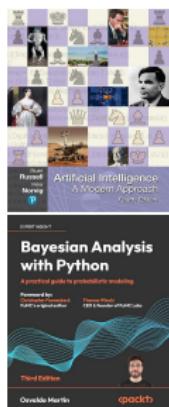


7.2: Posterior-based Decisions

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References:

- AIMA (Artificial Intelligence: a Modern Approach)
 - Chap 15: Probabilistic programming
- Martin, Bayesian Analysis with Python, 2018 (2e)



- ***Posterior-Based Decisions***
 - Chemical Shift: Example
 - Posterior Predictive Checks
 - Groups Comparison

Posterior-Based Decisions

- Sometimes describing the posterior is not enough
 - **Make decisions based on inference**
- E.g., is the coin fair ($\theta = 0.5$) or biased?
 - $\mathbb{E}[\hat{\theta}] = 0.324$ suggests bias
 - Can't rule out unbiased since:
 - $HPI = [0.03, 0.65]$
 - $0.5 \in HPI$
 - For **sharper decisions**:
 - Collect more data to reduce posterior spread
 - Define a more informative prior

Savage-Dickey Density Ratio

- **Savage-Dickey ratio** tests *point null-hypotheses* in Bayesian inference
- **Idea:** compare prior and posterior densities at a single point θ_0

$$BF_{01} = \frac{p(\theta_0 | H_1)}{p(\theta_0 | \mathcal{D}, H_1)}$$

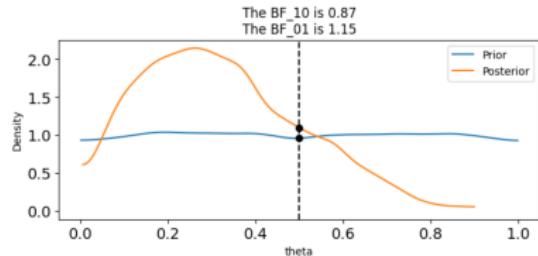
where:

- $p(\theta_0 | H_1)$ is *prior* density θ under alternative hypothesis H_1 , evaluated at θ_0
- $p(\theta_0 | \mathcal{D}, H_1)$ is *posterior* density θ under H_1 evaluated at θ_0
- **Intuition:** show how data changes belief about θ_0
 - If posterior density at θ_0 is much smaller than prior density, strong evidence against null hypothesis H_0
- **Cons:**
 - It's a point statistics (not considering the entire posterior)

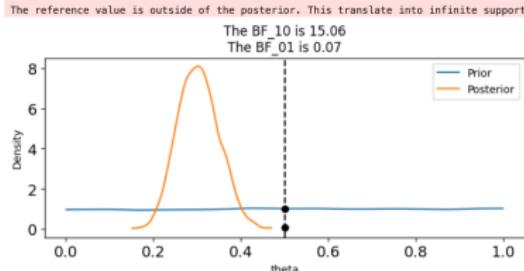
Bayes Factor (BF)	Interpretation
1 - 3	Not enough evidence
3 - 10	Substantial evidence
10 - 100	Strong evidence
> 100	Decisive evidence

Savage-Dickey Density Ratio: Example

```
[29]: az.plot_bf(idata1, var_name="theta", prior=np.random.uniform(0, 1, 10000), ref_val=0.5);
```



```
[30]: az.plot_bf(idata2, var_name="theta", prior=np.random.uniform(0, 1, 10000), ref_val=0.5);
```



- H_0 : “coin is fair”
 - The prior is 0.87
 - The posterior is 1.15
 - $BF_{10} = 0.76 \rightarrow$ **no evidence** to reject the null hypothesis
- In the other case
 - $BF_{10} = 15.06 \rightarrow$ **strong evidence** to reject the null hypothesis

Region of Practical Equivalence

- **Region of Practical Equivalence** (ROPE) = interval where values are “equivalent”
- **Example**
 - H_0 : “coin is fair” impractical if $\theta = 0.5$
 - ROPE: $\theta \in [0.45, 0.55]$ equivalent to 0.5
- **Hypothesis testing with ROPE and HPI**
 - Compare ROPE with HPI (Highest-Posterior Interval)
 - HPI within ROPE \rightarrow no effect, reject H_1
 - HPI outside ROPE \rightarrow effect present, reject H_0
 - HPI overlaps ROPE \rightarrow inconclusive
- **Decide ROPE before analysis based on domain knowledge**
 - Picking it after analysis is like choosing p-value threshold after seeing p-value

