Figure 9 Replication Code

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This demo intends to demonstrate overlapping scenarios between the Bayesian and frequentist settings assessing the assurance using credible interval based conditions. This demo focuses on the case when p_1 and p_2 are unknown.

```
library(bayesassurance)
```

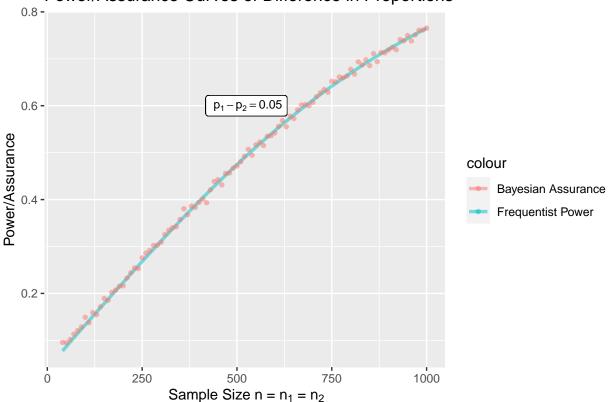
```
propdiffCI_classic <- function(n, p1, p2, alpha_1, beta_1, alpha_2, beta_2, sig_level){
    set.seed(1)
    if(is.null(p1) == TRUE & is.null(p2) == TRUE){
        p1 <- rbeta(n=1, alpha_1, beta_1)
        p2 <- rbeta(n=1, alpha_2, beta_2)
    }else if(is.null(p1) == TRUE & is.null(p2) == FALSE){
        p1 <- rbeta(n=1, alpha_1, beta_1)
    }else if(is.null(p1) == FALSE & is.null(p2) == TRUE){
        p2 <- rbeta(n=1, alpha_2, beta_2)
    }
    p <- p1 - p2

    power <- pnorm(sqrt(n / ((p1*(1-p1)+p2*(1-p2)) / (p)^2)) - qnorm(1-sig_level/2))
    return(power)
}</pre>
```

The following set of examples assign different values for p1 and p2 that adhere to different sets of critical differences while maintaining the same 0.5 throughout.

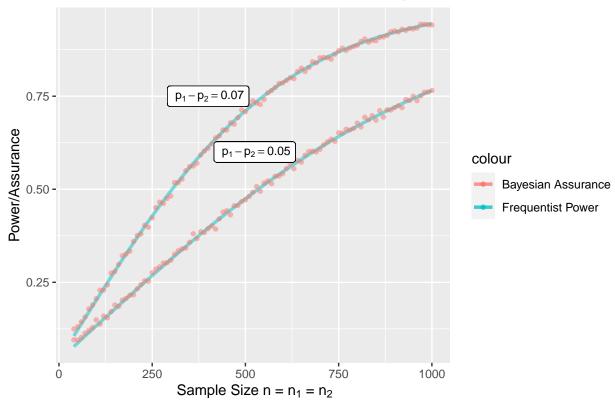
Case 1: p1 - p2 = 0.1

Power/Assurance Curves of Difference in Proportions



Case 2: p1 - p2 = 0.07

Power/Assurance Curves of Difference in Proportions



Case 3: p1 - p2 = 0.05

Power/Assurance Curves of Difference in Proportions

