

pomegranate

fast and flexible probabilistic modelling in python

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Acknowledgements



UNIVERSITY of WASHINGTON
eScience Institute
ADVANCING DATA-INTENSIVE DISCOVERY IN ALL FIELDS





Overview

pomegranate is **more flexible** than other packages, **faster**, is **intuitive to use**, and can do it all **in parallel**



Overview: this talk

Overview

Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Finale: Train a mixture of HMMs in parallel



Overview: supported models

Six Main Models:

1. Probability Distributions
2. General Mixture Models
3. Markov Chains
4. Hidden Markov Models
5. Bayes Classifiers / Naive Bayes
6. Bayesian Networks

Two Helper Models:

1. k-means++/kmeans||
2. Factor Graphs



Overview: model stacking in pomegranate

Distributions

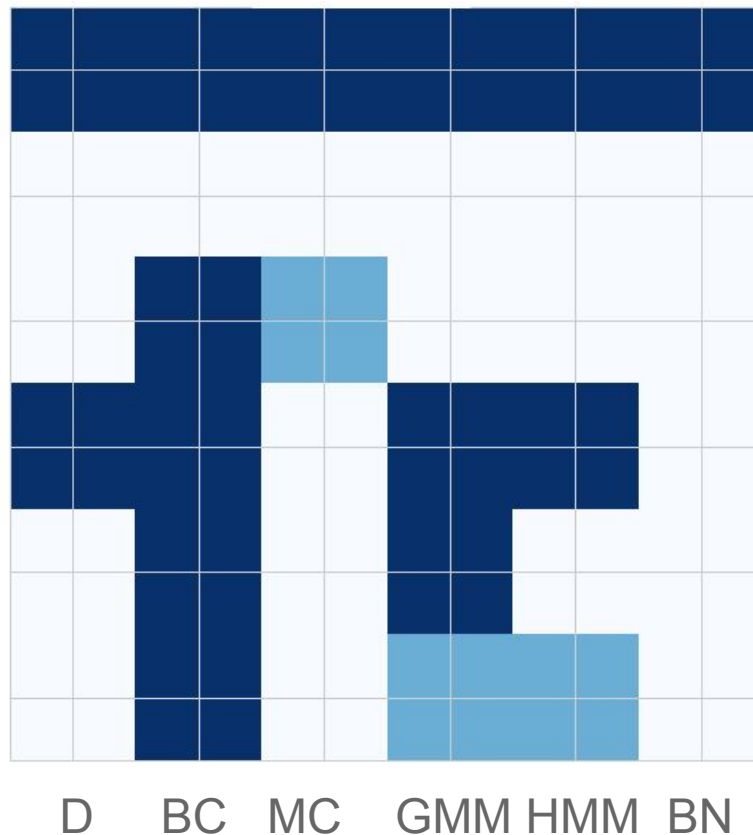
Bayes Classifiers

Markov Chains

General Mixture Models

Hidden Markov Models

Bayesian Networks





Overview: model stacking in pomegranate

Distributions

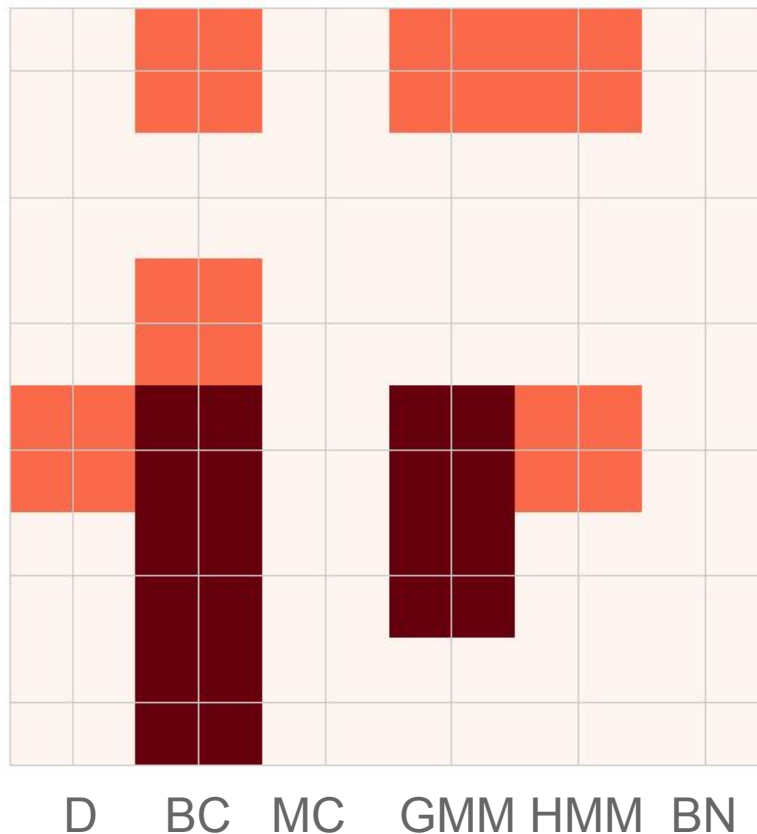
Bayes Classifiers

Markov Chains

General Mixture Models

Hidden Markov Models

Bayesian Networks





The API is common to all models

`model.log_probability(X) / model.probability(X)`

`model.sample()`

`model.fit(X, weights, inertia)`

All models have these methods!

`model.summarize(X, weights)`

`model.from_summaries(inertia)`

`model.predict(X)`

`model.predict_proba(X)`

`model.predict_log_proba(X)`

`Model.from_samples(X, weights)`

All models composed of distributions (like GMM, HMM...) have these methods too!



pomegranate supports many models

Univariate Distributions

1. UniformDistribution
2. BernoulliDistribution
3. NormalDistribution
4. LogNormalDistribution
5. ExponentialDistribution
6. BetaDistribution
7. GammaDistribution
8. DiscreteDistribution
9. PoissonDistribution

Kernel Densities

1. GaussianKernelDensity
2. UniformKernelDensity
3. TriangleKernelDensity

Multivariate Distributions

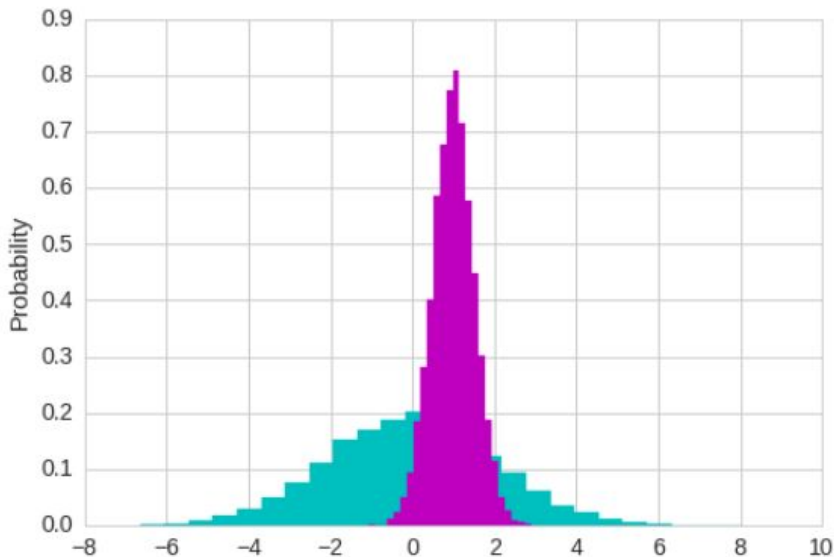
1. IndependentComponentsDistribution
2. MultivariateGaussianDistribution
3. DirichletDistribution
4. ConditionalProbabilityTable
5. JointProbabilityTable



Models can be created from known values

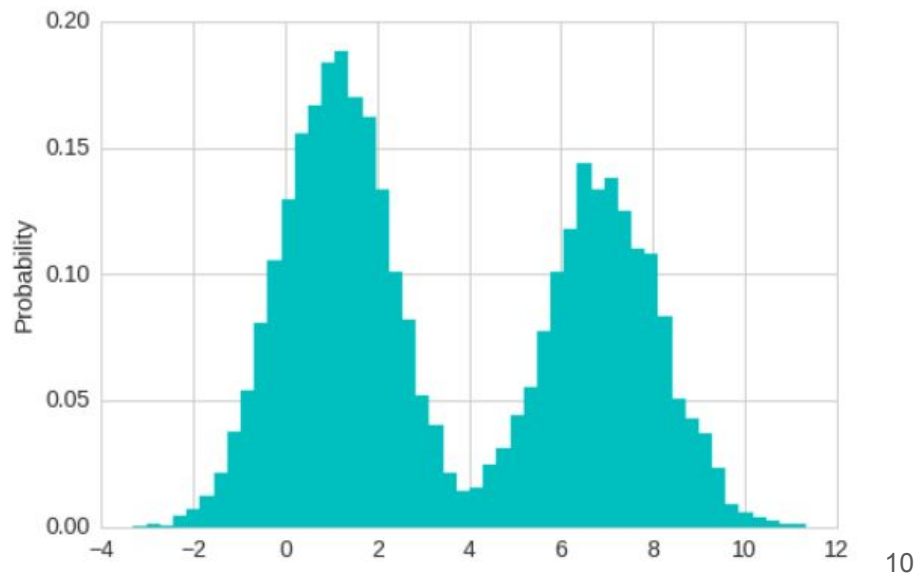
`mu, sig = 0, 2`

`a = NormalDistribution(mu, sig)`



`X = [0, 1, 1, 2, 1.5, 6, 7, 8, 7]`

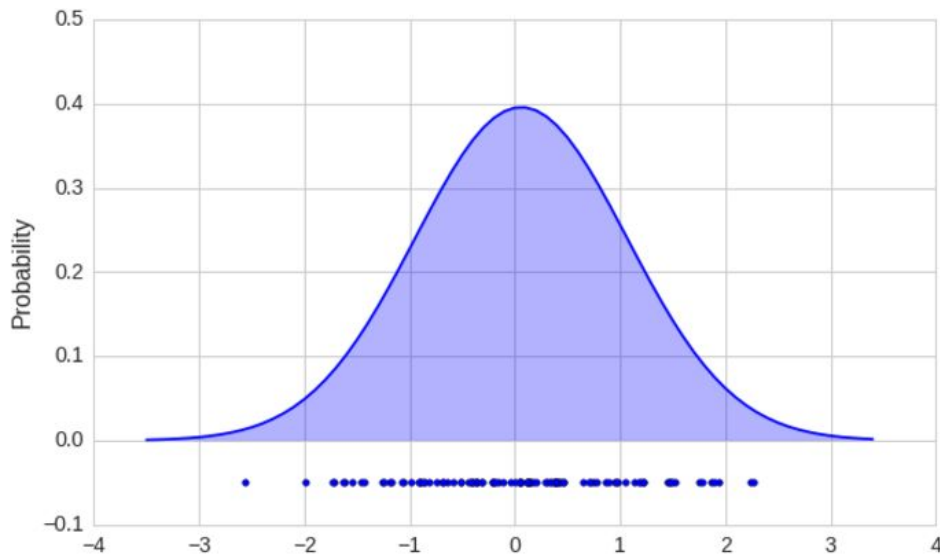
`a = GaussianKernelDensity(X)`





Models can be learned from data

```
X = numpy.random.normal(0, 1, 100)  
a = NormalDistribution.from_samples(X)
```





pomegranate can be faster than numpy

Fitting a Normal Distribution to 1,000 samples

```
data = numpy.random.randn(1000)

print "numpy time:"
%timeit -n 100 data.mean(), data.std()

print
print "pomegranate time:"
%timeit -n 100 NormalDistribution.from_samples(data)
```

numpy time:
100 loops, best of 3: 46.6 μ s per loop

pomegranate time:
100 loops, best of 3: 22.2 μ s per loop



pomegranate can be faster than numpy

Fitting Multivariate Gaussian to 10,000,000 samples of 10 dimensions

```
data = numpy.random.randn(10000000, 10)

print "numpy time:"
%timeit -n 10 data.mean(), numpy.cov(data.T)
print
print "pomegranate time:"
%timeit -n 10 MultivariateGaussianDistribution.from_samples(data)
```

numpy time:
10 loops, best of 3: 1.02 s per loop

pomegranate time:
10 loops, best of 3: 799 ms per loop



pomegranate uses BLAS internally

```
from scipy.linalg.cython_blas cimport dgemm
```

```
dgemm('N', 'T', &d, &d, &n, &alpha, y, &d, X, &d, &beta, pair_sum, &d)
```



pomegranate will soon have GPU support

[WIP] Add GPU support for models #270

 Open jmschrei wants to merge 1 commit into `master` from `GPU`

 Conversation 3

 Commits 1

 Files changed 3



jmschrei commented 12 days ago

Owner



This PR will add GPU support for all models, starting with multivariate gaussian distributions, GMMs, and HMMs, using the package `cupy`. Stay tuned!



pomegranate uses additive summarization

pomegranate reduces data to sufficient statistics for updates and so only has to go datasets once (for all models).

Here is an example of the Normal Distribution sufficient statistics

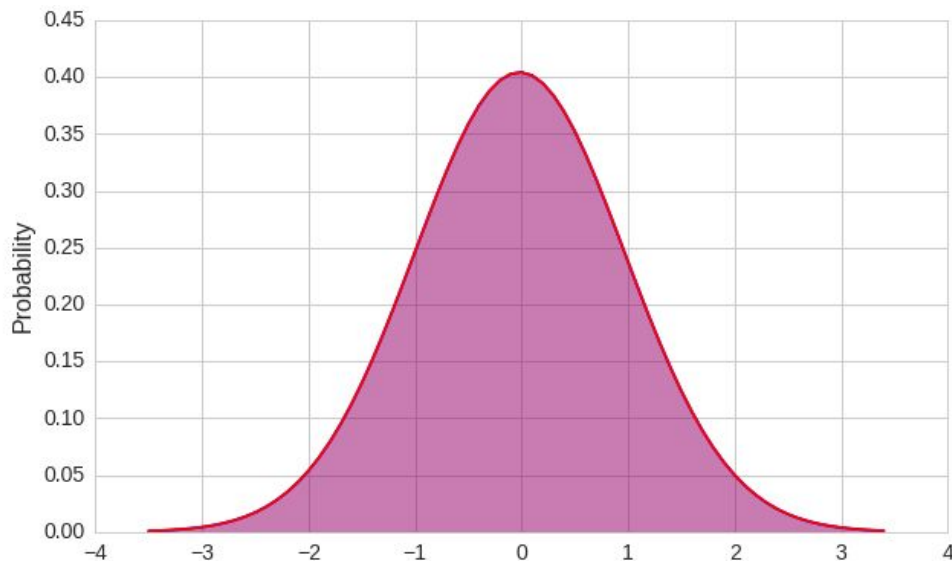
$$\sum_{i=1}^n w_i \quad \sum_{i=1}^n w_i x_i \quad \sum_{i=1}^n w_i x_i^2 \quad \longrightarrow \quad \begin{aligned} \mu &= \frac{\sum_{i=1}^n w_i x_i}{\sum_{i=1}^n w_i} \\ \sigma^2 &= \frac{\sum_{i=1}^n w_i x_i^2}{\sum_{i=1}^n w_i} - \frac{\left(\sum_{i=1}^n w_i x_i \right)^2}{\left(\sum_{i=1}^n w_i \right)^2} \end{aligned}$$



pomegranate supports out-of-core learning

Batches from a dataset can be reduced to additive summary statistics, enabling exact updates from data that can't fit in memory.

