

# Classification using Machine-Learning Algorithms (MALA)

Math 579

Dr. Jen-Mei Chang

Rodrigo Farnham

Tuyen Ly

Jason Wang



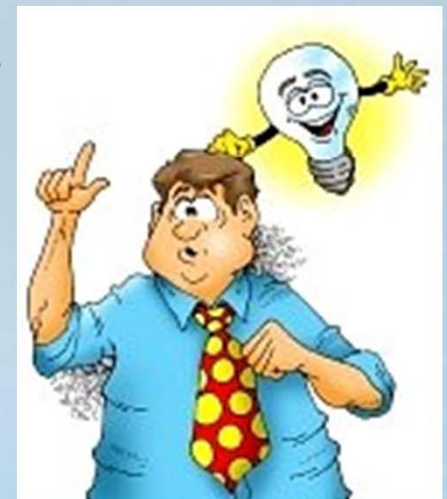


# Our Idea

*“The placenta is the crystal ball of the baby.”*

- Dr. Carolyn Salafia

- o Extract and identify features of the placenta
- o Extract relevant patient/mother data
- o Use above data to train a machine-learning system, hopefully to make predictions of a baby's future health



# Overview

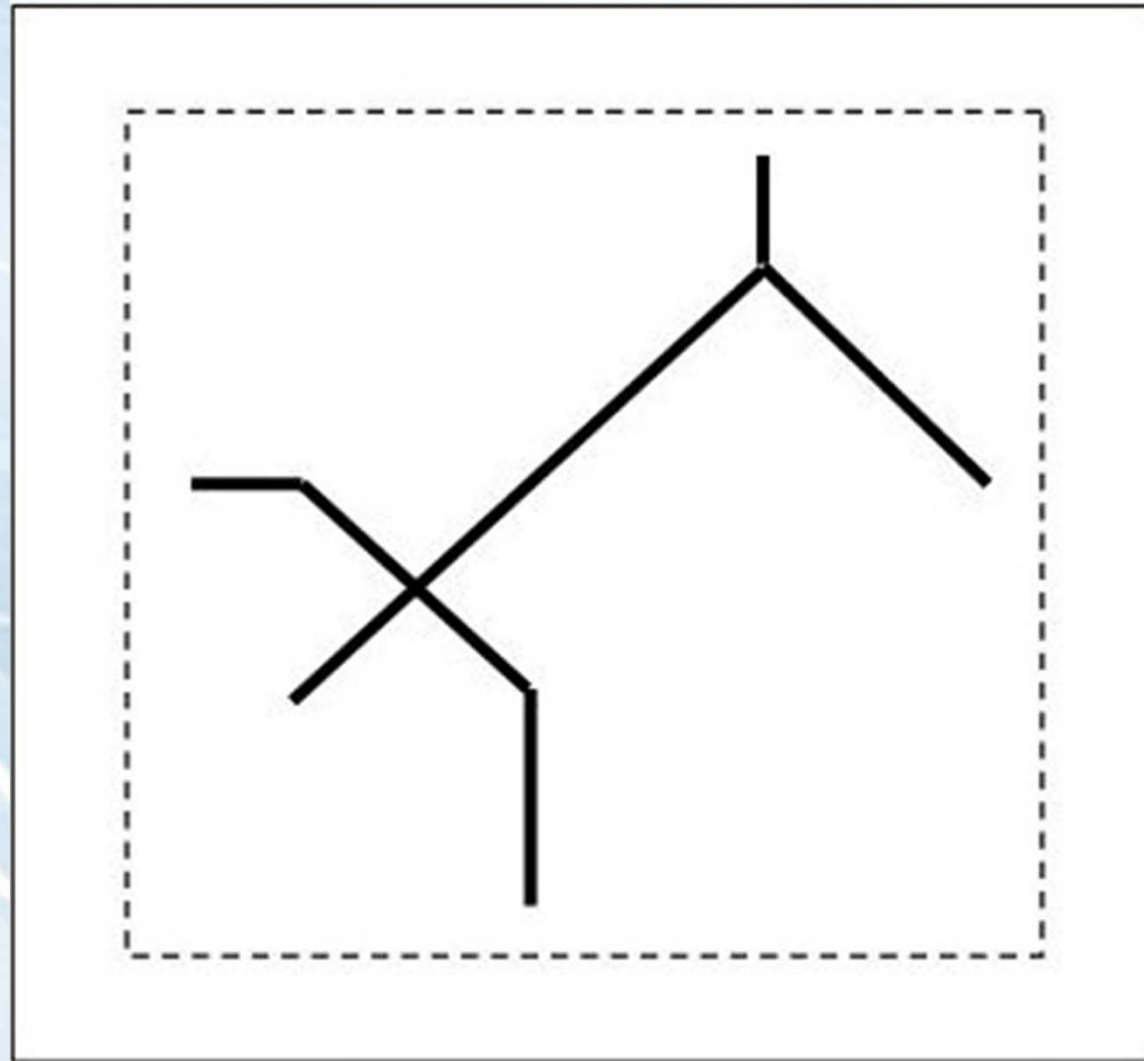
- 1) Analyze placental attributes
- 2) Maternal Attribute Classification
- 3) Learning Machine System: Weka

