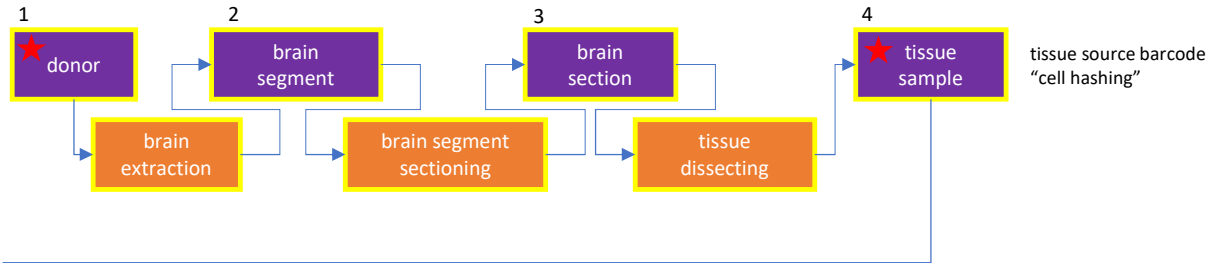
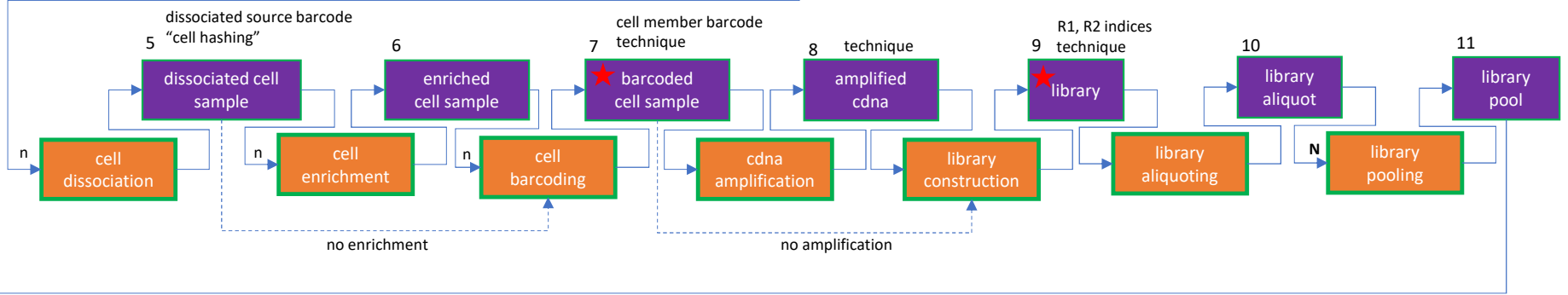


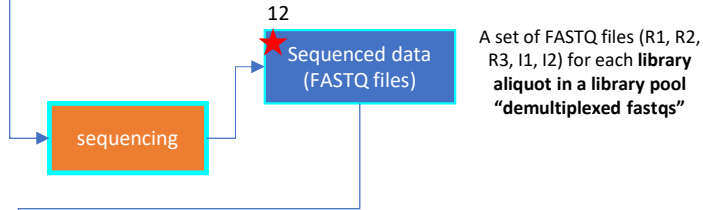
Anatomical level
Specimen Portal



Cellular level
Sequencing Portal



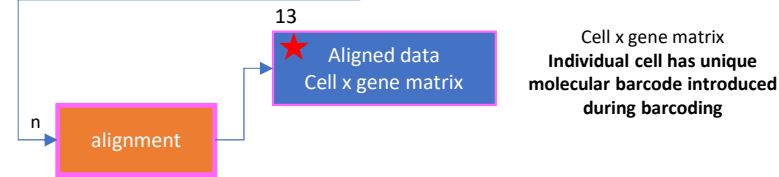
Seq Core



A set of FASTQ files (R1, R2, R3, I1, I2) for each library aliquot in a library pool "demultiplexed fastqs"

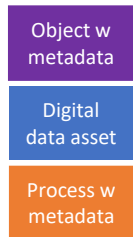
N : Multiple inputs possible WITHIN AND ACROSS species
n : Multiple inputs possible WITHIN the SAME species

NeMO



Cell x gene matrix
Individual cell has unique molecular barcode introduced during barcoding

Sequencing results from another library aliquot can be merged if it originates from the same barcoded cell sample



★ Present in all methods