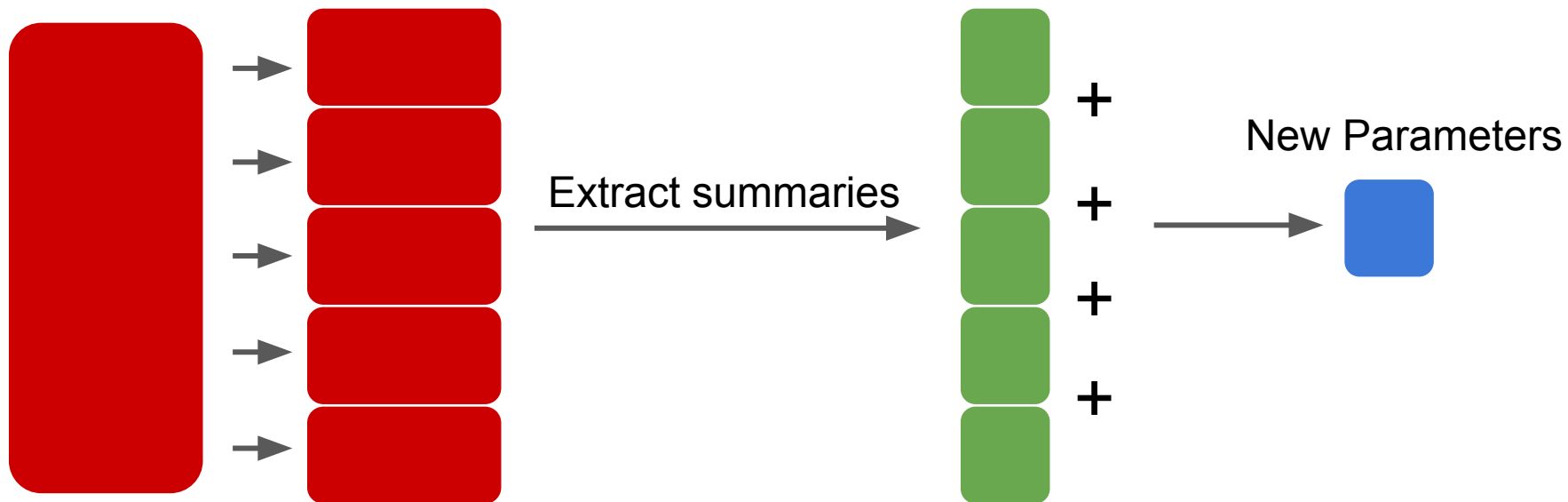
```
a.fit(data)
b.summarize(data[:1000])
b.summarize(data[1000:2000])
b.summarize(data[2000:3000])
b.summarize(data[3000:4000])
b.summarize(data[4000:])
b.from_summaries()
```



Fit Mean: -0.0174820965846, Fit STD: 0.986767322871
Summarize Mean: -0.0174820965846, Summarize STD: 0.986767322871



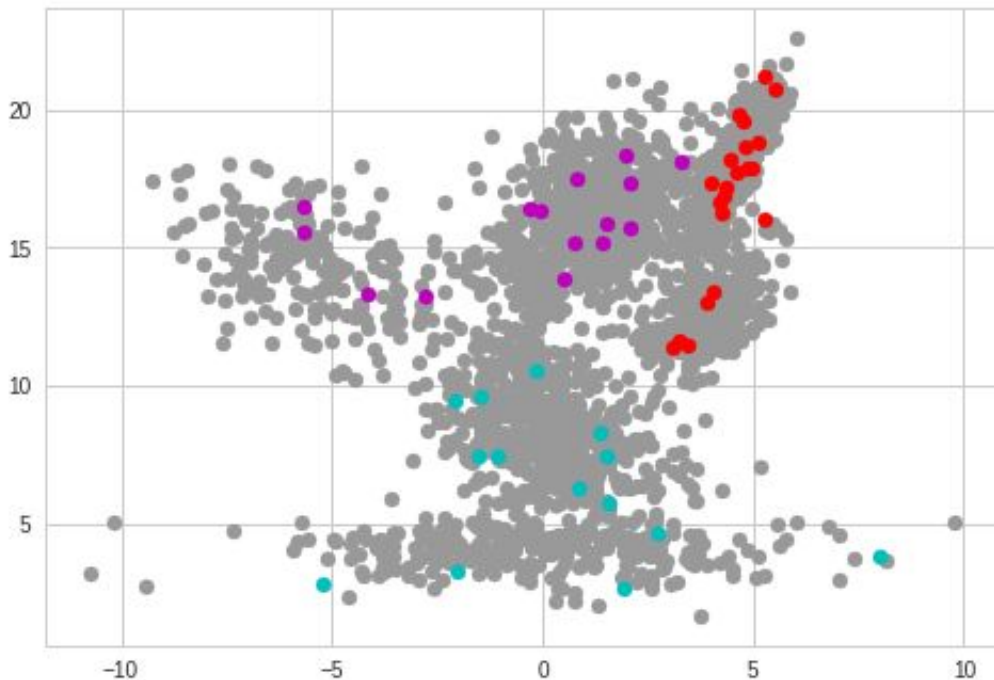
Parallelization exploits additive summaries





pomegranate supports semisupervised learning

Summary statistics from supervised models can be added to summary statistics from unsupervised models to train a single model on a mixture of labeled and unlabeled data.

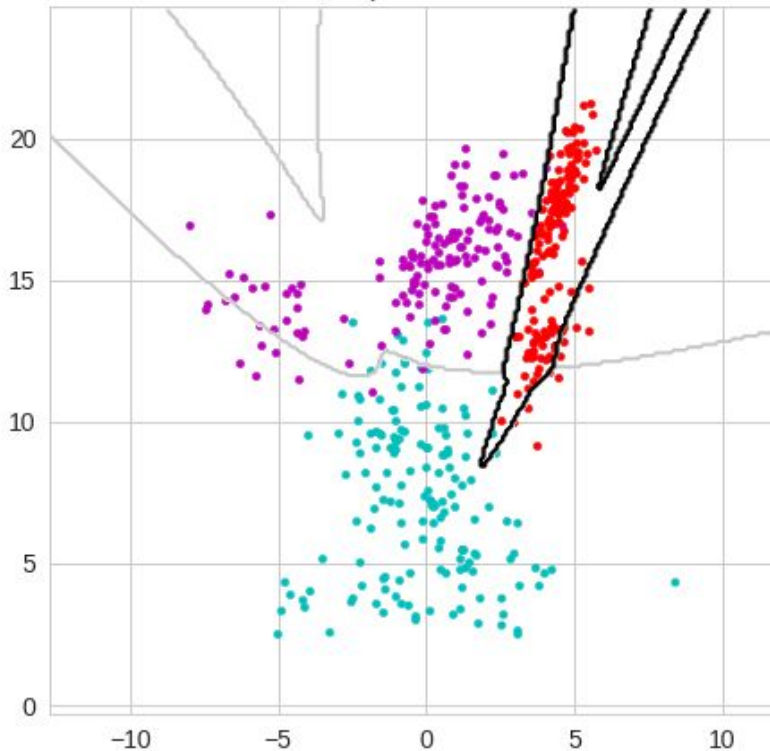




pomegranate supports semisupervised learning

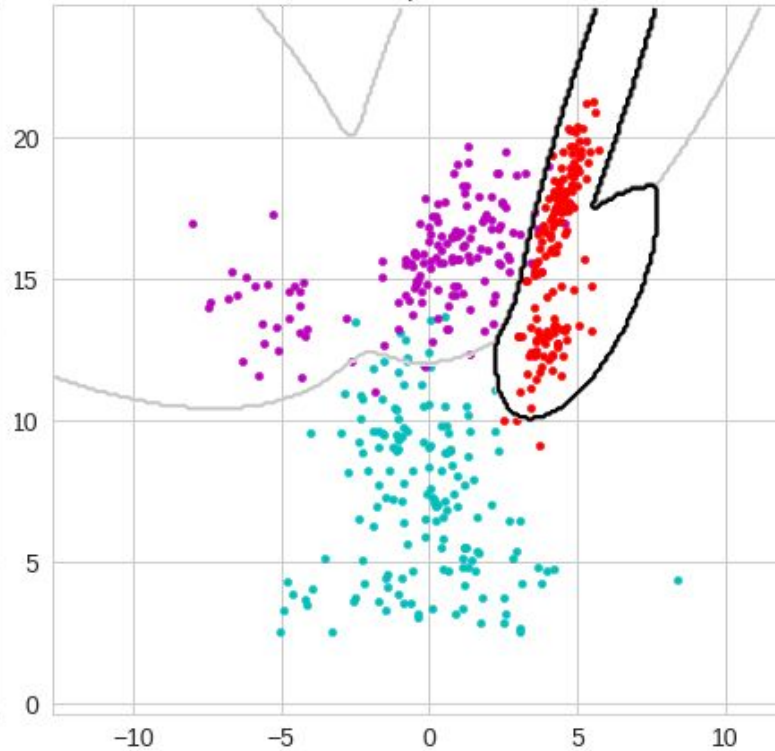
Supervised Accuracy: 0.93

Test Data, Supervised Boundaries



Semisupervised Accuracy: 0.96

Test Data, Semi-supervised Boundaries





pomegranate can be faster than scipy

```
mu, cov = numpy.random.randn(2000), numpy.eye(2000)
d = MultivariateGaussianDistribution(mu, cov)
X = numpy.random.randn(2000, 2000)
print "scipy time: ",
%timeit multivariate_normal.logpdf(X, mu, cov)
print "pomegranate time: ",
%timeit MultivariateGaussianDistribution(mu, cov).log_probability(X)
print "pomegranate time (w/ precreated object): ",
%timeit d.log_probability(X)
```

```
scipy time: 1 loop, best of 3: 1.67 s per loop
pomegranate time: 1 loop, best of 3: 801 ms per loop
pomegranate time (w/ precreated object): 1 loop, best of 3: 216 ms per loop
```



pomegranate uses aggressive caching

$$P(X|\mu, \sigma) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right)$$

$$\log P(X|\mu, \sigma) = -\log(\sqrt{2\pi}\sigma) - \frac{(x - \mu)^2}{2\sigma^2}$$

$$\log P(X|\mu, \sigma) = \alpha - \frac{(x - \mu)^2}{\beta}$$





Example 'blast' from Gossip Girl

Spotted: Lonely Boy. Can't believe the love of his life has returned. If only she knew who he was. But everyone knows Serena. And everyone is talking. Wonder what Blair Waldorf thinks. Sure, they're BFF's, but we always thought Blair's boyfriend Nate had a thing for Serena.



Example 'blast' from Gossip Girl

Why'd she leave? Why'd she return? Send me all the deets.
And who am I? That's the secret I'll never tell. The only one.
—XOXO. Gossip Girl.



How do we encode these 'blasts'?

Better lock it down with Nate, B. Clock's ticking.

+1 Nate

-1 Blair



How do we encode these 'blasts'?

This just in: S and B committing a crime of fashion. Who doesn't love a five-finger discount. Especially if it's the middle one.

