.3\right]$	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 λ
LossPositiveSample $\left[128\right]$	all Frame	Number of random positive samples for the loss com-
		putation
LossNegativeSample $[128]$	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
NbChannels=18
[scores.softmax]
Input=scores
Type=Softmax
NbOutputs=[scores] NbChannels
WithLoss=1
; RPN network: coordinates layer
[coordinates]
Input=...
Type=Conv
KernelWidth=1
KernelHeight=1
; 36 channels for 4 coordinates x 9 anchors
NbChannels=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
```

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

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
NbChannels=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[1].h

Convolutional layer.

Option [default value]	Description	
KernelWidth	Width of the kernel	
KernelHeight	Height of the kernel	
NbChannels	Number of output channels	
SubSampleX [1]	X-axis subsampling factor of the output feature maps	
SubSampleY [1]	Y-axis subsampling factor of the output feature maps	
SubSample $[1]$	Subsampling factor of the output feature maps	
	(mutually exclusive with SubSampleX and SubSampleY)	
StrideX $[1]$	X-axis stride of the kernels	
StrideY $[1]$	Y-axis stride of the kernels	
Stride $[1]$	Stride of the kernels	
	(mutually exclusive with StrideX and StrideY)	
PaddingX $[0]$	X-axis input padding	
PaddingY [0]	Y-axis input padding	
Padding $[0]$	Input padding	
	(mutually exclusive with Padding% and Padding%)	

Rectifier, Softplus, TanhLeCun, Linear, Saturation or Tanh WeightsFiller Weights initial values filler
NormalFiller(0.0, 0.05) BiasFiller
BiasFiller [NormalFiller(0.0, 0.05)] Mapping.SizeX [1] Mapping.SizeY [1] Mapping.SizeY [1] Mapping.Size [1] Mapping.StrideX [1] Mapping.StrideX [1] Mapping.StrideY [1] Mapping.StrideY [1] Mapping.StrideY [1] Mapping.StrideY [1] Mapping.StrideY [1] Mapping.Stride [1] Mapping.Stride [1] Mapping.Stride [1] Mapping.Stride [1] Mapping.OffsetX [0] Mapping.OffsetX [0] Mapping.OffsetY [0] Mapping.Offset [0] Mapping.Offset [0] Mapping.Offset [0] Mapping.NbIterations [0] Mapping canvas pattern default width height Mapping.OffsetX and Mapping.StrideY Mapping.OffsetX and Mapping.StrideY Mapping.Offset [0] Mapping canvas default Y-axis offset Mapping.Offset [0] Mapping.Offset [0] Mapping canvas default offset (mutually exclusive with Mapping.OffsetX and Mapping.OffsetY) 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
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Mapping(in).StrideY)
Mapping(in).OffsetX [0] Mapping canvas default X-axis offset for input layer in
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Mapping (in).Offset [0] Mapping canvas default offset for input layer in
(mutually exclusive with Mapping(in).OffsetX and
Mapping(in).OffsetY)
Mapping(in).NbIterations [0] 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 $[1]$	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, with O 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)
$\begin{tabular}{ll} Weights Export Transpose \\ [0] \end{tabular}$	all Frame	If true, import/export transposed 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 $\left[0.0\right]$	Spike, Spike_RRAM	Neural leak time constant τ_{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 $\left[1;0.1\right]$	Spike_RRAM	Mean minimum synaptic weight w_{min}
WeightsMaxMean $[100;10.0]$	Spike_RRAM	Mean maximum synaptic weight w_{max}
WeightsMinVarSlope $\left[0.0 ight]$	Spike_RRAM	OXRAM specific parameter
${\tt WeightsMinVarOrigin} \; [0.0]$	Spike_RRAM	OXRAM specific parameter
${\tt WeightsMaxVarSlope} \; [0.0]$	Spike_RRAM	OXRAM specific parameter
WeightsMaxVarOrigin $\left[0.0\right]$	Spike_RRAM	OXRAM specific parameter
WeightsSetProba $\left[1.0\right]$	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 $\left[1\right]$	Spike_RRAM	Synaptic redundancy (number of RRAM device per
		synapse)
BipolarWeights $\left[0 ight]$	Spike_RRAM	Bipolar weights
${\tt BipolarIntegration} \; [0]$	Spike_RRAM	Bipolar integration
${\tt LtpProba}\;[0.2]$	Spike_RRAM	Extrinsic STDP LTP probability (cumulative with in-
		trinsic SET switching probability P_{SET})
${\tt LtdProba}\;[0.1]$	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
		period

DigitalIntegration $\left[0 ight]$	Spike_RRAM	If false, the analog value of the devices is integrated,
		instead of their binary value

4.7.7 Deconv

Deconvolutionlayer.

Option [default value]	Description		
KernelWidth	Width of the kernel		
KernelHeight	Height of the kernel		
NbChannels	Number of output channels		