*“A suite of machine learning software written at the University of Waikato”*



# Placental Attribute: Heat Map

**Sample  
Vessel  
Network**





















# Placental Attribute: Heat Map

Matlab  
Output

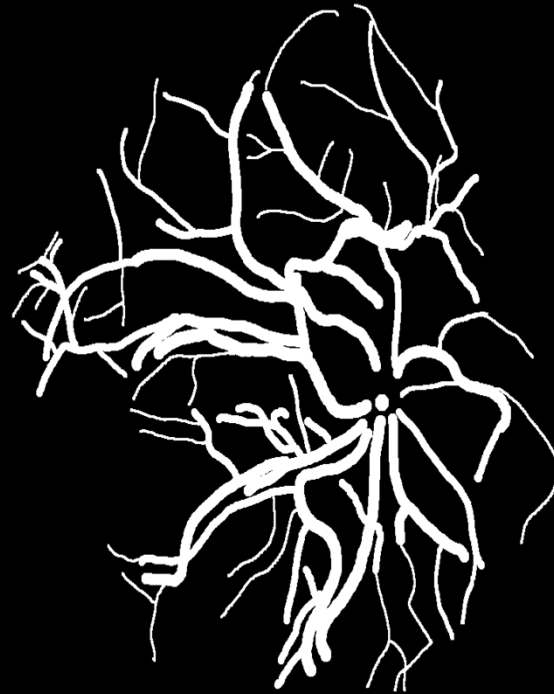
Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf
Inf	3	3	2.83	2	1	0	1	2	Inf
Inf	2	2	2.41	1.41	1	0	1	1.41	Inf
Inf	1	1	1.41	1	0	1	0	1	Inf
Inf	0	0	1	0	1	1.41	1	0	Inf
Inf	1	1	0	1	1.41	2.41	1.41	1	Inf
Inf	1	0	1	0	1	2	2.41	2	Inf
Inf	1.41	1	1	0	1	2	3	3	Inf
Inf	2.41	2	1	0	1	2	3	4	Inf
Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf

largest value:  
(largest distance from  
vessel network)