-1 Blair

-1 Serena

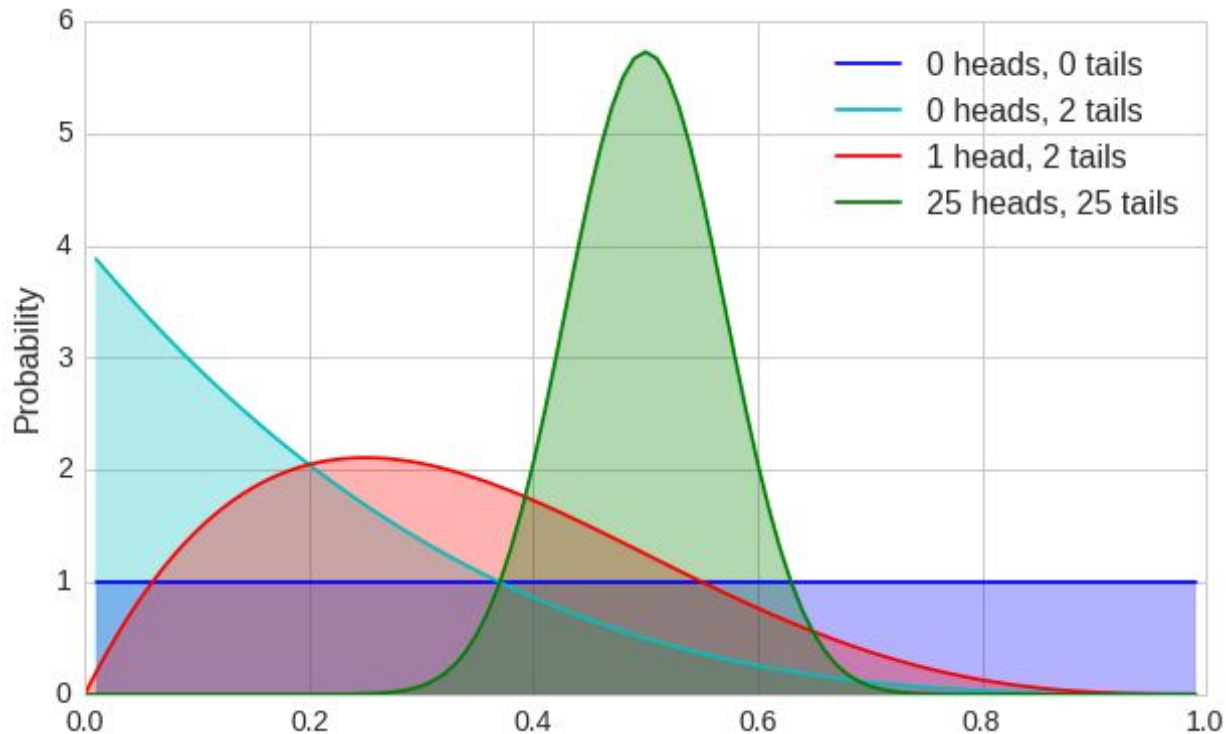


Simple summations don't work well



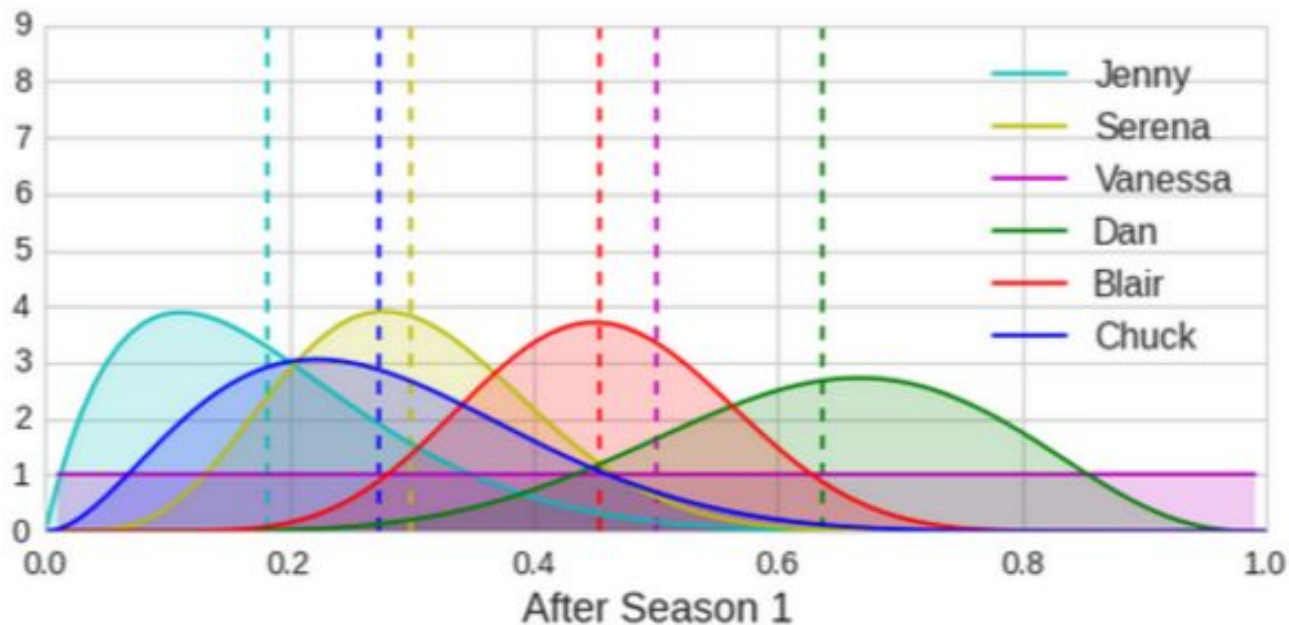


Beta distributions can model uncertainty



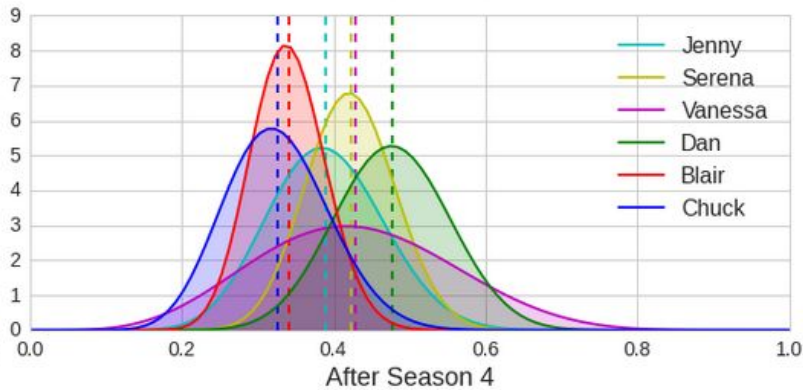
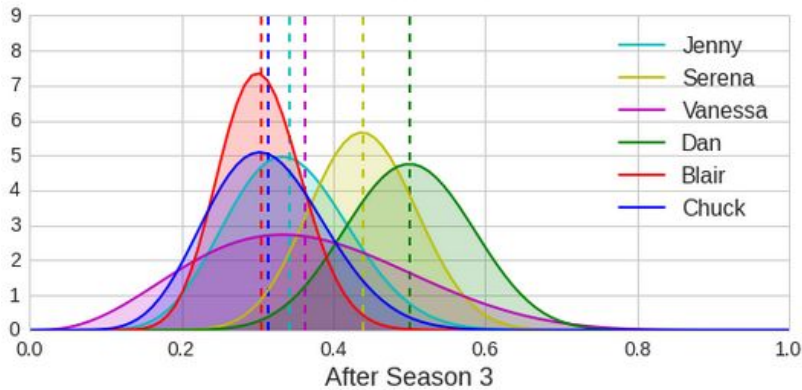
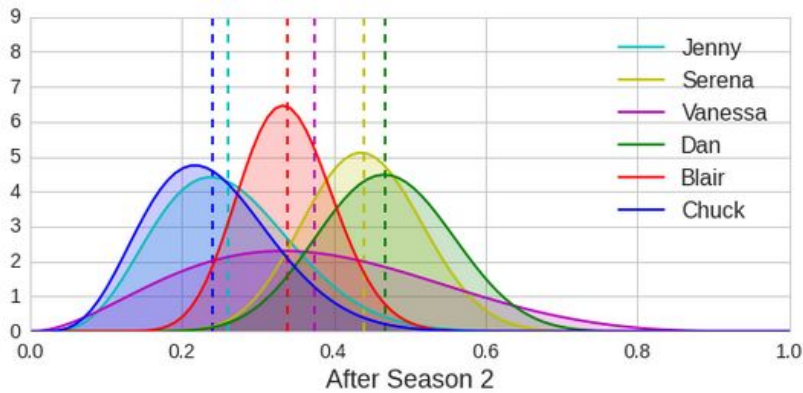
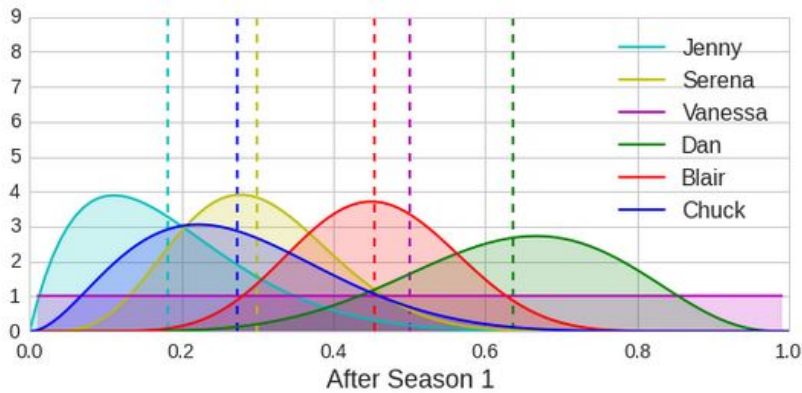


Beta distributions can model uncertainty





Beta distributions can model uncertainty





Overview: this talk

Overview

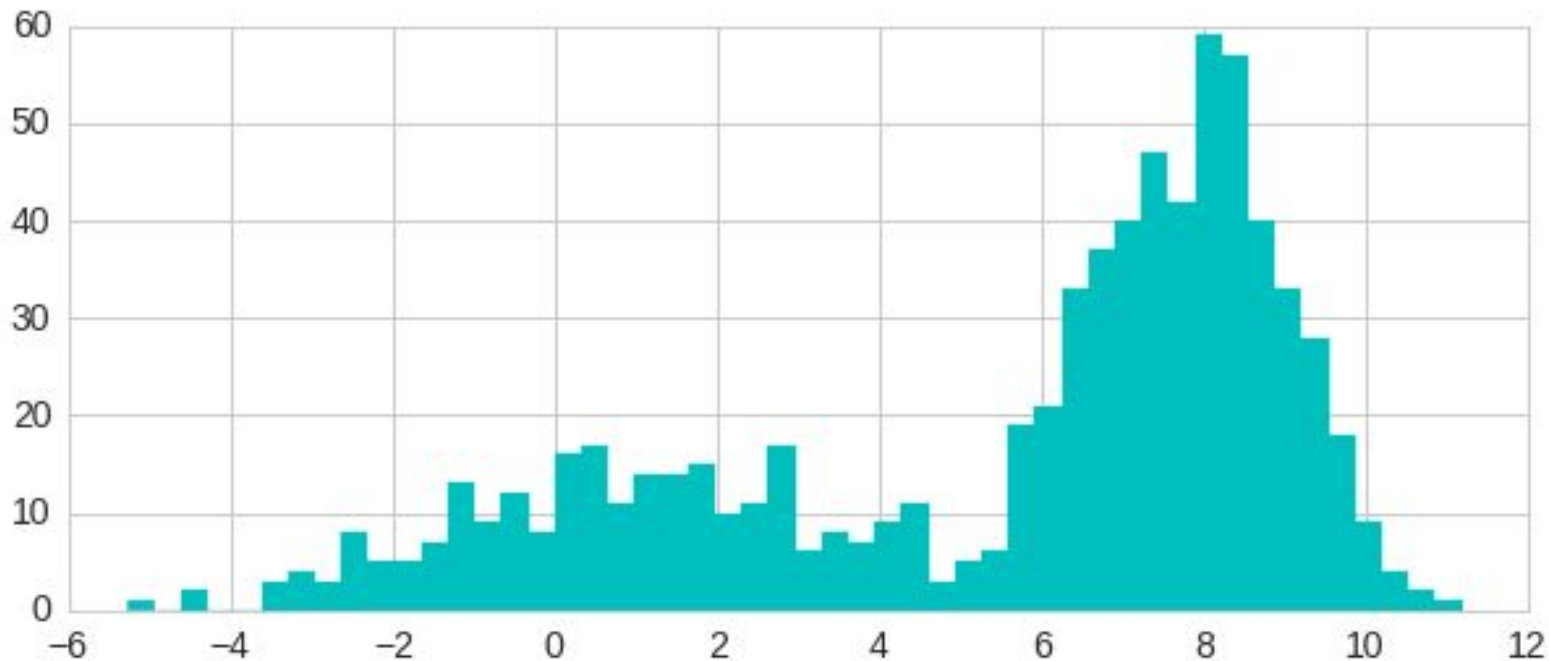
Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Finale: Train a mixture of HMMs in parallel

