

CS340

Bayesian concept learning cont'd

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Bayesian inference

- H : Hypothesis space of possible concepts:
- $X = \{x_1, \dots, x_n\}$: n examples of a concept C .
- Evaluate hypotheses given data using Bayes' rule:

$$p(h | X) = \frac{p(X | h) p(h)}{\sum_{h' \in H} p(X | h') p(h')}$$

- $p(h)$ [“prior”]: domain knowledge, pre-existing biases
- $p(X|h)$ [“likelihood”]: statistical information in examples.
- $p(h|X)$ [“posterior”]: degree of belief that h is the true extension of C .

Hypothesis space

- Mathematical properties (~50):
 - odd, even, square, cube, prime, ...
 - multiples of small integers
 - powers of small integers
 - same first (or last) digit
- Magnitude intervals (~5000):
 - all intervals of integers with endpoints between 1 and 100
- Hypothesis can be defined by its **extension**

$$h = \{x : h(x) = 1, x = 1, 2, \dots, 100\}$$

Likelihood $p(X|h)$

- Assume samples are iid, so $p(X|h) = \prod_{i=1}^n p(x_i|h)$
- **Size principle:** Smaller hypotheses receive greater likelihood, and exponentially more so as n increases.

$$p(X|h) = \begin{cases} \frac{1}{|size(h)|^n} & \text{if all } x_1, \dots, x_n \in h \\ 0 & \text{if any } x_i \notin h \end{cases}$$

- This is the likelihood of the *ordered sequence* x_1, \dots, x_n sampled randomly (with replacement) from h (**strong sampling assumption**).
- Captures the intuition of a representative sample.

Likelihood function

- Since $p(\vec{x}|h)$ is a distribution over vectors of length n , we require that, for all h , $\sum_{\vec{x}} p(x|h) = 1$
- It is easy to see this is true, e.g., for h =even numbers, $n=2$

$$\sum_{x_1=1}^{100} \sum_{x_2=1}^{100} p(x_1, x_2|h) = \sum_{x_1=1}^{100} \sum_{x_2=1}^{100} p(x_1|h)p(x_2|h) = \sum_{x_1 \in \text{even}} \sum_{x_2 \in \text{even}} \frac{1}{50} \frac{1}{50} = 1$$

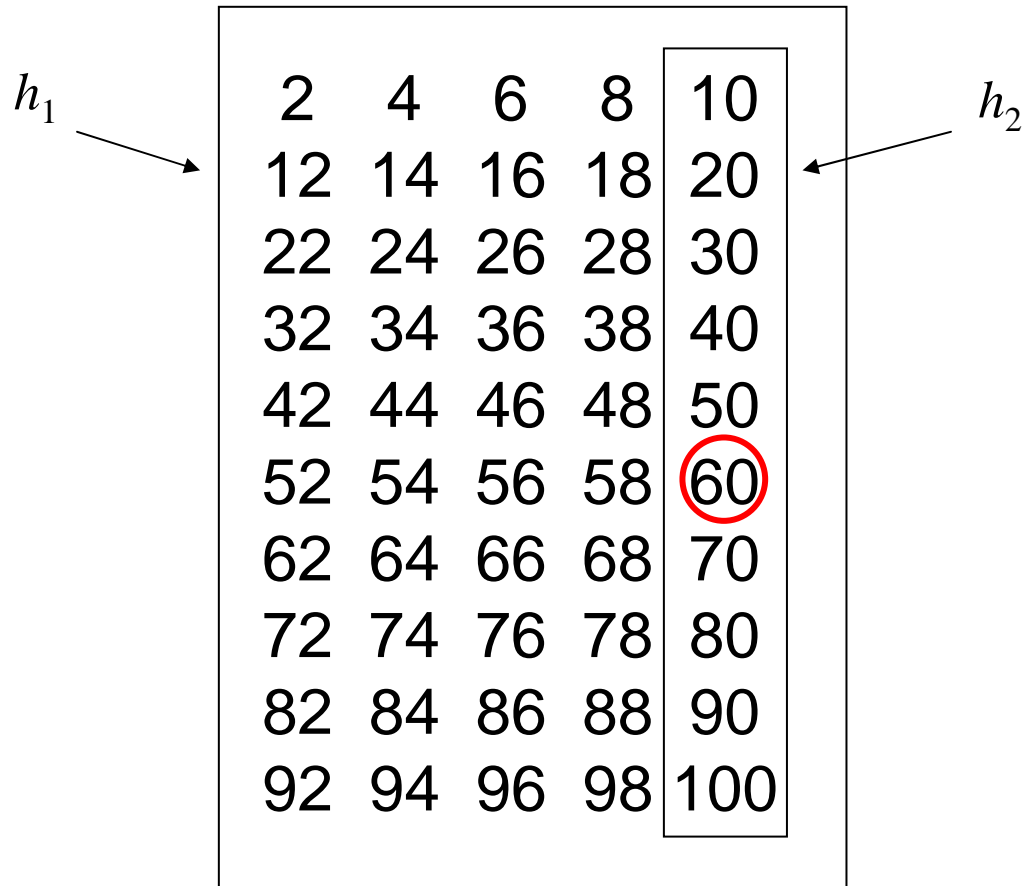
- If x is fixed, we do not require $\sum_h p(X|h) = 1$
- Hence we are free to multiply the likelihood by any constant independent of h

Illustrating the size principle

The diagram illustrates the size principle using a 10x5 grid of numbers. The numbers are arranged in columns of 10, starting from 2 in the top-left and ending at 100 in the bottom-right. A smaller box highlights the last column (10, 20, 30, 40, 50, 60, 70, 80, 90, 100). Two arrows, labeled h_1 and h_2 , point towards the grid from the left and right respectively.

