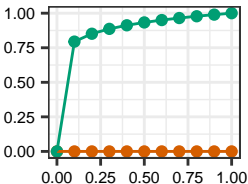
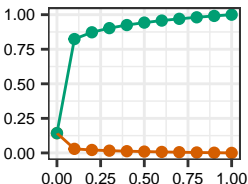


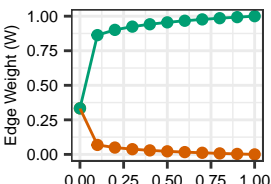
0% transcriptome neighbors



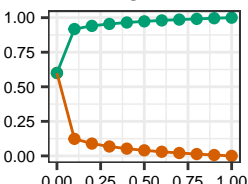
25% transcriptome neighbors



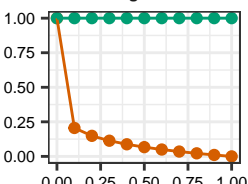
50% transcriptome neighbors



75% transcriptome neighbors



100% transcriptome neighbors



.. value

—●— between barcodes

—●— within barcode