StrideX [1]	X-axis stride of the kernels		
StrideY [1]	Y-axis stride of the kernels		
Stride [1]	Stride of the kernels		
	(mutually exclusive with StrideX and StrideY)		
PaddingX $[0]$	X-axis input padding		
PaddingY [0]	Y-axis input padding		
Padding [0]	Input padding		
	(mutually exclusive with PaddingX and PaddingY)		
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.SizeX $[1]$	Mapping canvas pattern default width		
Mapping.SizeY $[1]$	Mapping canvas pattern default height		
Mapping.Size $igl[1igr]$	Mapping canvas pattern default size		
	(mutually exclusive with Mapping.SizeX and Mapping.SizeY)		
Mapping.StrideX $\left[1 ight]$	Mapping canvas default X-axis step		
Mapping.StrideY $\left[1 ight]$	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 $\left[0 ight]$	Mapping canvas default X-axis offset		
Mapping.OffsetY $\left[0 ight]$	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)		
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 $\left[1 ight]$	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		

ig Mapping(in).OffsetY $[0]$	Mapping canvas default Y-axis offset for input layer in	
Mapping(in).Offset $[0]$	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 $[1]$	all Frame	If true, don't use bias
BackPropagate $\left[1 ight]$	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, with of the output feature map, of the input feature map (channel), he the kernel row and we the kernel column, in the order of the outermost dimension (in the leftmost position) to the innermost dimension (in the rightmost position)
$ \begin{array}{c} {\tt WeightsExportTranspose} \\ [0] \end{array} $	all Frame	If true, import/export transposed kernels

4.7.8 Pool

Pooling layer.

Option [default value]	Description	
Pooling	Type of pooling (Max or Average)	
PoolWidth	Width of the pooling area	
PoolHeight	Height of the pooling area	
NbChannels	Number of output channels	
StrideX [1]	X-axis stride of the pooling areas	
StrideY [1]	Y-axis stride of the pooling areas	
Stride [1]	Stride of the pooling areas	
	(mutually exclusive with StrideX and StrideY)	
PaddingX $[0]$	X-axis input padding	
PaddingY $[0]$	Y-axis input padding	
Padding [0]	Input padding	
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,	
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
$ \ \ Mapping.SizeX \left[1\right] $	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 $egin{bmatrix} 1 \end{bmatrix}$	Mapping canvas default Y-axis step
Mapping.Stride $[1]$	Mapping canvas default step
	(mutually exclusive with Mapping.StrideX and Mapping.StrideY)
Mapping.OffsetX $\left[0\right]$	Mapping canvas default X-axis offset
Mapping.OffsetY $\left[0\right]$	Mapping canvas default Y-axis offset
Mapping.Offset $\left[0\right]$	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)
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 $\left[1 ight]$	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 $[0]$	Mapping canvas default X-axis offset for input layer in
Mapping(in).OffsetY $[0]$	Mapping canvas default Y-axis offset for input layer in
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	
NbChannels	Number of output channels	

ArgMax	Name of the associated pool layer for the argmax (the pool	
	layer input and the unpool layer output dimension must	
5.3	match)	
StrideX [1]	X-axis stride of the pooling areas	
StrideY [1]	Y-axis stride of the pooling areas	
Stride $[1]$	Stride of the pooling areas	
	(mutually exclusive with StrideX and StrideY)	
ig PaddingX $ig[0ig]$	X-axis input padding	
PaddingY $[0]$	Y-axis input padding	
Padding [0]	Input padding	
ActivationFunction [Linear]	Activation function. Can be any of Logistic, LogisticWithLoss,	
	Rectifier, Softplus, TanhLeCun, Linear, Saturation Or Tanh	
Mapping.SizeX $[1]$	Mapping canvas pattern default width	
Mapping.SizeY $\begin{bmatrix} 1 \end{bmatrix}$	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	
	(mutually exclusive with Mapping.StrideX and Mapping.StrideY)	
Mapping.OffsetX $[0]$	Mapping canvas default X-axis offset	
Mapping.OffsetY $\left[0 ight]$	Mapping canvas default Y-axis offset	
Mapping.Offset $\left[0\right]$	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	
[4]	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	
	(mutually exclusive with Mapping(in).SizeX and	
[41]	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).Stride [1]	Mapping canvas default step for input layer in	
	(mutually exclusive with Mapping(in).StrideX and	
101	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	