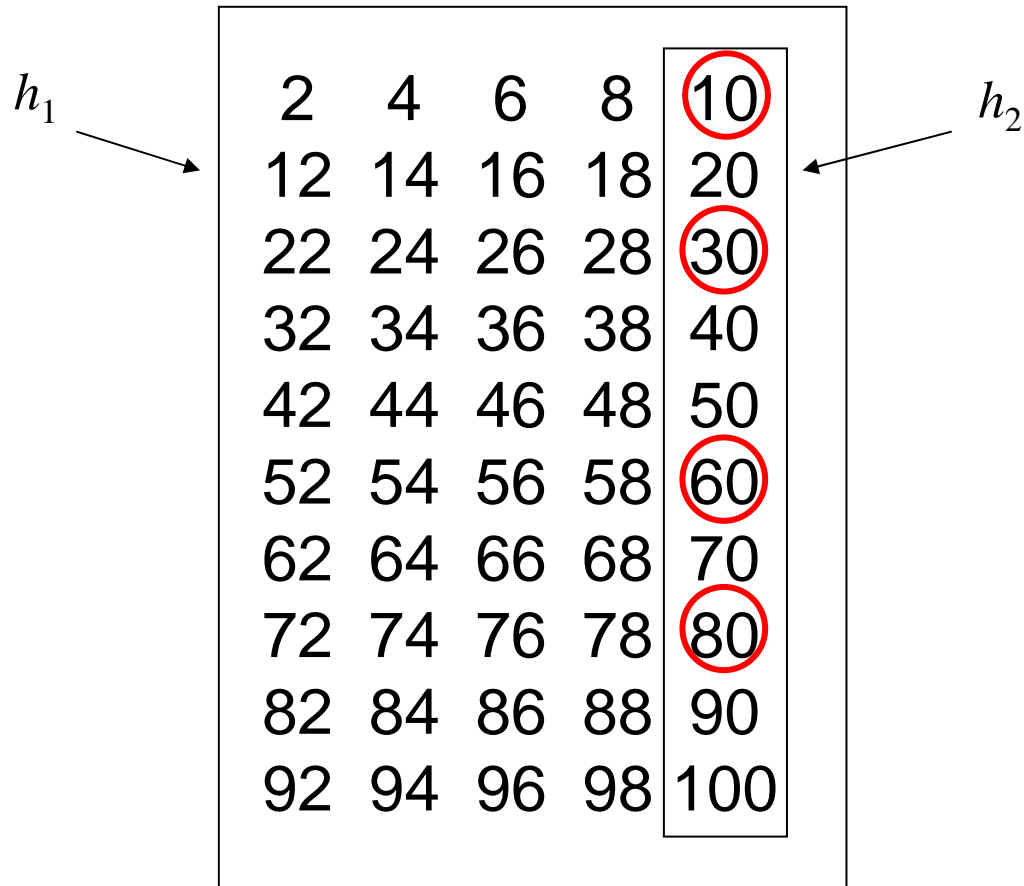
2	4	6	8	10
12	14	16	18	20
22	24	26	28	30
32	34	36	38	40
42	44	46	48	50
52	54	56	58	60
62	64	66	68	70
72	74	76	78	80
82	84	86	88	90
92	94	96	98	100

