

SciLifeLab

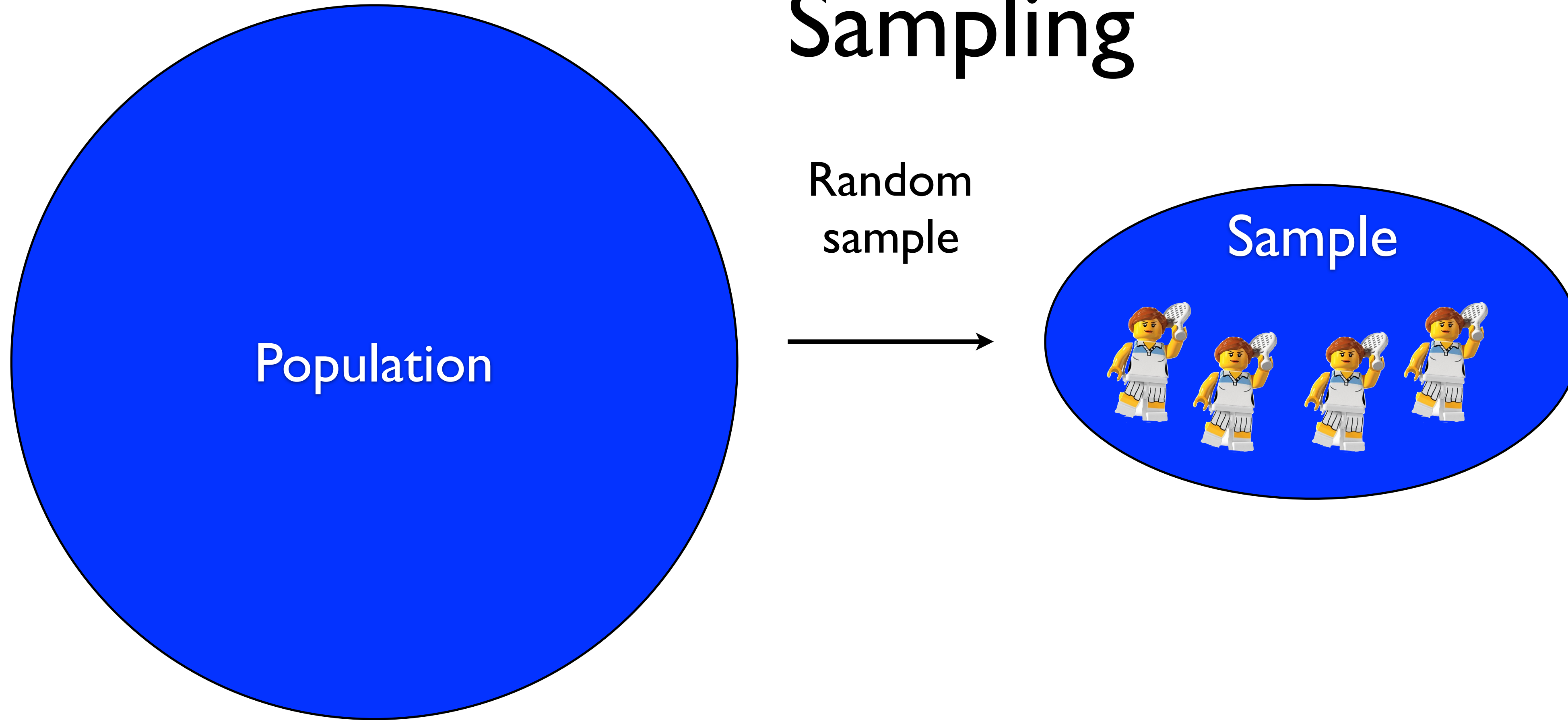
Hypothesis Testing

CB2030

Lukas Käll, KTH

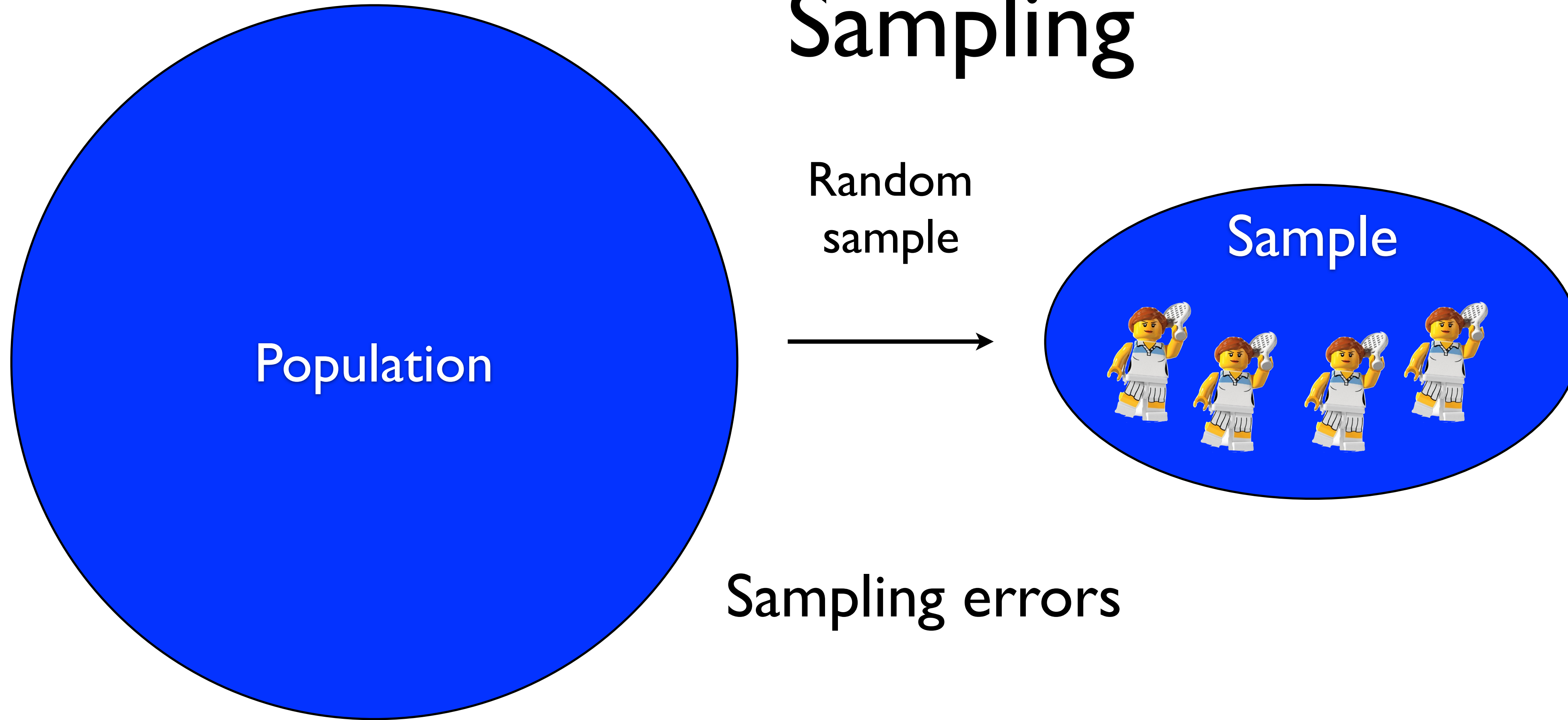


Sampling



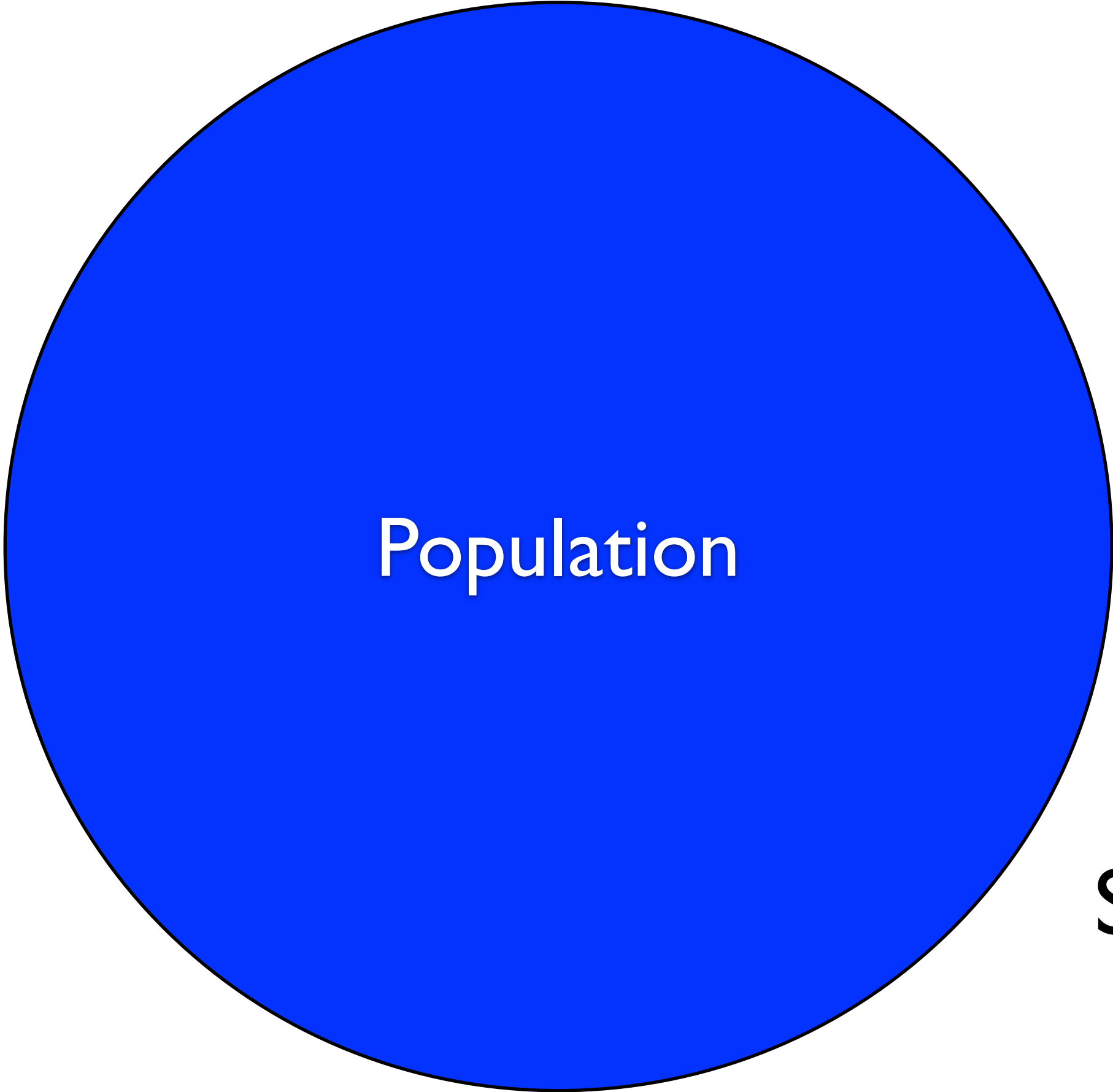
- We observe a population through a selected sample