Loss Function: Motivation

- For many problems:
 - **Make decisions based on inference**
 - **Decision cost is asymmetric**
 - Cost of a bad decision > or < benefit of a good decision
 - Vaccines may cause overreaction, but benefits outweigh risks
- Measure for **best decision**:
 - Benefits of a correct decision
 - Cost of a mistake
 - Trade-off between benefits and costs using a loss function
 - Use loss function for decisions

Loss Function

- Aka “cost function”
 - The inverse is known as “objective”, “fitness”, “utility function”
- **Loss function** quantifies “*how bad is an estimation mistake?*”
 - Larger loss indicates worse estimation
 - Loss is difference between:
 - The true value θ ; and
 - The estimated value $\hat{\theta}$

Loss	Expression	Point estimate
Quadratic loss	$(\theta - \hat{\theta})^2$	Mean of posterior
Absolute loss	$ \theta - \hat{\theta} $	Median of posterior
1-0 loss	$I(\theta \neq \hat{\theta})$	Mode of posterior

- Algorithm to make decisions in Bayesian statistics using loss function
 - Goal: pick a single value $\hat{\theta}$
 - True value θ is unknown
 - Estimate θ using posterior distribution
 - Find $\hat{\theta}$ minimizing expected loss function

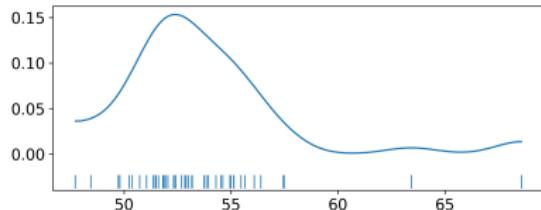
Tutorial

- Bayesian Coin

- Posterior-Based Decisions
 - *Chemical Shift: Example*
 - Posterior Predictive Checks
 - Groups Comparison

Chemical Shift

- **Nuclear magnetic resonance (NMR)**
 - Used to study molecules of living things
 - Measures observable quantities related to unobservable molecular properties (e.g., chemical shift)
- **Chemical shift** reveals local magnetic environment of atomic nuclei
 - Identifies molecular structure and functional groups
- **Example of chemical shift**
 - Data looks Gaussian with a couple of outliers



Use of Gaussians in Statistics

- Aka “normal” distribution
- **Gaussians** are:
 - easy to work with
 - abundant in nature
- **Pros:**
 - Average of large sample size tends to be Gaussian (Central Limit Theorem)
 - Many phenomena approximated using Gaussians (since they are average of effects)
 - Conjugate prior of Gaussian is Gaussian
- **Cons:**
 - Not robust to outliers
 - Important to relax assumption of Gaussianity

Chemical Shift: Example

- Assume Gaussian approximates data from example chemical shift
- Likelihood from normal distribution:

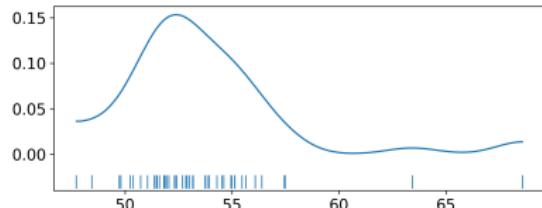
$$Y \sim N(\mu, \sigma)$$

- Weakly informative priors for mean and sigma of Y
 - Mean from uniform distribution

$$\mu \sim U(l = 40, h = 70)$$

- Larger than data range, e.g., [40, 70]
- Std dev from half-normal distribution

$$\sigma \sim HalfNormal(0, \sigma_\sigma = 10)$$



Chemical Shift: PyMC

```
[87]: with pm.Model() as model_g:  
    # The mean is Uniform in [40, 70] (which is larger than the data).  
    mu = pm.Uniform("mu", lower=40, upper=70)  
    # The std dev is half normal with a large value (which is a large value based on the data).  
    sigma = pm.HalfNormal("sigma", sigma=10)  
    # The model is  $N(\mu, \sigma)$ .  
    y = pm.Normal("y", mu=mu, sigma=sigma, observed=data)  
    # Sample.  
    idata_g = pm.sample(1000)
```

Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Multiprocess sampling (4 chains in 4 jobs)
NUTS: [mu, sigma]

Sampling 4 chains, 0 divergences  100% 0:00:00 / 0:00:00

Sampling 4 chains for 1_000 tune and 1_000 draw iterations (4_000 + 4_000 draws total) took 1 seconds.

