Adaptive Information Processing Model complexity and the MDL principle

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Signal Processing Group

Global overview

Part A: The Bayesian Information Criterion

Part B: Bayesian model estimation and the Context-tree model selection

Part C: Descriptive complexity

Prerequisites

Additional reading

Introduction

Bishop §1.2: Probability Theory

Bishop §1.3: Model Selection

Bishop §1.4: The Curse of Dimensionality

Probabilities

Bishop §2.1: Binary Variables

Bishop §2.2: Multinomial Variables

Part A

The Bayesian Information Criterion

Parameter and model estimation

Additional reading

Introduction

Bishop §3.3: Bayesian Linear Regression

Bishop §3.4: Bayesian Model Comparison

Parameter estimation

Define our variables!

Model \mathcal{M}_i model prior

Parameters θ_i parameter prior $p(\theta_i|\mathcal{M}_i)$

Data x^N

A-posteriori parameter distribution

$$p(heta_i|\mathcal{M}_i,x^N) = rac{p(heta_i|\mathcal{M}_i)p(x^N|\mathcal{M}_i, heta_i)}{p(x^N|\mathcal{M}_i)} \ p(x^N|\mathcal{M}_i) = \int_{\Omega} p(heta_i|\mathcal{M}_i)p(x^N|\mathcal{M}_i, heta_i) \,d heta_i$$

 $p(\mathcal{M}_i)$

Parameter estimation

Maximum Likelihood

We want a point estimate for θ_i (given \mathcal{M}_i).

$$\hat{ heta}_i = rg \max_{ heta_i} p(heta_i | \mathcal{M}_i, x^N) = rg \max_{ heta_i} p(x^N | \mathcal{M}_i, heta_i)$$

Where we assume a uniform prior or want to work without priors.

Model estimation

A-posteriori model distribution

$$egin{aligned} p(\mathcal{M}_i|x^N) &= rac{p(\mathcal{M}_i)p(x^N|\mathcal{M}_i)}{p(x^N)} \ p(x^N) &= \int_{\mathcal{M}_i} p(\mathcal{M}_i)p(x^N|\mathcal{M}_i) \, d\mathcal{M}_i \end{aligned}$$

Model estimation

Maximum Likelihood

We want a point estimate for \mathcal{M} .

$$\hat{\mathcal{M}} = rg \max_{\mathcal{M}_i} p(\mathcal{M}_i|x^N) = rg \max_{\mathcal{M}_i} p(x^N|\mathcal{M}_i)$$

Where we assume a uniform prior or want to work without priors.

Model estimation

We need to compute

$$p(x^N|\mathcal{M}_i) = \int_{\Theta_i} p(heta_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i) \, d heta_i$$

Often $p(\theta_i|\mathcal{M}_i,x^N)$ is sharply peaked and because

$$p(\theta_i|\mathcal{M}_i, x^N) \propto p(\theta_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i, \theta_i),$$

we might be able to approximate the integrand given above.

Maximum Likelihood and Overfitting

Additional reading

Overfitting

Bishop §1.1: Example: Polynomial Curve Fitting

Attempt 1 (Maximum Likelihood)

We approximate the integrand by its peak (θ_i^{MAP} or θ_i^{ML})

$$p(heta_i|\mathcal{M}_i)p(x^N|\mathcal{M}_i, heta_i)pprox \ \delta(heta_i- heta_i^{ extsf{ML}})p(heta_i|\mathcal{M}_i)p(x^N|\mathcal{M}_i, heta_i)$$

and find

$$p(x^N|\mathcal{M}_i) \propto p(heta_i^{\mathsf{ML}}|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i^{\mathsf{ML}})$$

So we end up with

$$egin{aligned} \mathcal{M}^{\mathsf{MAP}} &= rg \max_{\mathcal{M}_i} p(heta_i^{\mathsf{ML}}|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i^{\mathsf{ML}}) \ \mathcal{M}^{\mathsf{ML}} &= rg \max_{\mathcal{M}_i} p(x^N|\mathcal{M}_i, heta_i^{\mathsf{ML}}) \end{aligned}$$

Attempt 1: an example

Consider a linear regression model.

$$egin{align} y_n &= heta^T \underline{x}_n + n_n; \ y_n &\in \mathbb{R}; & heta \in \mathbb{R}^k; & \underline{x}_n \in \mathbb{R}^k; & n_n \sim \mathcal{N}(0, \sigma^2) \ \end{pmatrix}$$

Observe: $(y_1,\underline{x}_1),(y_2,\underline{x}_2),\ldots,(y_N,\underline{x}_N)$.

ML estimate: $\hat{\theta} = (X^T X)^{-1} X^T \underline{y}$.

Matrix: $oldsymbol{X} = \left[\underline{x}_1, \underline{x}_2, \dots \underline{x}_N \right]^T$.

Models: $\mathcal{M} \subset \{1, 2, \dots, k\}$. e.g.

$$\mathcal{M} = \{1,3\}; \qquad y_n = \theta_1 x_{n1} + \theta_3 x_{n3} + n_n$$

$$N=50; \quad \underline{x} \in [0,1]^3; \quad \theta = (0,0.6,0);$$
 $\sigma^2 = 1 \quad \text{actual } \sigma^2 = 0.799$
 $M \mid \hat{\theta}_1 \quad \hat{\theta}_2 \quad \hat{\theta}_3 \mid \hat{\sigma}^2 \mid \ln P_{ML} \mid \ln P_{ML} \mid \sigma^2 = \hat{\sigma}^2 \mid \sigma^2 = \hat{\sigma}^2$

0.766

0.766

0.686

-65.099

-65.101

-63.097

0 -0.522

 $0.752 \quad -0.940$

-0.472

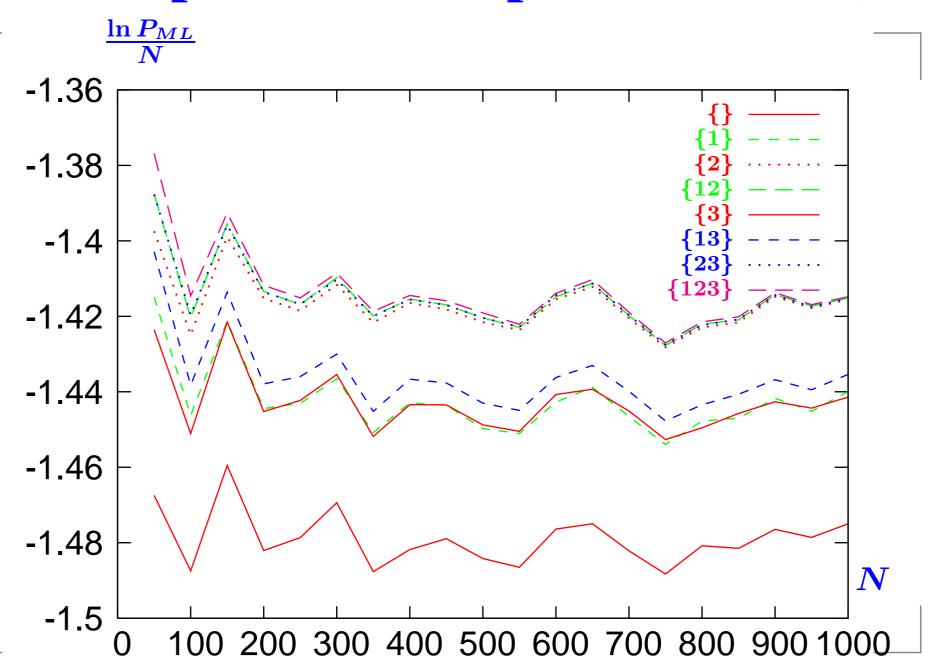
0.970

 $\{13\}$

{23}

-64.286

-64.287



From the graph we conclude that the "only noise" model $\mathcal{M} = \{\}$ has the worst performance, and that all models that include the actual parameter θ_2 , i.e. $\{\{2\}, \{1,2\}, \{2,3\}, \{1,2,3\}\}$ perform almost the same and the most complex of these, $\{1,2,3\}$, performs the best but is clearly an unwanted over-estimation.

Attempt 1: another example

A discrete data example.

Consider a binary second order Markov process:

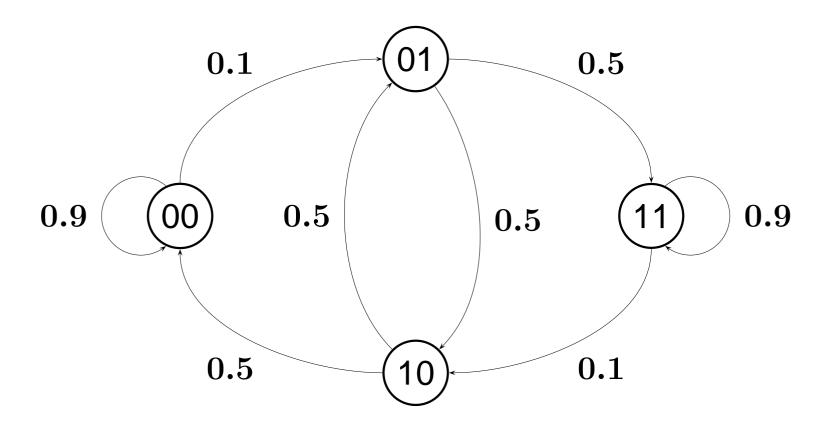
$$\Pr\{X_i = 1 | x^{i-1}\} = \Pr\{X_i = 1 | x_{i-2}x_{i-1}\}.$$

So, it is actually a set of four i.i.d. sub-sources.

ML estimate of an i.i.d. binary source:

$$n(s|x)=$$
 the number of times $s\in\mathcal{X}^*$ occurs in x $p(x^N| heta)=(1- heta)^{n(0|x^N)} heta^{n(1|x^N)}$ $rac{\partial}{\partial heta}\ln p(x^N| heta)=rac{n(1|x^N)-N heta}{ heta(1- heta)}=0$ $\hat{ heta}=rac{n(1|x^N)}{ heta}$

Attempt 1: another example (ctd.)



Attempt 1: another example (ctd.)

Let S be the state variable of an m-th order Markov source, so $S_i = X_{i-m} \dots X_{i-1}$ and $l(S_i) = m$ bits, then

$$\theta_s = \Pr\{X_i = 1 | S_i = s\}$$

The Maximum Likelihood estimator is

$$\hat{ heta}_s = rac{n(s1|x^N)}{n(s0|x^N) + n(s1|x^N)}$$

With this we find the ML probability for x^N (with initial state ς , see next slide)

$$p(x^N|m, \hat{ heta}, arsigma) = \prod_{s \in \{0,1\}^m} \left\{ \hat{ heta}_s^{n(s1|x^N)} (1-\hat{ heta}_s)^{n(s0|x^N)}
ight\}$$

Intermezzo: Initial state

The state variable of a m-th order Markov source is defined by the most recent m past symbols of the source. Initially, we cannot know the source state because we haven't seen m or more symbols yet.

