# Meeting 22/01/2020

Stephen Coleman

24/01/2020

**Summary**

- Move to MATLAB code entirely (suspicious of high Phi values and of command line code generally);
- Transfer Gaussian mixture component to Paul's MATLAB;
- Test this - check derivations in code and test onf MDItestdata from original MDI paper;
- Apply to TGCA data as manipulated for the BCC paper; and
- Apply to Pan-Cancer dataset (to receive from Paul).