- The PyMC code is **one-to-one with the model**:

$$\begin{cases} \mu \sim U(l = 40, h = 70) \\ \sigma \sim HalfNormal(0, \sigma_\sigma = 10) \\ Y \sim N(\mu, \sigma) \end{cases}$$

- Get 1000 samples from the posterior

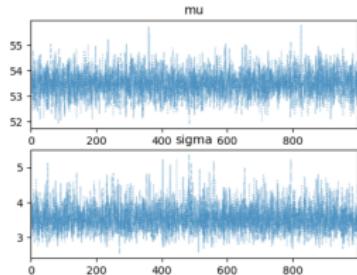
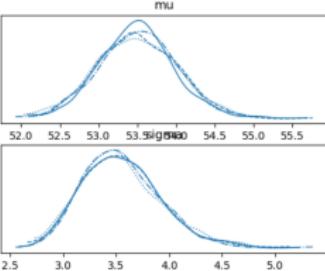
Chemical Shift: PyMC

- Compute 4 traces for 2 variables μ , σ

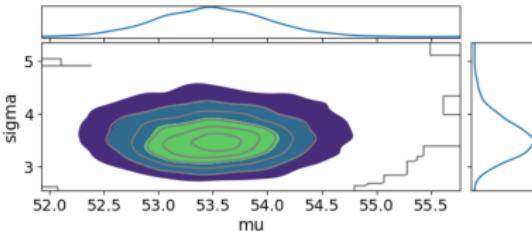
- Results are well-formed
- $\mu \in [52.55, 54.45]$
- $\sigma \in [2.86, 4.23]$

- Is the model good?

```
[92]: # There are 4 traces for 2 variables.  
az.plot_trace(idata_g);
```



```
[93]: # The posterior distribution of the params is bi-dimensional, since it has mu and sigma.  
az.plot_pair(idata_g, kind='kde', marginalize=True);
```



```
[94]: # Report a summary of the inference.  
az.summary(idata_g, kind='stats').round(2)
```

```
[94]:
```

	mean	sd	hdi_3%	hdi_97%
mu	53.50	0.51	52.55	54.46
sigma	3.55	0.38	2.86	4.23

- Posterior-Based Decisions
 - Chemical Shift: Example
 - ***Posterior Predictive Checks***
 - Groups Comparison

Samples from Posterior Distribution

- Given **posterior distribution** $\Pr(\theta|y)$, generate predictions \tilde{y} based on data y and estimated parameters $\hat{\theta}$:

$$\Pr(\tilde{y}|y) = \int_{\hat{\theta}} \Pr(\tilde{y}|\theta) \Pr(\theta|y) d\theta = \int \text{model} \times \text{posterior}$$

- This is called the “*posterior predictive distribution*” as it predicts *future data* using the *posterior distribution*
- Conceptually**
 - Sample a value of θ from the posterior $\Pr(\theta|y)$
 - Feed the value of θ to the likelihood $\Pr(\tilde{y}|\theta)$
 - Obtain \tilde{y}
- This process has **two sources of uncertainty**:
 - Parameter uncertainty
 - Captured by the posterior $\Pr(\theta|y)$
 - Sampling uncertainty
 - Captured by the likelihood $\Pr(\tilde{y}|\theta)$

Posterior Predictive Check (PPC)

- **PPC approach:**
 - Generate predictions \tilde{y} with observed data y from posterior distribution
 - Check consistency between predictions and observed data
- **Intuition:** can the model reproduce observed data?
 - Always check!
- **Differences** arise from:
 - Mistakes
 - Model limitations
 - E.g., works for average behavior but fails for rare values
 - Data limitations