Mapping(in).Offset $[0]$	Mapping canvas default offset for input layer in	
	(mutually exclusive with Mapping(in).OffsetX and	
M (() M 7 () [0]	Mapping(in).OffsetY)	
Mapping(in).NbIterations $[0]$	Mapping canvas pattern default number of iterations for	
	input layer in (0 means no limit)	

4.7.10 FMP

Fractional max pooling layer (Graham, 2014).

Option [default value]	Description
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NbChannels	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
${\tt PseudoRandom}\;[1]$	all Frame	If true, use pseudorandom sequences, else use random
		sequences

4.7.11 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)]	

Configuration parameters (Frame models)

Option [default value]	Model(s)	Description
NoBias $[1]$	all Frame	If true, don't use bias
BackPropagate $\left[1 ight]$	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}
$;100 \; {\tt TimeFs}]$		

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 $[0.0]$	Spike, Spike_RRAM	Neural leak time constant τ_{leak} (if 0, no leak)
Refractory $[0.0]$	Spike, Spike_RRAM	Neural refractory period T_{refrac}
TerminateDelta [0]	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 $\left[0.0 ight]$	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 $\left[1.0 ight]$	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 $\left[0\right]$	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})
$\texttt{StdpLtp} \; [1000 \; \texttt{TimePs}]$	Spike_RRAM	STDP LTP time window T_{LTP}
${\tt InhibitRefractory} \qquad [0$	Spike_RRAM	Neural lateral inhibition period $T_{inhibit}$
TimePs]		
EnableStdp $[1]$	Spike_RRAM	If false, STDP is disabled (no synaptic weight change)
${\tt RefractoryIntegration} \; [1]$	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

N2D2 IP only 4.7.12 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
ScalingSolver.*	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

4.7.13 Softmax

Softmax layer.

Option [default value]	Description
NbOutputs	Number of output neurons
WithLoss $[0]$	Softmax followed with a multinomial logistic layer

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:

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

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

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

4.7.14 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 $\left[0.75\right]$	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

4.7.15 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

4.7.16 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.

4.7.17 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 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
NbChannels=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
NbChannels=[conv1]NbChannels
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
NbChannels=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.

```
Map(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 1 1 1 0 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 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
NbChannels=[conv2]NbChannels
Stride=2
Pooling=Max
Mapping.Size=1

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

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

5.2 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]NbChannels
```

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.

5.2.1 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
NbChannels=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
```

5.2.2 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
NbChannels=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-2 0
. . .
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
```

5.2.3 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=TargetR0Is
MinOverlap=0.33; Min. overlap fraction to match the R0I to an annotation
FilterMinWidth=5; Min. R0I width
FilterMinHeight=5; Min. R0I height
FilterMinAspectRatio=0.5; Min. R0I aspect ratio
FilterMaxAspectRatio=1.5; Max. R0I 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
MinOverlap=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.

5.2.4 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 13.



Figure 13: Example of the target visualization helper tool.

5.3 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 5.1.

5.3.1 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]:

```
...

<del>[softmax]</del>

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...).

5.3.2 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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