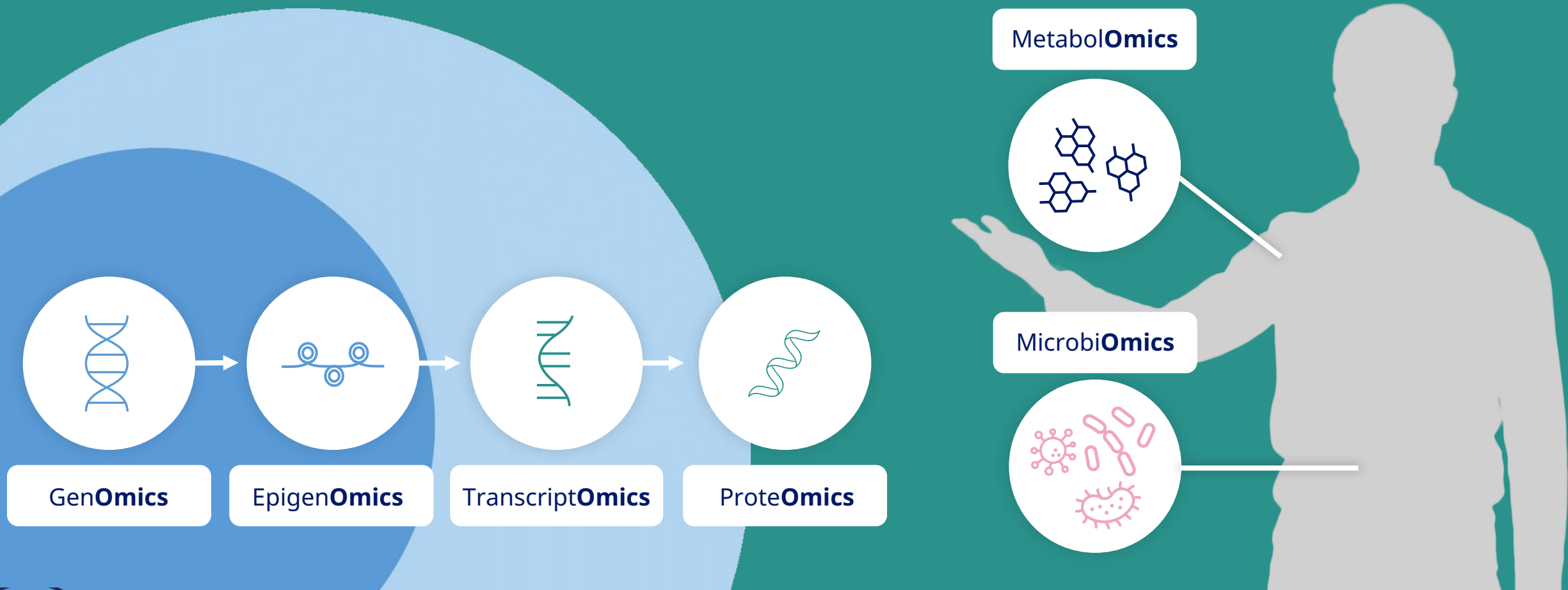




Alzheimer's disease in the omics era

Alzheimer's disease in the omics era

Using 'Omics' to identify, describe and quantify biomolecules and molecular processes



