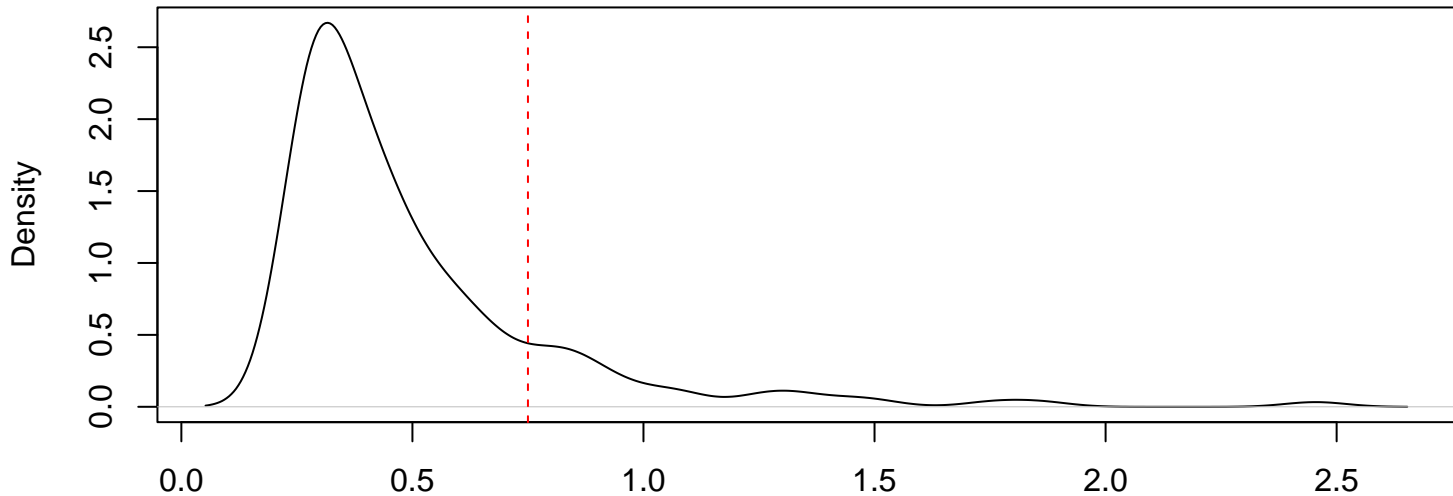
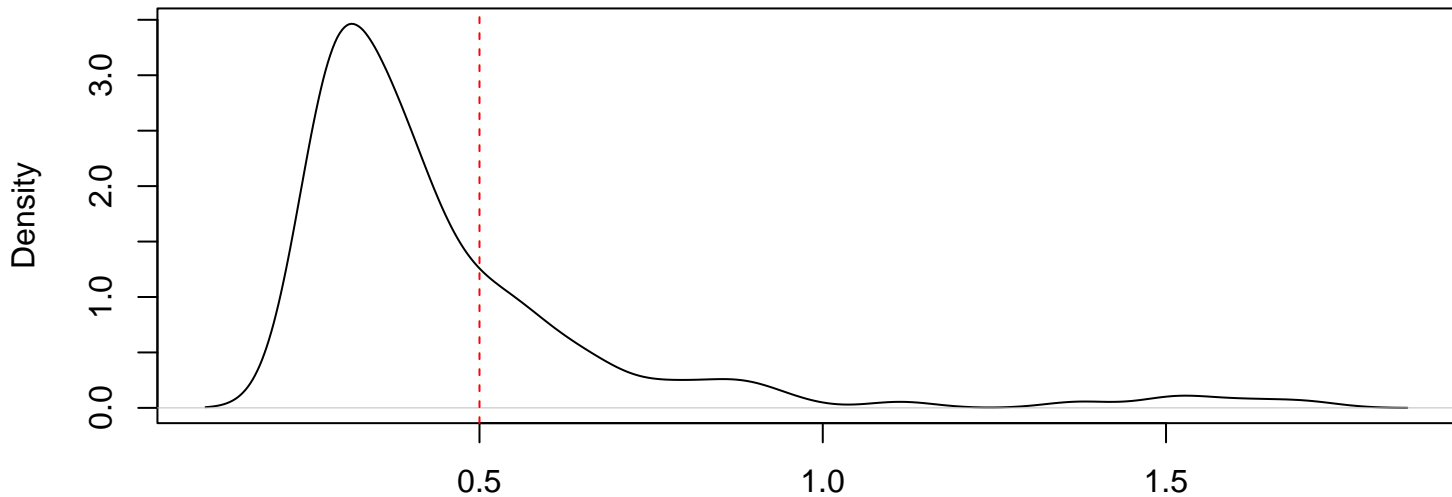


**density.default(x = Myeloids.markers.qc\$avg\_log2FC)**



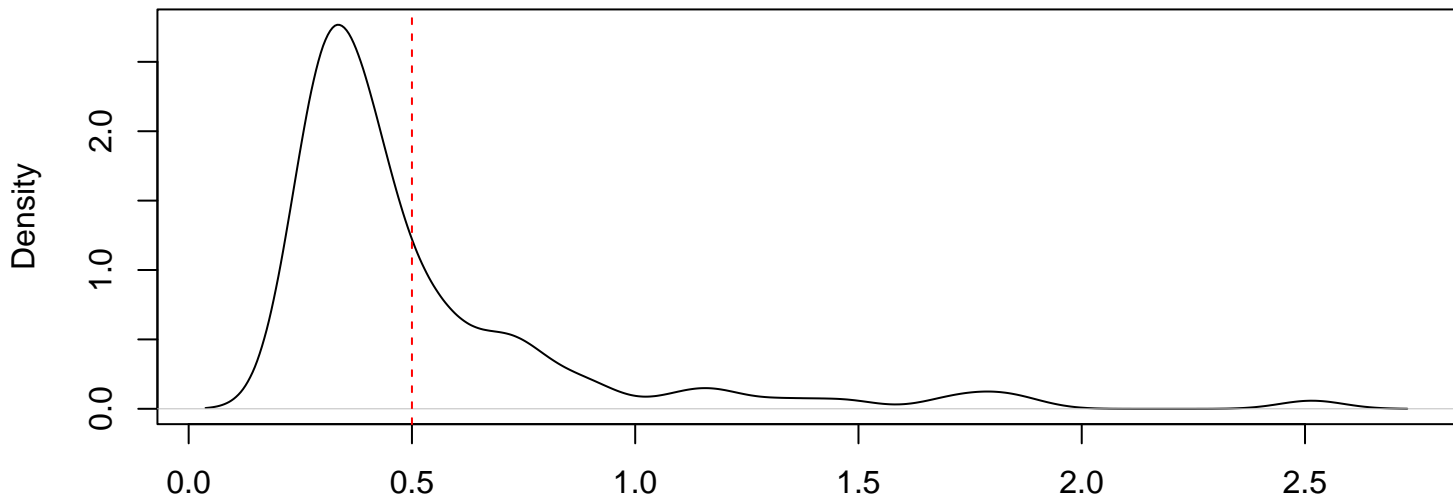
N = 185 Bandwidth = 0.06615

**density.default(x = dNKs.markers.qc\$avg\_log2FC)**



N = 148   Bandwidth = 0.04984

**density.default(x = tcells.markers.qc\$avg\_log2FC)**



N = 96 Bandwidth = 0.07133