So we assume knowledge of some initializing symbols that help in defining the first m values of the state variable. We will denote these initial symbols by ς and usually leave them unspecified.

Intermezzo: Initial state

This also implies that when we use the count function $n(s0|x^N)$ we imply the use of ς , e.g. let $x^5=10110$, m=2, and $\varsigma=01$. We wish to count the number of ones in state s=01.

$$n(s1|x^5) = 2$$

We consider the concatenation of ς and x^5 : 01 101100 and count the number of occurrences of s1=011 in that string.

Intermezzo: Initial state

s	$n(s0 x^5)$	$n(s1 x^5)$
00	0	0
01	0	2
10	0	1
11	2	0

So indeed, we count 2 zeros and 3 ones.

Attempt 1: another example (ctd.)

```
mytest(50, [0.1, 0.5, 0.5, 0.9], 4)
octave:1>
ML models sequence logprobs:
          logpr = -34.657359
Order 0:
          logpr = -14.546445
Order 1:
Order 2:
          logpr = -14.883390
          logpr = -15.185437
Order 3:
Order 4:
          logpr = -15.444986
          mytest(200,[0.1,0.5,0.5,0.9],4)
octave:2>
ML models sequence logprobs:
          logpr = -137.416984
Order 0:
          logpr = -102.949521
Order 1:
          logpr = -87.992931
Order 2:
          logpr = -84.732718
Order 3:
          logpr = -80.546002
Order 4:
```

Obviously, this method does not work.

Any model that includes the actual model assigns essentially the same probability to the data.

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- But high order models cannot predict well (too restricted).
- The higher order models are too well tuned.

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- We observe that (usually) the higher order models give higher probabilities to the sequence.
- But high order models cannot predict well (too restricted).
- The higher order models are too well tuned.

This is undesirable, the estimated model adapts itself to the noise and the resulting model is an over estimation of the actual model.

Preventing Overfitting

Additional reading

Laplace Approximation

Bishop §4.4: The Laplace Approximation

Attempt 2 (Laplace approximation)

We approximate the integrand by a Gaussian around the peak. The mean and variance of the Gaussian are determined by the integrand.

This approximation turns out to give more interesting results.

Suppose we have an arbitrary non-negative real function f(z), where z is a k-dimensional vector. We need an estimate of the normalizing constant Z_f .

$$Z_f = \int f(z)\,dz$$

Let z_0 be a maximum of f(z). Use the Taylor expansion.

$$\ln f(z) pprox \ln f(z_0) - rac{1}{2}(z-z_0)A(z-z_0)$$
 $A_{ij} = -rac{\partial^2}{\partial z_i \partial z_j} \ln f(z)igg|_{z=z_0}$

Approximate f(z) by the unnormalized Gaussian

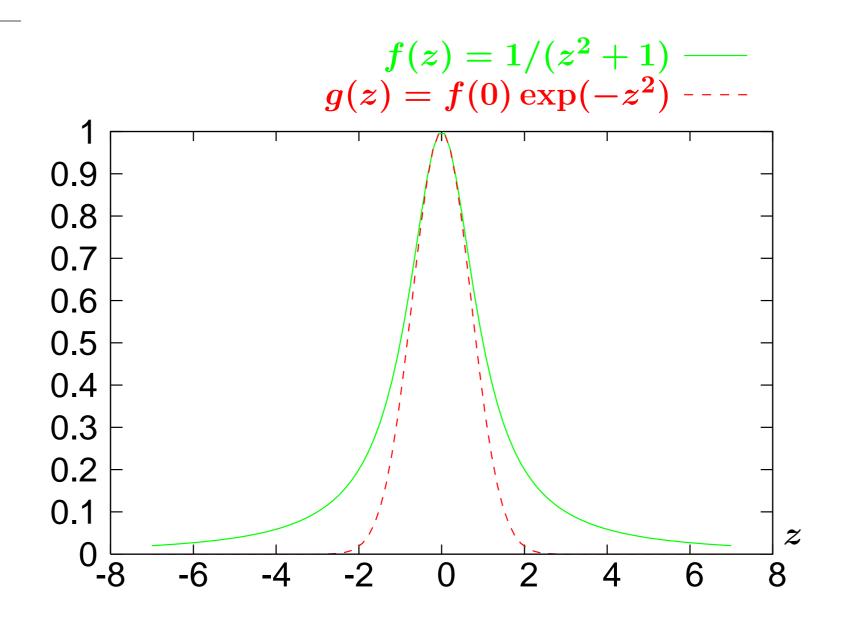
$$g(z) = f(z_0) \exp\left(-rac{1}{2}(z-z_0)A(z-z_0)
ight)$$

A, not necessarily good, approximation of Z_f is

$$Z_fpprox Z_g=\int g(z)\,dz=f(z_0)\sqrt{rac{(2\pi)^k}{\det A}}$$

Example 1:

$$f(z)=rac{1}{z^2+1}$$
 Has maximum at $z_0=0$. $Z_f=\pi$ $A=-rac{\partial^2}{\partial z^2}\ln f=-rac{f''f-f'^2}{f^2}$ $f(0)=1;\quad f'(0)=0;\quad f''(0)=-2; ext{ so } A=2$ $g(z)=f(0)\exp\left(-rac{1}{2}zAz
ight)=e^{-z^2}$ $Z_g=\sqrt{\pi}$



Attempt 2 (Laplace approximation)

Consider again $p(x^N|\mathcal{M}_i)$.

$$p(x^N|\mathcal{M}_i) = \int_{\Theta} p(heta_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i) \, d heta_i$$

We again use the fact that

$$p(heta_i|\mathcal{M}_i,x^N) arpropto p(heta_i|\mathcal{M}_i)p(x^N|\mathcal{M}_i, heta_i)$$

is often sharply peaked, say at $\hat{\theta}_i$. Using the Laplace approximation we may write

$$p(x^N|\mathcal{M}_i) pprox \sqrt{rac{(2\pi)^k}{\det A}} p(\hat{ heta}_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i,\hat{ heta}_i)$$

Comparing two models give

$$rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)}pprox rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)}rac{\sqrt{rac{(2\pi)^{k_i}}{\det A_i}}p(\hat{ heta}_i|\mathcal{M}_i)}{\sqrt{rac{(2\pi)^{k_j}}{\det A_j}}p(\hat{ heta}_i|\mathcal{M}_j)}rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)}$$

Comparing two models give

$$\frac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)} \approx \frac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)} \frac{\sqrt{\frac{(2\pi)^{k_i}}{\det A_i}} p(\hat{\theta}_i|\mathcal{M}_i)}{\sqrt{\frac{(2\pi)^{k_j}}{\det A_j}} p(\hat{\theta}_j|\mathcal{M}_j)} \frac{p(x^n|\mathcal{M}_i, \hat{\theta}_i)}{p(x^N|\mathcal{M}_j, \hat{\theta}_j)}$$

initial model preference

Comparing two models give

$$rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)}pprox rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)} rac{\sqrt{rac{(2\pi)^{k_i}}{\det A_i}}p(\hat{ heta}_i|\mathcal{M}_i)}{\sqrt{rac{(2\pi)^{k_j}}{\det A_j}}p(\hat{ heta}_i|\mathcal{M}_j)} rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)}$$

cost of (number of) parameters

Comparing two models give

$$rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)}pprox rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)} rac{\sqrt{rac{(2\pi)^{k_i}}{\det A_i}}p(\hat{ heta}_i|\mathcal{M}_i)}{\sqrt{rac{(2\pi)^{k_j}}{\det A_j}}p(\hat{ heta}_i|\mathcal{M}_j)} rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)}$$

likelihood ratio

Comparing two models give

$$rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)}pprox rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)}rac{\sqrt{rac{(2\pi)^{k_i}}{\det A_i}}p(\hat{ heta}_i|\mathcal{M}_i)}{\sqrt{rac{(2\pi)^{k_j}}{\det A_j}}p(\hat{ heta}_i|\mathcal{M}_j)}rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)}$$

Cost factors are: initial model preference, cost of (number of) parameters, likelihood ratio.

This is ML model estimation, it works because we consider the model complexity also!

BIC: Bayesian Information Criterion

A more refined approximation (Schwartz criterion or Bayesian Information Criterion) gives

$$\log rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)} pprox \log rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)} + \log rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)} + rac{1}{2}(k_i - k_j) \log N,$$

where k_i resp. k_j gives the number of free parameters in model \mathcal{M}_i or \mathcal{M}_j respectively.

This BIC is widely applied and turned out to be very usefull.

What happens when we apply the correction term $\frac{k}{2} \log N$? We shall revisit the two examples.

Example 1 with BIC correction

$$N=50; \quad \underline{x} \in [0,1]^3; \quad heta=(0,0.6,0);$$
 $\sigma^2=1 \quad ext{actual } \sigma^2=0.852$

\mathcal{M}	$\hat{\theta}_1$	$\hat{\theta}_{2}$	$\hat{\theta}_3$	$\hat{\sigma}^2$	$\ln P_{BIC}$
{}	0	0	0	1.068	-72.653
$\{1\}$	0.699	0	0	0.909	-70.632
$\{2\}$	0	0.773	0	0.841	-68.923
$\{3\}$	0	0	0.572	0.944	-71.491
$\{12\}$	0.159	0.662	0	0.837	-70.790
$\{13\}$	0.553	0	0.172	0.905	-72.478
$\{23\}$	0	0.811	-0.050	0.840	-70.869
$\{123\}$	0.240	0.728	-0.159	0.834	-72.670

Example 1 with BIC correction

$$N=1000;\quad \underline{x}\in[0,1]^3;\quad heta=(0,0.6,0);$$
 $\sigma^2=1\quad ext{actual }\sigma^2=0.977$

\mathcal{M}	$\hat{\theta}_1$	$\hat{\theta}_{2}$	$\hat{\theta}_3$	$\hat{\sigma}^2$	$\ln P_{BIC}$
{}	0	0	0	1.077	-1457.2
$\{1\}$	0.411	0	0	1.022	-1433.4
$\{2\}$	0	0.551	0	0.976	-1410.3
$\{3\}$	0	0	0.362	1.034	-1439.3
$\{12\}$	-0.017	0.564	0	0.976	-1413.7
$\{13\}$	0.315	0	0.128	1.020	-1435.6
$\{23\}$	0	0.637	-0.117	0.974	ullet -1412.7
$\{123\}$	0.040	0.620	-0.133	0.974	ullet -1416.1

Example 2 with BIC correction

```
octave:1> mytest(50,[0.1,0.5,0.5,0.9],4)
          scaled ML log probabilities:
Parameter
          logpr = -36.613371
Order 0:
          logpr = -18.458468
Order 1:
Order 2:
          logpr = -22.707436
          logpr = -30.833529
Order 3:
Order 4:
          logpr = -46.741170
          mytest(200,[0.1,0.5,0.5,0.9],4)
octave:2>
          scaled ML log probabilities:
Parameter
          logpr = -140.066143
Order 0:
Order 1:
          logpr = -108.247838
Order 2:
          logpr = -98.589565
          logpr = -105.925987
Order 3:
          logpr = -122.932541
Order 4:
```

BIC correction

The examples indicate that the correct model (order) is recovered, basically by using an ML selection criterion with an additional penalty term for the model complexity.

