#### Title Author\*

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# **Thanks**

To someone.

Began: July 16, 2012 Updated: Thursday 20<sup>th</sup> June, 2013

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### Part I

# **Intro Examples**

#### El mundo real

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#### Lists

Example of making list automated, saving space and reducing errors.

#### **Terms**

#### endocrine chemicals

- proteins
- steroid hormones
- amine hormones
- testosterone
- insulin
- thyroxine
- epinephrine
- tropic hormones
- thyrotropin
- luteinizing hormone
- follicle-stimulating

mone

- corticotropin
- growth hormone
- prolactin
- melanocyte-stimulating hormone
- enkephalins
- endorphins
- POMĊ

hor-

#### receptor tyrosine kinases

- dimerization
- autophosphorylation
- SH2 domains
- EGFR
- Ras
- Raf MEK
- ERK
- SOS
- Sevenless Drk/Grb2
- Ras-GAP
- Ras-GEF

- cAMP
- signal amplification
- cĞMP
- guanylyl cyclase
- phospholipids
- calmodulin
- protein-tyrosine phosphatase'
- serine/threonine kinase
- intracellular signal transduction
- NF-κB
- ΙκΒ

#### R

See proposed hierarchical neural networks pseudo-code block at end of paper for example implementation of this method in the R statistical language (Code 1)-error handling and background of creation of some parts of the neural networks are left out for sake of clarity.

#### Code 1: hierarchical neural networks in R

```
# Load neural net library
library(nnet)
library(neuralnet)
# Load trained neural nets
source("model.training.NN.hierarchical.v1")
trainedNeuralNets = NNtrained()
# Load data
source("data.run.NN.hierarchical")
this.data = NNdata()
# define the number of loops before an error has occured
error.loop.value = 3
neuralNet <- function(neuralNetFxn, this.loop.input){</pre>
       # this function runs specific level of a hierarchal neural network then recursively calls the
           next layer
       # base case is defined when a parent neural net has no children
       # run the initial neural net on the input
       # classifications is a tuple of probabilities for being in given class
       list(classifications, subfunctions) := neuralNetFxn(this.loop.input)
       # set the threshold
       threshold = this.loop.threshold.value
       # get index of next neural net function to be called
       subidx = max(find.col(classifications>threshold))
```

#### **Python**

A playlist maker for python (Code 2), see more concise implementation in bash (Code 3).

#### Code 2: python playlist maker

```
#!/Python27/env python
#Biafra Ahanonu
#2012.12.02
#Makes .m3u extended playlist at first level folders in a directory
#Modules used
import os,re,time
#Import settings from settings.py
```

```
from settings import *
#Help filter out duplicates
from sets import Set
def folderBrowser():
      #Opens a folder
      import Tkinter, tkFileDialog
      root = Tkinter.Tk()
      root.withdraw()
      dir = tkFileDialog.askdirectory(parent=root,initialdir=DEFAULT_DIR,title='Please select a
           directory')
      return dir
def fileTree(dir,relDir):
      #Crawls through a directory and finds audio files, returns list of files
      print dir
      #Variable to save files to
      filesToSave = []
      #Valid mp3 files
      validFiles = set(VALID_AUDIO_FORMATS)
```

#### bash

Improved playlist maker implemented in bash (Code 3).

#### Code 3: bash playlist maker

```
#!/bin/bash
# biafra ahanonu
# 2013.02.23
# script to make playlists
# Yes/No function
getYesNo(){
      select terminateSignal in "Yes" "No"
             case $terminateSignal in
                     "Yes" )
                            return 1;;
                     "No" )
                           return 0;;
             esac
       done
}
# Change file separator to allow use of files with spaces
oldIFS=$IFS
IFS=(echo -en "\n\b")
# Ask user for directory
echo "Directory? "
read userDir
echo $userDir
cd $userDir
# Ask to remove old .m3u files
```

#### **FASTA**

Example use of the fasta formatting for protein sequences.

#### **FASTA** sequences

 $D_1$ a dopamine receptor | Homo sapiens | Humans

MRTINTSAMDGTGLVVERDFSVRILTACFLSLILISTILGONTLVCAAVIRFRHIRSKVTNFFVISLAVSDLLVAVLVMPWKAVAEIAGFWPFGSFCNIWVAFDIMCSTASILNICVISVDRYWAISSPFRYERKMTPKAAFIUISVAWTLSVILSFIPVGLSWHKAKPTSP SDCNATSLAETIDNCDSSLSRTYAISSSVISFYIPVAIMIVTYTRIYRIAQKQIRRIAALERAAVHAKNCQTTTGNCKPVECSQPESSFKMSFKRETKVIKTLSVIMGVFVCCWLPFFLINCILPFCGSGETQPFCIDSNTFDVFVWFGWANSSLNPIIYAFNADFRK AFSTLLGCYRLCPATNNAIETVSINNNGAAMFSSHHEPRGSISKECNLVYLIPHAVGSSEDLKKEEAAGIARPLEKLSPALSVILDYDTDVSLEKIQPITQNGQHPT

#### **Problems**

If there is any confusion about the questions, shoot me an email or talk to me after class.

