### **Basic**

- There are 4 main parts in a layer program:
  - constants
  - data
  - networks
  - scripts

### **Constants**

- There are three different constants to define (default values):
  - threads: number of threads for parallelization (4)
  - batch size: size of the batch for the network (100)
  - log: log file where some messages are saved (netparser.log)

```
const{
  threads=4
  batch=10
  log="mnist.log"
}
```

### **Data**

Data can be read in ascii or binary. Data format is the following:

```
samples dim output

sample1_1 ... sample1_dim out1_1 .. out1_c

...

samplen_1 ... samplen_dim outn_1 .. outn_c
```

- samples is the number of samples (int)
- dim is the dimensionality of the samples (int)
- out is the number of output targets, classes in a classification problem (int)
- each row is a sample (float values)

Note: when samples are color images the channels appear separated, firts R, then G and then B and dim

must be  $3 \times rows \times cols$ 

• An example of data definition:

```
data {
   D1 [filename="../training", binary]
   D2 [filename="../test", binary]
   Dval [filename="../valid", ascii]
}
```

- D1, D2 and Dval are the name of the data variables
- Full path to file can be used

## **Networks**

- · Networks are the main part of a Layer program
- Networks are composed by three main parts:
  - Data
  - Layers
  - Connections
- The networks are defined like this:

```
network N1 {
...
}
```

• where N1 is the name of the network

## **Networks - Data**

- · The Networks have to defined at least a training data set
- Test and validation data sets are optional:

```
network N1 {
   //data
   data tr D1 //mandatory
   data ts D2 //optional
   data va Dval //optional
   ...
}
```

 When test or validation data are provided the error function of the net will be also evaluated for that data sets

# **Networks - Layers**

- There are different layers that can be created:
  - FI: Input Fully Connected layer
  - CI: Input Covolutional layer
  - F: Fully Connected layer
  - FO: Ouput layer
  - C: Convolutional layer
  - MP: MaxPooling layer
  - · CA: Cat layer

### **Networks - Layers - FI**

FI has not parameters just serve as a connection with the data

```
//NETWORK
network N1 {
   //data
   data tr D1

// Fully Connected Input
FI in
...
}
```

- in is the name of the layer
- The number of units is the dimensionality representation of the data

### **Networks - Layers - CI**

- Convolutional input has three mandatory parameters that indicate how the raw data has to be mapped into an input Map
- The parameters are:
  - o nz: number of channels
  - o nr: image rows
  - o nc: image cols

```
//NETWORK
network N1 {
   //data
   data tr D1

// Convolutional Input for MNIST
CI in [nz=1, nr=28, nc=28]
...
}
```

• in is the name of the layer

### **Networks - Layers -CI**

• Convolutional input has optional parameters:

```
cr: crop rowscc: crop cols
```

- Images will be randomly cropped on training phase
- Images will be center-cropped in test phase

```
//NETWORK
network N1 {
  //data
  data tr D1

// Convolutional Input for MNIST
  // Images will be randomly cropped to 24x24
  CI in [nz=1, nr=28, nc=28, cr=24, cc=24]
  ...
}
```

## **Networks - Layers - F**

· A fully connected layer

```
//NETWORK
network N1 {
   //data
   ...

// Fully connected
F f1 [numnodes=1024]
   ...
}
```

- f1 is the name of the layer
- numnodes is a mandatory parameter

### **Networks - Layers - FO**

- FO has a mandatory parameter {classification,regression} that define the cost error: cross-entropy or mse respectivelly
- For regression, optionally, we can define an autoencoder

```
network N1 {
    ...
    // Targets are the output in data (one-hot)
    F0 out1 [classification] // Cross-entropy

// Targets are the output in data (real values)
    F0 out2 [regression] // mse over output

// Targets are the same input in data
    F0 out3 [regression,autoencoder] // mse over input
    ...
}
```

## **Networks - Layers - C**

- Convolutional layers have three mandatory parameters:
  - nk: number of kernels
  - · kr: height of kernel
  - kc: width of kernel

```
network N1 {
    ...
    // Convolutional Layer
    C c1 [nk=32, kr=3, kc=3]
    ...
}
```

- c1 is the name of the layer
- Convolutional layers have the following optional parameters:
  - rpad: indicates if padding is done in the rows (0)
  - cpad: indicates if padding is done in the cols (0)
  - stride: stride value (1)

```
network N1 {
    ...
    // Convolutional Layer with padding
    // To keep the same size of Input Maps
    // use rpad=cpad=1
    C c1 [nk=32, kr=3, kc=3,rpad=1,cpad=1]

    // Stride=2 and only padding in cols
    C c2 [nk=32, kr=3, kc=3,cpad=1,stride=2]
    ...
}
```

