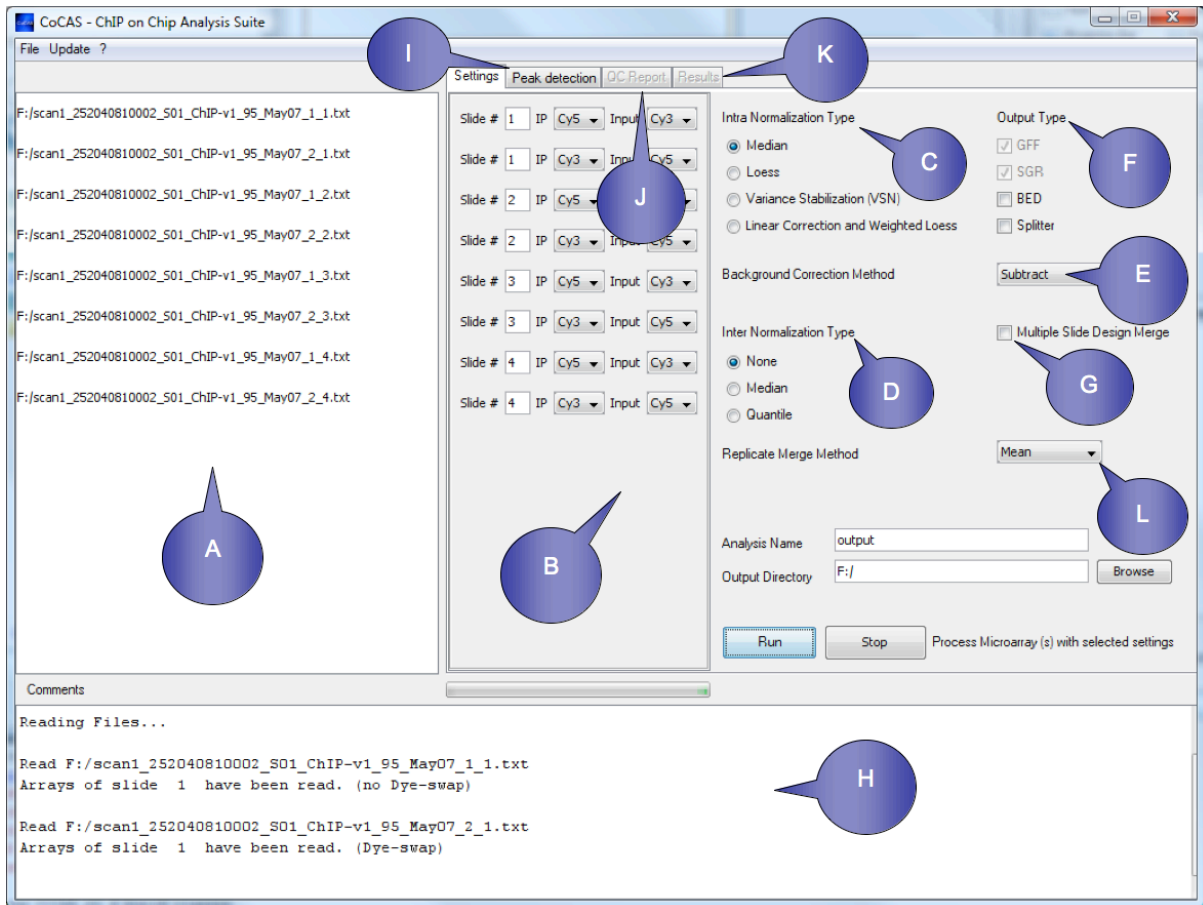


Sample Screenshot of CoCAS's main interface



Supplementary Fig. S3: Example screenshot of CoCAS's main interface. (A) Loaded microarray file list, (B) slide numbering and dye selection, (C) and (D) intra- and inter-normalisation parameterisation, respectively. Other parameters include background subtraction (E), output file format (F), and use of multiple array designs (G). Progress can be viewed in an R output console (H). Peak detection parameters and standalone peak detection are available in a separate tab (I). QC report and peak detection results are displayed in tabs (J) and (K) respectively. (L) Replicate merging method.