However, this BIC is derived as an approximation to the true Bayesian a-posteriori probability.

A better justification for the BIC should exist!

Part B

Bayesian model estimation and the Context-tree model selection

Additional reading

Bishop §14.1: Bayesian Model Averaging

Bishop §14.4: Tree-based Models

Additional notation

i^{th} order Markov Model	\mathcal{M}_i	The state is determined by the previous i symbols.	
Parameter vector	$ heta_i$	This vector describes all probabilities $P(x_n x_{n-i},x_{n-i+1},\ldots,x_{n-1}).$	
Parameter element	$ heta_{i,s}$	$\theta_{i,s} = P(x_n x_{n-i}^{n-1} = s).$	
Model state	s	s is a binary sequence of length i .	
Initial state	ς	ς is also a binary sequence of length i .	

Example 2: [Revisit first lecture]

Let \mathcal{M}_i be the *i*-th order binary Markov model (source).

Then $\Theta_i = [0,1]^{2^i}$.

Beta distribution for prior $p(\theta_i|\mathcal{M}_i)$, with $\alpha = \beta = \frac{1}{2}$ (Jeffreys prior).

$$egin{aligned} p(heta_i|\mathcal{M}_i,arsigma) &= \left(rac{\Gamma(lpha+eta)}{\Gamma(lpha)\Gamma(eta)}
ight)^{2^i} \prod_{s\in\{0,1\}^i} heta_{i,s}^{lpha-1} (1- heta_{i,s})^{eta-1} \ &= rac{1}{\pi^{2^i}} \prod_{s\in\{0,1\}^i} heta_{i,s}^{-1/2} (1- heta_{i,s})^{-1/2} \end{aligned}$$

$$egin{aligned} p(x^N|\mathcal{M}_i,arsigma) &= \int_{\Theta_i} p(heta_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i) \, d heta_i \ &= rac{1}{\pi^{2i}} \ &\int_{\Theta_i} \prod_{s \in \{0,1\}^i} heta_{i,s}^{n(s1|x^N)-1/2} (1- heta_{i,s})^{n(s0|x^N)-1/2} \, d heta_i \ &= rac{1}{\pi^{2i}} \ &\prod_{s \in \{0,1\}^i} \int_{[0,1]} heta_{i,s}^{n(s1|x^N)-1/2} (1- heta_{i,s})^{n(s0|x^N)-1/2} \, d heta_{i,s} \ &= \prod_{s \in \{0,1\}^i} rac{\Gamma(n(s0|x^N) + rac{1}{2})\Gamma(n(s1|x^N) + rac{1}{2})}{\pi\Gamma(n(s0|x^N) + n(s1|x^N) + 1)} \end{aligned}$$

So we must study the behaviour of

$$egin{aligned} P_e(a,b) & riangleq rac{\Gamma(a+rac{1}{2})\Gamma(b+rac{1}{2})}{\pi\Gamma(n+1)} \ a & riangleq n(0|x^N) \ b & riangleq n(1|x^N) \end{aligned}$$

It is a memoryless sub-sources of the Markov source. x^N is generated i.i.d. with parameter θ .

The actual probability of x^N under this source is

$$p(x^N|\mathcal{M}, \theta) = (1 - \theta)^a \theta^b$$

We can write

$$P_e(a,b) = rac{rac{1}{2}rac{3}{2}\cdots(a-rac{1}{2})\cdotrac{1}{2}rac{3}{2}\cdots(b-rac{1}{2})}{(a+b)!}$$

Again with the help of Stirling's approximation we can derive, for large a and b the following. (Excercise). Note: a+b=N.

$$\log_2 rac{p(x^N|\mathcal{M}, heta)}{P_e(a,b)} \leq rac{1}{2}\log_2 N + rac{1}{2}\log_2 rac{\pi}{2}$$

Actually, we can prove that for all $a \ge 0$ and $b \ge 0$

$$\log_2 \frac{p(x^N | \mathcal{M}, \theta)}{P_2(a, b)} \leq \frac{1}{2} \log_2 N + 1$$

Back to the *i*-th order Markov source.

$$egin{aligned} p(x^N|\mathcal{M}_i, heta_i,arsigma) &= \prod_{s\in\{0,1\}^i} heta_{i,s}^{n(s1|x^N)} (1- heta_{i,s})^{n(s0|x^N)} \ p(x^N|\mathcal{M}_i,arsigma) &= \prod_{s\in\{0,1\}^i} P_e(n(s1|x^N),n(s0|x^N)) \end{aligned}$$

With the previous bound we find

$$egin{aligned} \log_2 & rac{p(x^N | \mathcal{M}_i, eta_i, eta)}{p(x^N | \mathcal{M}_i, eta)} = \ & \log_2 rac{\prod_{s \in \{0,1\}^i} heta_{i,s}^{n(s1|x^N)} (1 - heta_{i,s})^{n(s0|x^N)}}{\prod_{s \in \{0,1\}^i} P_e(n(s1|x^N), n(s0|x^N))} \ & = \sum_{s \in \{0,1\}^i} \log_2 rac{ heta_{i,s}^{n(s1|x^N)} (1 - heta_{i,s})^{n(s0|x^N)}}{P_e(n(s1|x^n), n(s0|x^N))} \ & \leq \sum_{s \in \{0,1\}^i} rac{1}{2} \log_2 n(s|x^N) + 1 \stackrel{*1}{\leq} rac{2^i}{2} \log_2 rac{N - i}{2^i} + 2^i \end{aligned}$$

(*1): here we use Jensen's inequality.

So we conclude that for any parameter vector θ_i we have (approximately!)

[From now on we do not explicitly write ς anymore]

$$\log_2 p(x^N | \mathcal{M}_i) \approx \log_2 p(x^N | \mathcal{M}_i, \theta_i) - \frac{2^i}{2} \log_2 \frac{N-i}{2^i} - 2^i$$

Maximum Likelihood parameters (and $N\gg \max\{2^i,2^j\}$)

$$egin{split} \log_2 rac{p(\mathcal{M}_i|x^N)}{p(\mathcal{M}_j|x^N)} &pprox \log_2 rac{p(\mathcal{M}_i)}{p(\mathcal{M}_j)} + \log_2 rac{p(x^N|\mathcal{M}_i,\hat{ heta}_i)}{p(x^N|\mathcal{M}_j,\hat{ heta}_j)} \ &-rac{2^i-2^j}{2}\log_2 N \end{split}$$

So, again we observe the parameter cost!

Recap: Memoryless binary source: one parameter $\theta = \Pr\{X = 1\}$

Recap: Markov order-k: one parameter per state. There are 2^k states. The k symbols x_{i-k}, \ldots, x_{i-1} form the context of the symbol x_i .

Real world models: Length of context depends on its contents.

e.g. Natural language (English, Dutch, \cdots): if context starts with $x_{i-1} = q$ then no more symbols are needed.

A tree source is a nice concept to describe such sources.

A tree source consists of a set S of suffixes that together form a tree.

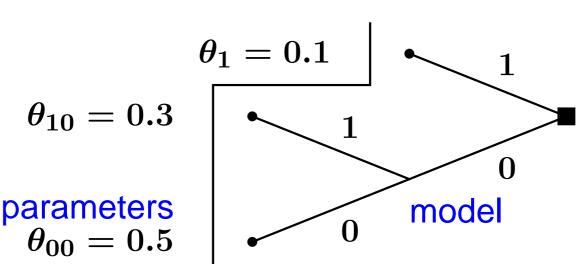
To each suffix (leaf) s in the tree there corresponds a parameter θ_s .

Some more notation: By $x_{|s}^N$ we denote the sub-sequence of symbols from x^N that are preceded by the sequence s.

Example: $x^8 = 01011010$; s = 01; then $x^8_{|01} = x_3 x_5 x_8 = 010$.

Example 3: Let $S \stackrel{\Delta}{=} \{00, 10, 1\}$ and $\theta_{00} = 0.5, \theta_{10} = 0.3$, and $\theta_1 = 0.1$ then

$$egin{aligned} \Pr\{X_i = 1| \cdots, x_{i-2} = 0, x_{i-1} = 0\} = 0.5, \ \Pr\{X_i = 1| \cdots, x_{i-2} = 1, x_{i-1} = 0\} = 0.3, \ \Pr\{X_i = 1| \cdots, x_{i-1} = 1\} = 0.1. \end{aligned}$$



$$\begin{array}{c} \text{parameters} \\ \theta_{00} = 0.5 \end{array}$$

Just as before ("Bayesian model estimation") we must estimate the sequence probabilities of the memoryless subsources that correspond to the leaves of the tree (states of the source).

Let the full sequence be x^N and the subsequence for state s be written as $x_{|s}^N$. Before we wrote

$$P_e(a,b) = rac{\Gamma(a+rac{1}{2})\Gamma(b+rac{1}{2})}{\pi\Gamma(a+b+1)}$$

We shall now use the shorthand notation for the estimated probability of the subsequence generated in state s given the full sequence x^i :

$$egin{aligned} P_e(a_s,b_s) &= rac{\Gamma(a_s+rac{1}{2})\Gamma(b_s+rac{1}{2})}{\pi\Gamma(a_s+b_s+1)} \ a_s &= n(s0|x^N) = n(0|x^N_{|s}) \ b_s &= n(s1|x^N) = n(1|x^N_{|s}) \end{aligned}$$

Example 4: Let
$$S = \{00, 10, 1\}$$
.