Sampling

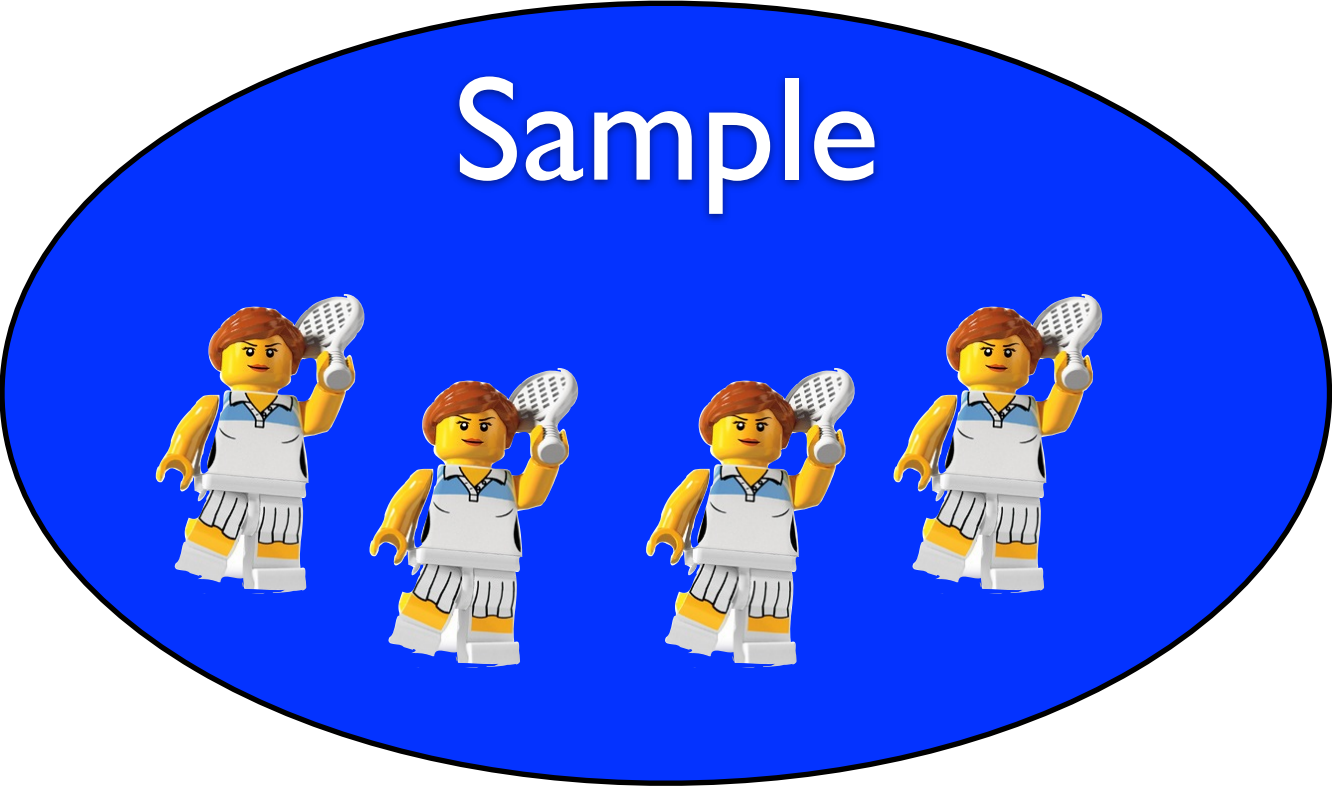


- We observe a population through a selected sample

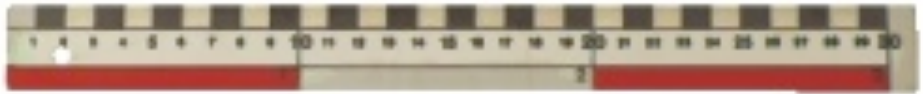
Sampling



Random
sample



Sampling errors

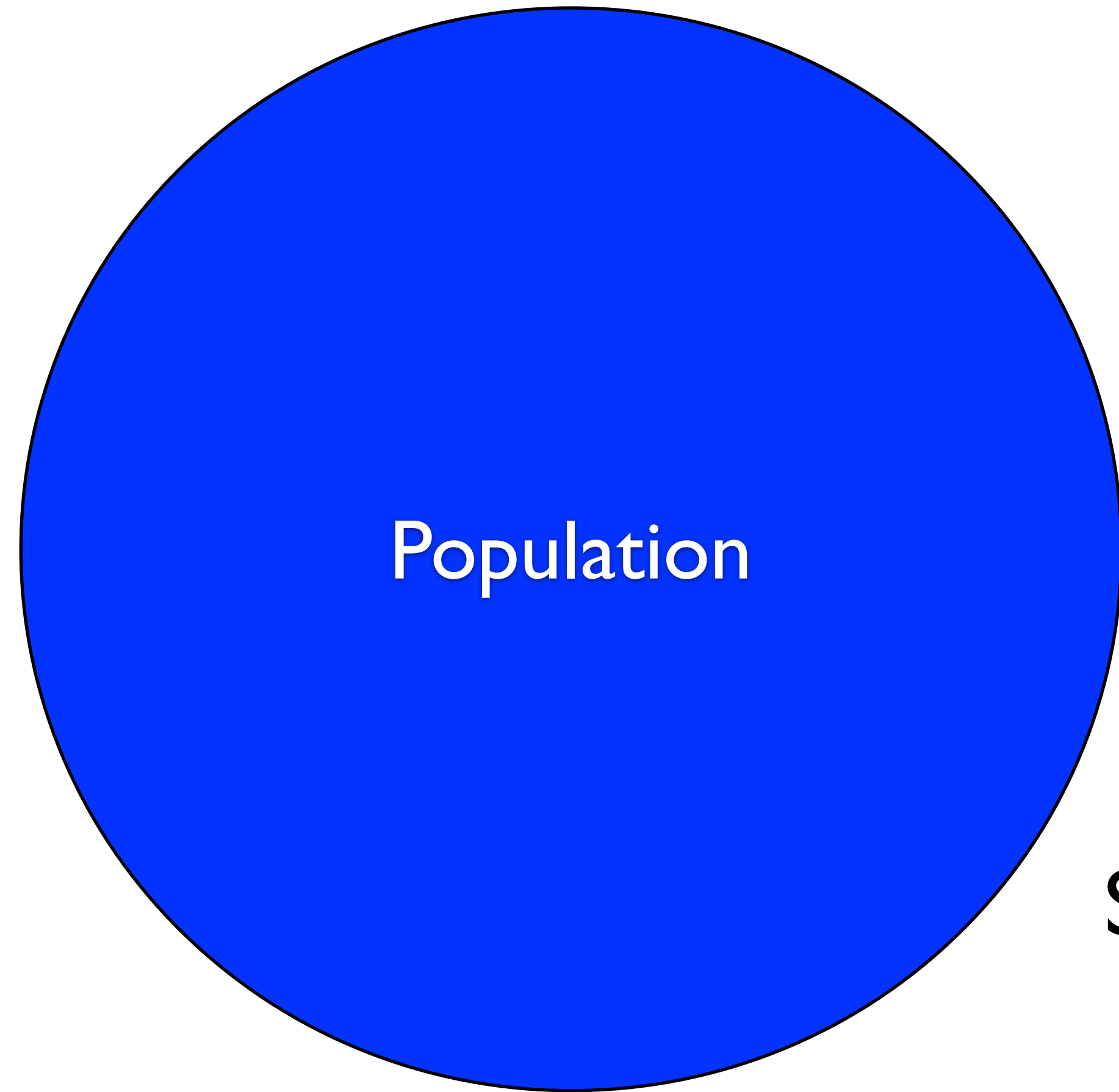


Measurement Errors

- 1. Systematic
- 2. Noise

- We observe a population through a selected sample