# Placental Attribute: Heat Map

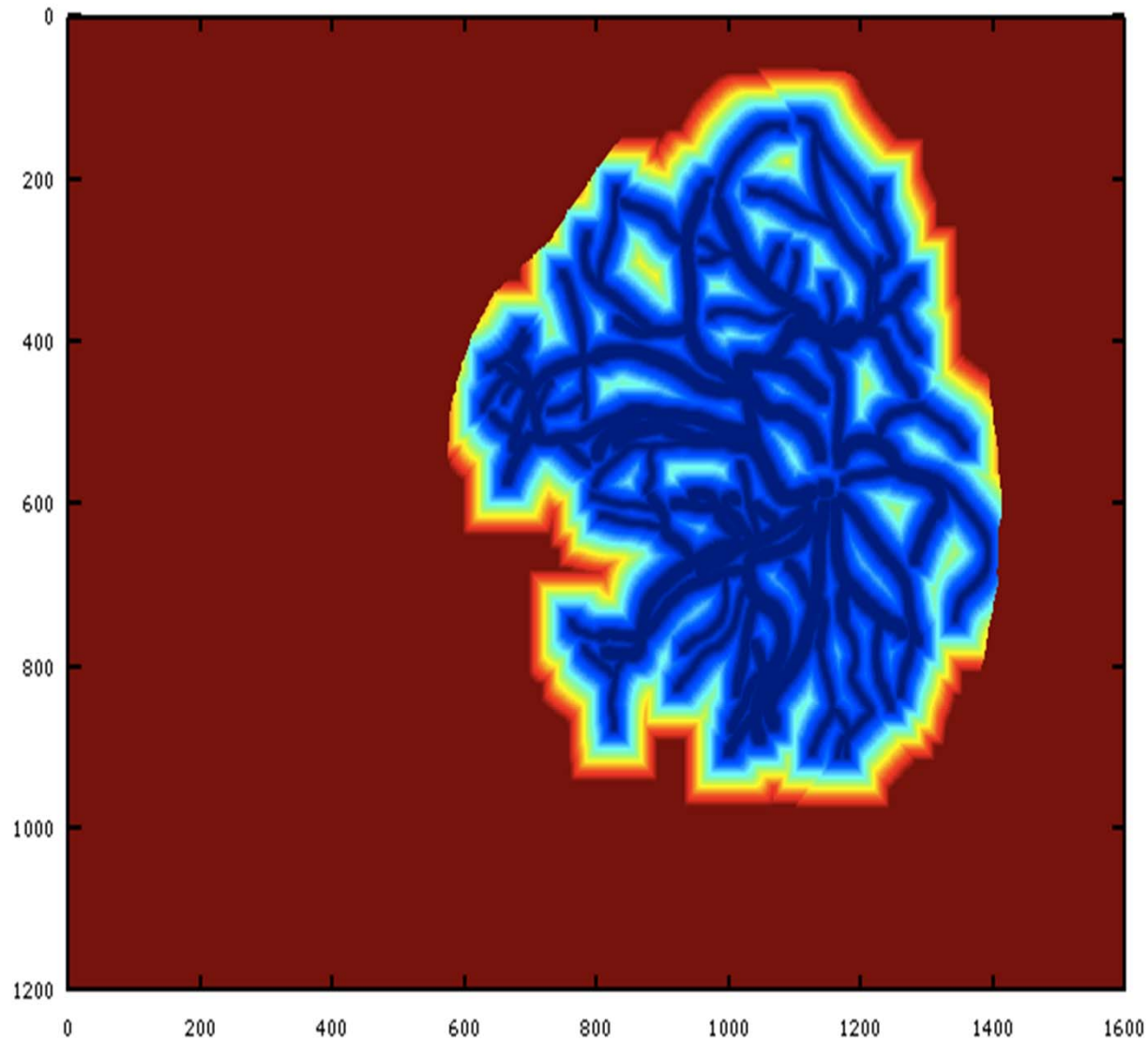
**Vessel  
Network**





# Placental Attribute: Heat Map

**Vessel  
Network**



# Maternal Attributes

## Strength:

- A whole lot of data, total of 209 attributes
  - mom's ethnicity
  - mom's height
  - mom's age at pregnancy start date
  - child's birth weight
  - mom's total weight gain
  - # of previous pregnancies
  - # of previous live births
  - etc.

## Weaknesses:

- *Too much* data
  - ↳ A lot of irrelevant data as well as dependent data





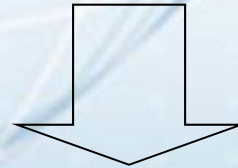
# Maternal Attributes

	A	B	C	D	E	F	G	H	
1	LABID	BirthWeight	GestationalDays	Weightbeforecuttingcordmembrane	Distancebetweenrupturesiteplacentalmargin	CordLength	CordWeight	Weightaftercuttingcordmembrane	DELTA
2	1,353	2,770	261		530 5-10	20	#NULL!		400
3	1,402	2,692	254		-99 -99	-99	#NULL!		-99
4	1,433	2,624	262		440 <5	45	#NULL!		350
5	1,462	4,018	284		420 <5	37	#NULL!		340
6	1,471	3,489	256		570 <5	51	#NULL!		450
7	1,472	3,338	278		460 5-10	23	#NULL!		370
8	1,473	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
9	1,474	3,421	286		560 <5	36	#NULL!		450
10	1,475	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
11	1,476	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
12	1,478	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
13	1,479	2,369	285		370 -99	48	#NULL!		320
14	1,480	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
15	1,482	-999	187		300 <5	19	#NULL!		240
16	1,489	3,790	274		600 <5	60	#NULL!		460
17	1,490	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
18	1,491	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
19	1,492	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
20	1,498	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
21	1,499	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
22	1,501	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
23	1,502	3,172	277		470 Unknown	33	#NULL!		400
24	1,503	3,042	268		480 <5	56	#NULL!		340
25	1,504	3,657	295		560 5-10	57	#NULL!		460
26	1,505	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	#NL
27	1,506	3,185	275		540 5-10	37	#NULL!		450
28	1,507	3,391	271		690 <5	34	#NULL!		600
29	1,508	2,556	259		570 <5	61	#NULL!		420
30	1,509	3,484	266		510 <5	51	#NULL!		420
31	1,510	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
32	1,511	#NULL!	#NULL!	#NULL!		#NULL!	#NULL!	#NULL!	
33	1,512	3,462	281		530 <5	42	#NULL!		430
34	1,513	2,220	243		440 5-10	15	#NULL!		330
35	1,514	2,885	270		530 >10	57	#NULL!		440
36	1,515	2,872	280		500 <5	50	#NULL!		420
37	1,516	2,827	263		500 <5	53	#NULL!		390
38	1,517	3,139	264		600 <5	35	#NULL!		470

# Maternal Attributes

## The attributes we chose to look at:

- “Reasonable” attributes
- Attributes whose data is “spread out”
- Results from research papers/publications



- I. Mother’s total weight gain<sup>1</sup>
- II. Number of previous pregnancies
- III. Summary Index # of Prenatal Care Adequacy<sup>2</sup>
- IV. Gestational Days
- V. Family’s Poverty Level Index<sup>3</sup>

<sup>1</sup> <http://www.telegraph.co.uk/health/healthnews/7926233/Putting-on-too-much-weight-in-pregnancy-risk-babys-health.html>

<sup>2</sup> <http://www.sjph.net.sd/files/vol4i4/SJPH-vol4i4-p403-410.pdf>

<sup>3</sup> [http://www.epi.umn.edu/mch/resources/hg/hg\\_childpoverty.pdf](http://www.epi.umn.edu/mch/resources/hg/hg_childpoverty.pdf)