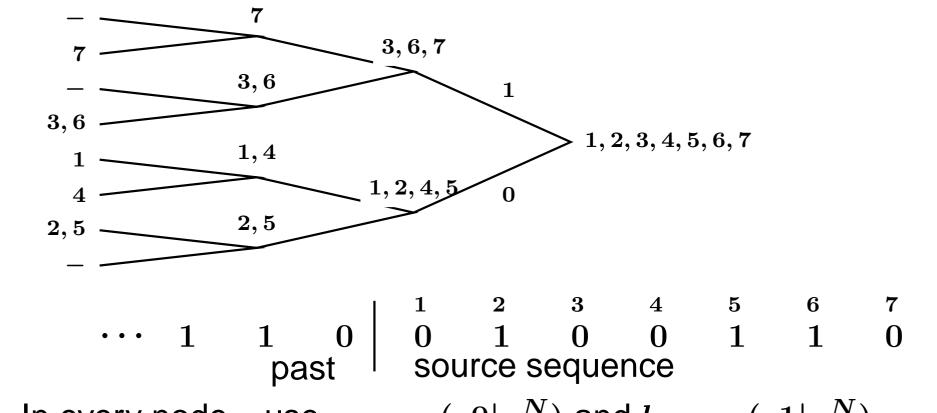
$$egin{aligned} p(0100110|\cdots 110) &= \underbrace{P_e(00)}_{10} \underbrace{P_e(11)}_{00} \underbrace{P_e(010)}_{1} \ &= rac{1}{2} \cdot rac{3}{4} \cdot rac{1}{2} \cdot rac{3}{4} \cdot rac{1}{2} \cdot rac{3}{4} \cdot rac{1}{6} = rac{9}{1024} \end{aligned}$$

See "Bayesian Estimation"

$$\log_2 rac{p(x^N|\mathcal{S}, heta)}{\prod_{s \in \mathcal{S}} P_e(a_s, b_s)} \leq rac{|\mathcal{S}|}{2} \log_2 rac{N}{|\mathcal{S}|} + |\mathcal{S}|$$

Problem: We do not know S!

Context tree (of depth D)



In every node s use $a_s=n(s0|x^N)$ and $b_s=n(s1|x^N)$.

We shall assign a probability to the subsequence $x_{|s|}^N$ for every state s in the context tree.

We shall do this in such a way that in the root of the tree we assign a probability to the whole sequence x^N that is a mixture of all possible tree sources.

We use the following observations to build, recursively, this probability.

The probability we build is written as follows

$$P_w^s = P_w(x_{|s}^N),$$

where P_w^s is the shorthand notation we shall use.

Later we return to this and make the notation more precise.

Suppose s is a leaf:

All we know are a_s and b_s so we assign the subsequence probability

$$P_w^s = P_e(a_s,b_s).$$

Now if s is an internal node, we have two options for the subsequence probability.

1: $P_e(a_s, b_s)$.

2: $P_w^{0s}P_w^{1s}$.

We must make a choice or better even, mix these options. So we set

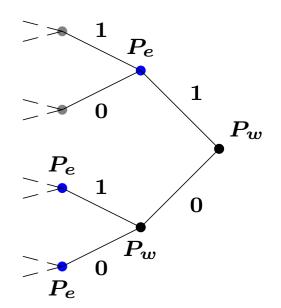
$$P_w^s = rac{P_e(a_s,b_s) + P_w^{0s} P_w^{1s}}{2}.$$

Analysis.

Let $S = \{00, 10, 1\}$ and we use a context tree with depth D > 2.

We look at the P_w 's for different nodes.

For the nodes $s \in \mathcal{S}$ we consider (in the analysis) only the P_e 's.



$$egin{split} P_w^{00} &\geq rac{1}{2} P_e(a_{00},b_{00}) \ P_w^{10} &\geq rac{1}{2} P_e(a_{10},b_{10}) \ P_w^1 &\geq rac{1}{2} P_e(a_1,b_1) \end{split}$$

Now we consider nodes nearer to the root and take only the $P_w^{0s}P_w^{1s}$ part.

$$egin{split} P_w^0 &\geq rac{1}{2} P_w^{00} P_w^{10} \ &\geq rac{1}{8} P_e(a_{00}, b_{00}) P_e(a_{10}, b_{10}) \ P_w^\lambda &\geq rac{1}{2} P_w^0 P_w^1 \ &\geq rac{1}{32} P_e(a_{00}, b_{00}) P_e(a_{10}, b_{10}) P_e(a_1, b_1) \end{split}$$

Here λ denotes the root of the tree.

For general trees (or suffix sets) S we find

$$P_w^{\lambda} \geq 2^{1-2|\mathcal{S}|} \prod_{s \in \mathcal{S}} P_e(a_s,b_s)$$

So

$$\log_2 P_w^{\lambda} \geq \log_2 p(x^N|\mathcal{S}, heta) - \left(2|\mathcal{S}| - 1 + rac{|\mathcal{S}|}{2}\log_2 N + |\mathcal{S}|
ight).$$

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Real sequence probability

For general trees (or suffix sets) S we find

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So

$$\log_2 P_w^{\lambda} \geq \log_2 p(x^N|\mathcal{S}, \theta) - \left(2|\mathcal{S}| - 1 + \frac{|\mathcal{S}|}{2}\log_2 N + |\mathcal{S}|\right).$$

Cost of describing the tree

For general trees (or suffix sets) S we find

$$P_w^{\lambda} \geq 2^{1-2|\mathcal{S}|} \prod_{s \in \mathcal{S}} P_e(a_s,b_s)$$

So

$$\log_2 P_w^{\lambda} \geq \log_2 p(x^N|\mathcal{S}, heta) - \left(2|\mathcal{S}| - 1 + rac{|\mathcal{S}|}{2} \log_2 N + |\mathcal{S}| \right).$$

Cost of the parameters

For general trees (or suffix sets) S we find

$$P_w^{\lambda} \geq 2^{1-2|\mathcal{S}|} \prod_{s \in \mathcal{S}} P_e(a_s,b_s)$$

So

$$\log_2 P_w^{\lambda} \geq \log_2 p(x^N|\mathcal{S}, \theta) - \left(2|\mathcal{S}| - 1 + \frac{|\mathcal{S}|}{2}\log_2 N + |\mathcal{S}|\right).$$

Contributions to the weighted probability are: Real sequence probability; Cost of describing the tree; Cost of the parameters

This algorithm achieves the (asymptotically) optimal log-likelihood ratio (not only on the average but also individually for every data sequence).

$$\log rac{p(x^N|\mathcal{S}, heta)}{P_w^\lambda} \leq 2|\mathcal{S}| - 1 + rac{|\mathcal{S}|}{2}\log_2 N + |\mathcal{S}|.$$

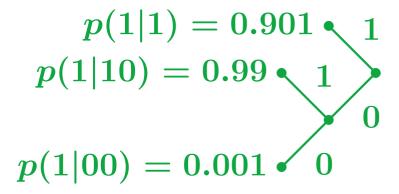
Another essential property of the "Context-Tree Weighting" (CTW) algorithm is its efficient implementation. The number of trees squares with every increment of D and yet the amount of work is at most linear in $D \cdot N$.

We can even write a stronger result when we realise that the method has no knowledge of a "real model". Let S_D be the set of all tree models with a maximal depth of atmost D.

$$egin{aligned} \log P_w^{\lambda} & \geq \max_{\mathcal{S} \in \mathcal{S}_D} \left\{ \log p(x^N | \mathcal{S}, heta) - \\ & \left(2|\mathcal{S}| - 1 + rac{|\mathcal{S}|}{2} \log_2 N + |\mathcal{S}|
ight)
ight\}. \end{aligned}$$

This algorithm is an instantiation of the MDL principle. It finds (in the class S_D) the model S that maximizes the sequence probability.

Example: Consider the following actual model.



We shall use the following models.

Example: Consider the following actual model.

$$p(1|1) = 0.901$$
 1 $p(1|10) = 0.99$ 1 0 $p(1|00) = 0.001$ 0

We shall use the following models.

$$p(1)$$
 •

Order-0

Example: Consider the following actual model.

$$p(1|1) = 0.901$$
 1 $p(1|10) = 0.99$ 1 0 $p(1|00) = 0.001$ 0

We shall use the following models.

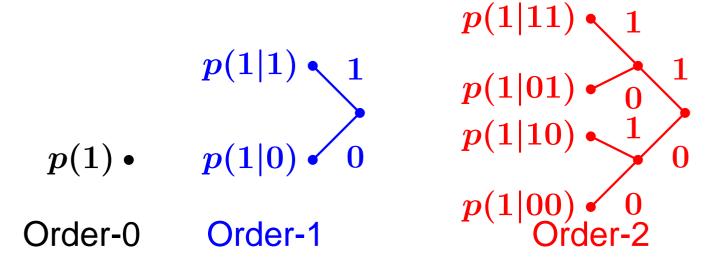
$$p(1|1) \cdot 1$$
 $p(1) \cdot p(1|0) \cdot 0$

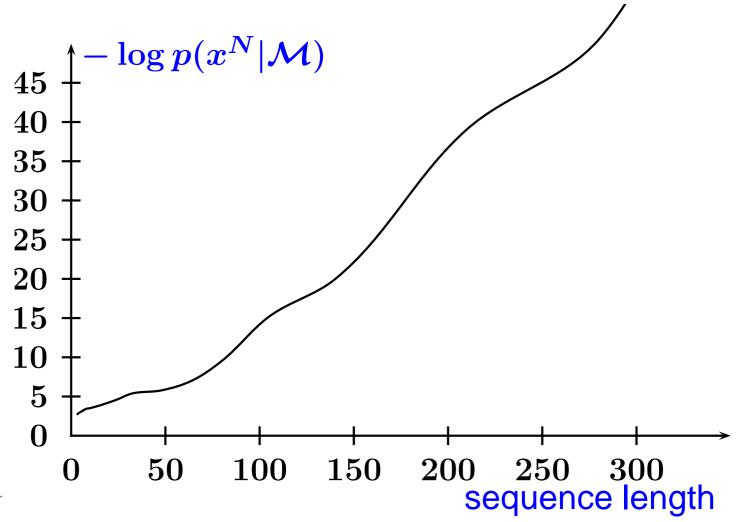
Order-0 Order-1

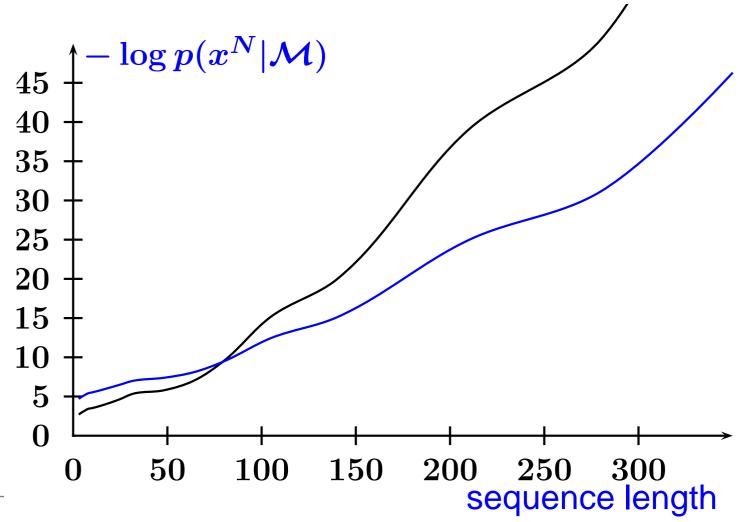
Example: Consider the following actual model.