Biological variation

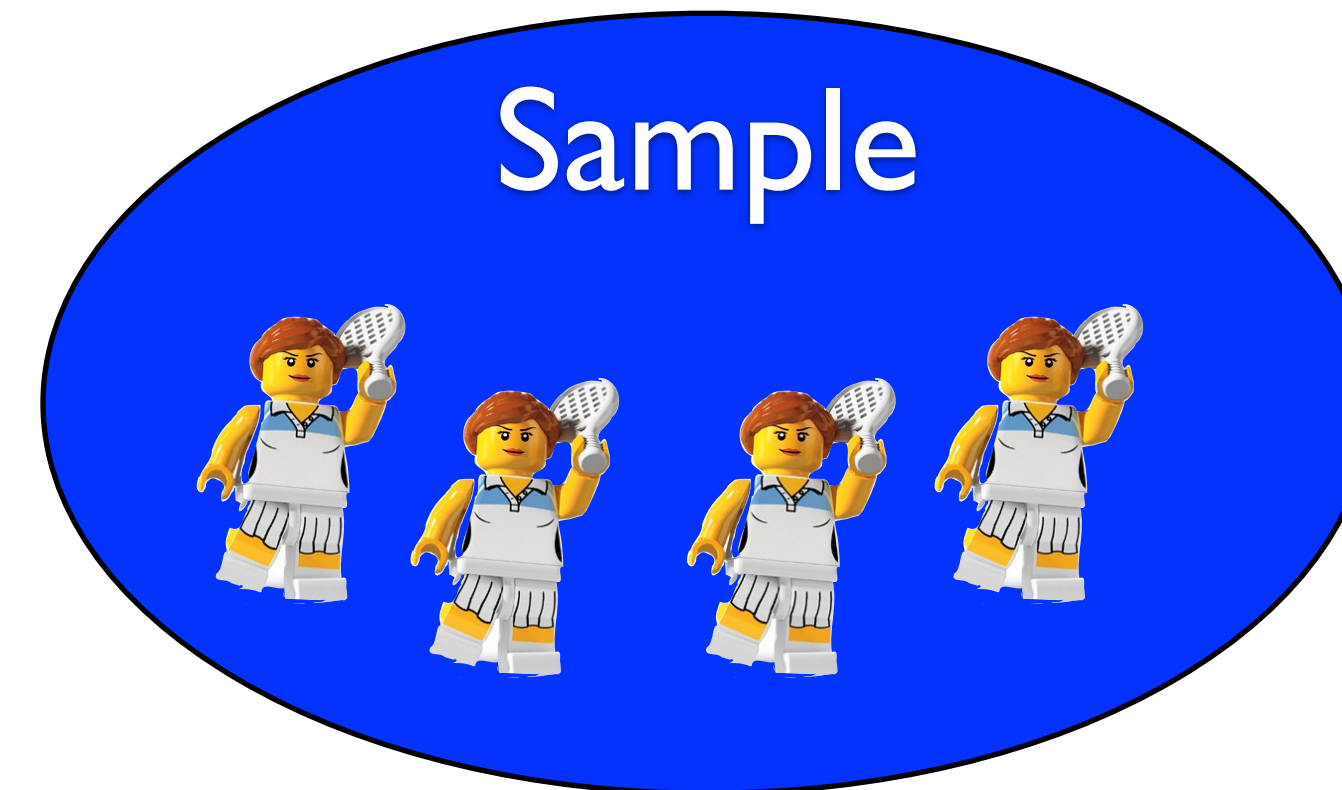


Sampling

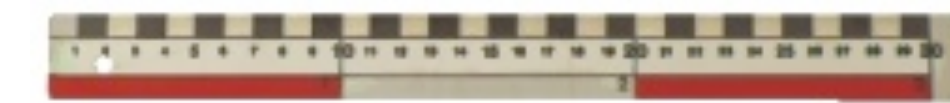
Random
sample



Technical variation



Sampling errors

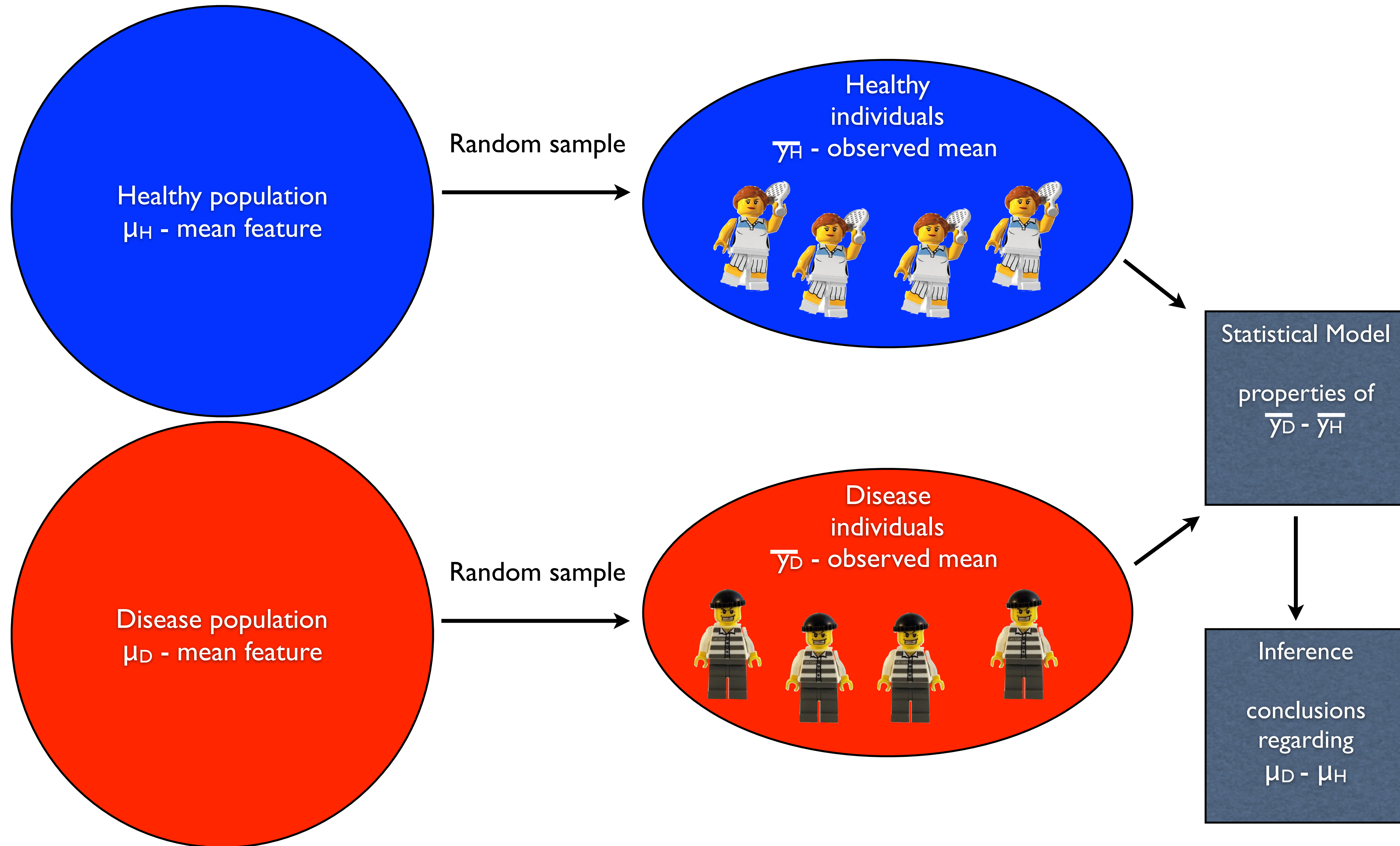


Measurement Errors

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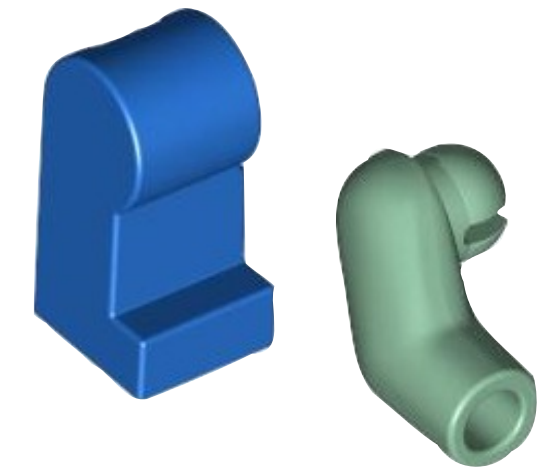