# Birth Weight & Beta Value

## The Importance of Birth Weight:

doi: 10.1111/j.1365-3016.2008.00935.x

229

Fetal growth correlates

### Placental characteristics and birthweight

Carolyn M. Salafia<sup>a,d</sup>, Jun Zhang<sup>a</sup>, Adrian K. Charles<sup>a</sup>, Michaeline Bresnahan<sup>a,b</sup>, Patrick Shrout<sup>c</sup>, Wenyu Sun<sup>a</sup> and Elizabeth M. Maas<sup>d</sup>

<sup>a</sup>Department of Epidemiology, Mailman School of Public Health, Columbia University College of Physicians and Surgeons, <sup>b</sup>New York State Psychiatric Institute, <sup>c</sup>Department of Psychology, New York University, New York, NY, <sup>d</sup>EarlyPath Clinical and Research Diagnostics, Larchmont, NY, <sup>e</sup>Epidemiology Branch, National Institute of Child Health and Human Development, National Institutes of Health, Bethesda, MD, USA, and <sup>f</sup>Department of Pathology, Princess Margaret Hospital, Perth, Western Australia

#### Summary

Salafia CM, Zhang J, Charles AK, Bresnahan M, Shrout P, Sun W, Maas EM. Prenatal characteristics and birthweight. *Paediatric and Perinatal Epidemiology* 2008; **22**: 229–239.

Standard gross placental measures capture dimensions relevant to specific placental functions. Our objective was to determine their accountability independent of placental weight for variance in birthweight, an important proxy for intrauterine 'adequacy' in fetal origins studies. The sample consisted of 24 152 singleton liveborn children of the Collaborative Perinatal Project delivered from 34 to 42 completed weeks gestation, with complete data for six placental measures (placental disc shape, umbilical cord length, distance from cord insertion to nearest margin, large diameter, small diameter, placental thickness) and placental weight. Associations between birthweight and placental measures were examined using multiple linear regression. Placental weight alone accounted for 36.6% of birthweight variation; the six other placental measures accounted for 28.1%. Combined, all placental measures accounted for 39.1% of birthweight variation. Seven maternal characteristics (age, height, weight, parity, socioeconomic status, cigarette use, and race) were investigated to determine whether their known associations with birthweight were mediated by placental markers. Analysis suggested that the impact of all maternal characteristics except smoking was consistent

#### Correspondence:

Carolyn M. Salafia, MD MS,  
Assistant Professor of  
Epidemiology, Mailman  
School of Public Health,  
Columbia University, 722  
West 168th Street, New York,  
NY 10032, USA.  
E-mail: salafiacm@aol.com



# Birth Weight & Beta Value

$$\beta = \frac{\log PW^*}{\log BW}$$



$\beta > 0.75$

**low functional  
efficiency**

$\beta < 0.75$

**high functional  
efficiency**

# Birth Weight & Beta Value

Kleiber's Law:

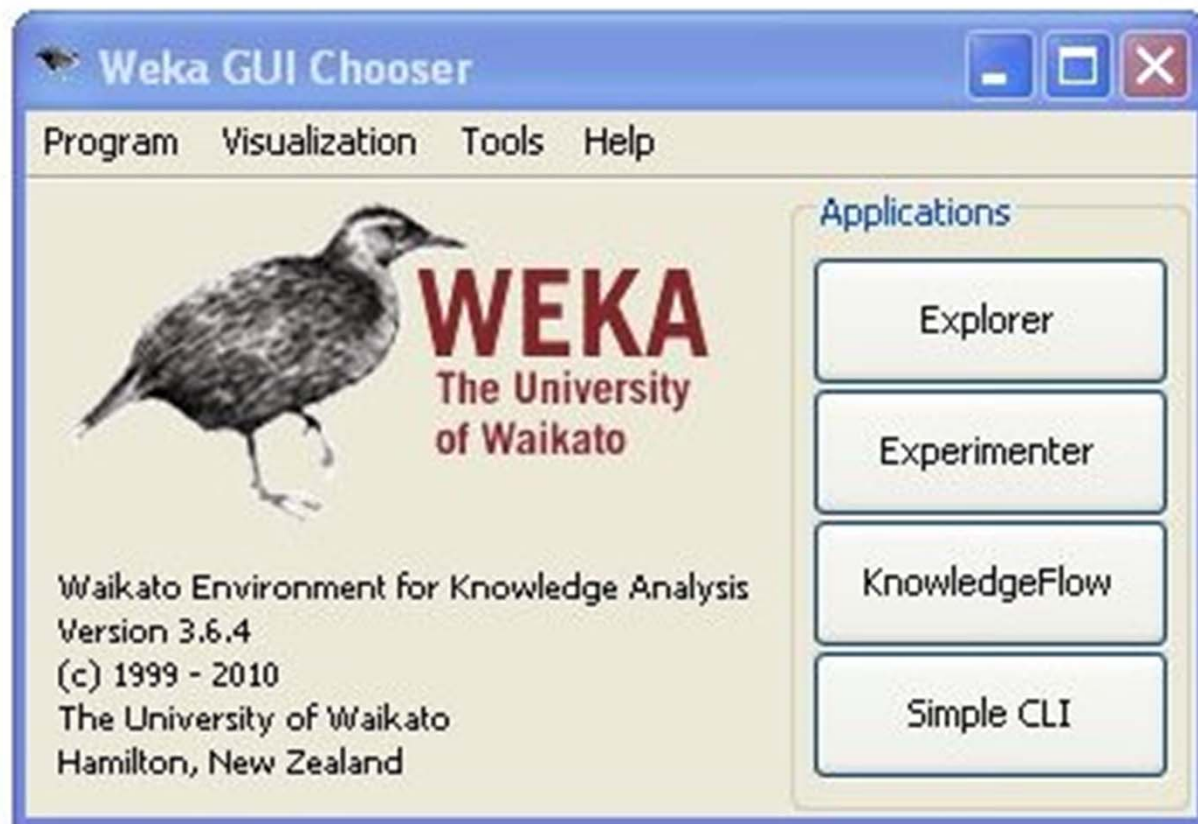
Metabolic rate ( $q_0$ ) is proportional to body mass ( $M$ )  
raised to  $3/4$  power