Illustrating the size principle



Data slightly more of a coincidence under h_1

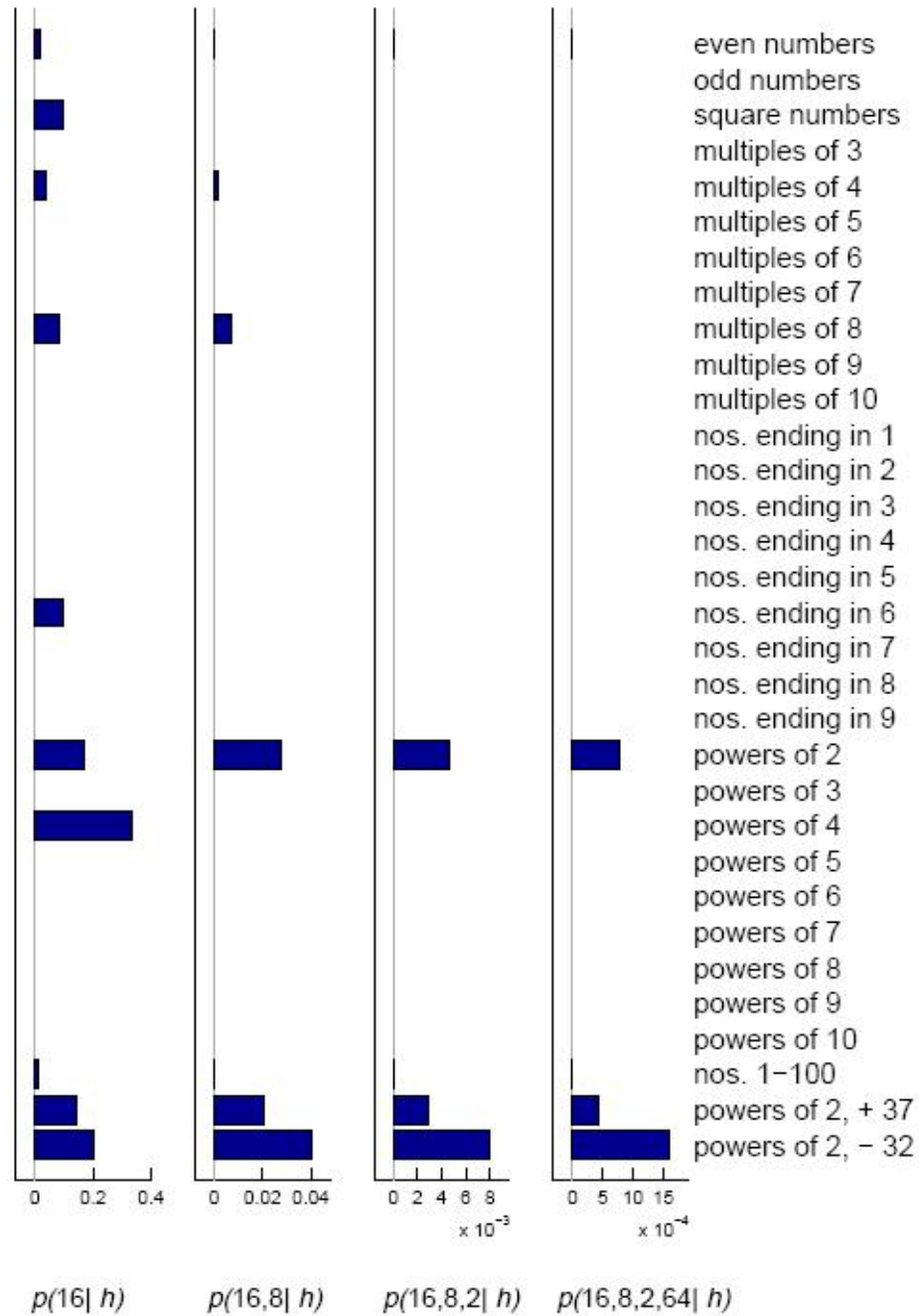
Illustrating the size principle



Data *much* more of a coincidence under h_1

Example of likelihood

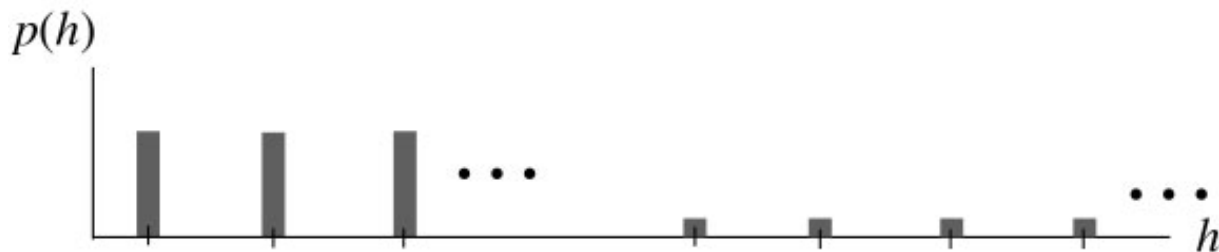
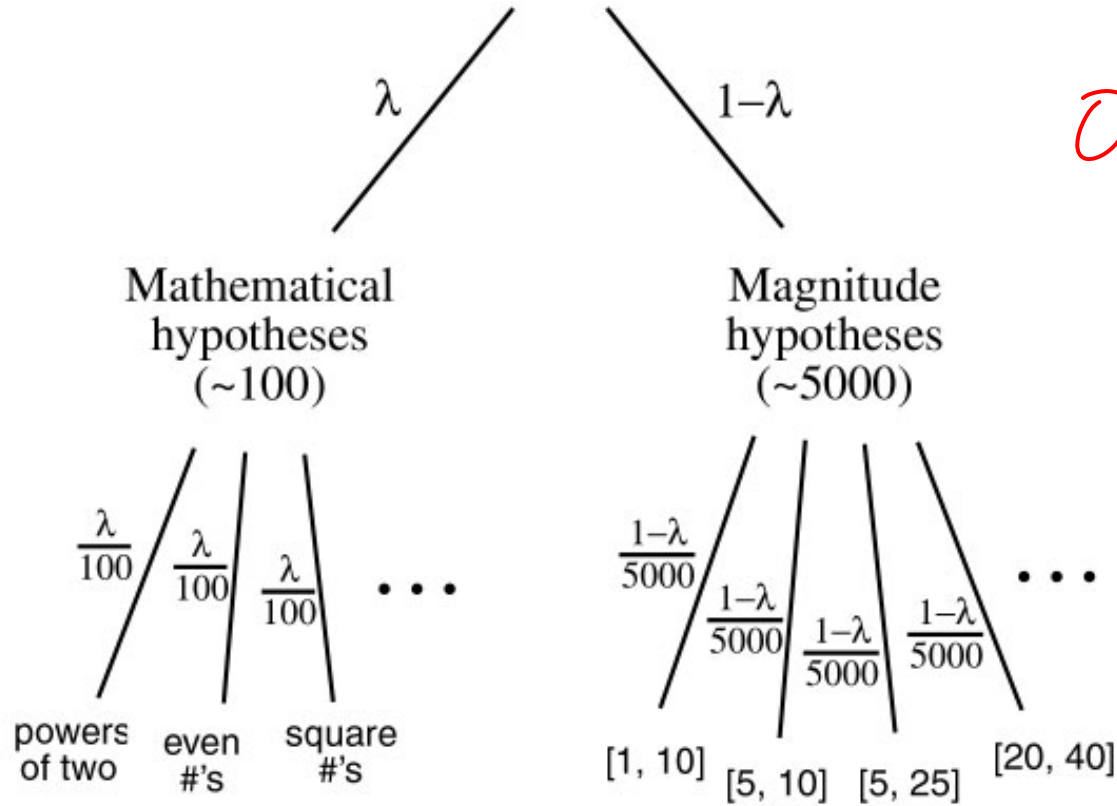
- $X = \{20, 40, 60\}$
- $H1 = \text{multiples of } 10 = \{10, 20, \dots, 100\}$
- $H2 = \text{even numbers} = \{2, 4, \dots, 100\}$
- $H3 = \text{odd numbers} = \{1, 3, \dots, 99\}$
- $P(X|H1) = 1/10 * 1/10 * 1/10$
- $p(X|H2) = 1/50 * 1/50 * 1/50$
- $P(X|H3) = 0$