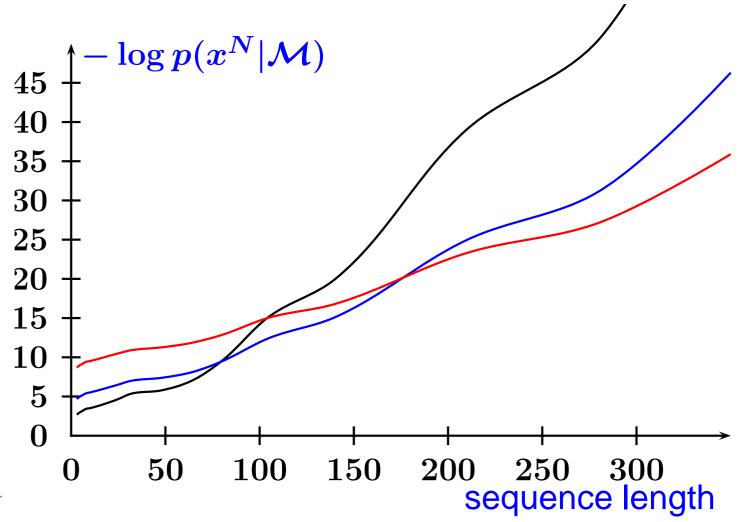
$$p(1|1) = 0.901$$
 1 $p(1|10) = 0.99$ 1 0 $p(1|00) = 0.001$ 0

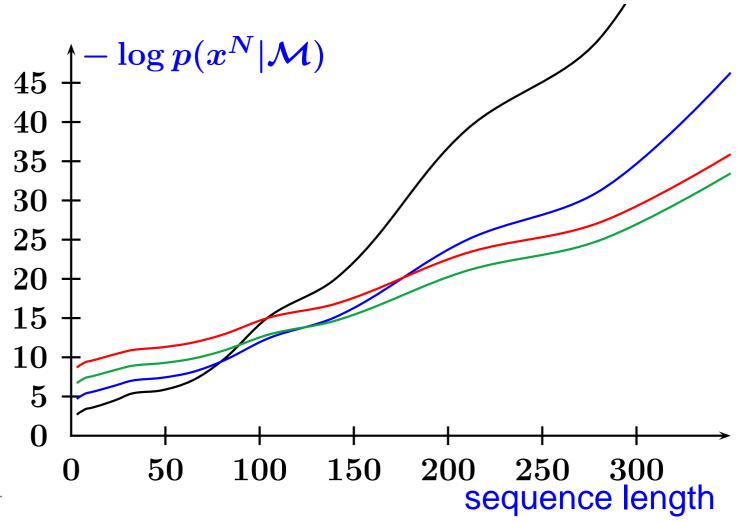
We shall use the following models.

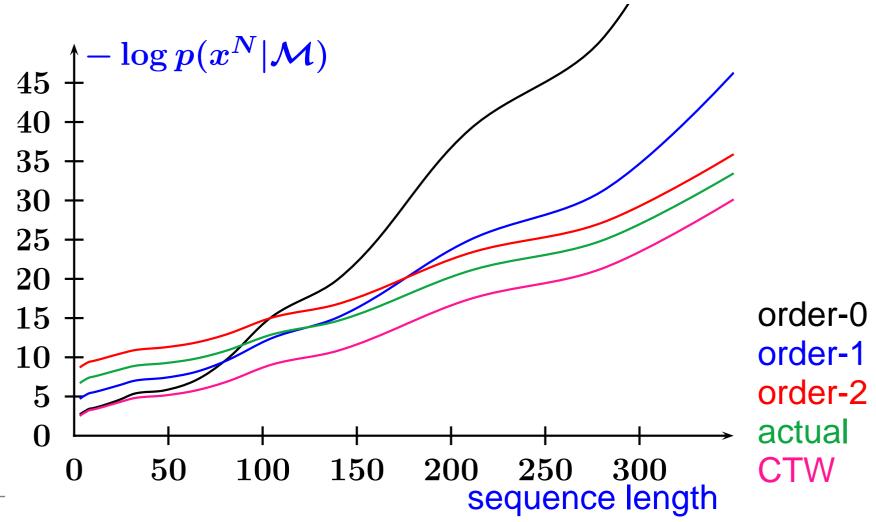












We see that initially the memoryless (order-0) model performs even better than the actual model.

After about 80 symbols the order-1 model becomes better than both the order-0 and the actual model.

From 120 symbols on the actual model is better than the simpeler models.

The order-2 model is always worse than the actual model. It describes the same probabilities but has too many parameters.

But the CTW model outperforms all models over the whole range of sequence lengths!

We shall now derive an expression, based on the previous method, for the a-posteriori model probability. We consider only binary data but the approach also works for arbitrary alphabets.

First we repeat our notation.

A model is described by a complete suffix set S.

The suffix set can be seen as the set of leaves of a binary tree. Our model class is the set of all complete binary trees whose depth is not more than D, for a given D. We write \mathcal{S}_D for the model class. So we say that $\mathcal{S} \in \mathcal{S}_D$. The depth of a tree is the length of the longest path from the root to a leaf.

Every model S has a set of parameters θ_s , one for every state $s \in S$ of the model. θ_s gives the probability of a 1 given that the previous symbols were s.

$$heta_s = \Pr\{X_t = 1 | X_{t-\ell}^{t-1} = s\}, \text{ where } \ell = |s|$$

The probability of a sequence, given a model S with parameters θ_s , $s \in S$ is

$$p(x^N|\mathcal{S}, heta) = \prod_{s \in \mathcal{S}} p(x^N_{|s}| heta_s)$$

and

$$p(x_{|s}^N| heta_s) = (1- heta_s)^{n(0|x_{|s}^N)} heta_s^{n(1|x_{|s}^N)}$$

Note (again) that $n(0|x_{|s}^N) = n(s0|x^N)$.

Actually, we silently assume that the first few symbols also have a "context". So we assume that there are some symbols preceding x^N .

We must define some prior distributions. First the prior on the parameters.

We use the beta distribution. (In a non-binary case this generalizes to the Dirichlet distribution.) As done before we select the parameters in the beta distribution to be $\frac{1}{2}$. So given a model $\mathcal S$ then for every $s\in\mathcal S$ we take

$$p(heta_s|\mathcal{S}) = rac{1}{\pi} heta_s^{-rac{1}{2}}(1- heta_s)^{-rac{1}{2}}$$

This results in the following sequence probability, first assuming one state s only

$$egin{aligned} p(x_{|s}^N) &= \int_0^1 p(heta_s|\mathcal{S}) heta_s^{n(s1|x^N)} (1- heta_s)^{n(s0|x^N)} \, d heta_s \ &= rac{\Gamma(n(s0|x^N) + rac{1}{2}) \Gamma(n(s1|x^N) + rac{1}{2})}{\pi \Gamma(n(s|x^N) + 1)} \end{aligned}$$

Now for any tree model \mathcal{S} we find

$$egin{align} p(x^N|\mathcal{S}) &= \prod_{s \in \mathcal{S}} \int_0^1 p(heta_s|\mathcal{S}) p(x_{|s}^N| heta_s) \, d heta_s \ &= \prod_{s \in \mathcal{S}} rac{\Gamma(n(s0|x^N) + rac{1}{2}) \Gamma(n(s1|x^N) + rac{1}{2})}{\pi \Gamma(n(s|x^N) rac{1}{AIP: \, ext{MODEL}})_{ ext{complexity and the MDL principle - p.70/12}} \end{aligned}$$

Next we need a prior on the tree models S in the set S_D . We wish to use the efficient CTW method of weighting so we choose the corresponding prior.

First define

$$\Delta_D(S) \stackrel{\Delta}{=} 2|\mathcal{S}| - 1 - |\{s \in \mathcal{S} : |s| = D\}|.$$

Then we take the prior

$$p(\mathcal{S}) = 2^{-\Delta_D(\mathcal{S})}$$

We prove that this is a proper prior probability.

Obviously, p(S) > 0 for all $S \in S_D$. We must show that it sums up to one.

We give a proof by induction.

Step 1: D = 0: $S_0 = \{\lambda\}$, the memoryless source.

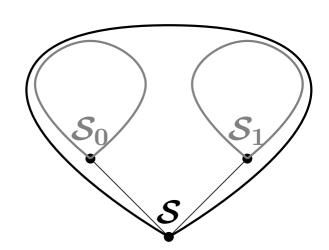
$$\Delta_0(\lambda) = 2 \cdot 1 - 1 - 1$$

Where the last -1 comes from the fact that the single state of λ is at level D=0 so $p(\lambda)=1$.

Induction: Assume it holds for $D \leq D^*$. Now if $\mathcal{S} \in \mathcal{S}_{D^*+1}$ then

- $\mathcal{S} = \lambda$, i.e. root node only.
- $m{\mathscr{S}}$ contains two trees on level 1, say $m{\mathscr{S}}_0\in S_{D^*}$ and $m{\mathscr{S}}_1\in S_{D^*}.$ We have

$$\Delta_{D^*+1}(\mathcal{S}) = 1 + \Delta_{D^*}(\mathcal{S}_0) + \Delta_{D^*}(\mathcal{S}_1)$$



• We repeat: \mathcal{S} contains two trees on level 1, say $\mathcal{S}_0 \in \mathcal{S}_{D^*}$ and $\mathcal{S}_1 \in \mathcal{S}_{D^*}$. We have

 $\Delta_{D^*+1}(\mathcal{S}) = 1 + \Delta_{D^*}(\mathcal{S}_0) + \Delta_{D^*}(\mathcal{S}_1)$

$$egin{aligned} \sum_{\mathcal{S} \in \, \mathbb{S}_{D^*+1}} 2^{-\Delta_{D^*+1}(\mathcal{S})} &= 2^{-1} + \ & \sum_{\mathcal{S}_0 \in \, \mathbb{S}_{D^*}} \sum_{\mathcal{S}_1 \in \, \mathbb{S}_{D^*}} 2^{-1-\Delta_{D^*}(\mathcal{S}_0) - \Delta_{D^*}(\mathcal{S}_1)} \ &= 2^{-1} + 2^{-1} \sum_{\mathcal{S}_0 \in \, \mathbb{S}_{D^*}} 2^{-\Delta_{D^*}(\mathcal{S}_0)} \sum_{\mathcal{S}_1 \in \, \mathbb{S}_{D^*}} 2^{-\Delta_{D^*}(\mathcal{S}_1)} \ &= 2^{-1} + 2^{-1} = 1 \end{aligned}$$

We now show that the weighted sequence probability

$$p(x^N) = \sum_{\mathcal{S} \in \, \mathbb{S}_D} p(\mathcal{S}) p(x^N | \mathcal{S}),$$

is produced by the weighting procedure of CTW, so

$$p(x^N) = P_w^{\lambda}$$
.

We shall prove this using (mathematical) induction.