$$q_0 \sim M^{3/4}$$





# Learning Machine: Weka



**Weka:** <http://www.cs.waikato.ac.nz/ml/weka/>

# Weka

Weka Explorer

Preprocess | Classify | Cluster | Associate | Select attributes | Visualize

Open file... | Open URL... | Open DB... | Generate... | Undo | Edit... | Save...

Filter: Choose **None** Apply

Current relation: Relation: sdv\_vs\_GA Instances: 139 Attributes: 361

Attributes: All | None | Invert | Pattern

No.	Name
351	A351
352	A352
353	A353
354	A354
355	A355
356	A356
357	A357
358	A358
<b>359</b>	<b>A359</b>
360	A360
361	preterm(1)

Remove

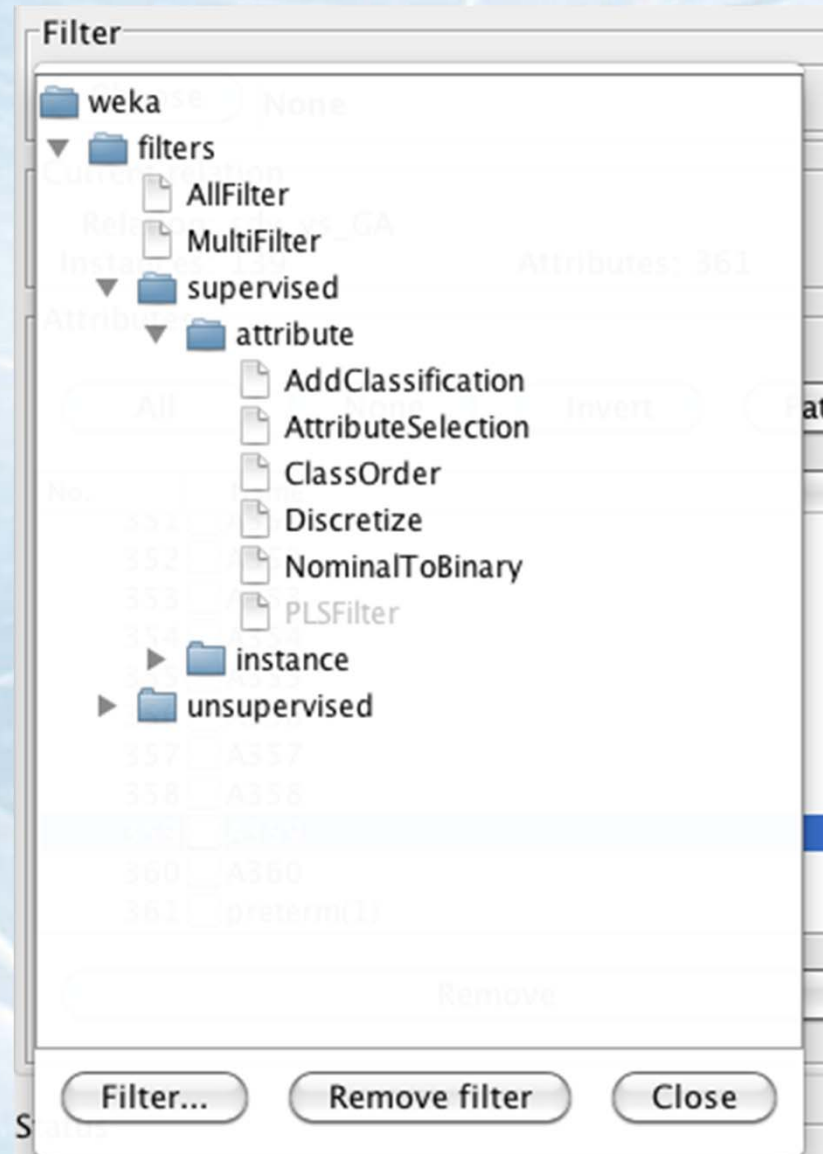
Selected attribute: Name: A359 Type: Numeric Missing: 0 (0%) Distinct: 111 Unique: 87 (63%)

Statistic	Value
Minimum	-155.08
Maximum	201.12
Mean	-0.456
StdDev	71.818

Class: preterm(1) (Nom) Visualize All

Bin	preterm(1) (Blue)	Other (Red)
1	13	0
2	17	0
3	38	0
4	24	0
5	7	0
6	2	0

# Weka





# Weka

The screenshot shows the Weka Explorer application window. The title bar reads "Weka Explorer". The main menu bar includes "Preprocess", "Classify", "Cluster" (which is selected), "Associate", "Select attributes", and "Visualize".