- 1. Look at ??, why is the heart rate of the iguana different at the same temperature? In the first case the body temperature is decreasing while in the second case the iguana raises its heart rate to increase blood flow, which leads to increased body temperature.
- 2. How does increasing heart rate while in direct sunlight help warm the body? More blood can flow to the skin and be warmed.
- 3. What does it mean for the Q10 of the metabolic rate of an animal to be 2? To be 1? the animal consumes half as much oxygen per hour at 20C as it does at 30C. In the case of 1, it doesn't change.

### **Equations**

#### **Equations**

Example of several equations chosen at random from different works. Illustrates the use of the **lbpeq** macro.

$$E = \frac{1}{2} \sum_{q=1}^{n} \sum_{k=1}^{K} [y_k(x^q, w) - t_k^q]^2$$
(1)

$$y_j(x) = \sum_{i=0}^n w_{i,j} \phi(x)_i$$
 (2)  $\phi(x)_i = exp(-\frac{|x - \mu_i|^2}{2\omega_i^2})$ 

$$M + Q_{abs} = \epsilon \sigma T_r^4 + h_c (T_r - T_a) + E + C$$
(4)

# **Appendix**

Additional figures and tables are included. References and index found at the end.

attenuation	laserPower	attenuationDirection
2	126.3	down
1. <i>7</i>	118. <i>7</i>	down
1.6	114.1	down
1.5	100.4	down
1.4	71.7	down
1.3	44.2	down
1.2	26.5	down
1.1	15.3	down
1	8.52	down
0.9	4.61	down
0.8	2.17	down
0.7	0.64	down
2	127	up
1.7	119	up
1.6	112.6	up
1.5	97.5	up
1.4	70.5	up
1.3	44	up
1.2	25	up
1.1	14.59	up
1	8.26	up
0.9	4.62	up
0.8	2.16	up
0.7	0.49	up

Name	Туре	$\%\Delta$	Onset	Localization
YAP	protein	60	E18	Nuclear- >Cytoplasm
Hippo	protein	40	E18	Cytoplasm
agmatine	metaboliote	20	Ś	Ex->Cytoplasm
Lim 1	protein	-50	E9.5	Cytoplasm
FoxC1/2	protein	-10	E11.5	Cytoplasm

Table 1 | Hypothetical list of kidney proteins and metabolites These were identified after removal of housekeeping and injury genes.  $\%\Delta$  is protein or metabolite level change from **onset** to cessation of kidney growth. Onset time is part hypothetical, part from the literature.

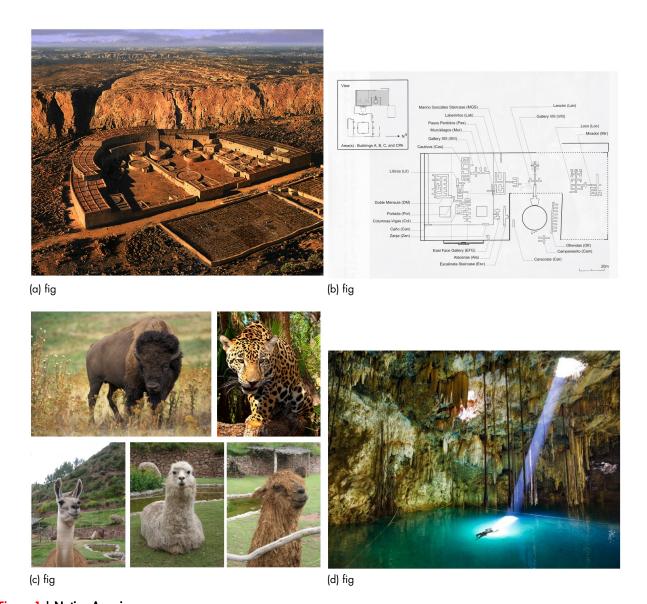


Figure 1 | Native Americans (a) Pueblo Bonito was an.... (b) Located in Peru. (c) The different type of game hunted in the Americas. (d) The cenote, which were seen as sacred by the Maya.

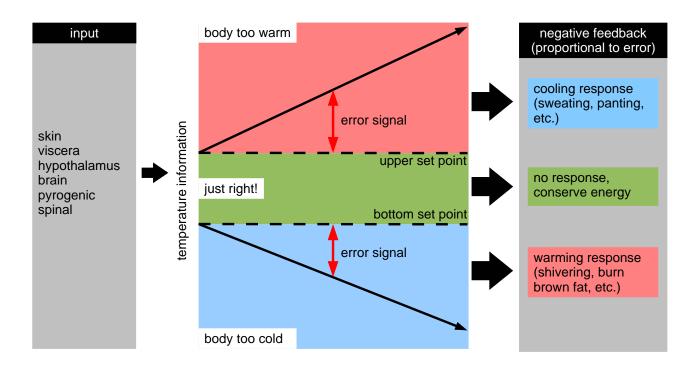


Figure 2 | Response regulation General concept of integrating temperature information and formulating a response.

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