**Can you explain how you know which c(1,2,1) vs (2,2,2) etc. to use again? Are we just trying out different ones to see which works best?**

* Any model that you should pick, should/must have d =2 .
* Iterate starting from p = 1, q =1 (given that both the ACF and PACF of the differenced data did not cut of nicely).
* Increment either p or q by 1 until we have a p-value > 0.05 (this would mean that we would not reject the null i.e. residuals are uncorrelated)