

Meeting 05/02/2020

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Summary

- Check how NAs are handled in pre-processing of BCC TCGA data.
- Mention `mclust` and `tabulation` to Eckart; look at Bayesian Hierarchical Clustering (and `BHC` the Bioconductor package). Also, log transform count data.
- Latex derivation of collapsed gibbs sampler for mixture allocation for the general model and the gaussian case.
- Test Gaussian mixtures in MDI on a simple simulation of standard normal data with a diagonal covariance matrix.
- Look at Bayesian hierarchical clustering and associated `BHC` package (on Bioconductor).