- We observe a population through a selected sample

Statistical inference procedure



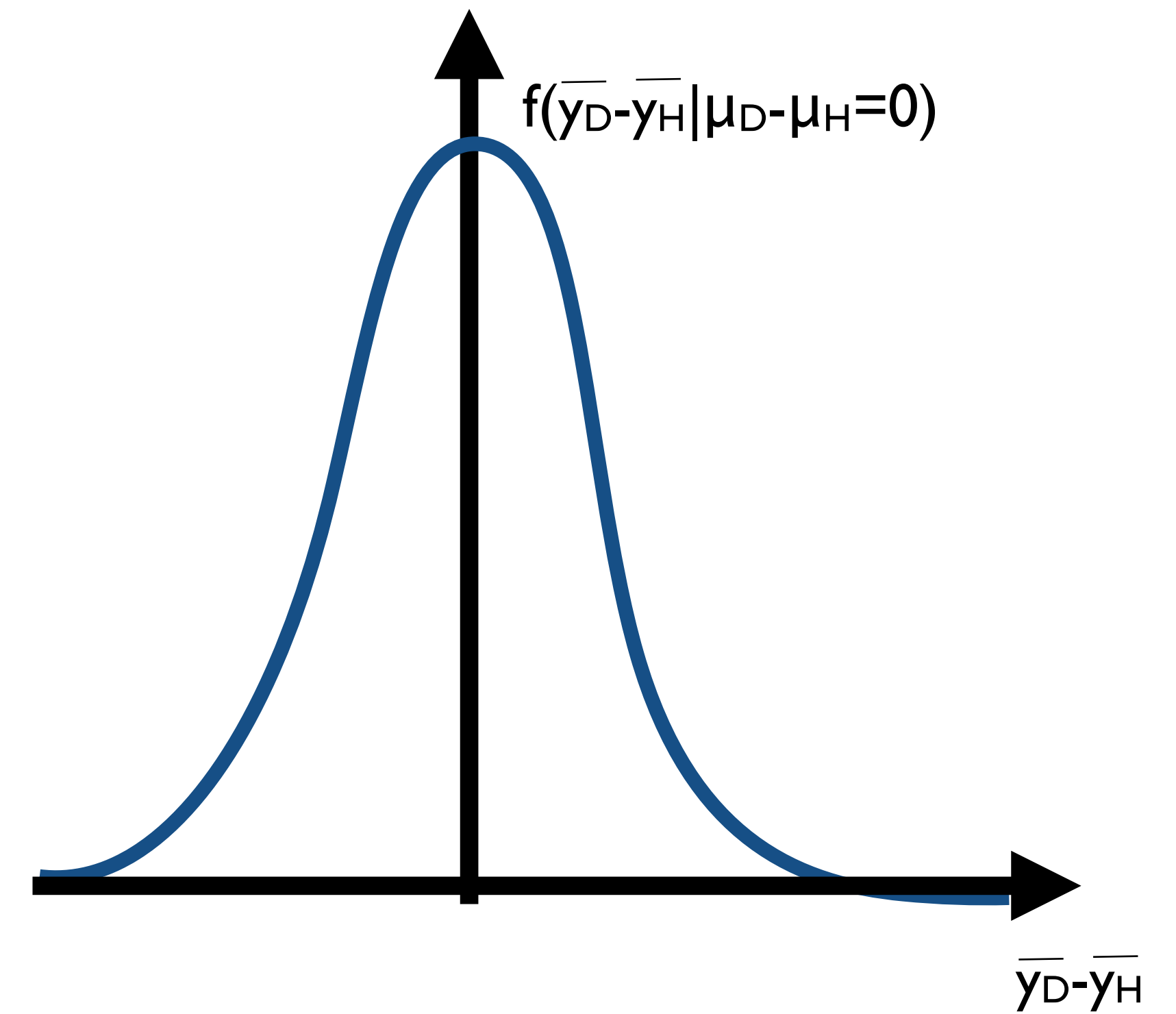
Our Hypotheses

- H_0 : The *null* hypothesis. The situation we are not interested in (typically $\mu_D - \mu_H = 0$)
- H_1 : The *alternative* hypothesis. The situation we want to detect (typically $\mu_D - \mu_H \neq 0$)



