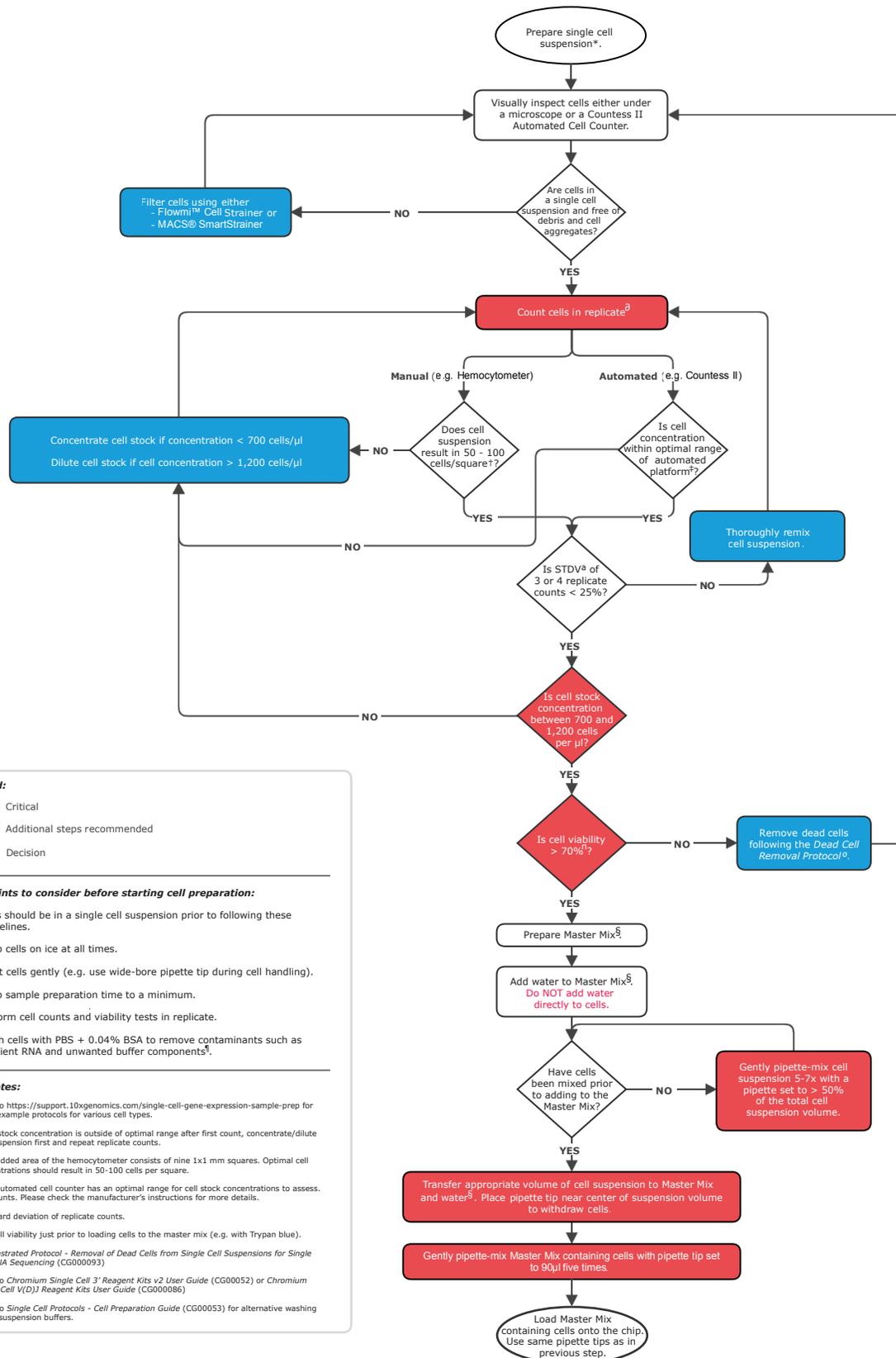


Chromium™ Single Cell Applications

Guidelines for Optimal Sample Preparation



Legend:

- Critical
- Additional steps recommended
- ◇ Decision

Key points to consider before starting cell preparation:

1. Cells should be in a single cell suspension prior to following these guidelines.
2. Keep cells on ice at all times.
3. Treat cells gently (e.g. use wide-bore pipette tip during cell handling).
4. Keep sample preparation time to a minimum.
5. Perform cell counts and viability tests in replicate.
6. Wash cells with PBS + 0.04% BSA to remove contaminants such as ambient RNA and unwanted buffer components[¶].

Footnotes:

- * Refer to <https://support.10xgenomics.com/single-cell-gene-expression/sample-prep-for-more-example-protocols-for-various-cell-types>.
- ^ If cell stock concentration is outside of optimal range after first count, concentrate/dilute cell suspension first and repeat replicate counts.
- † The gridded area of the hemocytometer consists of nine 1x1 mm squares. Optimal cell concentrations should result in 50-100 cells per square.
- ‡ Each automated cell counter has an optimal range for cell stock concentrations to assess cell counts. Please check the manufacturer's instructions for more details.
- § Standard deviation of replicate counts.
- ¶ Test cell viability just prior to loading cells to the master mix (e.g. with Trypan blue).
- ‡ Demonstrated Protocol - Removal of Dead Cells from Single Cell Suspensions for Single Cell RNA Sequencing (CG000093)
- § Refer to Chromium Single Cell 3™ Reagent Kits v2 User Guide (CG00052) or Chromium Single Cell V(D)J Reagent Kits User Guide (CG00086)
- ¶ Refer to Single Cell Protocols - Cell Preparation Guide (CG00053) for alternative washing and resuspension buffers.