'Bulk' versus single-cell transcriptomics

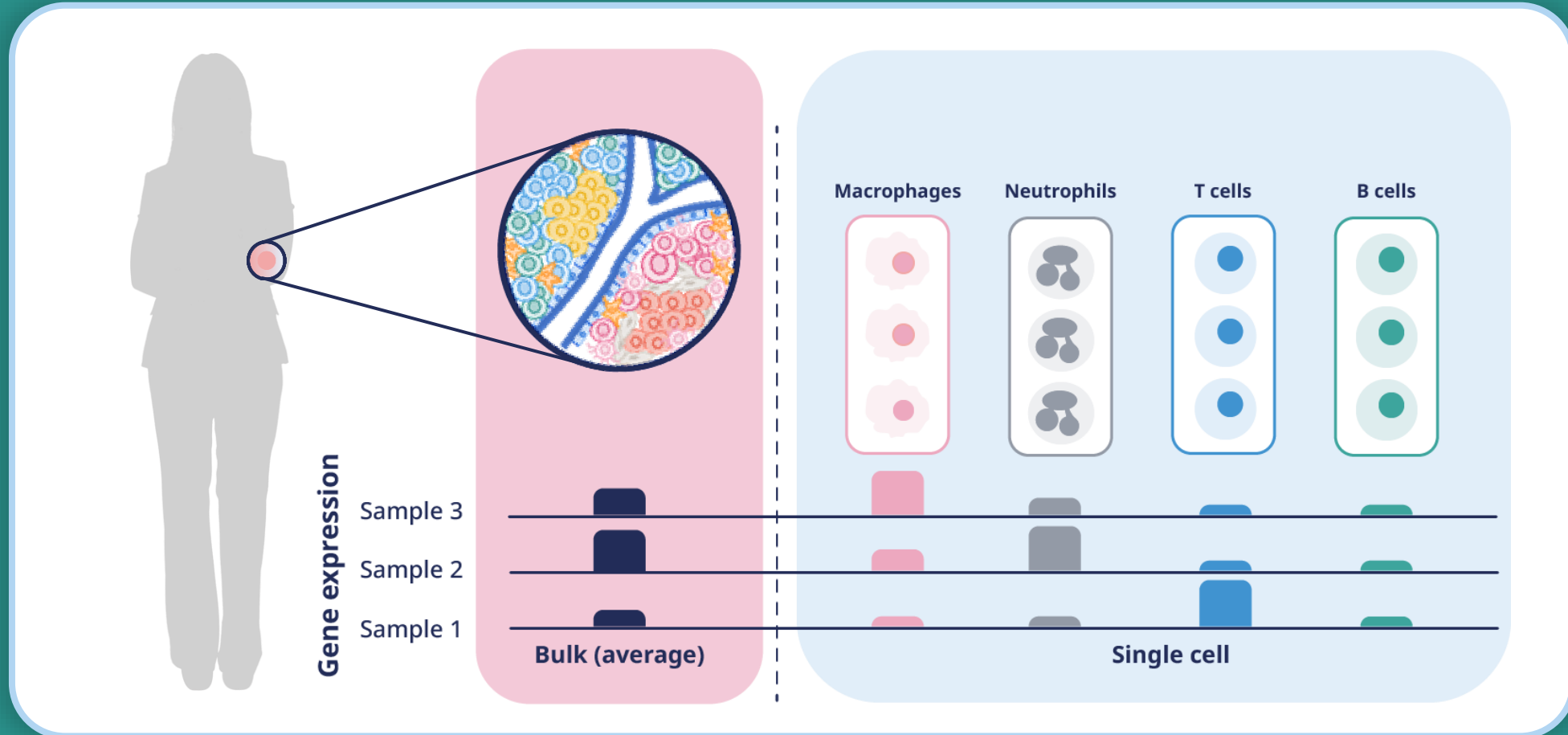