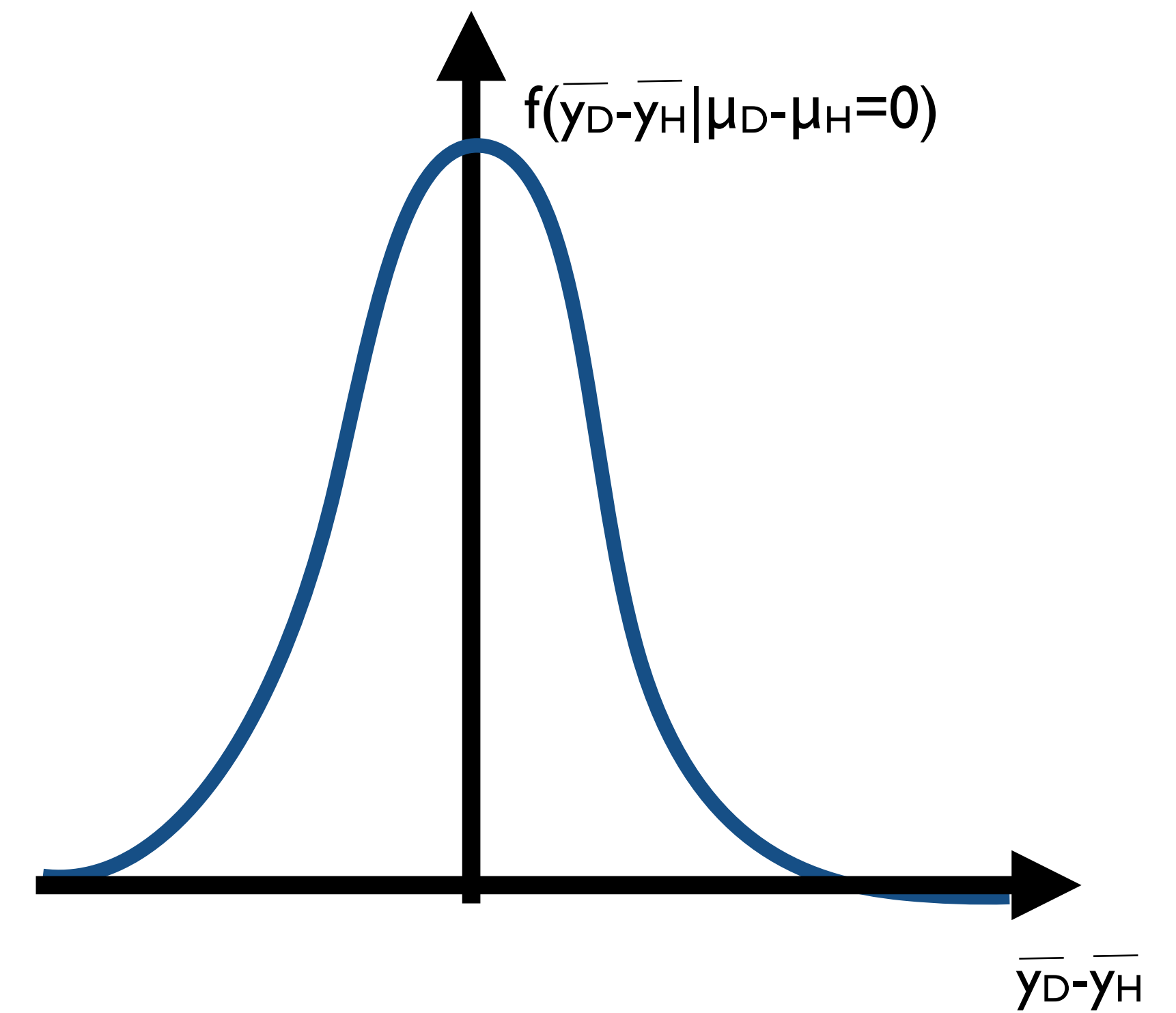
Sampling Distribution

- The distribution of any statistic of measurements in a sample, e.g. its mean, is known as a sampling distribution

