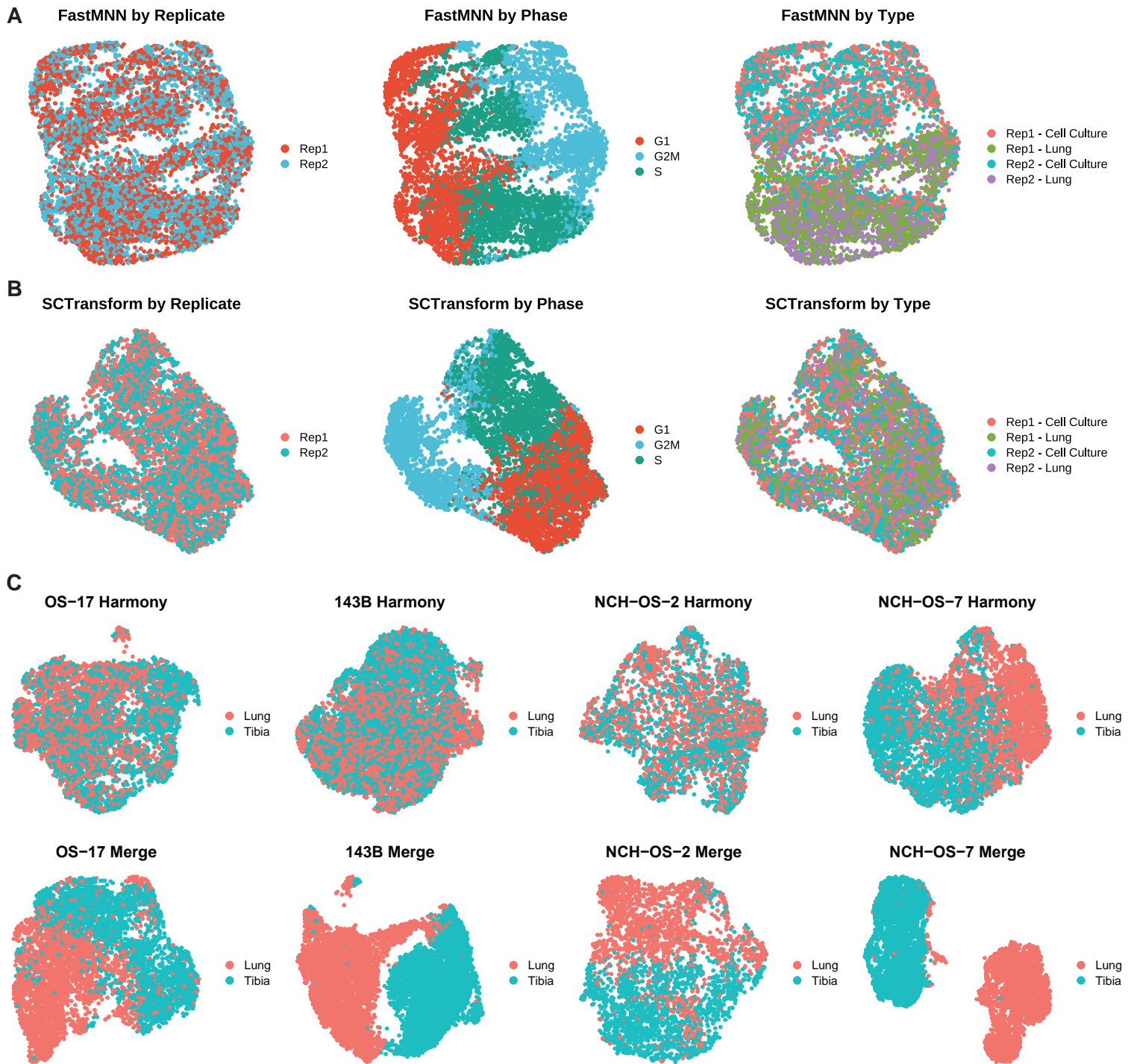Figure S8. Integrative analysis for within-model comparison of primary and metastatic lesions. A, B) Application of FastMNN or SCTransform between replicate samples demonstrated complete overlay, however cells separated largely by state of cell cycle (OS-17). C) Application of Harmony or merging to Seurat objects. Application of Harmony demonstrated nearly complete overlay. Merged objects displayed some overlay while maintaining a slight distinction between samples. Cells did not cluster based on state of cell cycle (data not shown).