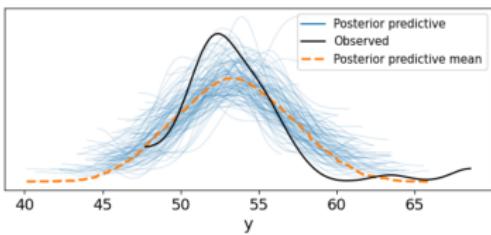
Bayesian Workflow Using PPC

1. Given a process, **true distribution** is unknown / unknowable
2. **Sample the process**
 - Get finite sample y
 - E.g., experiment, survey, simulation
3. **Inference**
 - Build probabilistic model using prior $\Pr(\theta)$ and likelihood $\Pr(y|\theta)$ to get posterior $\Pr(\theta|y)$
 - Posterior is distribution of model parameters θ given data
4. **Predictive distribution**
 - Compute predictions from posterior (posterior predictive distribution)
 - Posterior predictive is distribution of predicted samples averaged over posterior
5. **Validation**
 - Validate model by comparing original vs predicted samples

Chemical Shift Example: PPC

```
[95]: # Compute 100 posterior predictive samples.  
y_pred_g = pm.sample_posterior_predictive(idata_g, model=model_g)  
  
Sampling: [y]  
  
Sampling ... ━━━━━━━━━━━━━━━━ 100% 0:00:00 / 0:00:00
```

```
[96]: # Black: KDE of the data (observed)  
# Blue: KDEs of the posterior predictive samples  
# Orange: KDE of the posterior predictive mean  
az.plot_ppc(y_pred_g, mean=True, num_pp_samples=100);
```



• Apply the Bayesian workflow

- Sample posterior
- Apply model
- Get predictive posterior distribution (dashed orange)
- Compare to data (black)

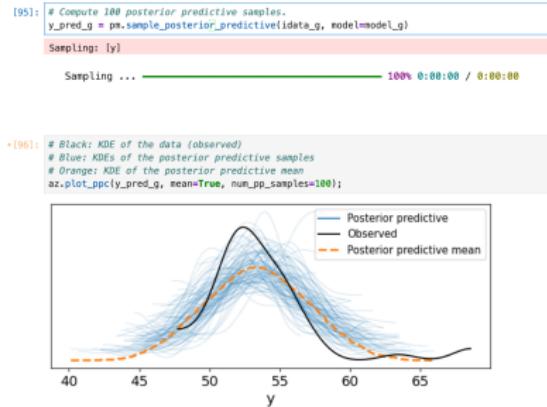
• Is the PPC model good?

- No!
- Posterior mean is right of data
- Posterior std dev is larger
- You can't generate the data from the model!

Chemical Shift: Model Critique

- **Problem**

- Two data points on distribution tails
- Normal distribution:
 - is “surprised” by these points
 - “reacts” by adjusting mean towards them and increasing the standard deviation

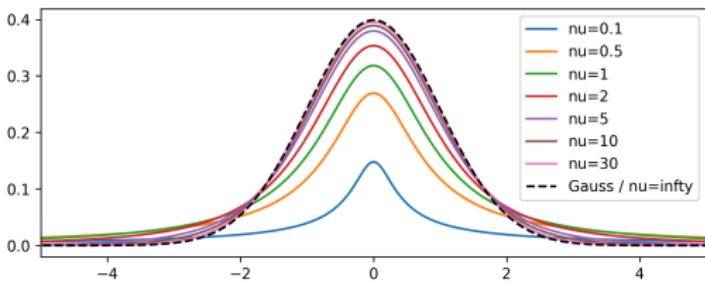


- **Possible solutions**

1. Declare and discard outliers
 - E.g., equipment malfunction (requires evidence)
2. Change the model
3. **Bayesian philosophy**
 - Encode assumptions into the model (e.g., priors, likelihoods)
 - Avoid ad-hoc heuristics (e.g., outlier removal rules)

Student's t-distribution: Recap

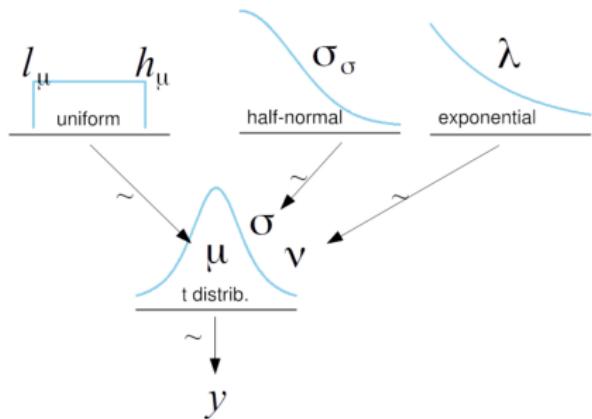
- Student's t-distribution has 3 params:
 1. Mean μ
 - Doesn't always exist
 2. Scale σ
 - Similar to std dev, but doesn't always exist
 3. Degrees of freedom $\nu \in [0, \infty]$
 - "Normality parameter": controls distribution normality
 - $\nu = 1$: heavy tails, no mean (Cauchy)
 - $\nu \rightarrow \infty$: Gaussian
- Student's t has **heavy tails** (high kurtosis)
 - Values more likely far from mean compared to Normal