Hierarchical prior

$$\text{Total probability mass} = \sum_h p(h) = 1$$

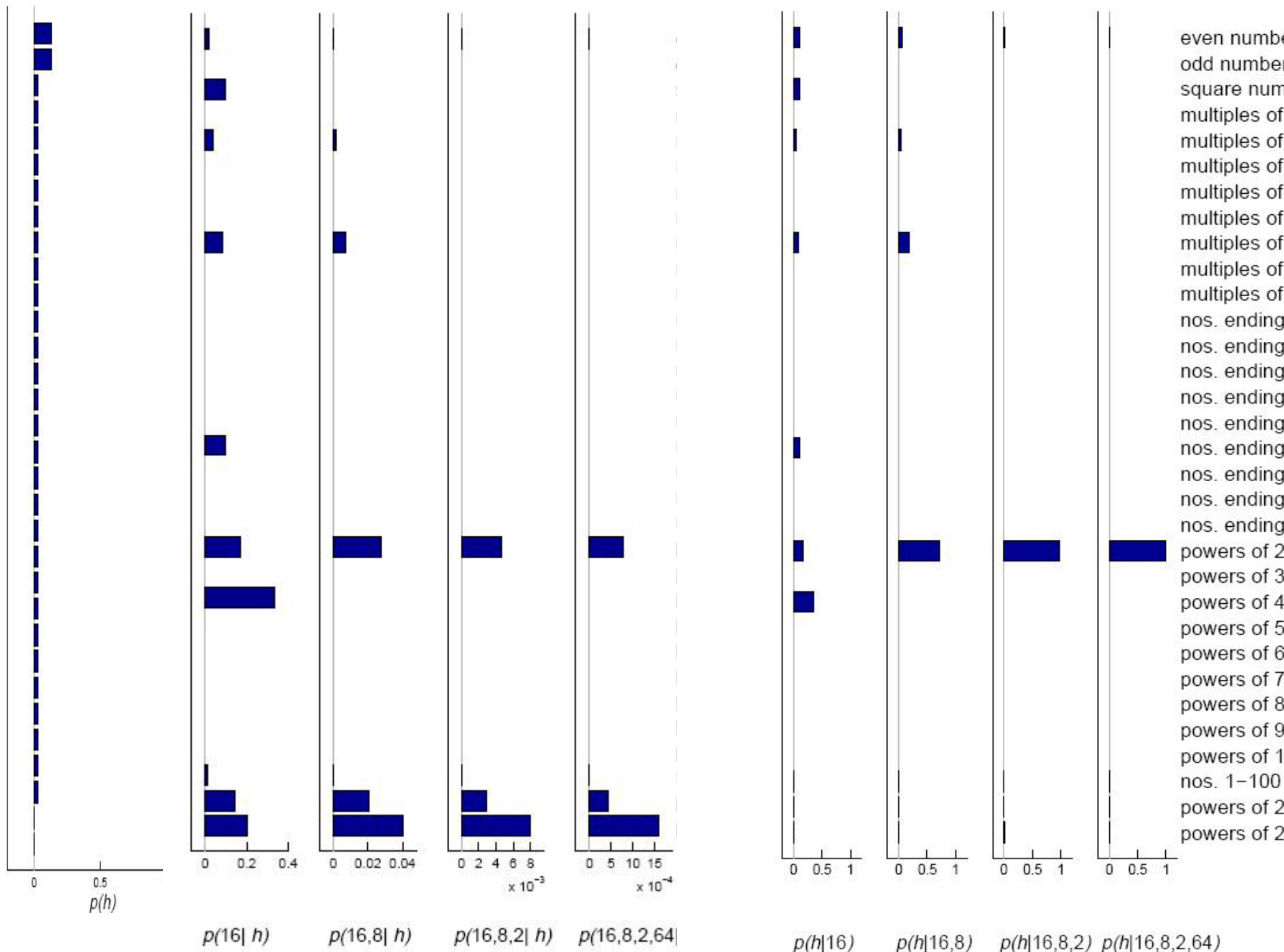
$$0 < \lambda < 1$$



Computing the posterior

- In this talk, we will not worry about computational issues (we will perform brute force enumeration or derive analytical expressions).

$$p(h | X) = \frac{p(X | h) p(h)}{\sum_{h' \in H} p(X | h') p(h')}$$



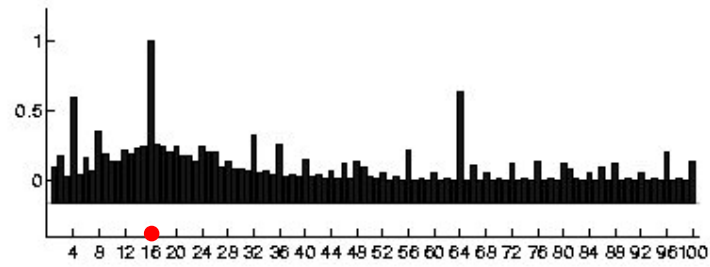
Generalizing to new objects

Given $p(h|X)$, how do we compute $p(y \in C | X)$, the probability that C applies to some new stimulus y ?