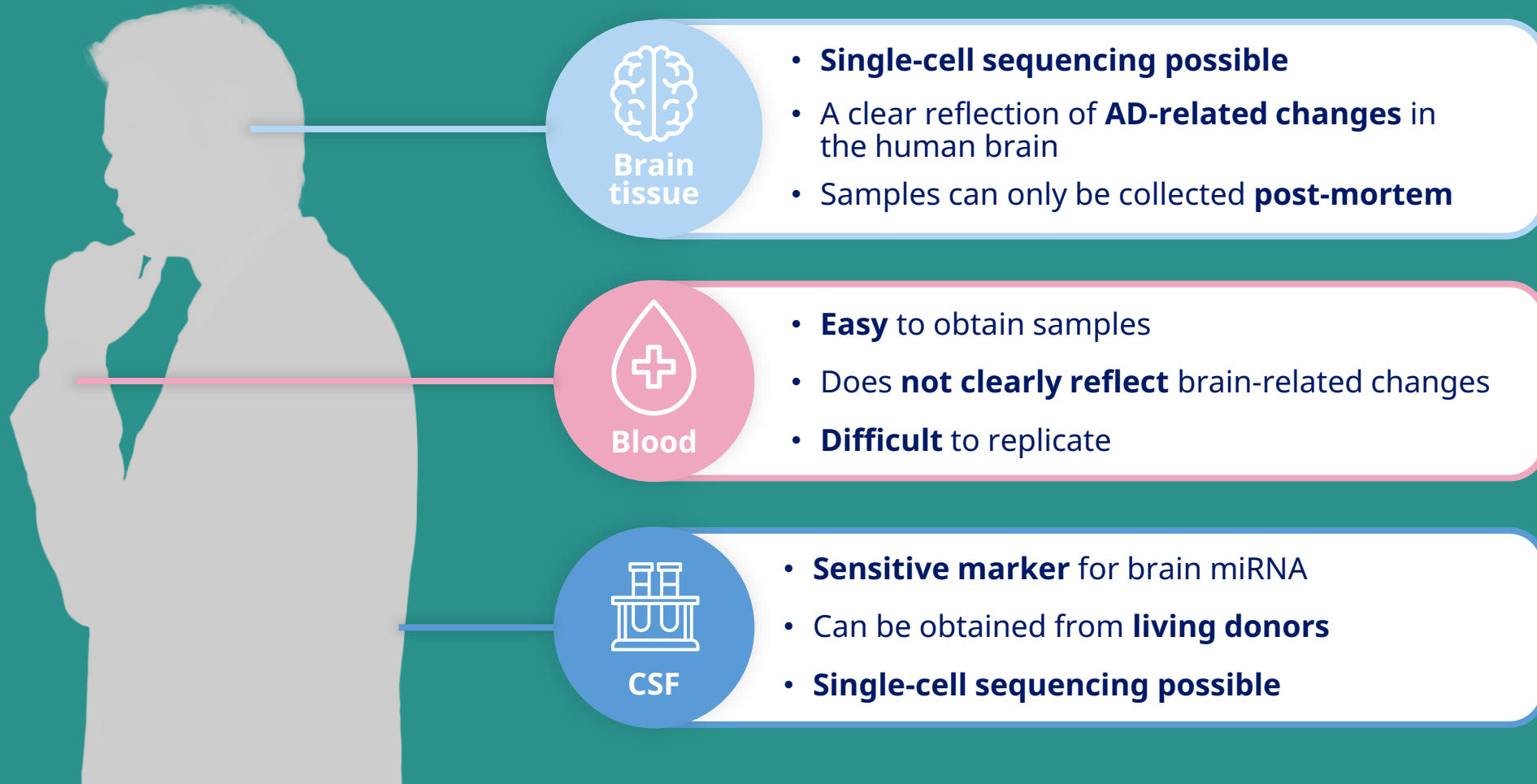