First assume D=0: $S_0=\{\lambda\}$, so the only tree in the set consists of a root only. Therefor $\Delta_0(\lambda)=0$. So,

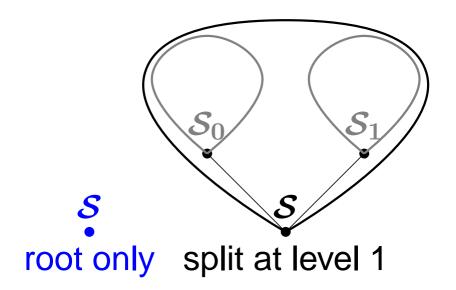
$$egin{aligned} p(x^N) &= p(\lambda) p(x^N | \lambda) \ &= 2^0 P_e(n(0|x^N), n(1|x^N)) \ &= P_w^\lambda, \end{aligned}$$

because λ is also a leaf and in a leaf $P_w = P_e$.

Now assume that for all $D < D^*$

$$\sum_{\mathcal{S} \in \, \$_D} p(\mathcal{S}) p(x^N | \mathcal{S}) = P_w^{\lambda}$$

The tree S is either the root only or it consists of a root plus two trees, S_0 and S_1 , on level one.



$$egin{aligned} \sum_{\mathcal{S} \in \, \mathbb{S}_{D^*+1}} & p(\mathcal{S})p(x^N|\mathcal{S}) = \ & = 2^{-1}P_e(n(0|x^N), n(1|x^N)) + \ & \sum_{\mathcal{S} \in \, \mathbb{S}_{D^*+1}: \mathcal{S}
eq \lambda} & p(\mathcal{S})p(x^N|\mathcal{S}) \end{aligned}$$

$$egin{aligned} \sum_{\mathcal{S} \in \, \$_{D^*+1} : \mathcal{S}
eq \lambda} p(\mathcal{S}) p(x^N | \mathcal{S}) &= \ &\sum_{\mathcal{S}_0 \in \, \$_{D^*}} \sum_{\mathcal{S}_1 \in \, \$_{D^*}} rac{1}{2} 2^{-\Delta_{D^*}(\mathcal{S}_0)} 2^{-\Delta_{D^*}(\mathcal{S}_1)} imes \ &p(x_{|0}^N | \mathcal{S}_0) p(x_{|1}^N | \mathcal{S}_1) \ &= rac{1}{2} \sum_{\mathcal{S}_0 \in \, \$_{D^*}} 2^{-\Delta_{D^*}(\mathcal{S}_0)} p(x_{|0}^N | \mathcal{S}_0) imes \ &\sum_{\mathcal{S}_1 \in \, \$_{D^*}} 2^{-\Delta_{D^*}(\mathcal{S}_1)} p(x_{|1}^N | \mathcal{S}_1) \ &= rac{1}{2} P_w^0 P_w^1 \ &= rac{1}{2} P_w^0 P_w^1 \end{aligned}$$

And so we find

$$egin{aligned} \sum_{\mathcal{S} \in \, \mathbb{S}_{D^*+1}} & p(\mathcal{S}) p(x^N | \mathcal{S}) = \ &= rac{1}{2} P_e(n(0|x^N), n(1|x^N)) + rac{1}{2} P_w^0 P_w^1 \ &= P_w^{\lambda} \end{aligned}$$

$$p(\mathcal{S}|x^N) = rac{p(\mathcal{S})p(x^N|\mathcal{S})}{p(x^N)}$$

$$p(\mathcal{S}|x^N) = rac{p(\mathcal{S})p(x^N|\mathcal{S})}{p(x^N)}$$

$$p(\mathcal{S}|x^N) = rac{\mathbf{2}^{-oldsymbol{\Delta}_D(\mathcal{S})}p(x^N|\mathcal{S})}{p(x^N)}$$

$$p(\mathcal{S}|x^N) = rac{2^{-\Delta_D(\mathcal{S})} \prod_{s \in \mathcal{S}} P_e(n(s0|x^N), n(s1|x^N))}{p(x^N)}$$

$$p(\mathcal{S}|x^N) = rac{2^{-\Delta_D(\mathcal{S})}\prod_{s\in\mathcal{S}}P_e(n(s0|x^N),n(s1|x^N))}{P_w^{\lambda}}$$

Thus we can compute the a-posteriori model probability.

$$p(\mathcal{S}|x^N) = rac{2^{-\Delta_D(\mathcal{S})}\prod_{s\in\mathcal{S}}P_e(n(s0|x^N),n(s1|x^N))}{P_w^{\lambda}}$$

So, we can use the same computations as in the CTW.

An efficient way to find the Bayesian MAP model exists, but its discussion is not a part of this course.

Part C

Descriptive complexity

Descriptive complexity

Simple sequences are "easy" to describe, complex ones must be described symbol by symbol.

Shannon complexity

Can the (Shannon) entropy be considered as a measure of complexity?

Yes, but the entropy depends on the probability of a sequence given an underlying source or stochastic data generating process.

Assuming that a source assigns probabilities $\Pr\{X = x\}$ the entropy of the source is defined as

$$H(X) = -\sum_{x \in \mathcal{X}} \Pr\{X = x\} \log_2 \Pr\{X = x\}.$$

This is the expected number of bits needed to represent X.

Shannon complexity

For a sequence x a corresponding notion is the ideal code wordlength given as

$$I(x) = -\log_2 \Pr\{X = x\}.$$

This can be interpreted as the most favorable representation length.

A disadvantage of Shannon's measures seems to be the fact that the complexity of a sequence depends on the probability of the sequence and not on the sequence itself.

Shannon complexity

Example 5: [of the 'unreasonable' interpretation]

Let $\mathcal{X} = \{01101010000010011110, 0011011100100100100\}$ and let the source select between the two sequences with equal probability $(\frac{1}{2}, \frac{1}{2})$.

The entropy of the source is 1 bit per sequence (of 20 symbols)! However, the two strings each appear much more complex than 1 bit!!

The complexity is hidden in the source description, namely in \mathcal{X} , which is already known by the receiver. We shall see that universal data compression gives a more fundamental answer to this problem.

Is it also possible to find a more meaningful measure using Shannon's information measure?

Because we do not know the model and its parameter values, we must consider data compression for parametrized classes of sources.

Example 6:

Parametrized binary source (I.I.D. source class)

Alphabet: $\mathcal{X} = \{0, 1\};$

Sequence: $x^N = x_1 \dots x_N$;

(N is the block length)

Probabilities: $\Pr\{X_i = 1\} = 1 - \Pr\{X_i = 0\} = \theta$.

 $0 \le \theta \le 1$.

Code: $C:\mathcal{X}^N o \{0,1\}^*$

Code word: $c(x^N) = c_1 \dots c_j \in C$

Length: $l_C(x^N) = l(c_1 \dots c_j) = j$

Ideal code wordlength

The best possible code wordlengths come from Huffman's algorithm, but these are hard to compute.

The task: minimize over the choice of lengths $l_C(x^N)$

$$\sum_{x^N \in \mathcal{X}^N} p(x^N) l_C(x^N)$$

where the lengths must satisfy Kraft's inequality

$$\sum_{x^N \in \mathcal{X}^N} 2^{-l_C(x^N)} \le 1$$

Ideal code wordlength

Ignoring the requirement that code wordlengths are integer, we find that the optimal code wordlengths are

$$l_C(x^N) = -\log_2 p(x^N)$$

The upward rounded version of these lengths still satisfy Kraft's inequality and the resulting code achieves Shannon's upper bound.

We write $l_C^*(x^N)$ for these ideal code wordlengths.

$$egin{aligned} l_C^*(x^N) &= \left\lceil -\log_2 p(x^N)
ight
ceil \\ &< -\log_2 p(x^N) + 1 \end{aligned}$$

Remember $n(a|x^N)$ is the number of times the symbol a occurs in x^N .

Sequence probability: $p(x^N) = (1- heta)^{n(0|x^N)} heta^{n(1|x^N)}$

Expected code word length: $ar{l_C} = \sum_{x^N \in \mathcal{X}^N} p(x^N) l_C(x^N)$

(Expected) code rate: $R_N=rac{ar{l_C}}{N}$

(Expected) code redundancy: $r_N = R_N - h(\theta)$

First assume that we know that $\theta=\theta_1=0.2$ or $\theta=\theta_2=0.9$ but we don't know which θ generated x^N .

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First assume that we know that $\theta=\theta_1=0.2$ or $\theta=\theta_2=0.9$ but we don't know which θ generated x^N .

We design a code C_1 assuming that $\theta = \theta_1$.

And a code C_2 assuming $\theta = \theta_2$.

We also create the code C_{12} which uses the smallest code word from C_1 and C_2 with a '0' or '1' prepended to indicate from which code the word comes.

First assume that we know that $\theta=\theta_1=0.2$ or $\theta=\theta_2=0.9$ but we don't know which θ generated x^N .

We design a code C_1 assuming that $\theta = \theta_1$.

And a code C_2 assuming $\theta = \theta_2$.

We also create the code C_{12} which uses the smallest code word from C_1 and C_2 with a '0' or '1' prepended to indicate from which code the word comes.

In all cases the code words are created using the ideal code wordlengths $l_C^*(x^N)$.

First assume that we know that $\theta=\theta_1=0.2$ or $\theta=\theta_2=0.9$ but we don't know which θ generated x^N .

We design a code C_1 assuming that $\theta = \theta_1$.

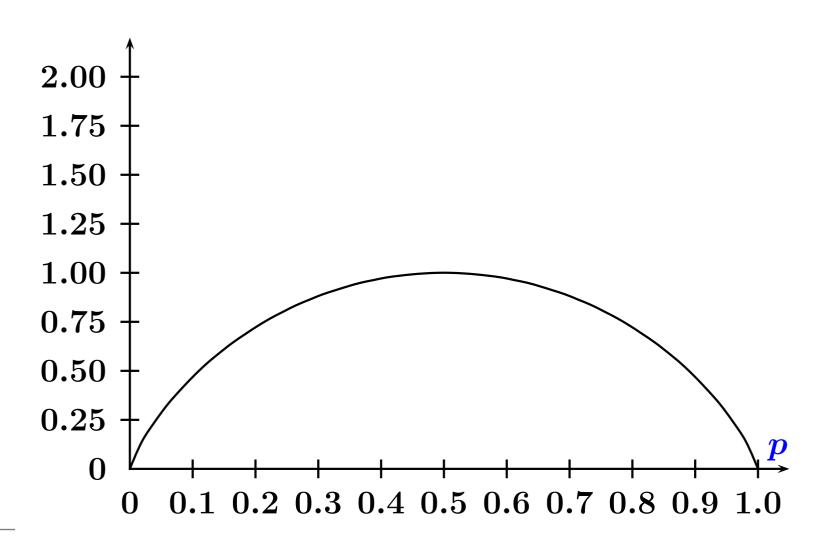
And a code C_2 assuming $\theta = \theta_2$.