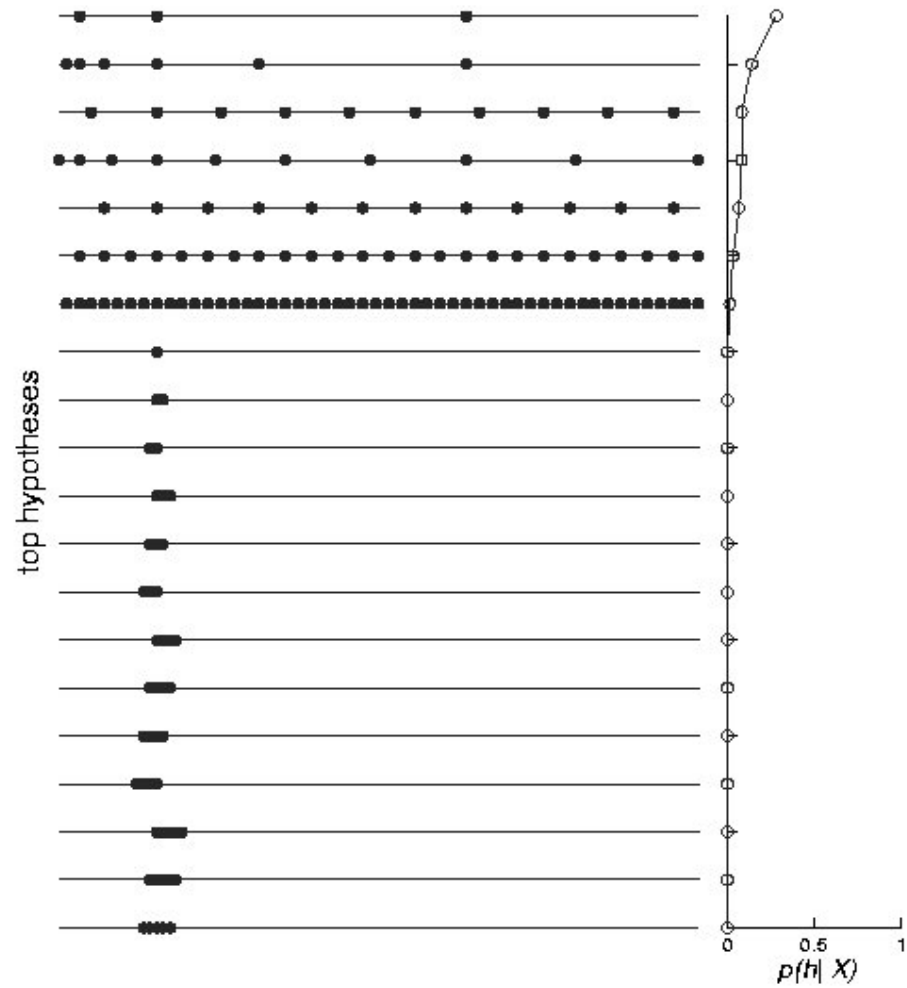
Posterior predictive distribution

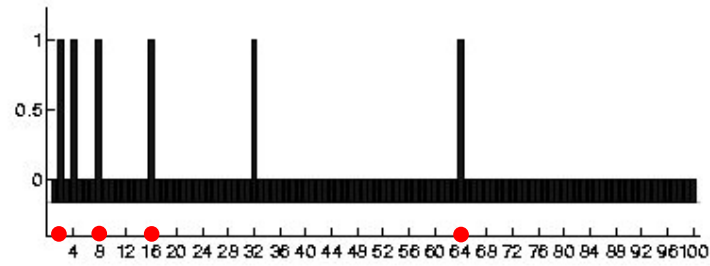
Compute the probability that C applies to some new object y by averaging the predictions of all hypotheses h , weighted by $p(h|X)$
(Bayesian model averaging):

$$\begin{aligned} p(y \in C | X) &= \sum_{h \in H} p(y \in C | h) p(h | X) \\ &= \sum_{h \supset \{y, X\}} p(h | X) \end{aligned}$$



Examples:
16





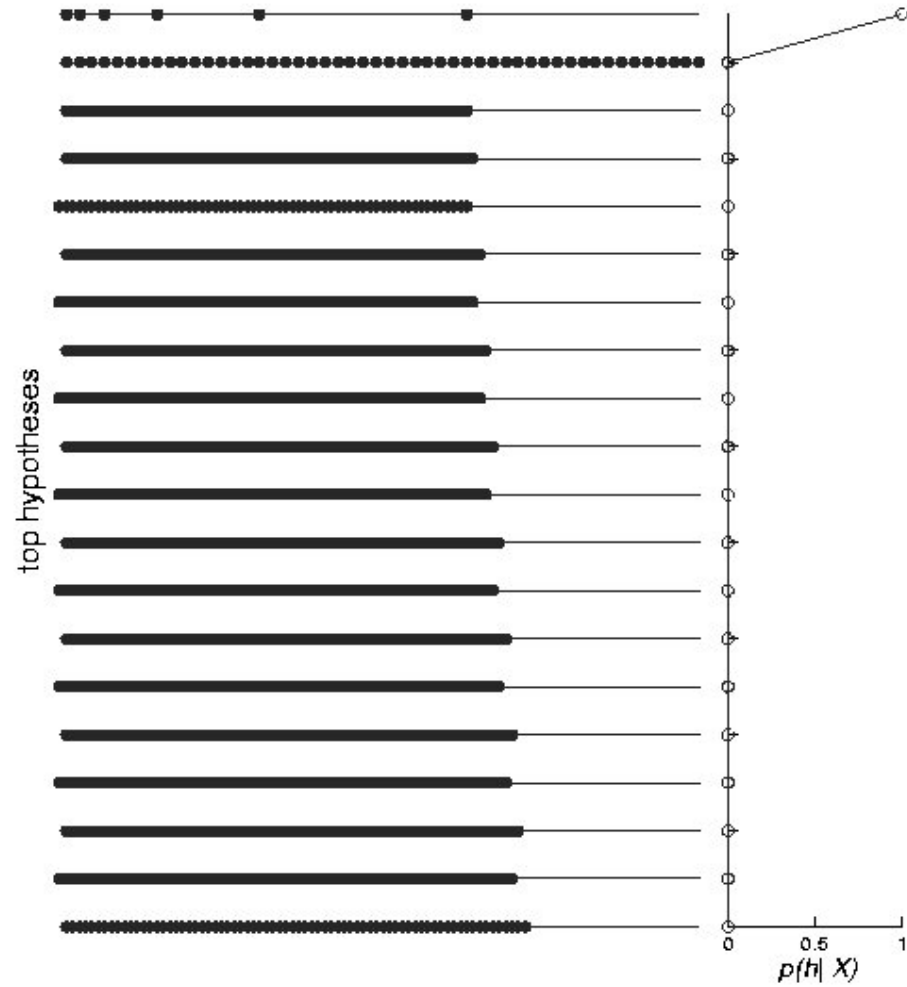
Examples:

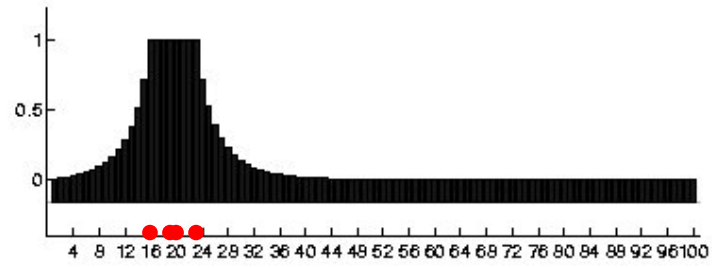
16

8

2

64





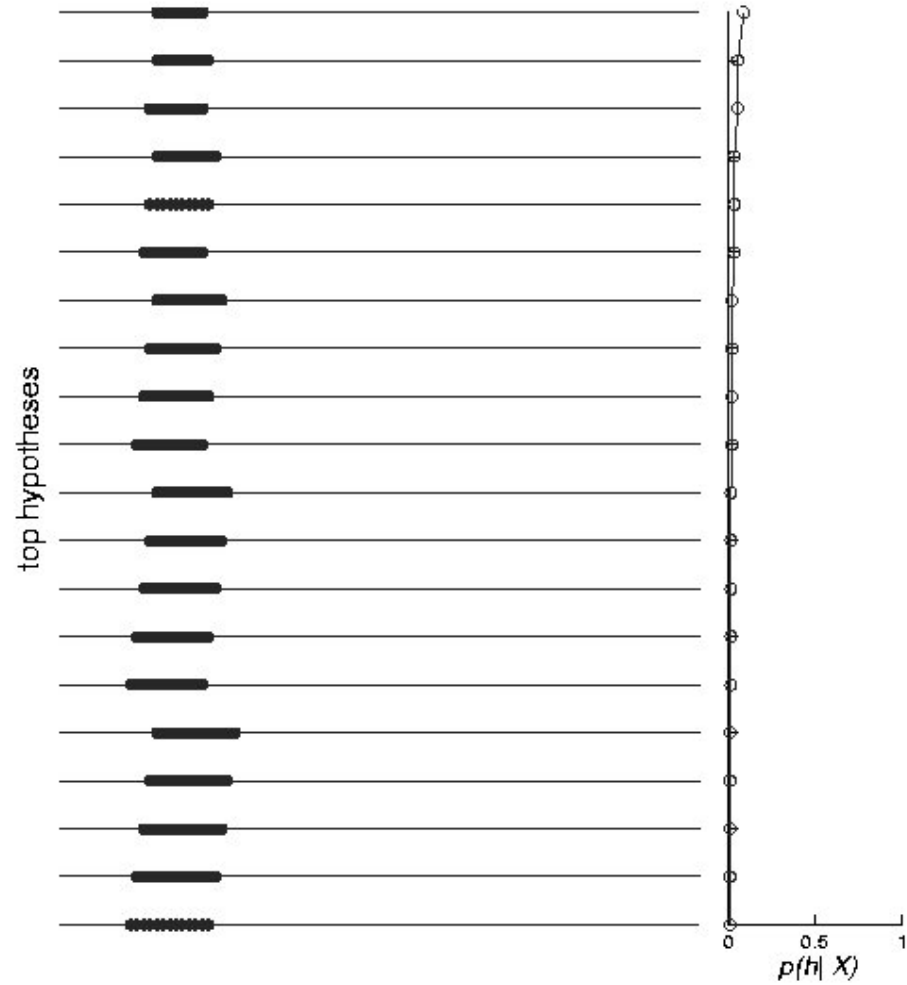
Examples:

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20

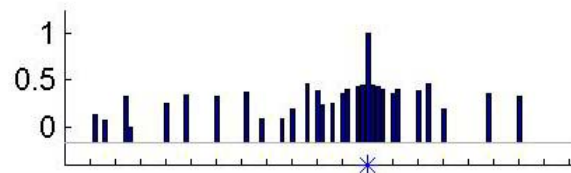
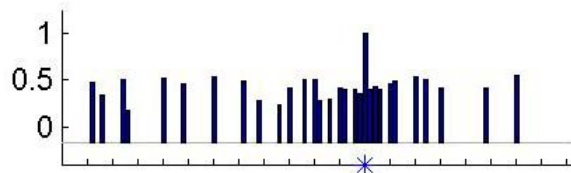


+ Examples

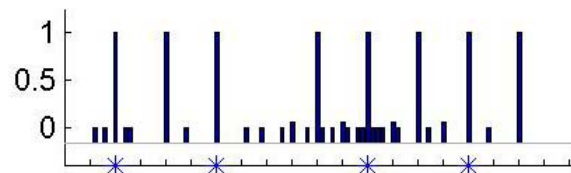
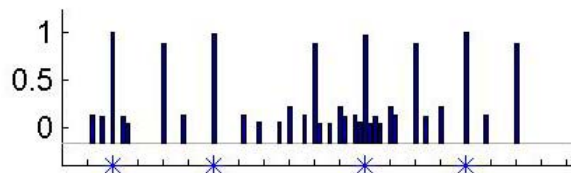
Human generalization