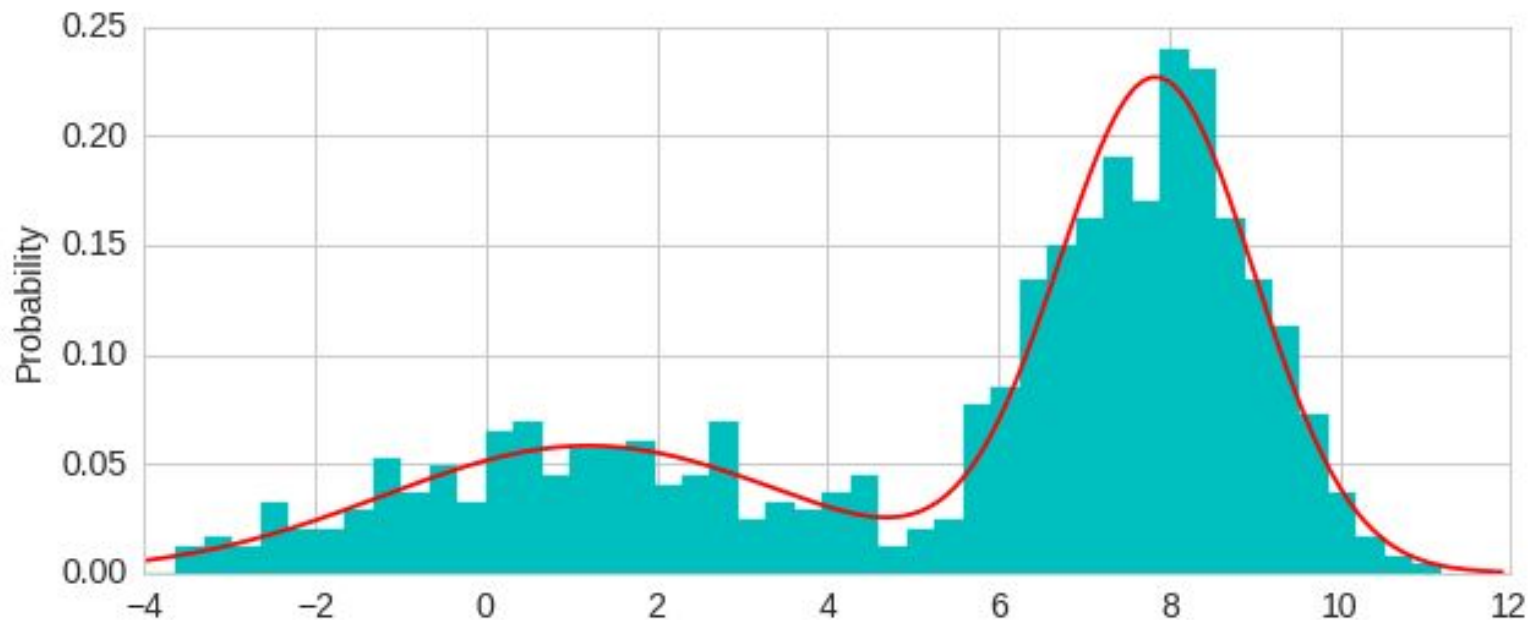


GMMs can model complex distributions





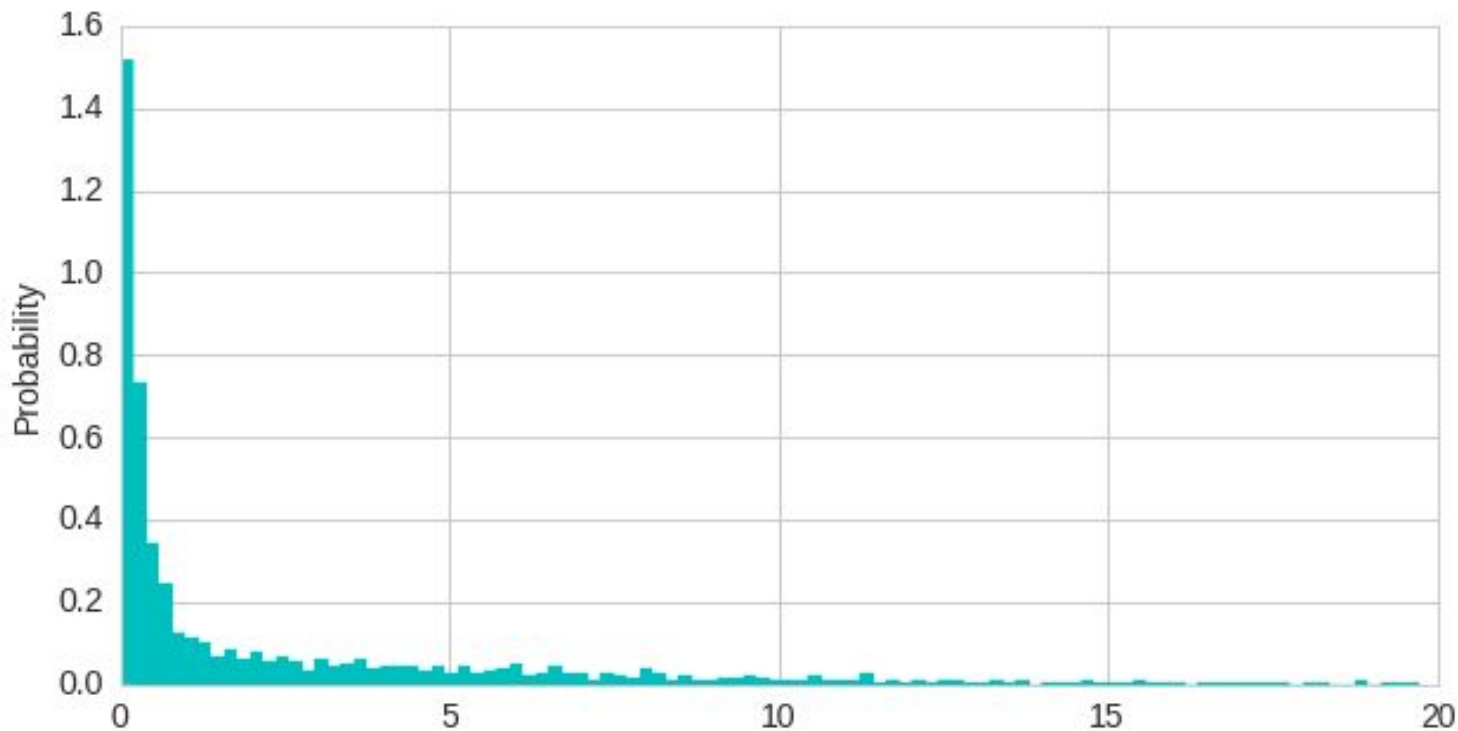
GMMs can model complex distributions



```
model = GeneralMixtureModel.from_samples(NormalDistribution, 2, X)
```

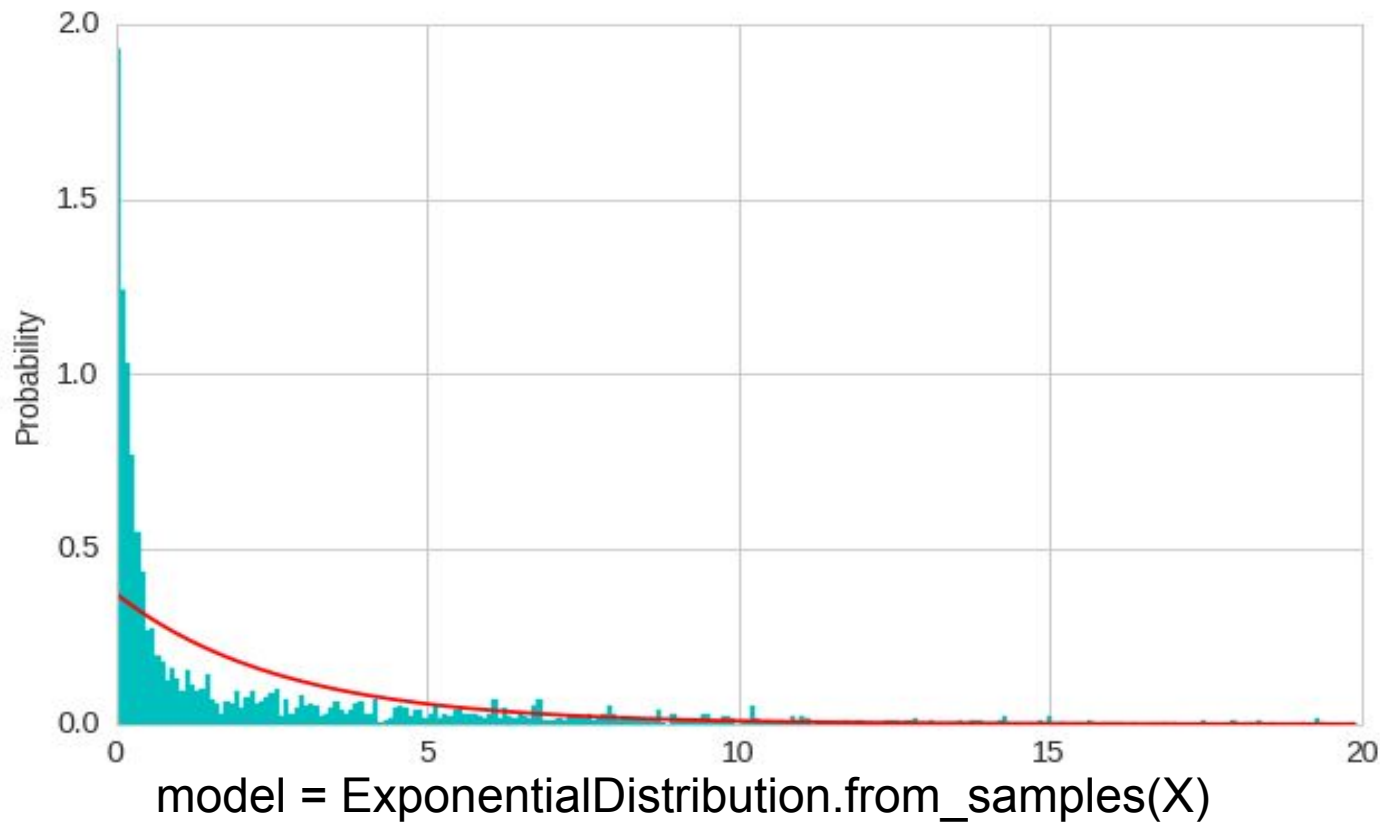


GMMs can model complex distributions



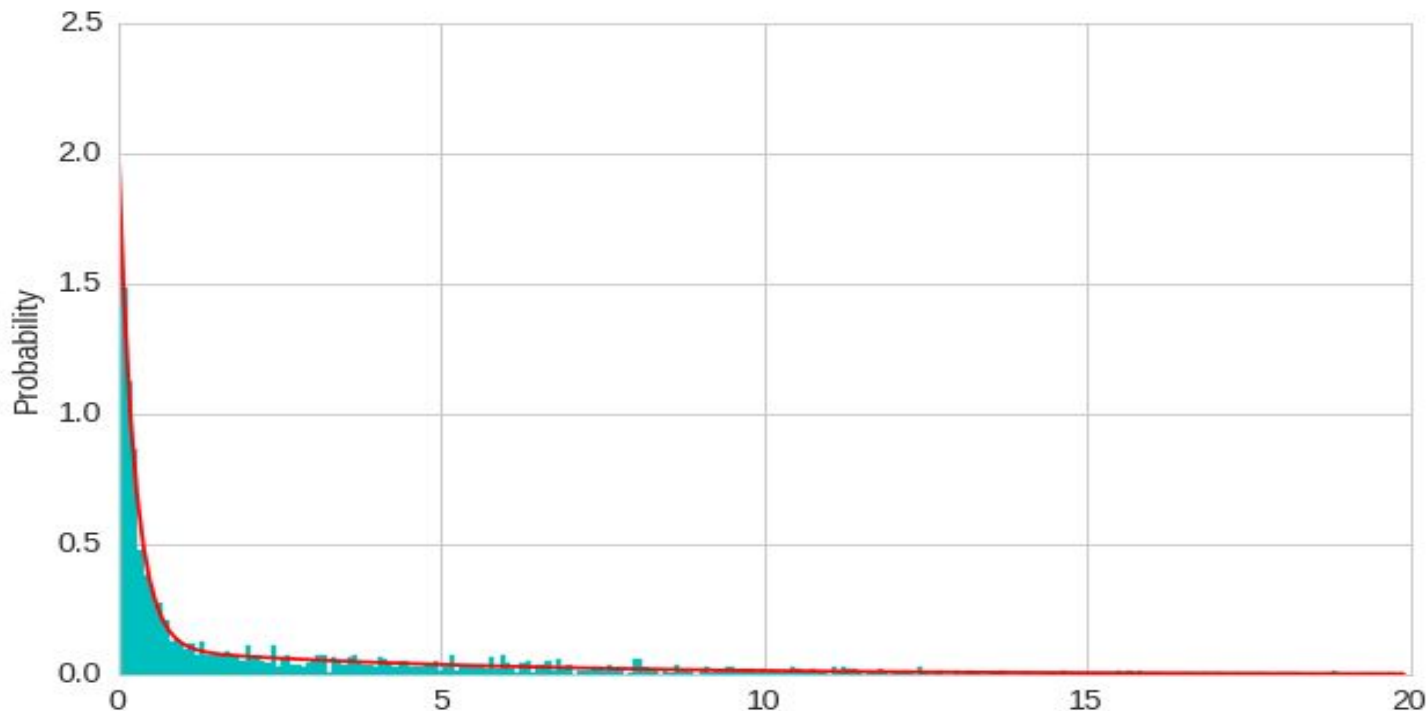


An exponential distribution is not right





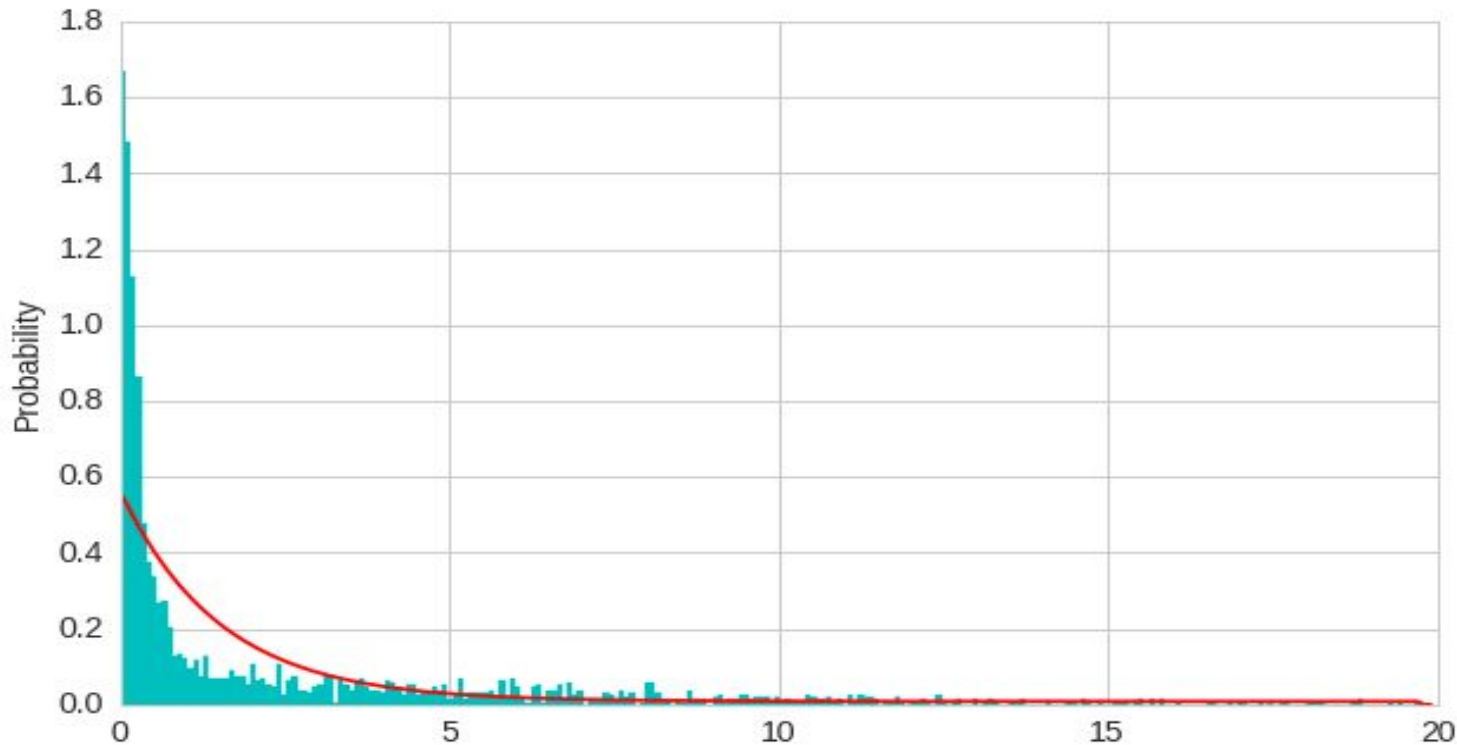
A mixture of exponentials is better



```
model = GeneralMixtureModel.from_samples(ExponentialDistribution, 2, X)
```



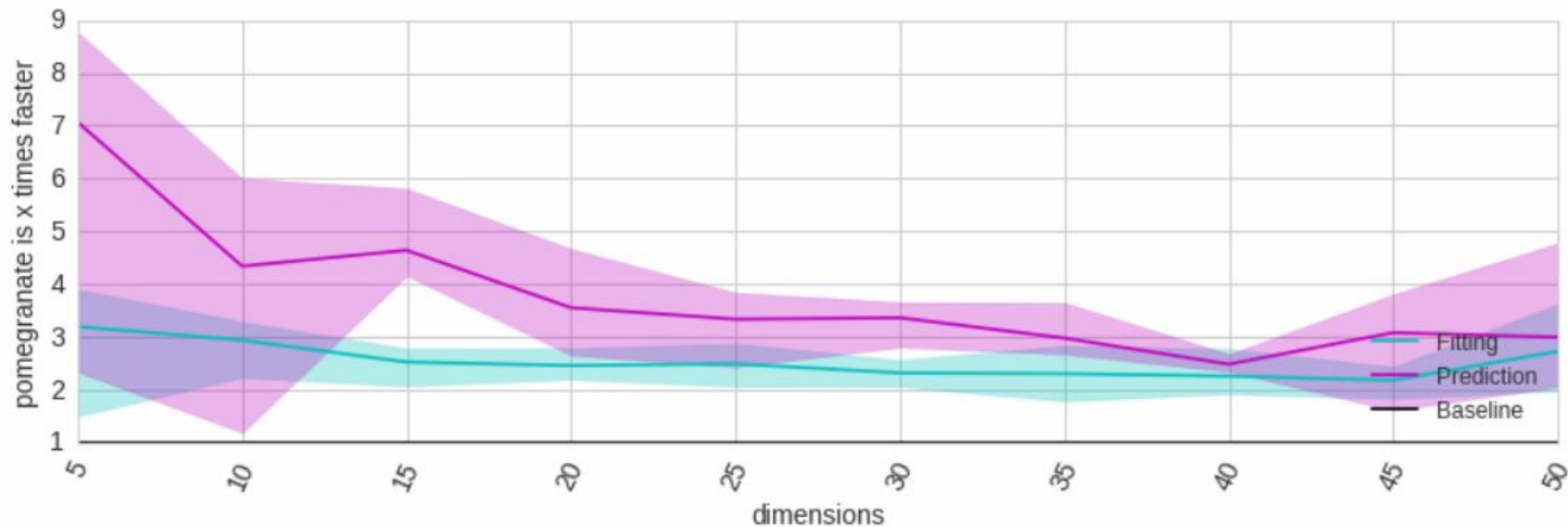
Heterogeneous mixtures natively supported



```
model = GeneralMixtureModel.from_samples([ExponentialDistribution, UniformDistribution], 2, X)
```