The "Clusterer" section contains a "Choose" button and a text field with the command: `EM -I 100 -N -1 -M 1.0E-6 -S 100`.

The "Cluster mode" section has several options:

- Use training set
- Supplied test set (with a "Set..." button)
- Percentage split (with a percentage field set to 66)
- Classes to clusters evaluation (with a dropdown menu showing "(Nom) preterm(1)")
- Store clusters for visualization

Below these options are buttons for "Ignore attributes", "Start", and "Stop".

The "Clusterer output" section displays the following text:

```
=== Run information ===  
Scheme:      weka.clusterers.EM -I 100 -N -1 -M 1.0E-6 -S 100  
Relation:    sdv_vs_GA  
Instances:   139  
Attributes:  361  
              [list of attributes omitted]  
Test mode:   split 66% train, remainder test
```

The "Result list (right-click for options)" section shows a list of results:

- 02:02:05 - Cobweb
- 02:02:35 - EM (highlighted)

The "Status" bar at the bottom indicates "Building model on training data...". To the right of the status bar are a "Log" button, a small bird icon, and the text "x 1".

# Weka

Weka Explorer

Preprocess | Classify | Cluster | Associate | **Select attributes** | Visualize

Attribute Evaluator  
Choose CfsSubsetEval

Search Method  
Choose BestFirst -D 1 -N 5

Attribute Selection Mode  
 Use full training set  
 Cross-validation Folds 10 Seed 1

(Nom) preterm(1)

Start Stop

Result list (right-click for options)  
02:10:05 - BestFirst + CfsSubsetEval

Attribute selection output

```
=== Run information ===
Evaluator:   weka.attributeSelection.CfsSubsetEval
Search:     weka.attributeSelection.BestFirst -D 1 -N 5
Relation:   sdv_vs_GA
Instances:  139
Attributes: 361
            [list of attributes omitted]
Evaluation mode: 10-fold cross-validation

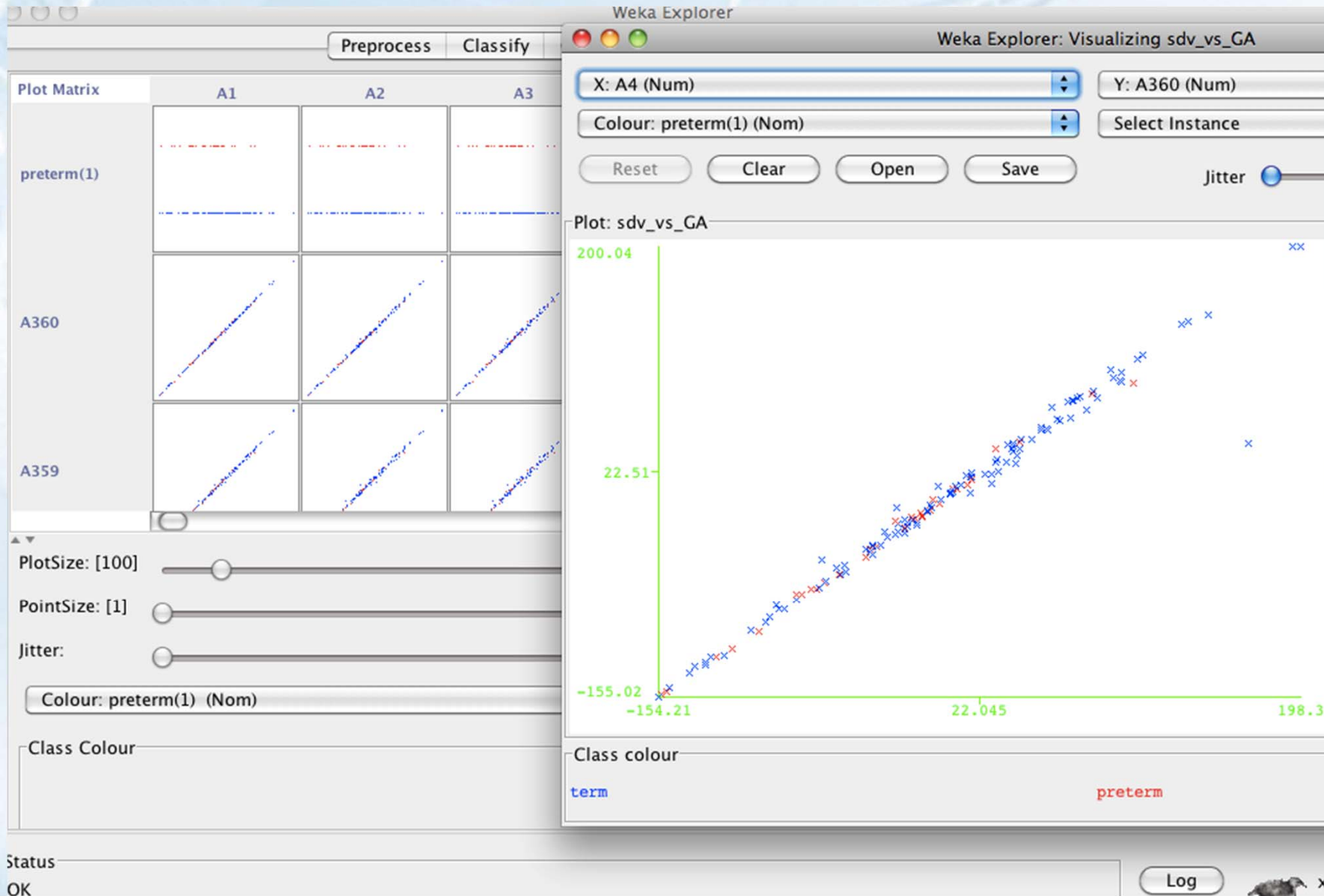
=== Attribute selection 10 fold cross-validation (stratified), seed: 1 ===
```