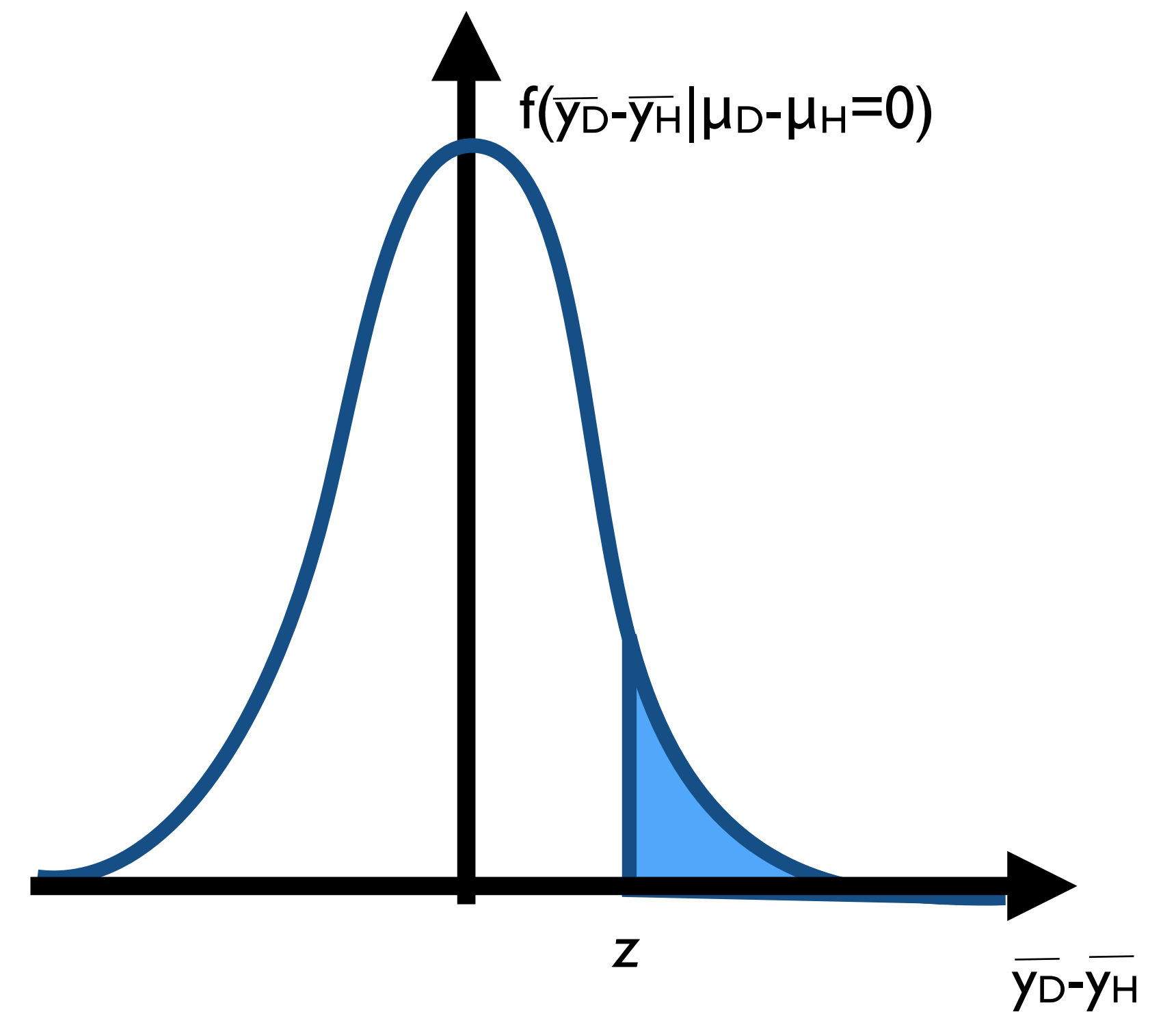


Sampling Distribution

- The distribution of any statistic of measurements in a sample, e.g. its mean, is known as a sampling distribution
- If we know the sampling distribution of the difference in sample mean under H_0 we can calculate how extreme the difference is.