GMMs faster than sklearn





Overview: this talk

Overview

Major Models/Model Stacks

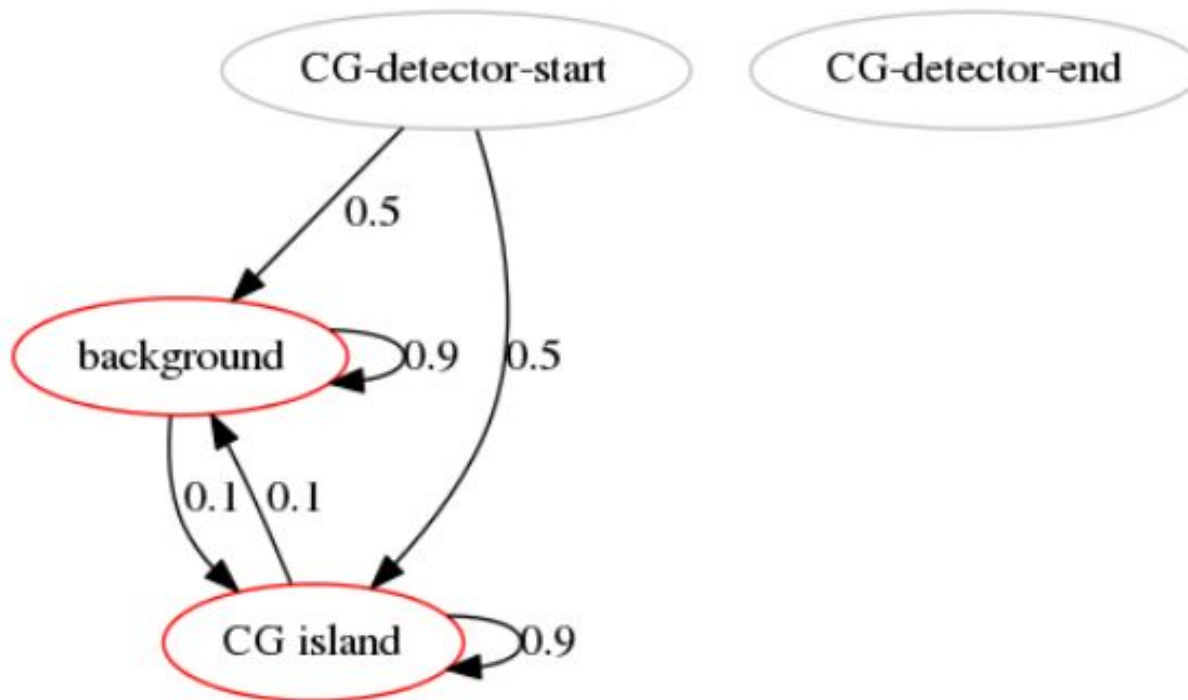
1. General Mixture Models
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Finale: Train a mixture of HMMs in parallel



CG enrichment detection HMM

GACTACGACT**CGCGCTCGCACGTCGCTCG**ACATCATCGACA





CG enrichment detection HMM

GACTACGACTCGCGCTCGCACGTCGCTCGACATCATCGACA

```
d1 = DiscreteDistribution({'A': 0.25, 'C': 0.25, 'G': 0.25, 'T': 0.25})
d2 = DiscreteDistribution({'A': 0.10, 'C': 0.40, 'G': 0.40, 'T': 0.10})

s1 = State(d1, name="background")
s2 = State(d2, name="CG island")

hmm = HiddenMarkovModel("CG-detector")
hmm.add_states(s1, s2)
hmm.add_transition(hmm.start, s1, 0.5)
hmm.add_transition(hmm.start, s2, 0.5)
hmm.add_transition(s1, s1, 0.9)
hmm.add_transition(s1, s2, 0.1)
hmm.add_transition(s2, s1, 0.1)
hmm.add_transition(s2, s2, 0.9)
hmm.bake()
```



pomegranate HMMs are feature rich

Feature	pomegranate	hmmlearn
Graph Structure		
Silent States	✓	
Optional Explicit End State	✓	
Sparse Implementation	✓	
Arbitrary Emissions Allowed on States	✓	
Discrete/Gaussian/GMM Emissions	✓	✓
Large Library of Other Emissions	✓	
Build Model from Matrices	✓	✓
Build Model Node-by-Node	✓	
Serialize to JSON	✓	
Serialize using Pickle/Joblib	✓	✓

Algorithms		
Priors		✓
Sampling	✓	✓
Log Probability Scoring	✓	✓
Forward-Backward Emissions	✓	✓
Forward-Backward Transitions	✓	
Viterbi Decoding	✓	✓
MAP Decoding	✓	✓
Baum-Welch Training	✓	✓
Viterbi Training	✓	
Labeled Training	✓	
Tied Emissions	✓	
Tied Transitions	✓	
Emission Inertia	✓	
Transition Inertia	✓	
Emission Freezing	✓	✓
Transition Freezing	✓	✓
Multi-threaded Training	✓	



GMM-HMM easy to define

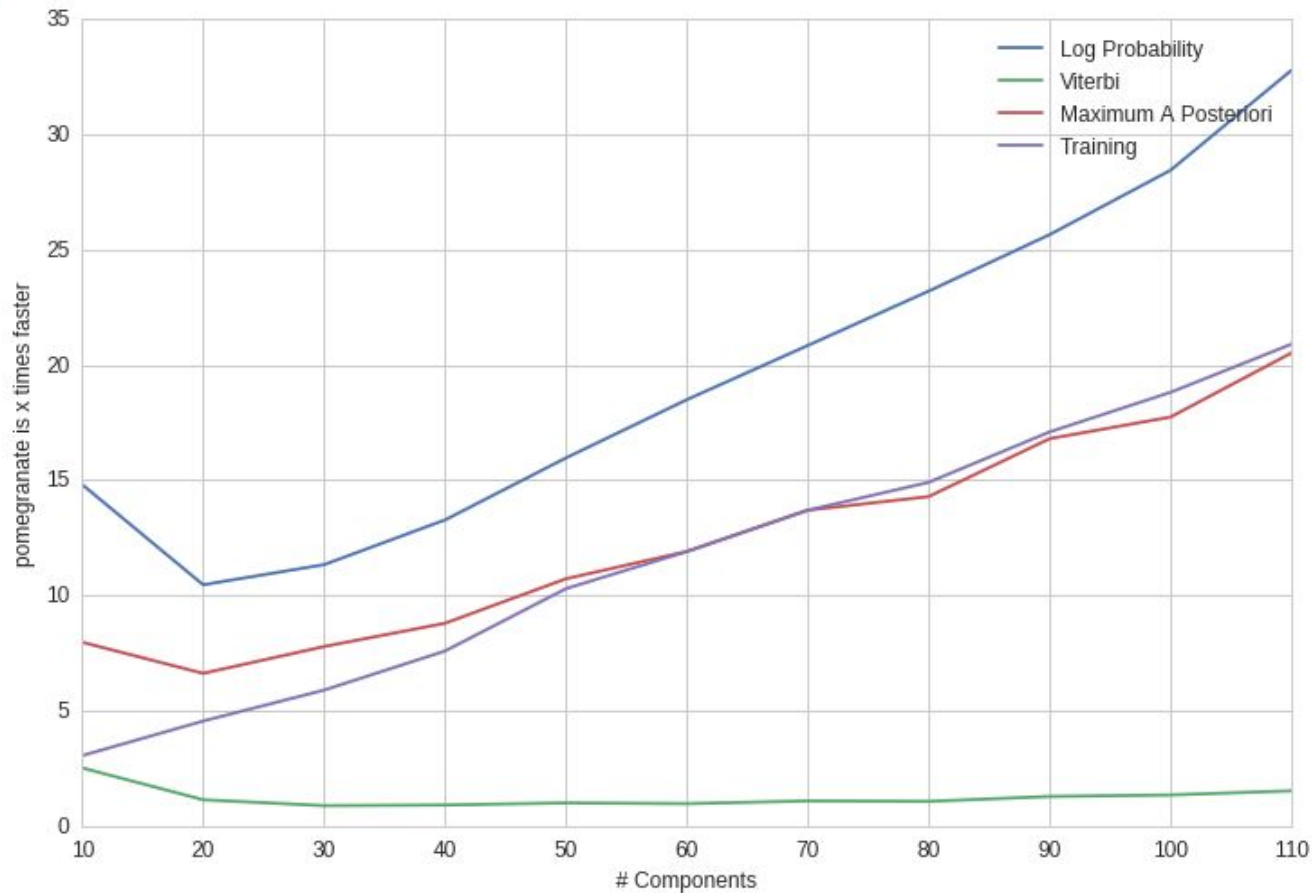
```
d1 = GeneralMixtureModel([NormalDistribution(5, 2), NormalDistribution(5, 4)])
d2 = GeneralMixtureModel([NormalDistribution(15, 1), NormalDistribution(15, 5)])

s1 = State(d1, name="GMM1")
s2 = State(d2, name="GMM2")

model = HiddenMarkovModel()
model.add_states(s1, s2)
model.add_transition(model.start, s1, 0.75)
model.add_transition(model.start, s2, 0.25)
model.add_transition(s1, s1, 0.85)
model.add_transition(s1, s2, 0.15)
model.add_transition(s2, s2, 0.90)
model.add_transition(s2, s1, 0.10)
model.bake()
```



HMMs are faster than hmmlearn





Overview: this talk

Overview

Major Models/Model Stacks

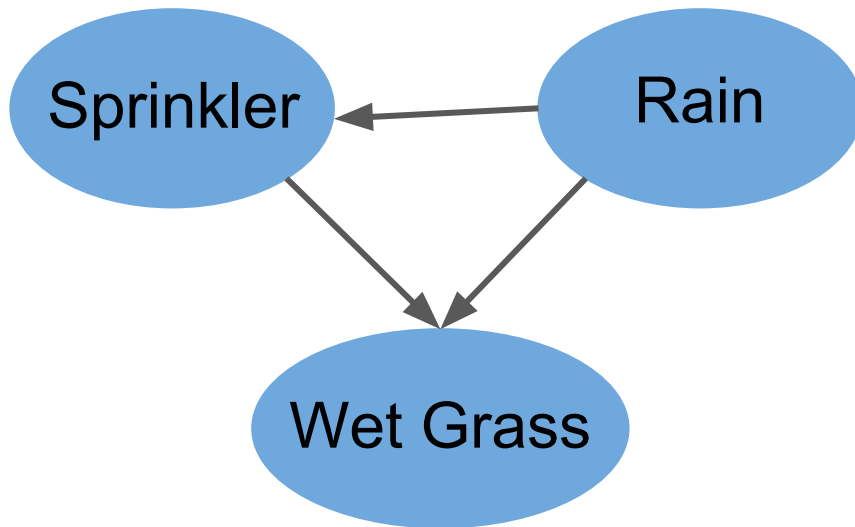
1. General Mixture Models
2. Hidden Markov Models
3. **Bayesian Networks**
4. Bayes Classifiers

Finale: Train a mixture of HMMs in parallel



Bayesian networks

Bayesian networks are powerful inference tools which define a dependency structure between variables.

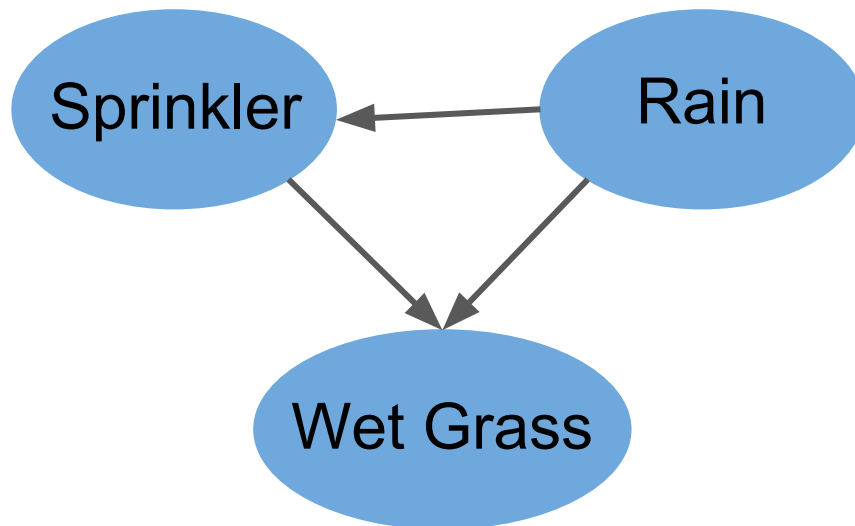




Bayesian networks

Two main difficult tasks:

- (1) Inference given incomplete information
- (2) Learning the dependency structure from data





Bayesian network structure learning

???

Three primary ways:

- “Search and score” / Exact
- “Constraint Learning” / PC
- Heuristics



Bayesian network structure learning

???

