Clustering evaluation: SNF vs baselines
(log-rank p annotated below method name)

Legend:
- Stability target (0.8)
- ARI vs PAM50
- NMI vs PAM50
- Stability ARI

Methods and p-values:
- SNF (proposed): p=0.144
- Single-omics RNA: p=0.022
- Single-omics CNV: p=0.697
- Single-omics Methylation: p=0.728
- Early Integration (concat PCA): p=0.111