lecture 2.2 example 1

preliminaries

```
# clear environment
rm(list = ls())
# set random seed
set.seed(123)
```

define params

```
# define true parameters of the data generating process
pTrue = 0.7

# define data set size
nData = 100

# prior params
aPrior = 1
bPrior = 1
```

simulate dataset

```
data = rbinom(n = nData, size = 1, prob = pTrue)
nWater = sum(data)
```

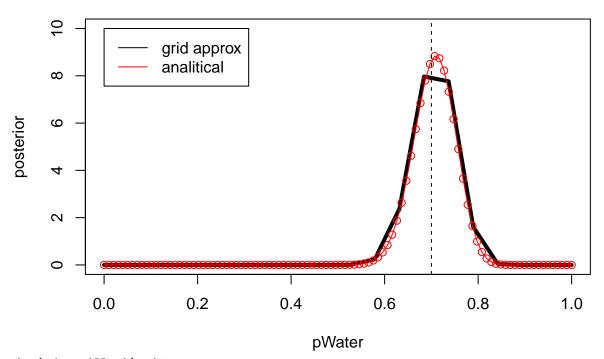
build simulation functions

```
prior <- function(nGridPoints) {
   pGrid = seq(from = 0, to = 1,length.out = nGridPoints)
   prior = dbeta(x = pGrid, shape1 = aPrior, shape2 = bPrior)
}

compute_posterior <- function(nGridPoints) {
   #build grid
   pGrid = seq(from = 0, to = 1,length.out = nGridPoints)
   gridSize = 1 / nGridPoints
   # compute likelihood
   likelihood = dbinom(x = nWater, size = nData, prob = pGrid)
   # compute posterior
   postGrid = likelihood * prior(nGridPoints)
   postGrid = postGrid / ( sum(postGrid) * gridSize )
   #
   return(postGrid)
}</pre>
```

simulations: 20 grid points

20 grid points



simulations: 100 grid points

##

100 grid points