Chemical Shift: Use Student's t-dist (1/3)

- Use Student's t-distribution instead of Normal in model for chemical shift

$$\begin{cases} \mu \sim U(l, h) \\ \sigma \sim HalfNormal(0, \sigma) \\ \nu \sim Exp(\lambda) \\ y \sim StudentT(\mu, \sigma, \nu) \end{cases}$$



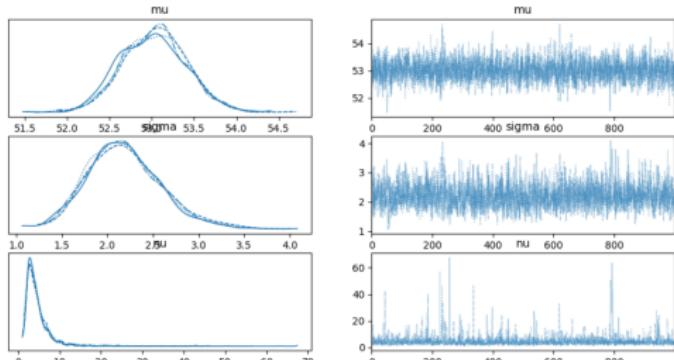
Chemical Shift: Use Student's t-dist (2/3)

- Decrease ν (**less Gaussian**) instead of increasing mean and standard deviation
 - μ similar to Gaussian estimate
 - σ smaller
 - $\nu \approx 5$ (not very Gaussian)
- Estimation more robust**
 - Outliers have less effect

```
[102]: # Use a Student-T model.  
with pm.Model() as model_t:  
    mu = pm.Uniform("mu", 40, 75)  
    sigma = pm.HalfNormal("sigma", sigma=10)  
    # A student with nu = 30 is close to a Gaussian.  
    nu = pm.Exponential("nu", 1/30)  
    #  
    y = pm.StudentT("y", mu=mu, sigma=sigma, nu=nu, observed=data)  
idata_t = pm.sample(1_000)
```

Auto-assigning NUTS sampler... ***

```
az.plot_trace(idata_t);***
```



```
[104]: az.summary(idata_t, kind="stats").round(2)
```

	mean	sd	hdi_3%	hdi_97%
mu	53.03	0.38	52.35	53.76
sigma	2.19	0.40	1.45	2.95
nu	4.65	3.92	1.20	9.20

Chemical Shift: Use Student's t-dist (3/3)

```
[105]: # Compute 100 posterior predictive samples.
```

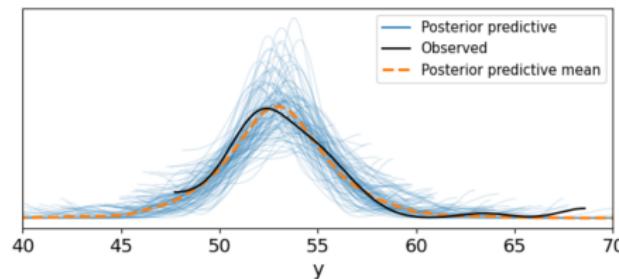
```
y_ppc_t = pm.sample_posterior_predictive(idata_t, model_t);
```

```
Sampling: [y]
```

```
Sampling ... 100% 0:00:00 / 0:00:00
```

```
[106]: ax = az.plot_ppc(y_ppc_t, num_pp_samples=100, mean=True)
```

```
ax.set_xlim(40, 70);
```



- PPC fits better than Normal model
- Plot is “hairy” because KDE is estimated only in data interval and 0 outside

Tutorial

- Robust modeling

- Posterior-Based Decisions
 - Chemical Shift: Example
 - Posterior Predictive Checks
 - ***Groups Comparison***