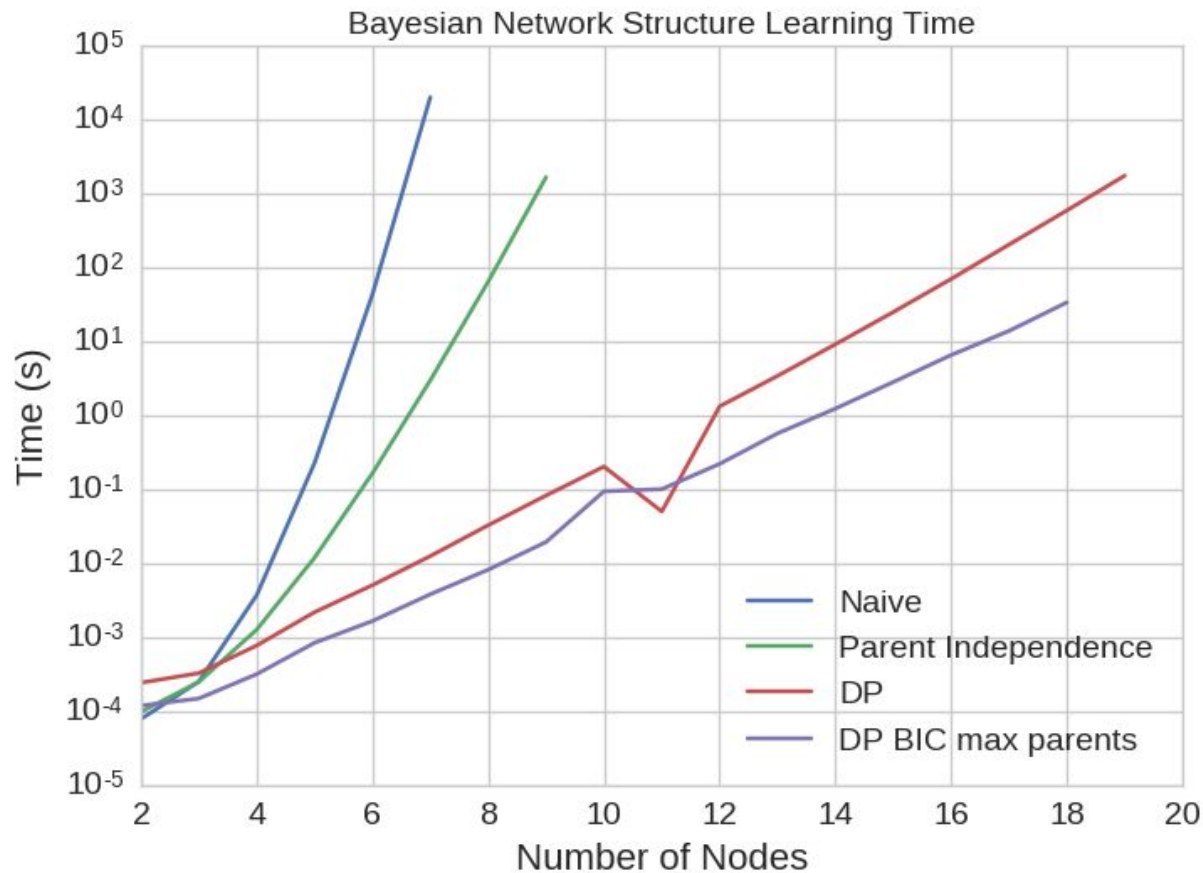
pomegranate supports:

- “Search and score” / Exact
- “Constraint Learning” / PC
- Heuristics



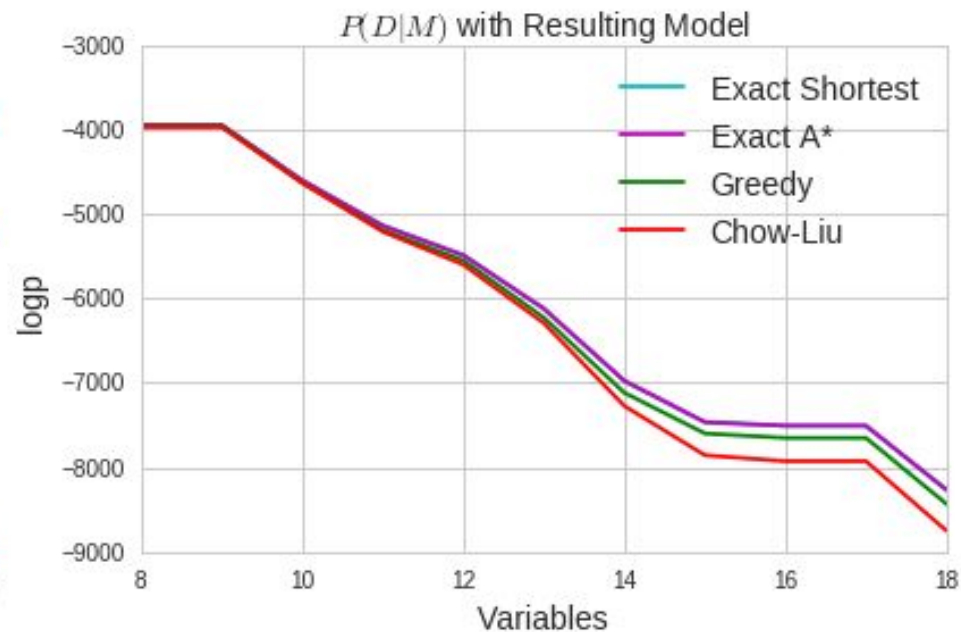
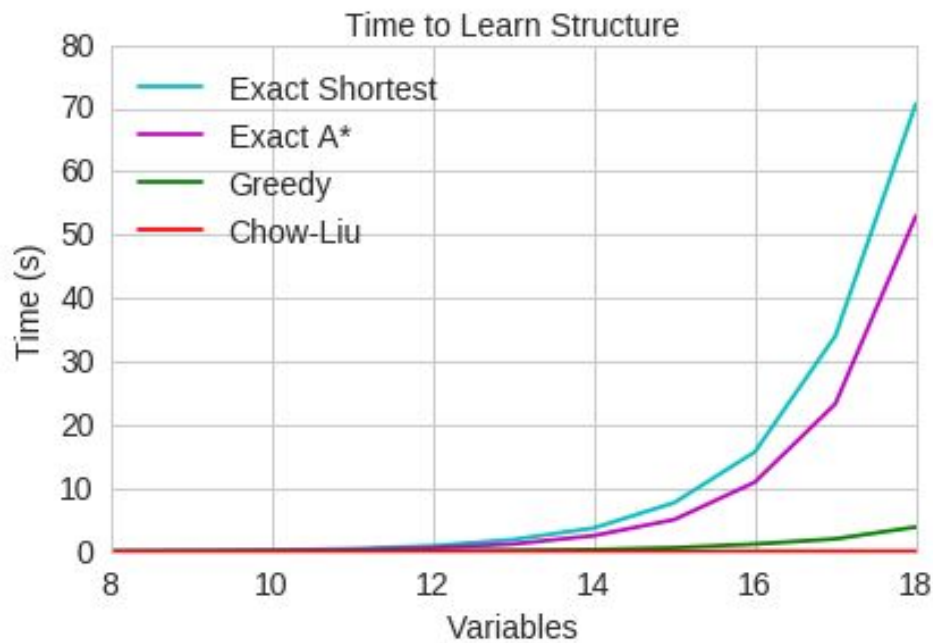
Exact structure learning is intractable

???



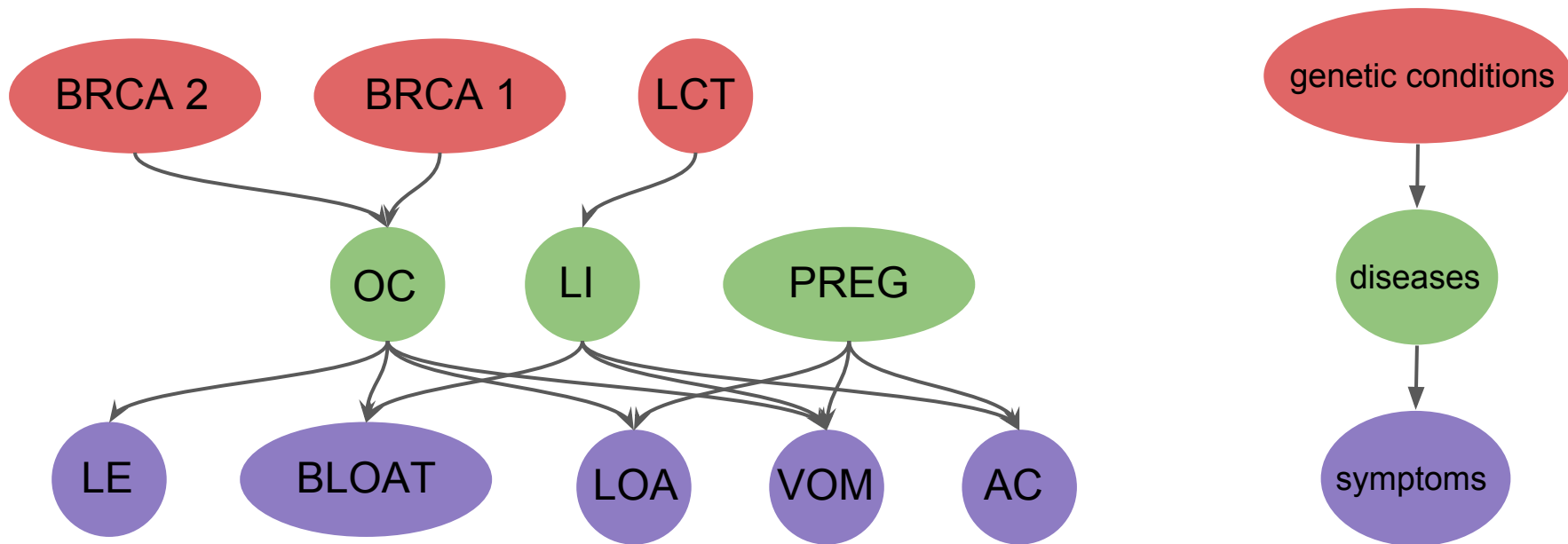


pomegranate supports four algorithms



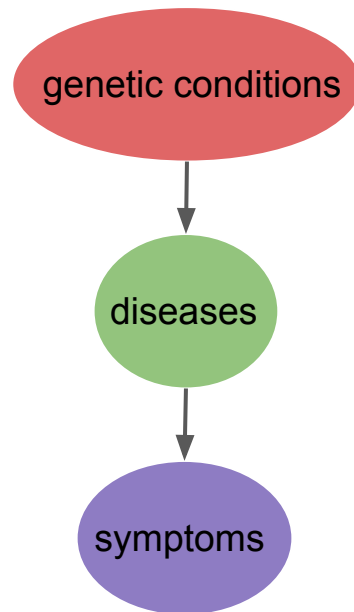
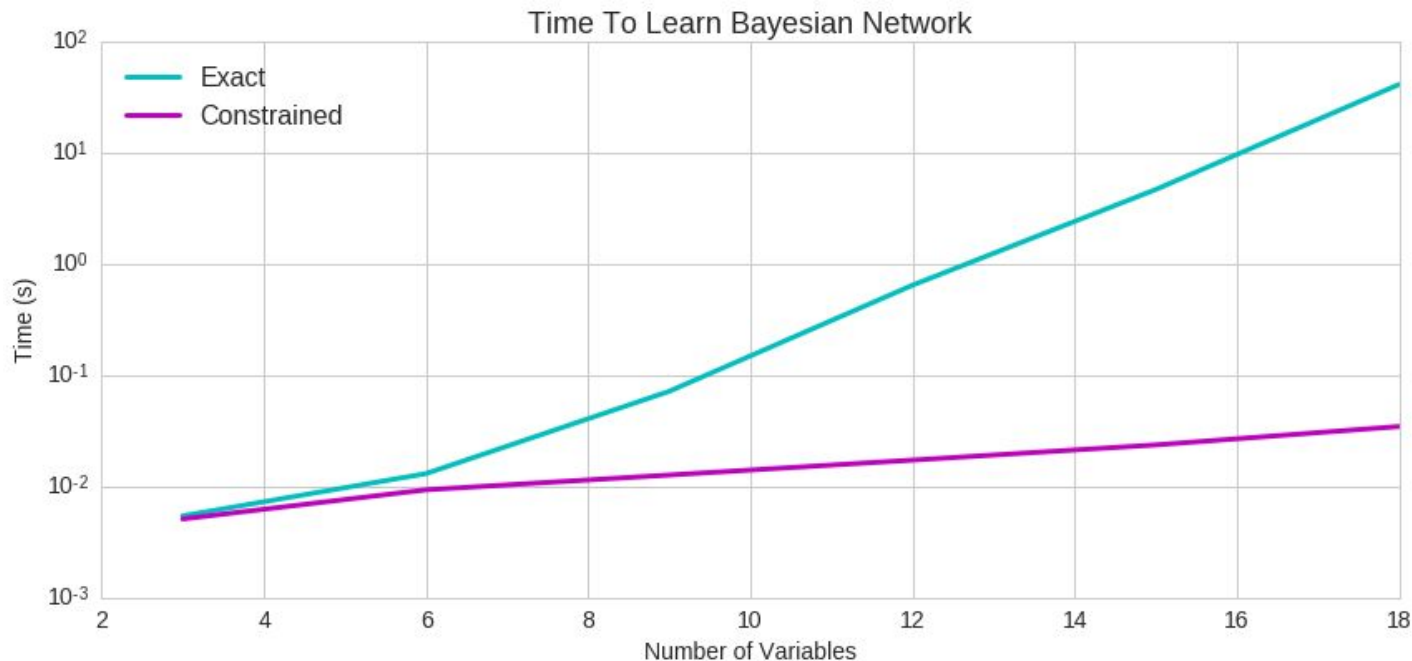


Constraint graphs merge data + knowledge



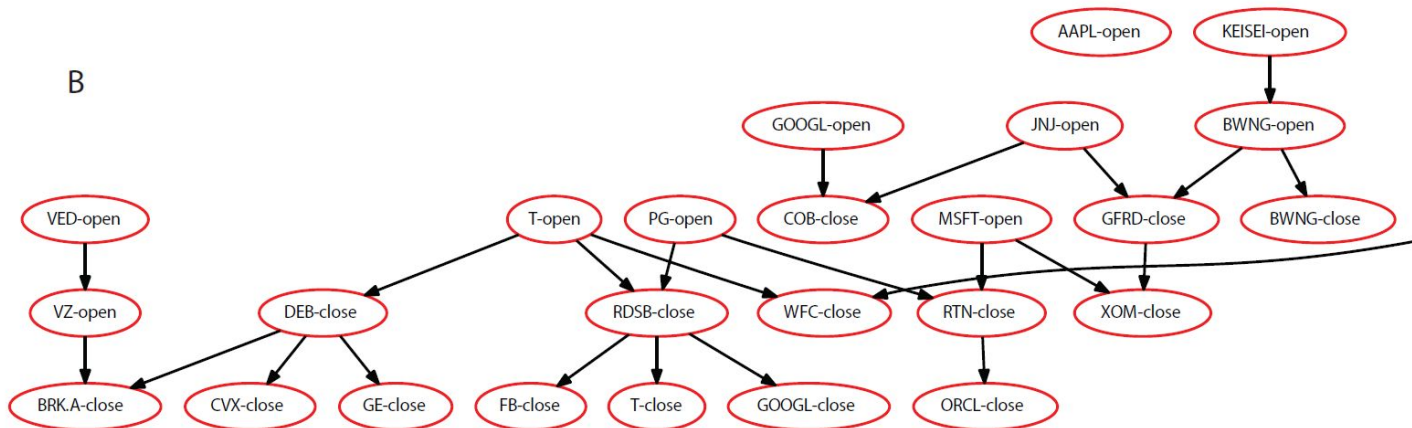
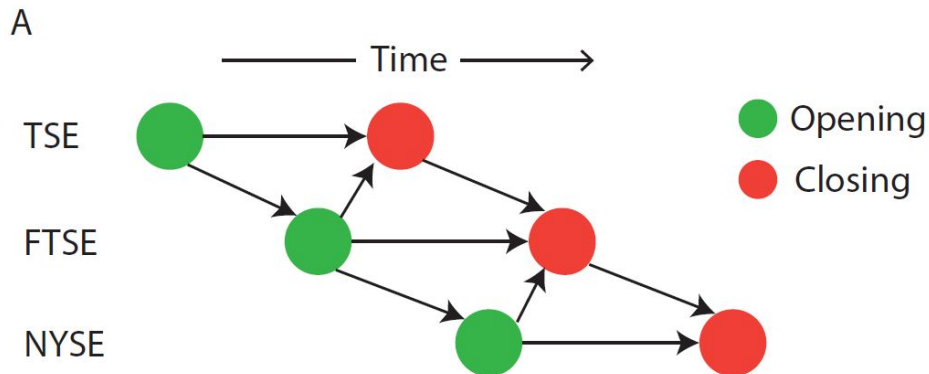