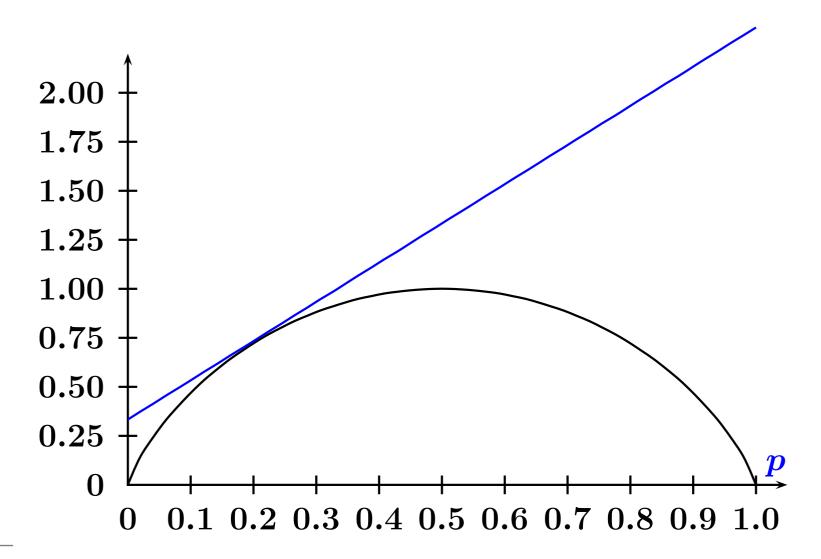
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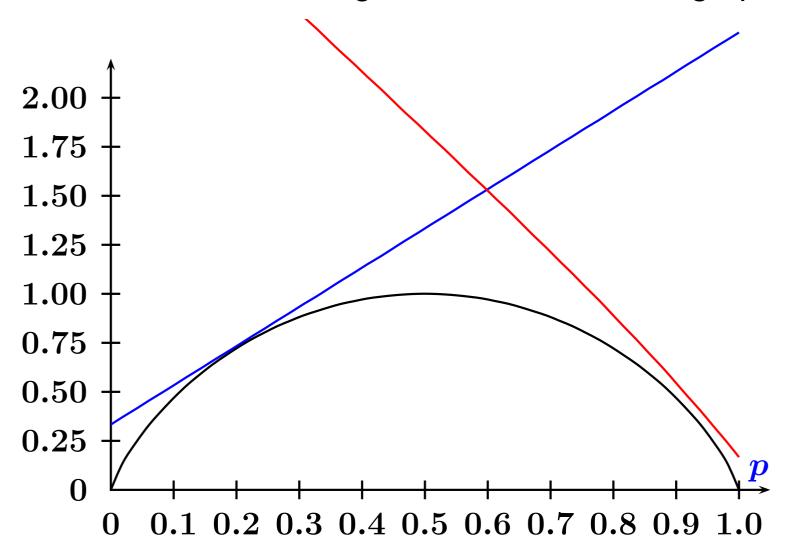
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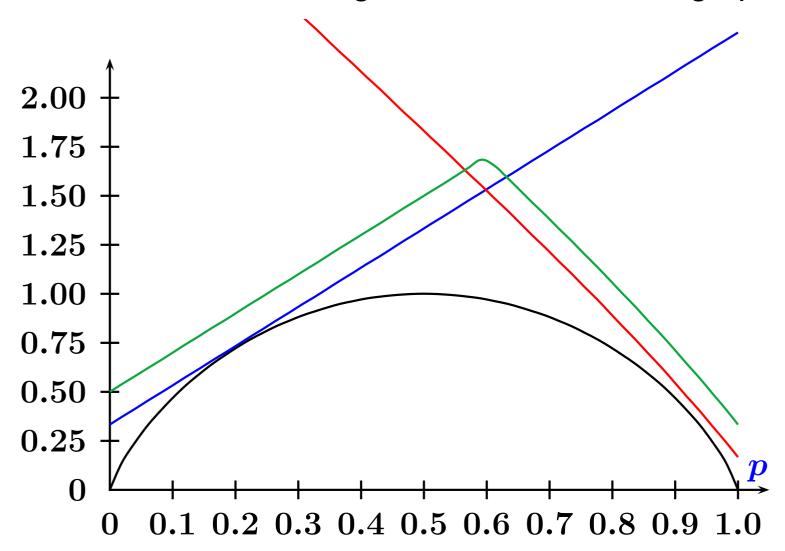
The code C_{mix} is make using the mixed (weighted) probabilities

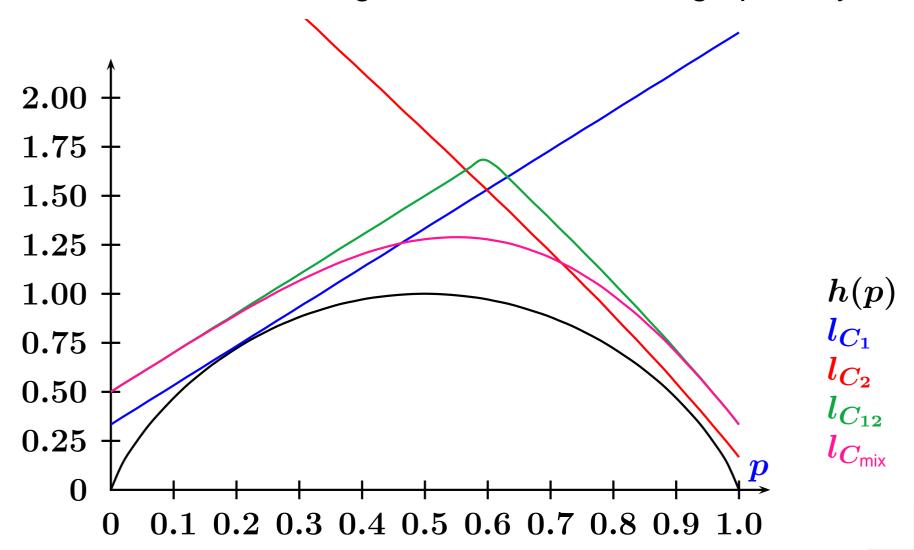
$$p_{\mathsf{mix}}(x^N) = rac{p(x^N| heta_1) + p(x^N| heta_2)}{2}$$











We conclude that

Using an ordinary source code only works (well) if we are accurate in predicting the source probabilities.

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We conclude that

- Using an ordinary source code only works (well) if we are accurate in predicting the source probabilities.
- That a two-part code works for more than one source. First part: description of the source (parameters). Second part: the compressed version of the sequence assuming the given source.
- Mixing (weighting) probabilities works at least as good as the two-part code and can be performed in one run through the data.

Theorem 1 [Optimal number of sources] For a sequence x^N generated by an binary i.i.d. source with unknown $\Pr\{X=1\}=\theta$ the optimal number of alternative sources is of order \sqrt{N} and the achieved redundancy of the resulting code C^* , relative to any i.i.d. source, is bounded as

$$r_N(C^*) < rac{\log_2 N}{2N} \left(1 + \epsilon
ight),$$

and also

$$r_N(C^*) > rac{\log_2 N}{2N} \left(1 - \epsilon
ight),$$

for any $\epsilon > 0$ and N sufficiently large. We shall not prove this theorem here.

Discussion:

For the binary i.i.d. source which is described by one parameter θ , the optimal redundancy is $\frac{\log_2 N}{2N}$.

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For the binary i.i.d. source which is described by one parameter θ , the optimal redundancy is $\frac{\log_2 N}{2N}$.

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The next result will explain some of these observations.

In the Bayesian Model estimation problem we looked at the log-regret criterion:

$$\log rac{p(x^N|\mathcal{M}_i, heta_i)}{p(x^N|\mathcal{M}_i)},$$

regret from not knowing the parameters.

or the criterion

$$\log rac{p(x^N|\mathcal{M}_i, heta_i)}{p(x^N)},$$

regret from not knowing the model plus parameters.

Remember

$$p(x^N|\mathcal{M}_i) = \int_{\Theta_i} p(heta_i|\mathcal{M}_i) p(x^N|\mathcal{M}_i, heta_i) \, d heta_i$$

and

$$p(x^N) = \int_{\mathfrak{M}} p(\mathcal{M}) p(x^N | \mathcal{M}) \, d\mathcal{M},$$

or more often when the model class is discrete

$$p(x^N) = \sum_{\mathcal{M} \in \, \mathfrak{M}} p(\mathcal{M}) p(x^N | \mathcal{M}).$$

If \mathcal{M}_i , θ_i has actually generated x^N then

$$-\log p(x^N|\mathcal{M}_i, heta_i)$$

is the ideal codeword length.

And

$$-\log p(x^N|\mathcal{M}_i)$$
 resp. $-\log p(x^N)$

is the actual codeword length of a good code using these 'estimated' probabilities.

Thus

$$\log rac{p(x^N|\mathcal{M}_i, heta_i)}{p(x^N|\mathcal{M}_i)}$$
 resp. $\log rac{p(x^N|\mathcal{M}_i, heta_i)}{p(x^N)}$

can be seen as

Data compression: The excess codeword length (individual redundancy).

Machine learning: The individual log-regret.

So now we see that the expected redundancy of a code C on sequences x^N from a source \mathcal{M}_i , θ_i , given by

$$r_N(C) = \sum_{x^N \in \mathcal{X}^N} p(x^N | \mathcal{M}_i, heta_i) \log rac{p(x^N | \mathcal{M}_i, heta_i)}{p(x^N | \mathcal{M}_i)}$$

resp.

$$r_N(C) = \sum_{x^N \in \mathcal{X}^N} p(x^N | \mathcal{M}_i, heta_i) \log rac{p(x^N | \mathcal{M}_i, heta_i)}{p(x^N)}$$

can also be seen as the expected log-regret with respect to \mathcal{M}_i and θ_i .

Redundancy-capacity theorem

We again take a Bayesian approach.

But we are also dealing with variable-length codes C. We first discuss how we can relate codeword lengths $l_C(x^N)$ to probabilities $p(x^N|\mathcal{M},\theta)$.

