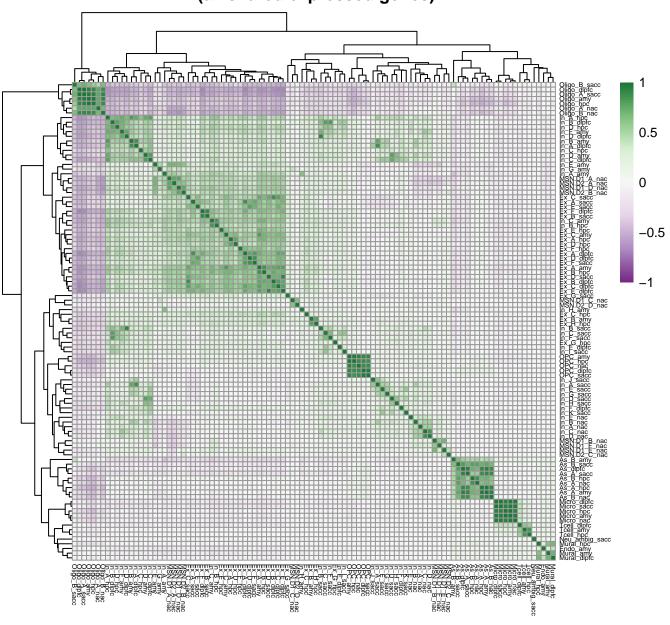
Correlation of cluster-specific t's from all regions (all shared expressed genes)



Correlation of cluster–specific t's from all regions (top 100 cluster genes space)