Constraint graphs merge data + knowledge





Modeling the global stock market





Finding the optimal Bayesian network given a constraint graph

Jacob M. Schreiber¹ and William S. Noble²

¹Department of Computer Science, University of Washington, Seattle, WA, United States of America

²Department of Genome Science, University of Washington, Seattle, WA, United States of America

ABSTRACT

Despite recent algorithmic improvements, learning the optimal structure of a Bayesian network from data is typically infeasible past a few dozen variables. Fortunately, domain knowledge can frequently be exploited to achieve dramatic computational savings, and in many cases domain knowledge can even make structure learning tractable. Several methods have previously been described for representing this type of structural prior



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Bayes classifiers rely on Bayes' rule

$$P(M|D) = \frac{P(D|M)P(M)}{\sum_M P(D|M)P(M)}$$

$$\textit{Posterior} = \frac{\textit{Likelihood} * \textit{Prior}}{\textit{Normalization}}$$



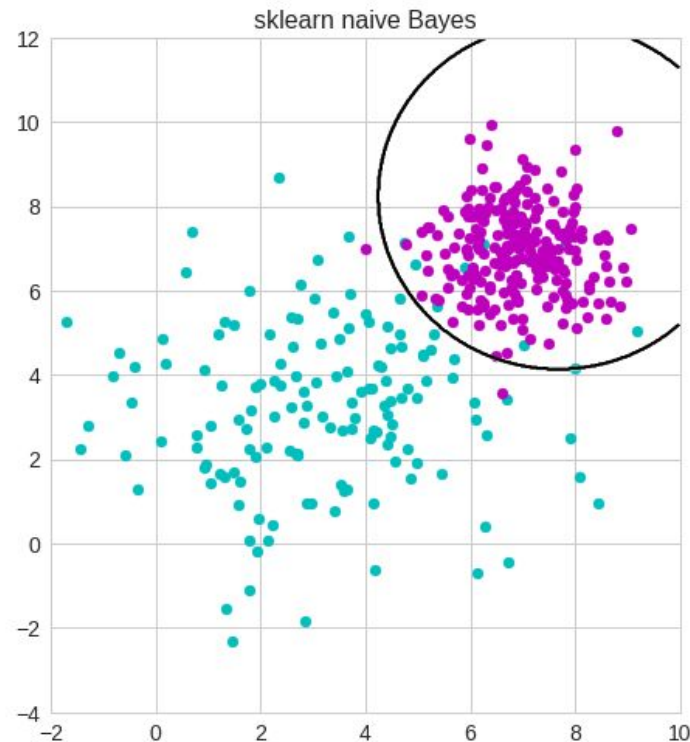
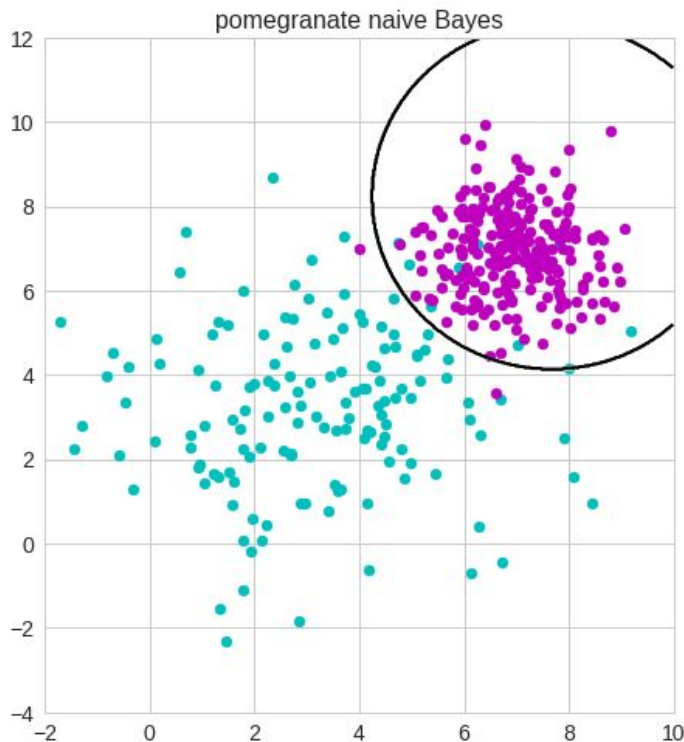
Naive Bayes assumes independent features

$$P(M|D) = \frac{\prod_{i=1}^d P(D_i|M)P(M)}{\sum_M \prod_{i=1}^d P(D_i|M)P(M)}$$

$$\textit{Posterior} = \frac{\textit{Likelihood} * \textit{Prior}}{\textit{Normalization}}$$



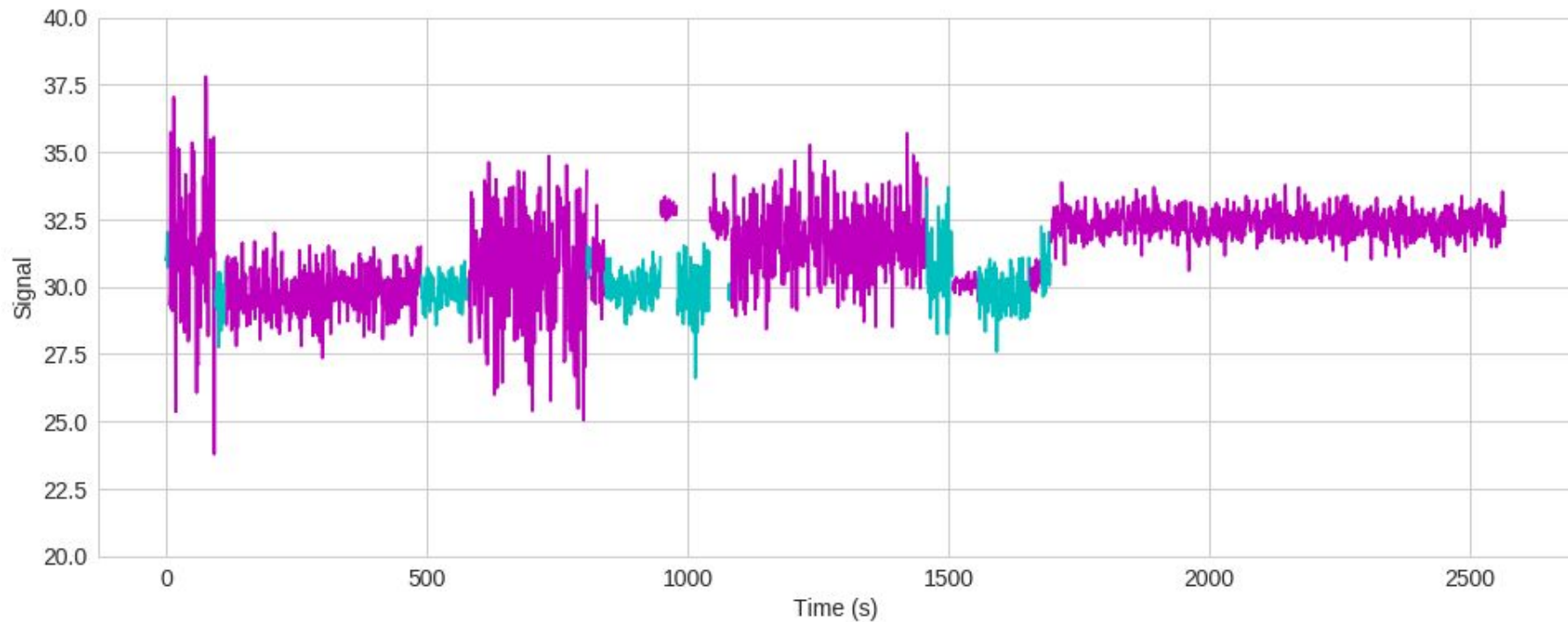
Naive Bayes produces ellipsoid boundaries



`model = NaiveBayes.from_samples(NormalDistribution, X, y)`



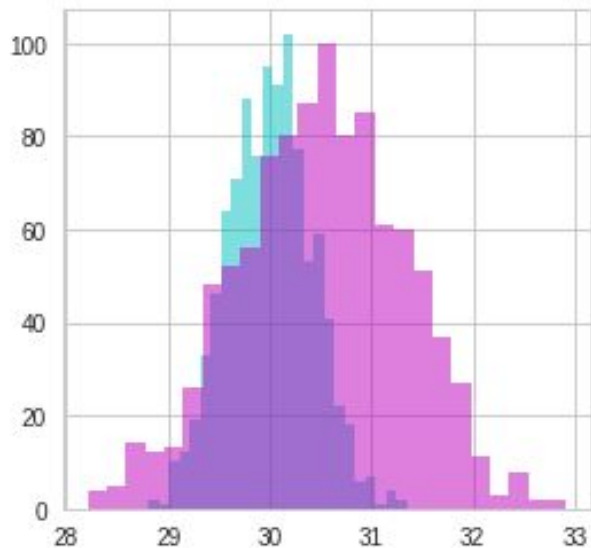
Naive Bayes can be heterogenous



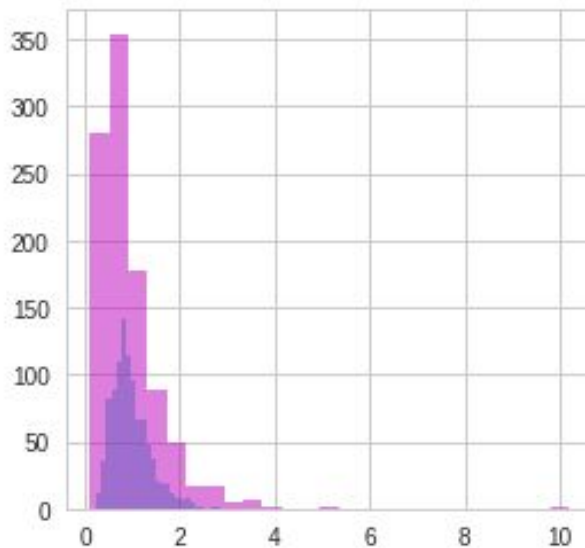


Data can fall under different distributions

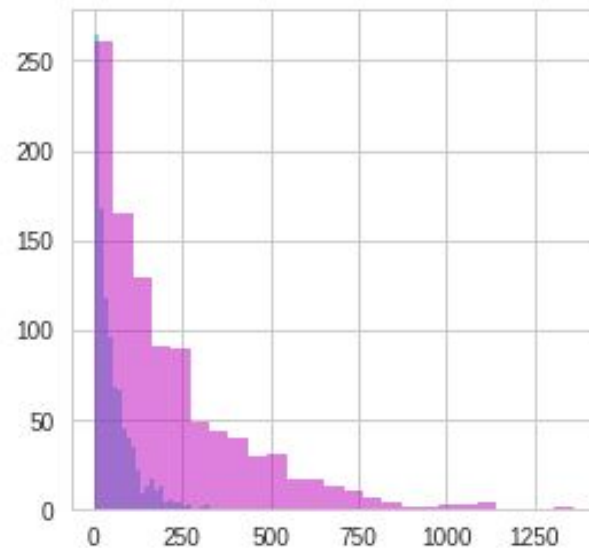
Mean



Standard Deviation



Duration





Using appropriate distributions is better

```
model = NaiveBayes.from_samples(NormalDistribution, X_train, y_train)
print "Gaussian Naive Bayes: ", (model.predict(X_test) == y_test).mean()
```

```
clf = GaussianNB().fit(X_train, y_train)
print "sklearn Gaussian Naive Bayes: ", (clf.predict(X_test) == y_test).mean()
```

```
model = NaiveBayes.from_samples([NormalDistribution, LogNormalDistribution,
ExponentialDistribution], X_train, y_train)
print "Heterogeneous Naive Bayes: ", (model.predict(X_test) == y_test).mean()
```