number of folds (%)	attribute
0( 0 %)	1 A1
0( 0 %)	2 A2
0( 0 %)	3 A3
0( 0 %)	4 A4
0( 0 %)	5 A5
0( 0 %)	6 A6
0( 0 %)	7 A7
4( 40 %)	8 A8
0( 0 %)	9 A9
0( 0 %)	10 A10
0( 0 %)	11 A11
0( 0 %)	12 A12
4( 40 %)	13 A13
0( 0 %)	14 A14
0( 0 %)	15 A15
0( 0 %)	16 A16
0( 0 %)	17 A17
0( 0 %)	18 A18
0( 0 %)	19 A19

Status  
OK

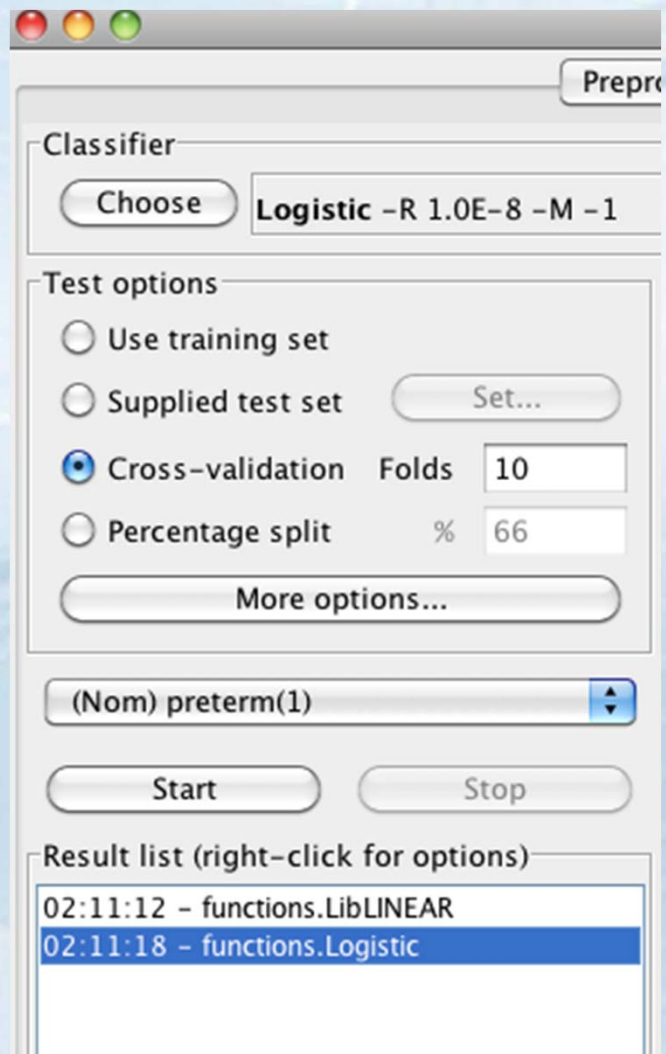
Log x 0

# Weka





# Weka



# Naïve Bayesian Classifier

Using Conditional Probability using Bayes's Theorem

$$p(C | F_1, \dots, F_n) = \frac{p(C) p(F_1, \dots, F_n | C)}{p(F_1, \dots, F_n)}.$$

in words:

$$\text{posterior} = \frac{\text{prior} \times \text{likelihood}}{\text{evidence}}.$$

Calculating the Conditional Probabilities

$p(F_1, \dots, F_n)$

$$p(C) p(F_1 | C) p(F_2 | C, F_1) p(F_3 | C, F_1, F_2) \dots p(F_n | C, F_1, F_2, F_3, \dots, F_{n-1}).$$