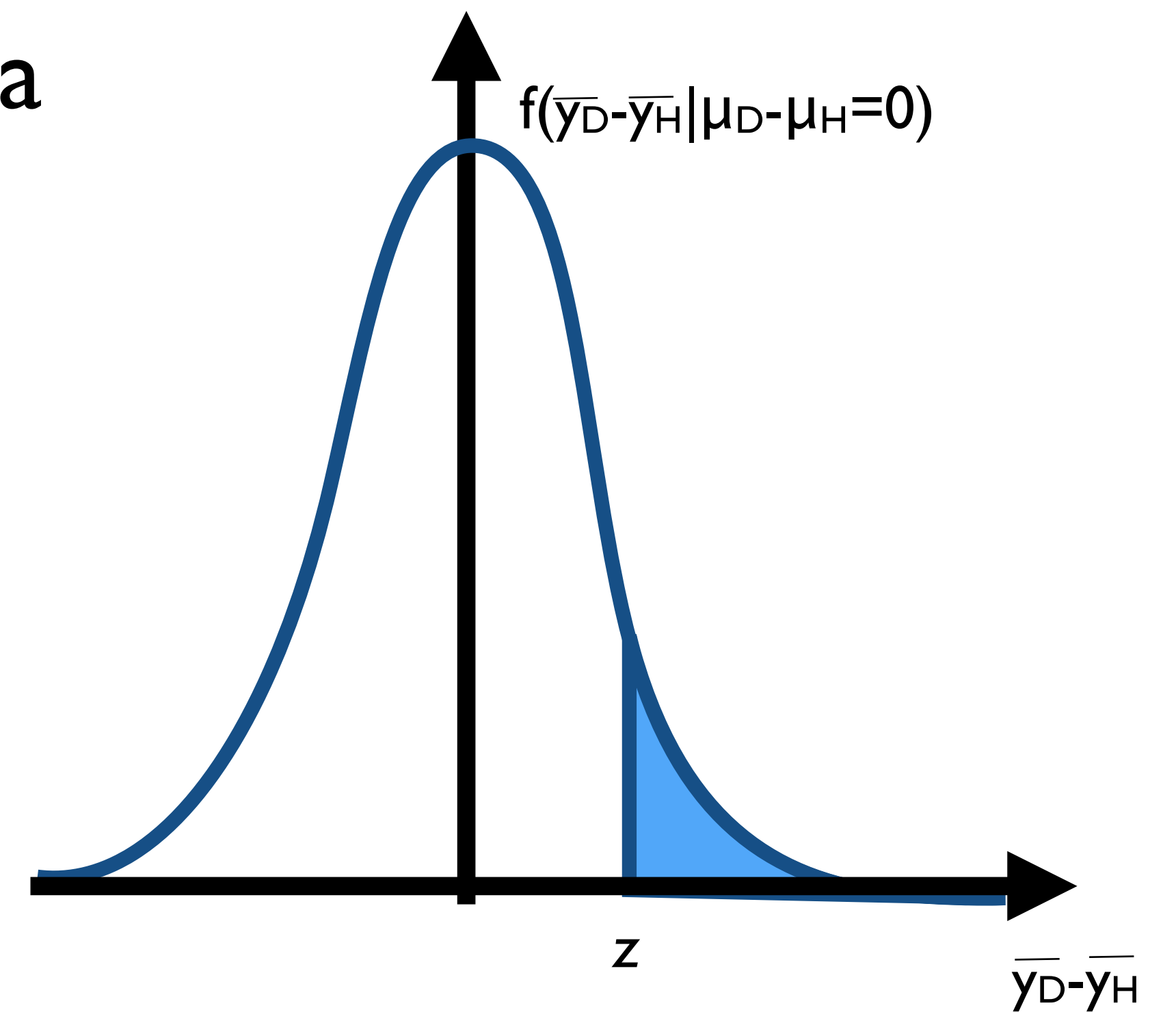


p value



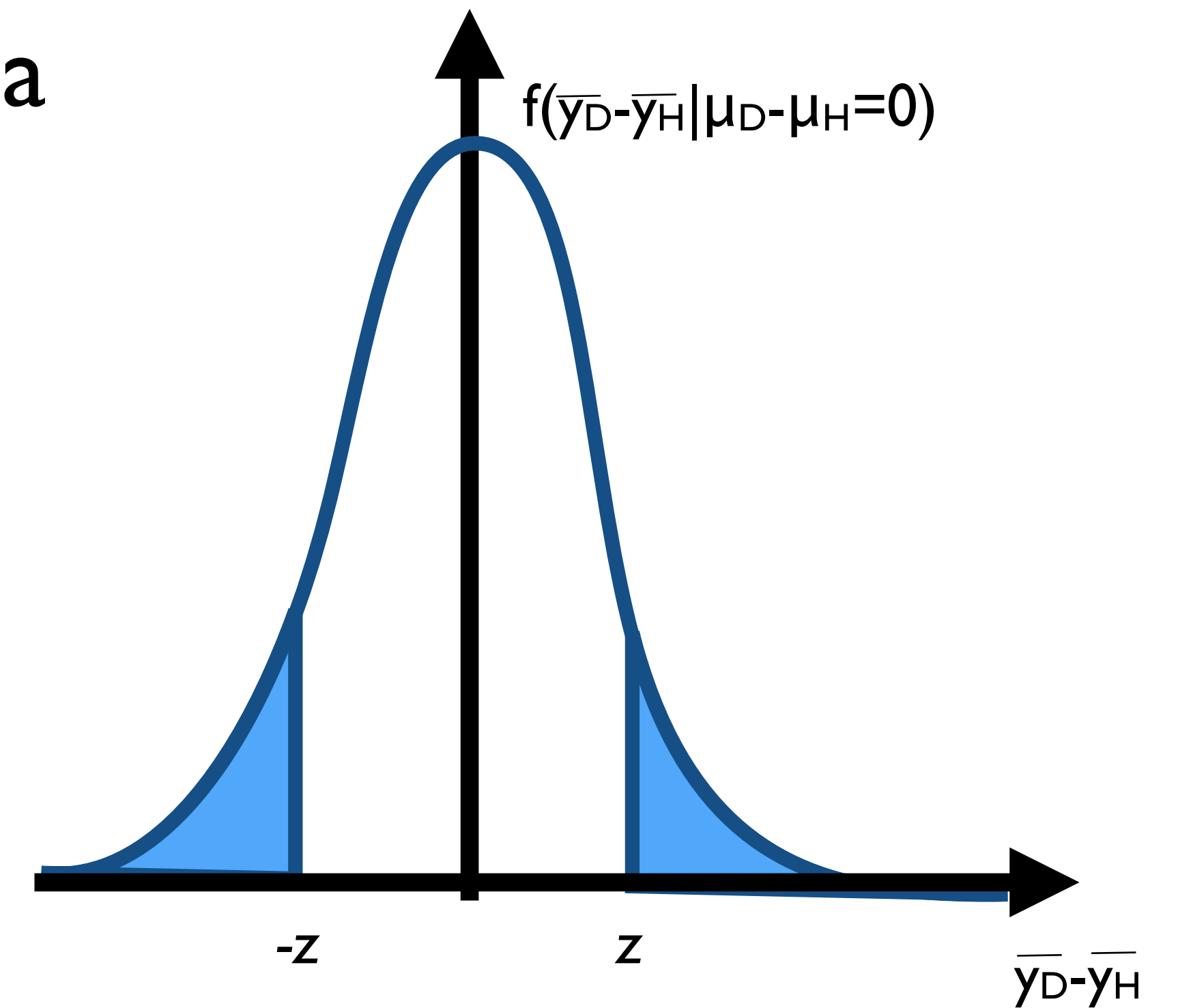
p value

- $\Pr(\bar{y}_D - \bar{y}_H \geq z | \mu_D - \mu_H = 0)$, *i.e.* the probability of a result at least as extreme as the one that was