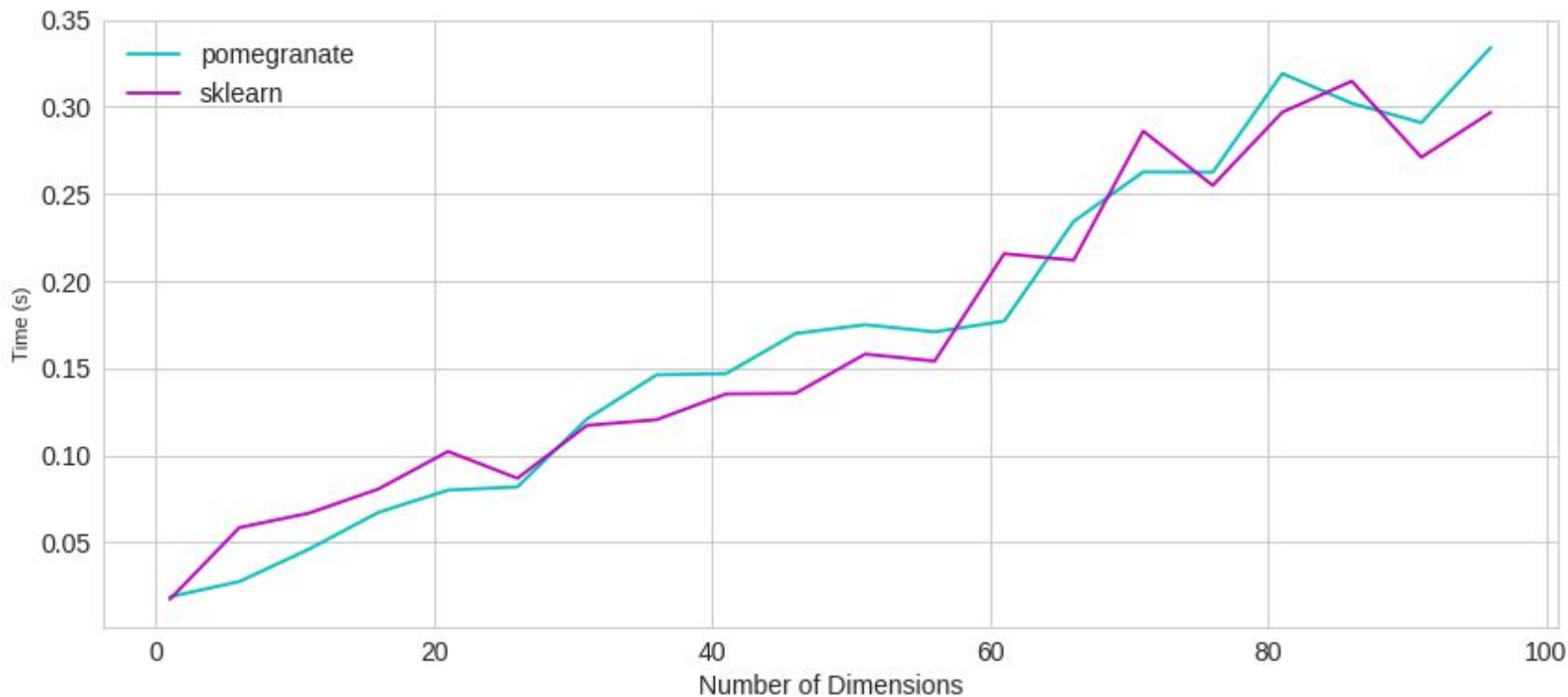
Gaussian Naive Bayes: 0.798

sklearn Gaussian Naive Bayes: 0.798

Heterogeneous Naive Bayes: 0.844



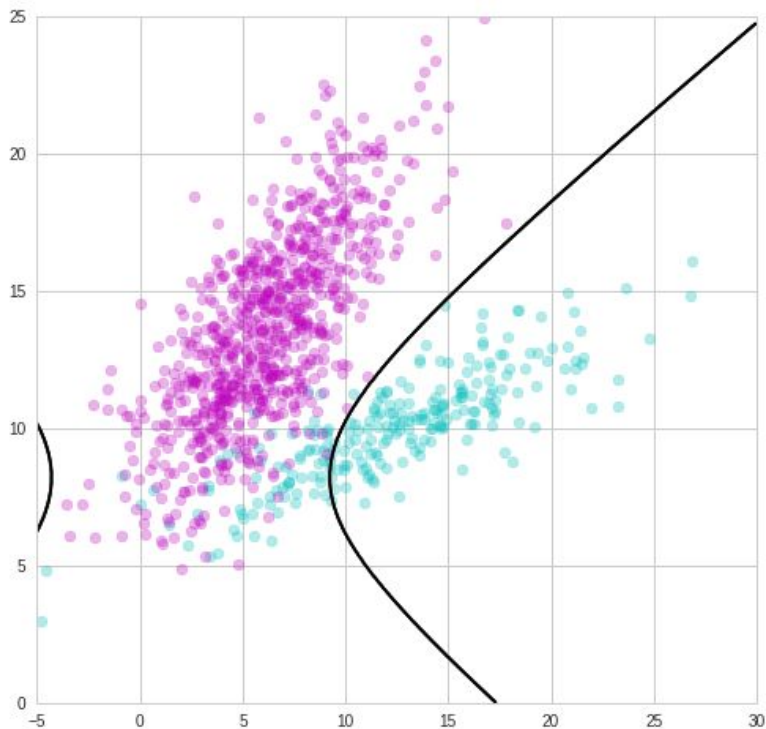
This additional flexibility is just as fast



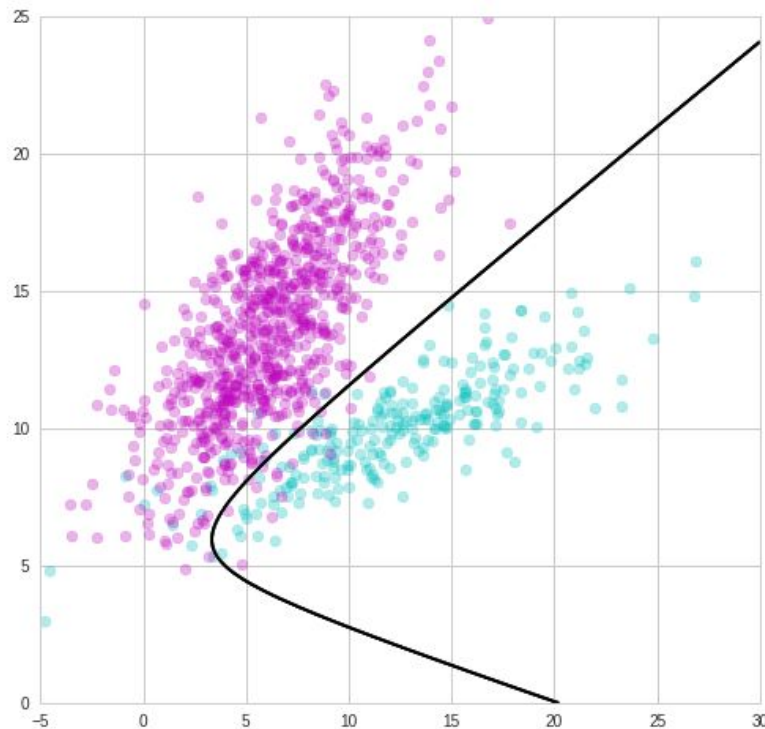


Bayes classifiers don't require independence

naive accuracy: 0.929

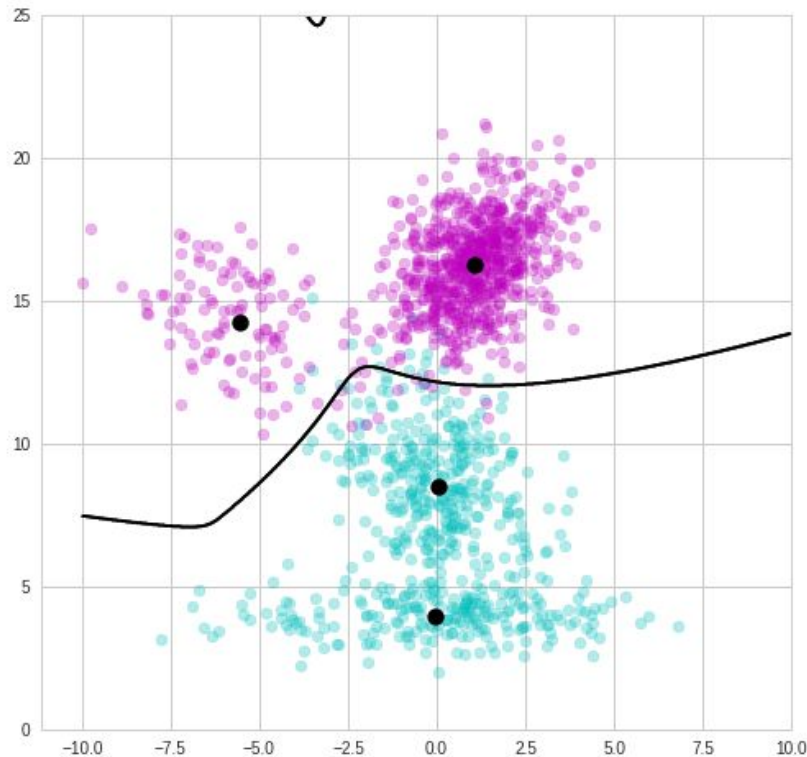
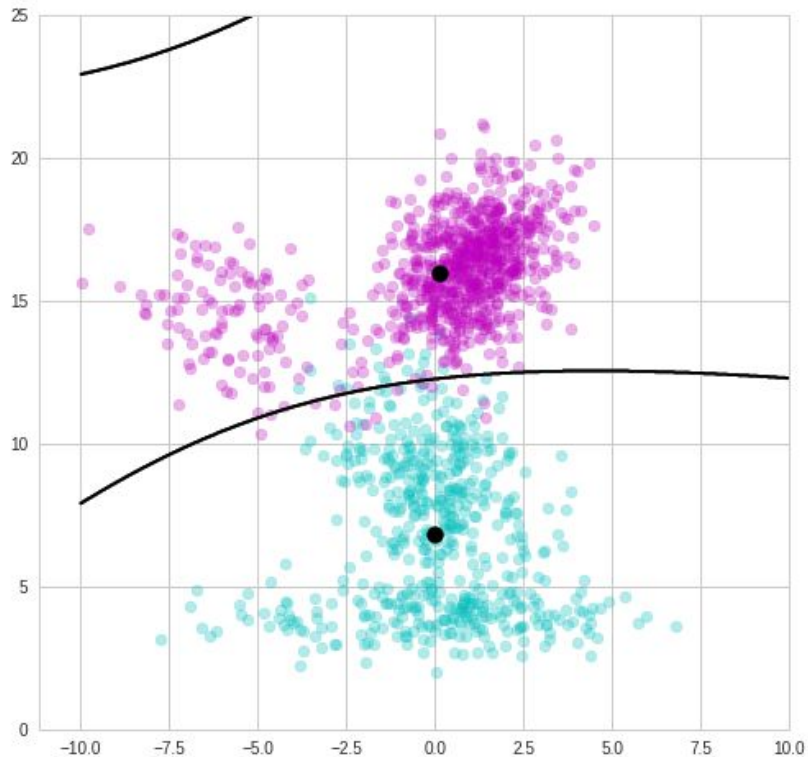


bayes classifier accuracy: 0.966





Gaussian mixture model Bayes classifier





Creating complex Bayes classifiers is easy

```
gmm_a = GeneralMixtureModel.from_samples(MultivariateGaussianDistribution, 2, X[y == 0])  
gmm_b = GeneralMixtureModel.from_samples(MultivariateGaussianDistribution, 2, X[y == 1])  
model_b = BayesClassifier([gmm_a, gmm_b], weights=numpy.array([1-y.mean(), y.mean()]))
```



Creating complex Bayes classifiers is easy

```
mc_a = MarkovChain.from_samples(X[y == 0])
mc_b = MarkovChain.from_samples(X[y == 1])
model_b = BayesClassifier([mc_a, mc_b], weights=numpy.array([1-y.mean(), y.mean()]))

hmm_a = HiddenMarkovModel.from_samples(X[y == 0])
hmm_b = HiddenMarkovModel.from_samples(X[y == 1])
model_b = BayesClassifier([hmm_a, hmm_b], weights=numpy.array([1-y.mean(), y.mean()]))

bn_a = BayesianNetwork.from_samples(X[y == 0])
bn_b = BayesianNetwork.from_samples(X[y == 1])
model_b = BayesClassifier([bn_a, bn_b], weights=numpy.array([1-y.mean(), y.mean()]))
```



Overview: this talk

Overview

Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Finale: Train a mixture of HMMs in parallel



Training a mixture of HMMs in parallel

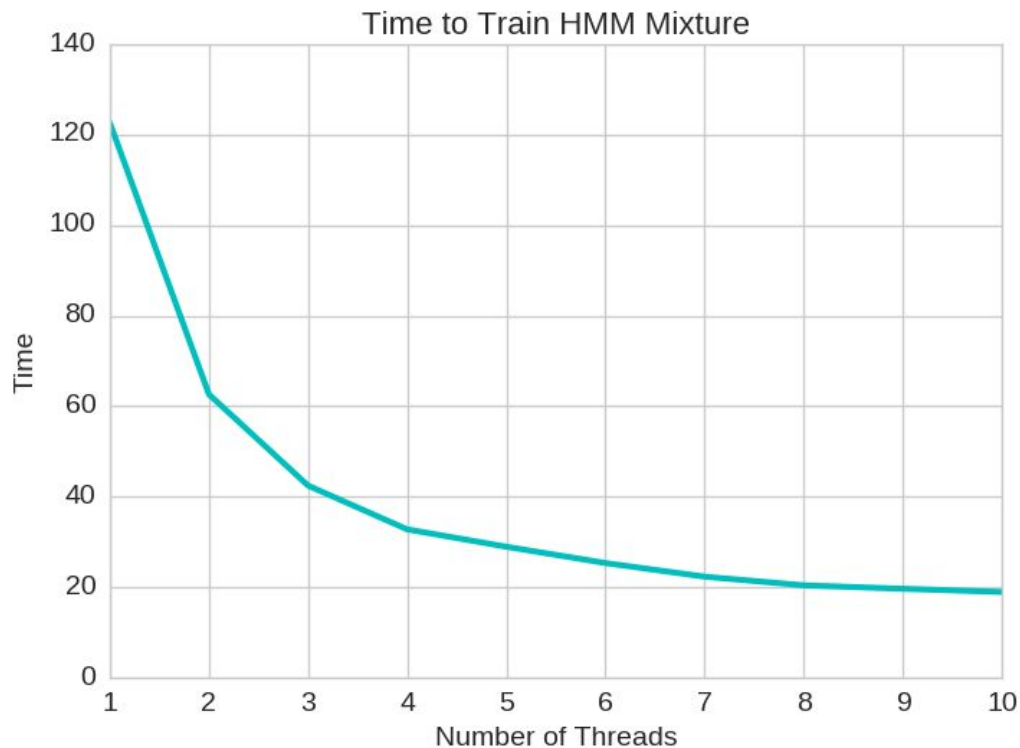
Creating a mixture of HMMs is just as simple as passing the HMMs into a GMM as if it were any other distribution

```
model_C = create_profile_hmm(dC, I)
model_mC = create_profile_hmm(dmC, I)
model_hmC = create_profile_hmm(dhmC, I)

model = GeneralMixtureModel([model_C, model_mC, model_hmC])
return model
```



Training a mixture of HMMs in parallel



`fit(model, X, n_jobs=n)`




Overview

pomegranate is **more flexible** than other packages, **faster**, is **intuitive to use**, and can do it all **in parallel**



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Bayes Classifiers and Naive Bayes

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pomegranate is a python package which implements fast, efficient, and extremely flexible probabilistic models ranging from probability distributions to Bayesian networks to mixtures of hidden Markov models. The most basic level of probabilistic modeling is the a simple probability distribution. If we're modeling language, this may be a simple distribution over the frequency of all possible words a person can say.



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<https://github.com/jmschrei/pomegranate/tree/master/tutorials>

Thank you for your time.