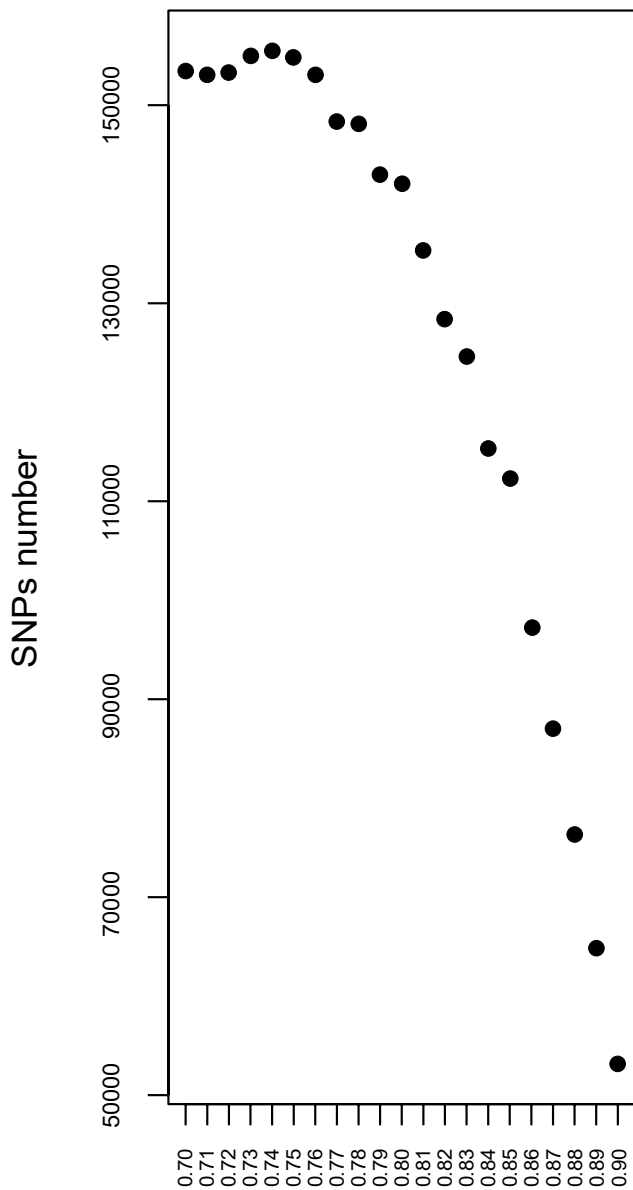


max_barcode_mismatch=1, min_samples_locus=4

max_SNPs_locus=0.6, max_Indels_locus=8



| CT | SNP number |
|------|------------|
| 0.70 | 153455 |
| 0.71 | 153025 |
| 0.72 | 153267 |
| 0.73 | 154953 |
| 0.74 | 155298 |
| 0.75 | 154955 |
| 0.76 | 153068 |
| 0.77 | 148363 |
| 0.78 | 148084 |
| 0.79 | 142908 |
| 0.80 | 142062 |
| 0.81 | 135355 |
| 0.82 | 128351 |
| 0.83 | 124538 |
| 0.84 | 114904 |
| 0.85 | 113339 |
| 0.86 | 97177 |
| 0.87 | 86941 |
| 0.88 | 76282 |
| 0.89 | 64823 |
| 0.90 | 53188 |

Clustering thresholds