# Naïve Bayesian Classifier

Independence Assumption:

$$p(F_i|C, F_j) = p(F_i|C)$$

Deriving:

$$p(C, F_1, \dots, F_n) = p(C) p(F_1|C) p(F_2|C) p(F_3|C) \dots$$

$$= p(C) \prod_{i=1}^n p(F_i|C).$$

**Normalization constant (1/Z)**  
is the same for all C,  
so we can ignore it

$$p(F_1, \dots, F_n) = \frac{1}{Z} p(C) \prod_{i=1}^n p(F_i|C)$$

Classification Rule

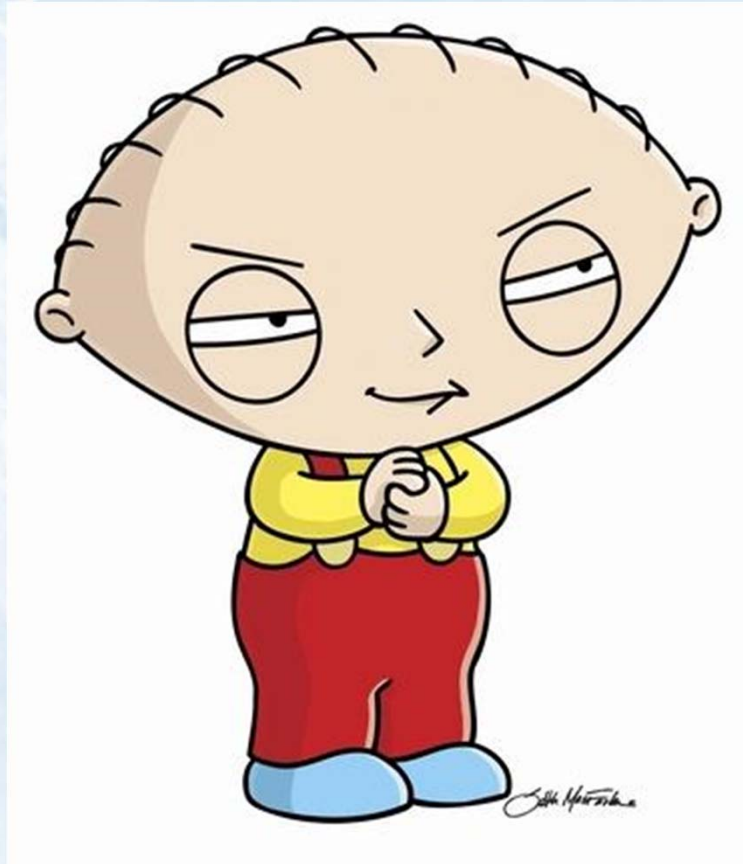
$$y(f_1, \dots, f_n) = \operatorname{argmax}_c p(C = c) \prod_{i=1}^n p(F_i = f_i|C = c).$$



# Naïve Bayesian Classifier

```
1 from math import log, exp
2
3 class BayesianClassifier(object):
4     def __init__(self):
5         self.observation_total = 0
6         self.observations = {}
7         self.observations_dual = {}
8         self.labels = {}
9         self.feats = {}
10
11     def train(self, event, evidence):
12         self.observations.setdefault(event, {})
13         self.labels.setdefault(event, 0)
14         self.observation_total += 1
15         self.labels[event] += 1
16         for v in evidence:
17             self.observations[event].setdefault(v, 0)
18             self.observations_dual.setdefault(v, {})
19             self.observations_dual[v].setdefault(event, 0)
20             self.feats.setdefault(v, 0)
21             self.observations[event][v] += 1
22             self.feats[v] += 1
23             self.observations_dual[v][event] += 1
24
25     def classify(self, evidence, complement=False):
26         estimates = [(self.cond_comp(event, evidence, complement), event)
27                     for event in self.labels]
28         highest_prob, likely_event = max(estimates)
29         estimates = [(exp(prob - highest_prob), event)
30                     for prob, event in estimates]
31         highest_prob = 1 / sum(prob for prob, event in estimates)
32         estimates = [(highest_prob * prob, event)
33                     for prob, event in estimates]
34
35         return likely_event, highest_prob
36
37     def cond_comp(self, event, evidence, complement):
38         if event not in self.labels:
39             return 0
40         else:
41             #Using log probabilities to prevent underflow
42             P_event = log(self.labels[event]) - log(self.observation_total)
43             P_condt = 0
44             for Bk in evidence:
45                 if Bk not in self.feats: continue
46                 P_condt += ( log(self.observations[event].get(Bk,0) + 1) -
47                             log(self.labels[event]) )
48             if complement:
49                 for Bk in self.feats:
50                     if Bk in evidence: continue
51                     P_condt += ( log(self.labels[event] -
52                                 self.observations[event].get(Bk, 0) + 1) -
53                                 log(self.labels[event]) )
54             return P_event + P_condt
55
```

# Questions? Comments?



STEWIE GRIFFIN WALLPAPER  
CREATED BY ZAC MARTIN

VERSION 1 - INCLUDES "WHAT THE DEUCE" TEXT  
VERSION 2 - DOES NOT INCLUDE "WHAT THE DEUCE" TEXT

WHAT THE DEUCE?