Group Comparison

- **Group comparison** tests for statistically significant results between *treatment and control group*
- E.g.,
 - *How well do patients respond to a new drug vs a placebo?*
 - *Is there a reduction in car accidents after new traffic regulation?*
 - *Does college student performance improve without cellphones at school?*
- **Effect size** quantifies the difference between two groups
 - From “does it work?” (hypothesis testing) to “how well does it work?” (estimate effect size)

Bogus Control Groups

- **Always ask for the baseline** used for comparison, when something is claimed to be harder/better/faster/stronger
 - E.g., sell sugary yogurts to boost the immune system by comparing it to milk
 - Better control group: less sugary yogurt
- **Placebo** is a psychological phenomenon where a patient experiences improvements after receiving an inactive treatment
 - Using a placebo is better than “no treatment”
 - Shows difficulty in accounting for all factors in an experiment

Group Comparison Bayesian-Style

- **Frequentist approach**
 - Compare p-value of difference of means in each group
- **Bayesian approach**
 - Compare posterior distribution of means between groups using:
 - Plot of posterior
 - Cohen's d
 - Probability of superiority

Sample Size Effect

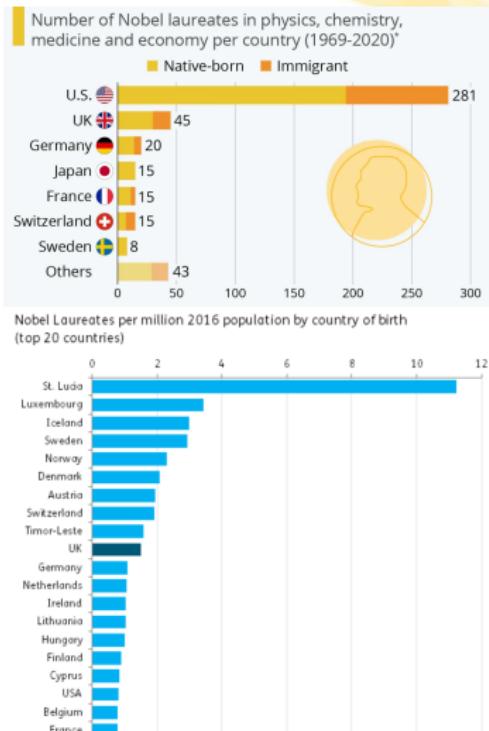
- **Sample size effect** is the impact of the number of observations on statistical results (e.g., p-values, confidence intervals)

- **Small sample effect**

- Large mean difference might not be statistically significant (low p-value)
- Estimates (e.g., means, proportions, correlations) fluctuate widely due to high sampling variability
- Outliers have a disproportionate influence
- Inference is unstable: results may not replicate

- **Large sample:**

- Tiny mean difference can be highly significant (small p-value) but meaningless
- E.g., Cohen's d, probability of superiority

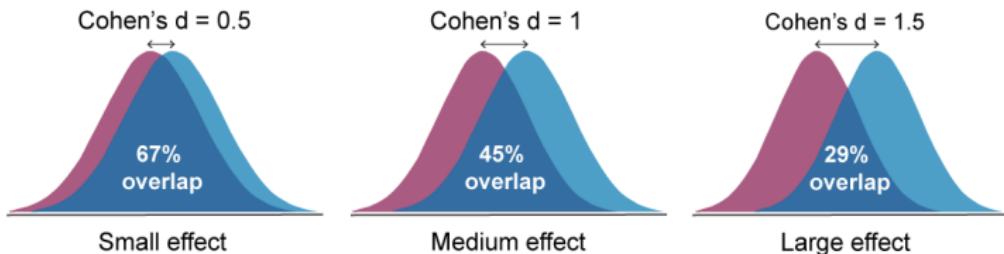


Cohen's d

- Cohen's d is the difference of means relative to pooled standard deviation

$$\frac{\mu_2 - \mu_1}{\sqrt{(\sigma_1^2 + \sigma_2^2)/2}}$$

- Normalizes effect by variability for pooled std dev
- Variability of each group normalizes mean difference
- Similar to a Z-score, number of std dev values differ



- Bayesian approach

- Compute posterior distribution of means and std → formula
- Compute distribution of Cohen's d → summary statistics