### **Networks - Layers - MP**

- MaxPool layers have two mandatory parameters:
  - · sizer: height of the pooling region
  - sizec: width of the pooling region

```
network N1 {
    ...
    // Maxpool layer sizer=sizec=2
    MP p1[sizer=2, sizec=2]
    ...
}
```

## **Networks - Layers - CA**

Cat layers does not require any parameter

# **Networks - Connections**

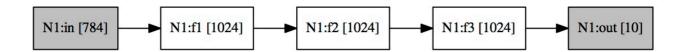
• To connect two layers use the symbols "->"

# **Networks - Examples**

**Example 1 - MLP MNIST** 

```
// A typical MLP for MNIST
// 784->1024->1024->10
//CONSTANTS
const{
  threads=8
  batch=10
  log="mnist.log"
}
// DATA FILES
data {
  D1 [filename="../training", binary]
  D2 [filename="../test", binary]
}
//NETWORK
network N1 {
  //data
  data tr D1 //mandatory
  data ts D2 //optional
  // Fully Connected Input
  FI in
  // Fully connected
  F f1 [numnodes=1024]
  F f2 \[ \text{numnodes} = \frac{1024}{\text{}} \]
  F f3 [numnodes=1024]
  // Fully Connected Output
  FO out [classification]
  // Links
  in->f1
  f1->f2
  f2->f3
  f3->out
```

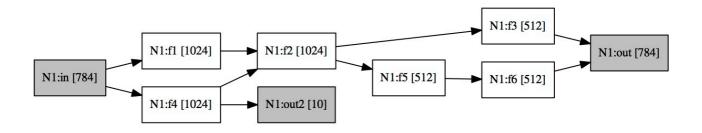
- layers generates a dot file with the name of each network
- Use "dot -T pdf N1.dot > N1.pdf" to generate a PDF with the network:



• but yes, more complicated networks can be defined

## **Example 2 - Hybrid MLP MNIST**

```
//NETWORK
network N1 {
  //data
  data tr D1
  data ts D2
  // Fully Connected Input
  FI in
  // Fully connected
  F f1 \[ \text{numnodes} = \frac{1024}{\text{}} \]
  F f2 [numnodes=1024]
  F f3 [numnodes=512]
  // Fully connected
  F f4 [numnodes=1024]
  F f5 \[ \text{numnodes=512} \]
  F f6 [numnodes=512]
  // Output
  FO out [regression, autoencoder]
  FO out2 [classification]
  // Connections
  in->f1
  f1->f2
  f2->f3
  f3->out
  in->f4
  f4->f2
  f4->out2
  f2->f5
  f5->f6
  f6->out
}
```



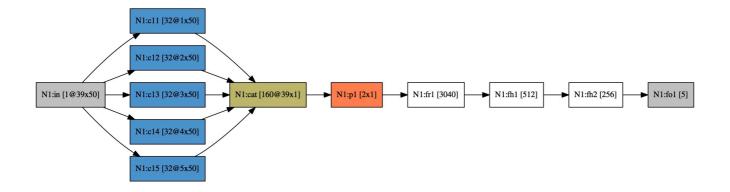
**Example 3 - Convolutional CIFAR** 

```
network N1 {
  data tr D1
  data ts D2
  CI in [nz=3, nr=32, nc=32]
  C c0 [nk=16, kr=3, kc=3]
  MP p0[sizer=2,sizec=2]
  C c1 [nk=32, kr=3, kc=3]
  MP p1 [sizer=2, sizec=2]
  C c2 [nk=64, kr=3, kc=3]
  MP p2 [sizer=2, sizec=2]
  // FC reshape
  F f0 []
  // FC hidden
  F f1 [numnodes=128]
  // FC output
  FO f2 [classification]
  // Conecctions
  in->c0
  c0->p0
  p0->c1
  c1->p1
  p1->c2
  c2->p2
  //reshape
  p2->f0
  f0->f1
  f1->f2
}
```



**Example 4 - CAT layers** 

```
network N1 {
  data tr D1
  data ts D2
  CI in [nz=1, nr=39, nc=50]
  C c11 \lceil nk=32, kr=1, kc=50, rpad=1 \rceil
  C c12 [nk=32, kr=2, kc=50,rpad=1]
  C c13 [nk=32, kr=3, kc=50,rpad=1]
  C c14 [nk=32, kr=4, kc=50,rpad=1]
  C c15 [nk=32, kr=5, kc=50, rpad=1]
  MP p1 [sizer=2, sizec=1]
  CA cat
  // FC reshape
  F fr1 □
  // FC hidden
  F fh1 \[ \text{numnodes=512} \]
  F fh2 [numnodes=256]
  // FC output
  FO fo1 [regression]
  // Connections
  in->c11
  in->c12
  in->c13
  in->c14
  in->c15
  c11->cat
  c12->cat
  c13->cat
  c14->cat
  c15->cat
  cat->p1
  p1->fr1
  fr1->fh1
  fh1->fh2
  fh2->fo1
}
```