Figure adapted from Fan J et al. *Exp Mol Med* 2020;52:1452–65.



Transcriptomics sample types in Alzheimer's disease



'Omics' approaches have huge potential in drug development and disease understanding

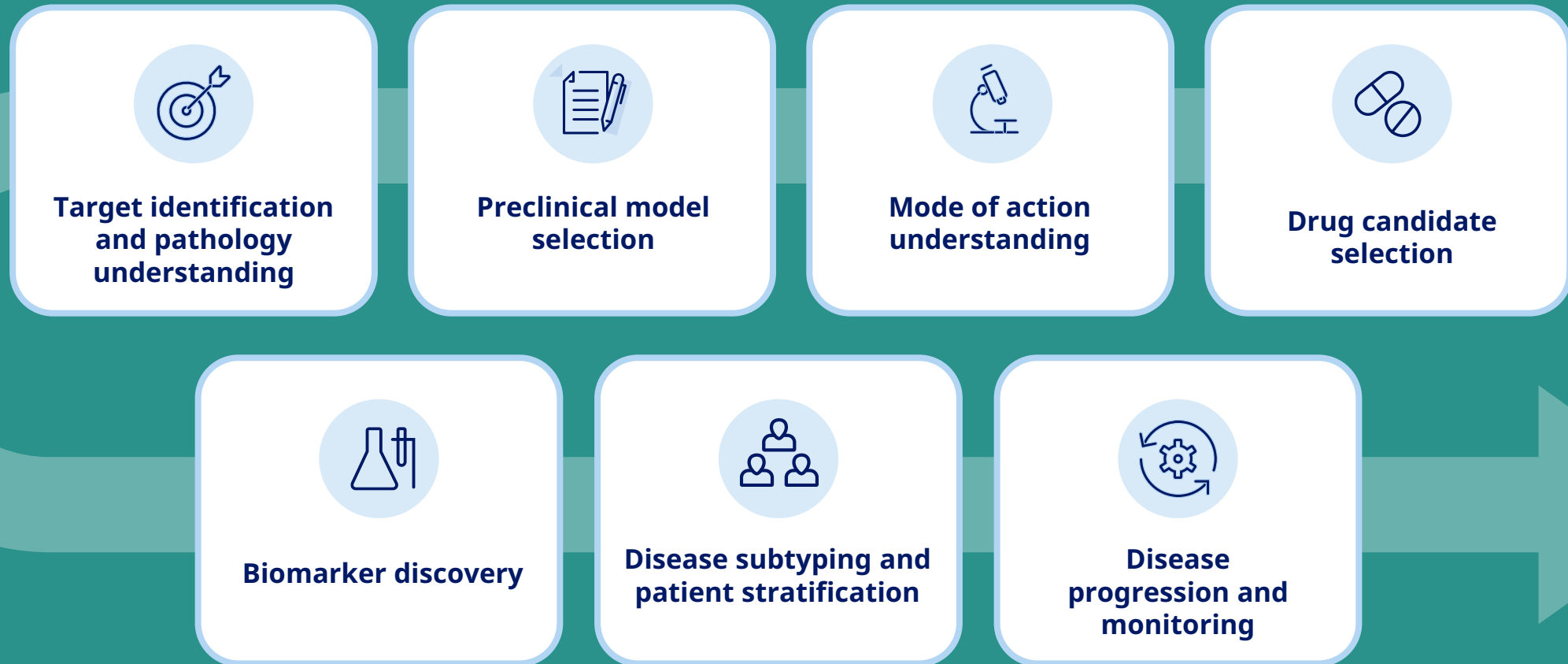
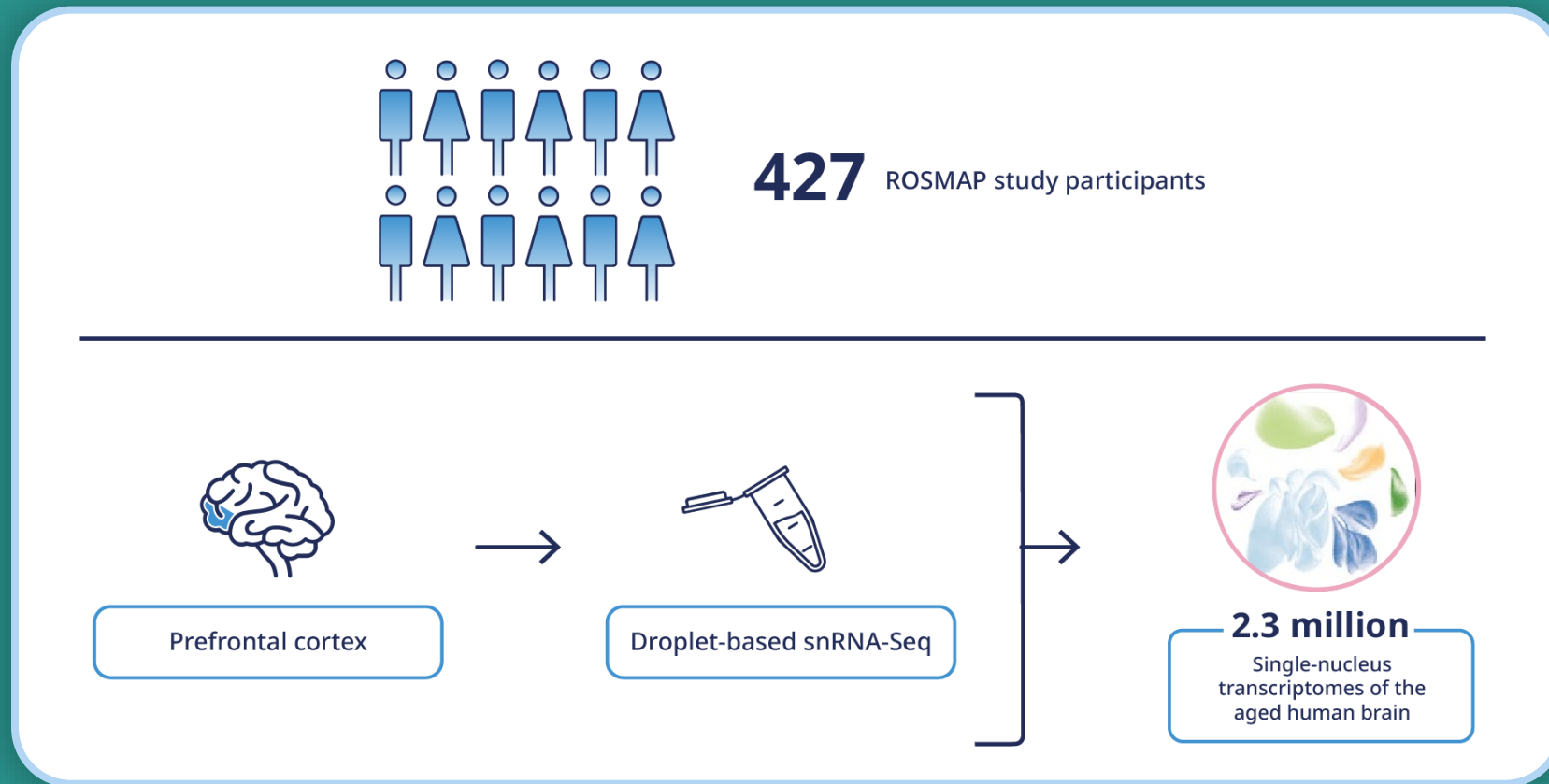


Figure modified from Van de Sande B et al. *Nat Rev Drug Discov* 2023;22:496–520.



Single-cell atlas of the aged human prefrontal cortex across 427 individuals



Can omics lead to precision medicine in AD?

3 key learnings

1

“Omics” (transcriptomics, proteomics) can aid in identifying novel drug targets, gaining a better understanding of pathophysiology, assessing potential treatment effects, and discovering novel biomarkers.

2

In contrast to bulk omics, single-cell techniques (performed on either post-mortem brain tissue or CSF/blood samples) provide knowledge on which specific cells and pathways are being modified throughout the disease course or by a drug or clinical intervention.

3

Single-cell atlas of the human brain are unravelling genes and transcriptomic changes associated with cognitive decline as well as preserved cognitive function (resilience factors).

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