observed, given H_0 , is known as a one-sided p value.



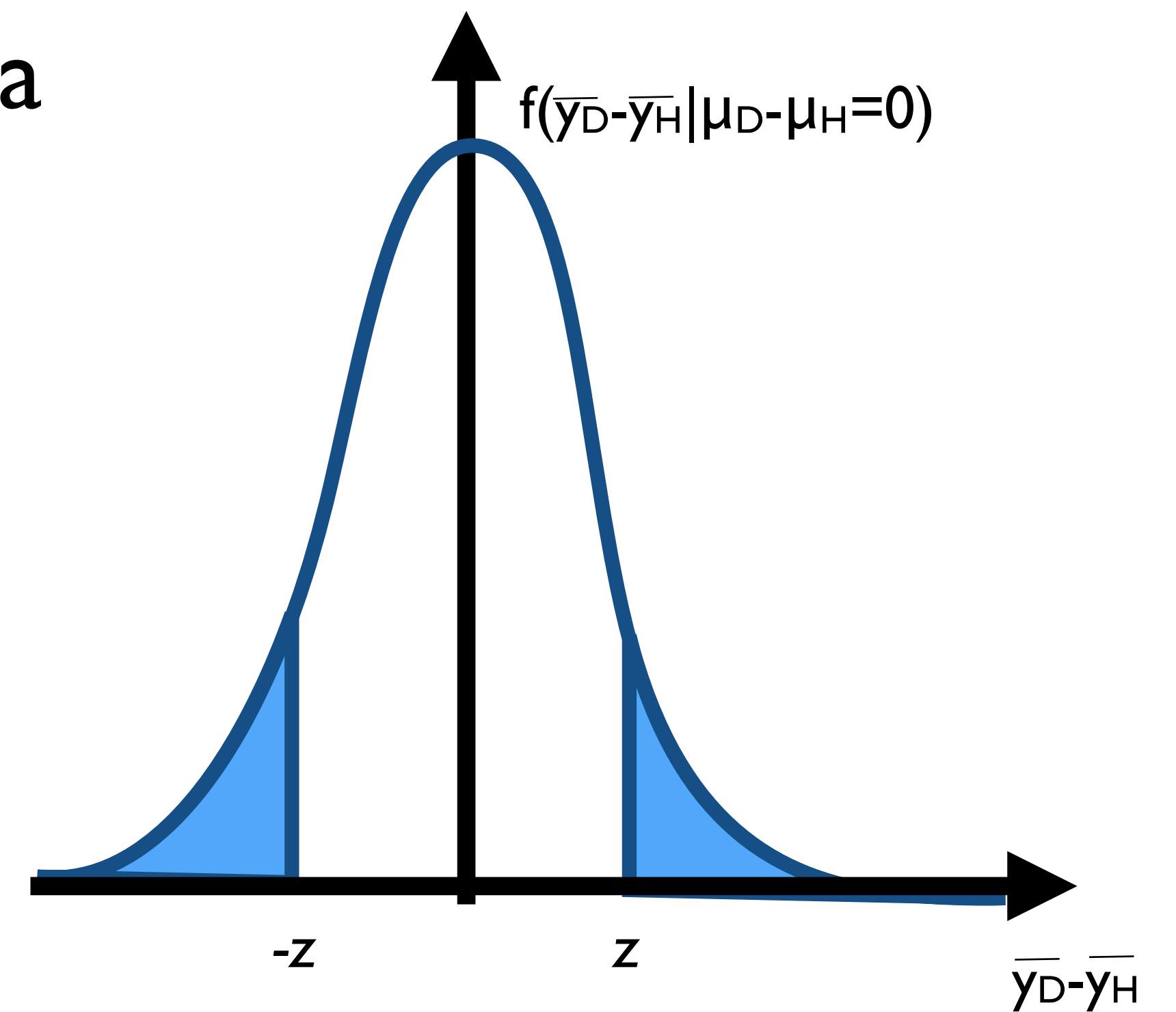
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- p values are uniformly distributed under H_0 .



Statistical inference procedure