# **Scripts**

- Script is a block of actions. There are two different types of actions:
  - functions
  - parameters modifications

# **Scripts Functions**

- There are several functions that can be applied to the defined objects:
  - For Data:
    - Zscore(): To normalize Data
    - yuv(): to convert RGB Maps to YUV maps
  - For Networks:
    - train(epochs): to train networks

More functions will be soon implemented

```
script {
   D1.yuv()
   D2.yuv()

   D1.zscore()
   D2.zscore(D1) //normalize D2 with D1 statistics

   N1.train(100)
}
```

# **Scripts Parameters**

• There are several parameters that can be modified:

```
mu: learning rate
mmu: momentum rate
l2: l2 regularization (weight decay)
l1: l1 regularization
maxn: maxnorm regularization
noiser: noise ratio after activation function
noisesd: standard deviation of noise (N(0.0, σ))
drop: dropout (0<drop<1)</li>
bn: batch normalization ({0,1})
act: activation (0 Linear, 1 Relu, 2 Sigmoid, 3 ELU)
flip: to flip input images ({1,0})
shift: to shift randomly input images
```

• Parameters can be modified for one particular layer:

```
script{
...
N1.c2.drop=0.5
N1.in.flip=1
N1.in.shift=2
...
}
```

• Or, parameters can be modified for all the layers of the same network:

```
script{
...
N1.mu=0.001
N1.12=0.0005
// noise in all the layers of the network:
N1.noiser=0.5
N1.noiser=0.3
...
}
```

· Some examples of scripts:

```
script {
  D1.yuv()
  D2.yuv()
  D1.zscore()
  D2.zscore(D1)
  N1.in.flip=1.0
  N1.in.noiser=0.5
  N1.in.noisesd=1.0
  N1.f1.drop=0.5
  N1.12=0.0005
  N1.bn=1.0
  // Learning rate annealing every 15 epochs
  N1.mu = 0.01
  N1.train(15)
  N1.mu = 0.005
  N1.train(15)
  N1.mu = 0.0025
  N1.train(15)
}
```

```
script {
    D1.zscore()
    D2.zscore(D1)

    N1.noiser=0.0
    N1.noisesd=1.0
    N1.in.noiser=0.5
    N1.mu=0.001
    N1.maxn=3.0

    N1.train(100)
}
```

# The whole picture

```
// A typical MLP
// 784->1024->1024->10
// Check N1.dot and get a pdf of the net with:
// dot -T pdf N1.dot > N1.pdf
//CONSTANTS
const{
  threads=8
  batch=10
  log="mnist.log"
}
// DATA FILES
data {
  D1 [filename="../training", binary]
  D2 [filename="../test", binary]
}
//NETWORK
network N1 {
  //data
  data tr D1 //mandatory
  data ts D2 //optional
  // Fully Connected Input
  FI in
  // Fully connected
  F f1 \[ \text{numnodes} = \frac{1024}{\text{}} \]
  F f2 \[ \text{numnodes=1024} \]
  F f3 [numnodes=1024]
  // Fully Connected Output
  FO out [regression]
  // Links
  in->f1
  f1->f2
  f2->f3
  f3->out
}
```

```
//RUN SCRIPT
script {

D1.zscore()
D2.zscore(D1)

//N1.noisesd=1.0
//N1.noiser=0.5
N1.mu=0.01
//N1.maxn=3.0

N1.train(10)

}
```

```
const {
threads=4
batch=100
data {
  D1 [filename="training", binary]
  D2 [filename="test", binary]
}
network N1 {
  data tr D1
  data ts D2
  // Covolutional Input
  CI in [nz=3, nr=32, nc=32]
  C c0 [nk=16, kr=3, kc=3]
  MP p0[sizer=2, sizec=2]
  C c1 [nk=32, kr=3, kc=3]
  MP p1 [sizer=2, sizec=2]
  C c2 [nk=64, kr=3, kc=3]
  MP p2 [sizer=2, sizec=2]
  // FC reshape
  F f0 []
  // FC hidden
  F f1 [numnodes=128]
```

```
// FC output
  FO f2 [classification]
  // Connections
  in->c0
  c0->p0
  p0->c1
  c1->p1
  p1->c2
  c2->p2
  //reshape
  p2->f0
  f0->f1
  f1->f2
}
// RUN
script {
  D1.zscore()
  D2.zscore(D1)
  N1.mu=0.1
  N1.in.noiser=0.5
  N1.in.noisesd=1.0
  //N1.bn=1.0
  N1.train(1000)
}
```

# **ADVANCED ISSUES**

## **Networks sharing layers**

TODO

## **Training several networks**

**TODO** 

...