The answer is through the ideal codeword length

$$l_C(x^N) \sim -\log_2 p(x^N|\mathcal{M}, \theta).$$

Let C be any binary, prefix-free code with K words, where l_i denotes the length of the i^{th} code word, that satisfies the Kraft inequality with equality, i.e.

$$\sum_{i=1}^{K} 2^{-l_i} = 1.$$

We see that 2^{-l_i} plays the role of a probability value, namely

$$2^{-l_i} > 0$$
, because $0 < l_i < \infty$,

$$\sum_{i=1}^{K} 2^{-l_i} = 1, \quad \text{total probability is 1.}$$

A probability vector $Q=\{q_1,q_2,\ldots,q_K\}$ is called a dyadic probability vector, if for all $i,1\leq i\leq K$, there exists integers n_i such that

$$q_i=2^{-n_i}.$$

An example

$$Q = \left\{ \frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{8}, \frac{1}{16}, \frac{1}{32}, \frac{1}{32} \right\}$$
 is dyadic, namely

$$q_1=q_2=q_3=2^{-2}, \qquad q_4=2^{-3}, \ q_5=2^{-4}, \qquad q_6=q_7=2^{-5}, \ \sum^7 q_i=1.$$

The code that corresponds to this probability vector has codeword lengths

$$l_1 = l_2 = l_3 = 2, \quad l_4 = 3$$
 $l_5 = 4$
 $l_6 = l_7 = 5.$

An example of such a code can have the following 7 words

$$c_1 = 00$$
 $c_2 = 01$ $c_3 = 10$ $c_4 = 110$ $c_5 = 1110$ $c_6 = 11110$ $c_7 = 11111$

If C is a code that satisfies the Kraft inequality with equality, then we denote the corresponding, unique dyadic probability vector by Q_C .

The set of all dyadic probability vectors of the same length K will be denoted by $\mathcal{Q}_{\mathcal{C}}$, where the vector length K is not specified explicitely. We also write \mathcal{Q} for the set of all probability vectors of the same length K.

Obviously any $Q_C \in \mathcal{Q}_{\mathcal{C}}$ is also a member of \mathcal{Q} , so

$$Q_{\mathcal{C}} \subset Q$$
.

So again, let $Q_{\mathcal{C}}$ be the set of all dyadic probabilities and Q be the set of all probabilities.

 \mathcal{S} is the set of all sources parametrized by a vector θ that takes values in a parameter space Θ .

We have seen the example of the binary i.i.d. source with a one dimensional parameter $\theta = \Pr\{X = 1\}$ and $\Theta = [0, 1]$.

If $Q_C \in \mathcal{Q}_{\mathcal{C}}$ then the redundancy of of the corresponding code C is given by

$$egin{aligned} r_N(C) &= \sum_{x^N \in \mathcal{X}^N} p(x^N | heta) \log_2 rac{p(x^N | heta)}{Q_C(x^N)} \ &= D(p(X^N | heta) \|Q_C(X^N)) \end{aligned}$$

Maximizing over the parameter values we get the maximum expected redundancy of a given code C.

$$r_N^+(C) = \sup_{ heta \in \Theta} D(p(X^N| heta) \|Q_C(X^N))$$

Now we can look for the best possible code that minimizes the maximum expected redundancy.

$$r_N^+ = r_N^+(C^*) = \min_C r_N^+(C).$$

So C^* is the code that minimizes the worst-case expected redundancy over all parameter values.

 r_N^+ is the resulting minimax expected redundancy.

Instead of the worst-case redundancy we can also consider weighted redundancies.

Let $w(\theta)$ be a prior distribution over θ . The Bayes redundancy is given by

$$\mathcal{D}(w;Q_C) = \int_{\Theta} D(p(X^N| heta)\|Q_C(X^N))w(heta)\,d heta$$

Because the maximum is never smaller than the average, we have

$$r_N^+(C) \geq \mathcal{D}(w;Q_C),$$

and likewise we obtain for the best possible code

$$r_N^+ \geq \min_C \mathcal{D}(w;Q_C).$$

If we allow all probabilities, not only dyadic ones, we obtain:

$$\mathcal{oldsymbol{\mathcal{D}}}(oldsymbol{w};oldsymbol{Q}) = \int_{oldsymbol{\Theta}} w(heta) D(p(X^N| heta) \|Q(X^N)) \, d heta$$

and because we can minimize over a larger set

$$r_N^+ \geq \min_Q \mathcal{D}(w;Q).$$

It turns out that the Q^* that realizes the minimum is the $w(\theta)$ weighted probability

$$Q^* = \int_{m{\Theta}} p(x^N| heta) w(heta) \, d heta.$$

And thus we can observe:

$$\mathcal{D}(w;Q^*) = \int_{\Theta} w(heta) D(p(X^N| heta) \|Q^*(X^N)) \, d heta$$

And thus we can observe:

$$egin{aligned} \mathcal{D}(oldsymbol{w}; oldsymbol{Q}^*) &= \int_{\Theta} w(heta) D(p(X^N| heta) || oldsymbol{Q}^*(X^N)) \, d heta \ &= \int_{\Theta} \sum_{x^N \in \mathcal{X}^N} oldsymbol{w}(heta) p(x^N| heta) \log_2 rac{p(x^N| heta)}{oldsymbol{Q}^*(x^N)} \, d heta \end{aligned}$$

Channel input heta probabilities w(heta)

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Channel transition probabilities $p(x^N|\theta)$

And thus we can observe:

$$egin{align} \mathcal{D}(oldsymbol{w};oldsymbol{Q}^*) &= \int_{oldsymbol{\Theta}} w(heta) D(p(X^N| heta) \| Q^*(X^N)) \, d heta \ &= \int_{oldsymbol{\Theta}} \sum_{x^N \in \mathcal{X}^N} w(heta) p(x^N| heta) \log_2 rac{p(x^N| heta)}{oldsymbol{Q}^*(oldsymbol{x}^N)} \, d heta \ \end{split}$$

Channel output x^N probabilities $Q^*(x^N)$

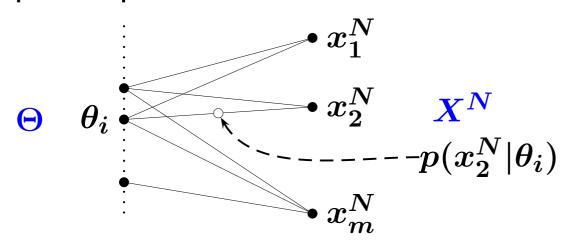
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Because this last bound is independent of the prior $w(\theta)$, we can tighten the bound by maximizing over all possible priors $w(\theta)$ and find

$$r_N^+ \geq \max_{w(heta)} I(heta; X^N) = C_{ heta o X^N}.$$

So the redundancy is lower bounded by (often it is equal to) the capacity of the channel from the source parameters to the source output sequence x^N .



Redundancy: learning source parameters from data

- Source coding: we don't want this, because it causes extra codeword length, but it is unavoidable.
- Machine learning: this is what's it about, but we cannot learn faster then the channel capacity.

The meaning of model information

Efficient description of data can be split into two parts:

- Information about the 'model'
 - Universal compression redundancy: The description of the parameters of the data generating process.
- Selection of one of the 'possible' sequences.
 - Universal compression: One of the "typical sequences" selected and described with $NH(P_x)$ bits.

The meaning of model information

The first part describes what the model 'can do'.

bits of π : Almost zero complexity. The model is easy to describe and can only generate this sequence. Easy to predict bits.

bits from an i.i.d. source $\theta = \frac{1}{2}$: Highly complex. The model is very simple but the set of possible sequences is large. Hard to predict bits.

The meaning of model information

Occam's razor:

One should not increase, beyond what is necessary, the number of entities required to explain anything.

The most useful statement of the principle for scientists is:

When you have two competing theories which make exactly the same predictions, the one that is simpler is the better.

Universal source coding:

Take the simplest model that describes your data.

The two-part description separates model information from random selection

Universal coding: There is a certain unavoidable cost for parameters in a model. It is the price for learning the parameters.

Distinguishable models (parameter values): For a sequence of length N we can use (selection or weighting) about \sqrt{N} distinct values.

Occam's razor: Take the simplest explanation that explains the observations.

This results in the notion of stochastic complexity

$$-\log_2 p(x^N|\mathcal{M})$$
 $p(x^N|\mathcal{M}) = rac{p(x^N|\mathcal{M},\hat{ heta}(x^N))}{\sum_{x^N \in \mathcal{X}^N} p(x^N|\mathcal{M},\hat{ heta}(x^N))}$

is known as the NML (Normalized Maximum Likelihood). That is must be normalized is reasonable because

$$\sum_{x^N \in \mathcal{X}^N} p(x^N | \mathcal{M}, \hat{ heta}(x^N)) \geq 1$$

And the normalizing constant determines the model cost. Note that we assume here that the model priors $p(\mathcal{M})$ are all equal!

Suppose I have two model classes, \mathcal{M}_1 and \mathcal{M}_2 , for my data x^N and the stochastic complexity $-\log_2 p(x^N|\mathcal{M}_1)$ is smaller than $-\log_2 p(x^N|\mathcal{M}_2)$.

Because the model information part is proportional to $\log_2 N$ and the "noise" part is proportional to N, a smaller complexity means "less noise". So \mathcal{M}_1 explains more of the data.

This leads to the Minimum Description Length Principle.

The best model for the data is is the model that results in the smallest stochastic complexity.

Stochastic complexity ≈ ideal codeword length. Coding interpretation:

$$L(\theta) = O(\log N); L(\mathsf{noise}) = O(N)$$

$$oxed{L(heta_2)}$$
 $oxed{\mathsf{noise_2}}$

$$oxed{L(heta_1)}$$
 $oxed{\mathsf{noise}_1}$

Say x^N with N=1000. $L(\mathsf{noise}_2) + L(\theta_2) = 500 + 5k_2$.

With model \mathcal{M}_1 , x^N has smaller stochastic complexity:

 $L(\mathsf{noise}_1) {>} L(\mathsf{noise}_2)$ hardly possible because $L(\theta_2) - L(\theta_1)$ cannot be large.

Stochastic complexity ≈ ideal codeword length. Coding interpretation:

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With model \mathcal{M}_1 , x^N has smaller stochastic complexity:

 $L(\text{noise}_1) < L(\text{noise}_2)$ very likely.

So, \mathcal{M}_1 explains more of the data (less noise)

Stochastic Complexity (MDL)

"Real data model": binary 1th order Markov,

$$\theta_0 = \Pr\{X_i = 1 | x_{i-1} = 0\} = \frac{1}{4},$$

$$\theta_1 = \Pr\{X_i = 1 | x_{i-1} = 1\} = \frac{1}{2}$$

Then: $\Pr\{X_i = 1\} = \frac{1}{3}$.

 \mathcal{M}_1 is i.i.d. with $\hat{ heta}_1 pprox rac{1}{3}$.

 \mathcal{M}_2 is 1th order Markov with $\hat{\theta}_2 \approx (\frac{1}{4}, \frac{1}{2})$.

$$H(X|\mathcal{M}_1,\hat{ heta}_1)=0.918$$
 bit.

$$H(X|\mathcal{M}_2,\hat{ heta}_2)=0.874$$
 bit.

Stochastic Complexity (MDL)

Stochastic complexity

$$S.C._1 \sim rac{\log_2 N}{2} + 0.918N.$$
 $S.C._2 \sim \log_2 N + 0.874N.$

For $N < 70: S.C._1 < S.C._2$ and for $N > 70: S.C._1 > S.C._2$.

So if there is not enough data the MDL selects a smaller model than the "true" model.

This is good!

There is not enough data to estimate properly a complex model.