Bayesian Model

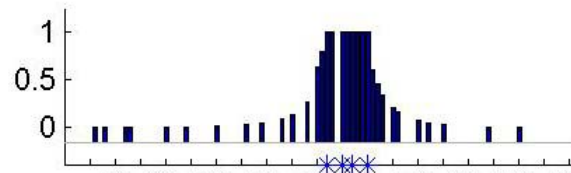
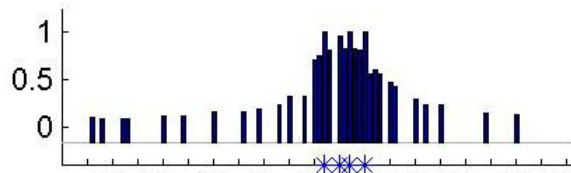
60



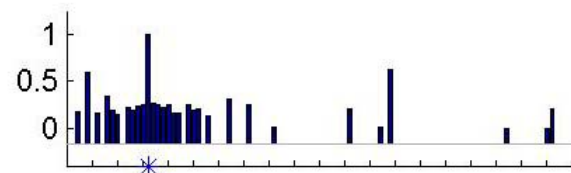
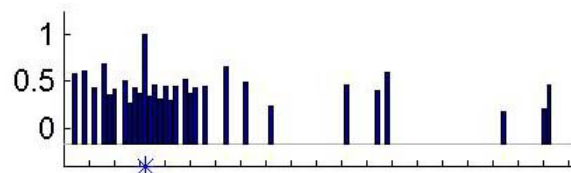
60 80 10 30



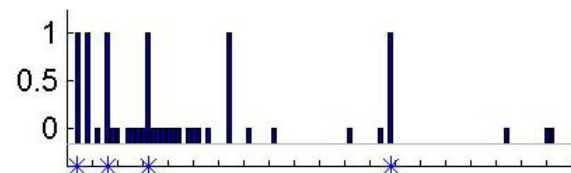
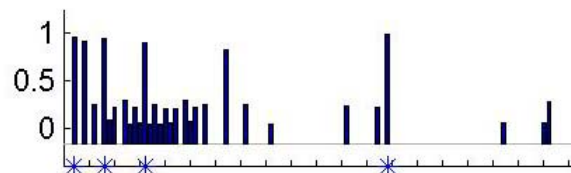
60 52 57 55



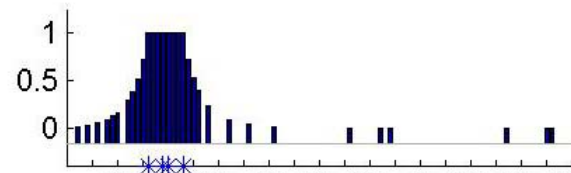
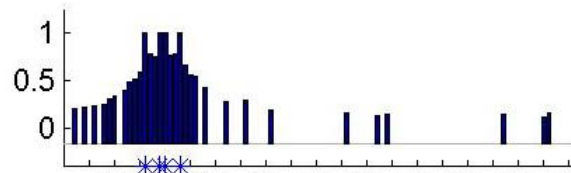
16



16 8 2 64



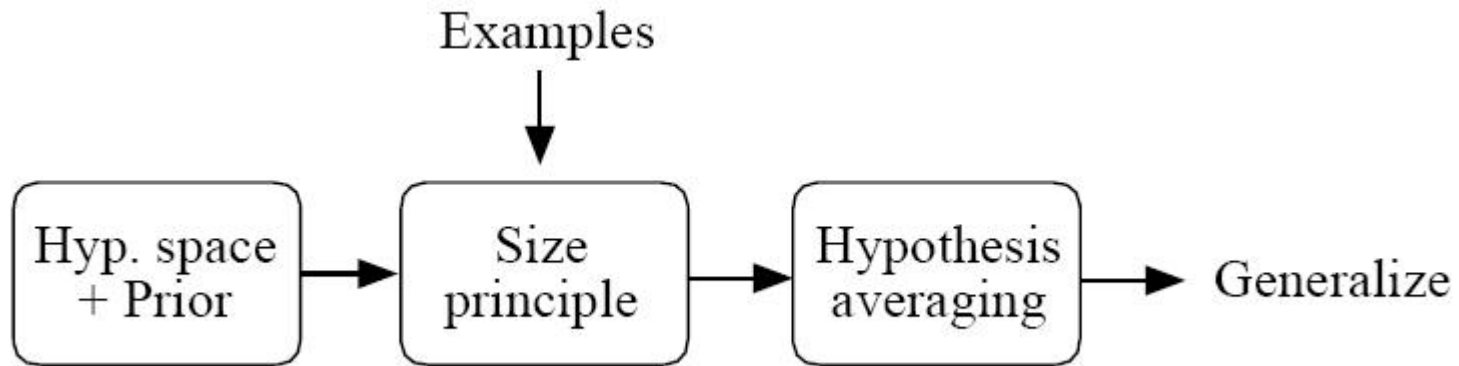
16 23 19 20



10 20 30 40 50 60 70 80 90 100

10 20 30 40 50 60 70 80 90 100

Summary of the Bayesian approach



1. Constrained hypothesis space H
2. Prior $p(h)$
3. Likelihood $p(X|h)$
4. Hypothesis (model) averaging:

$$p(y \in C | X) = \sum_h p(y \in C | h) p(h | X)$$