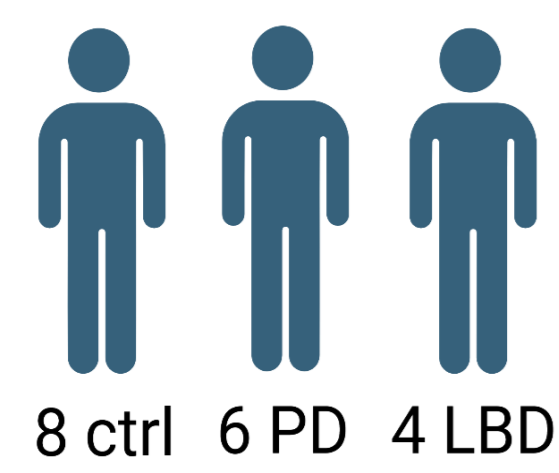


SINGLE-CELL RNA SEQUENCE ANALYSIS OF POST-MORTEM MIDBRAIN TISSUE REVEALS ALTERED GLIAL CELL FUNCTION IN PARKINSON'S DISEASE

ABSTRACT

Synucleinopathy marked by aberrant misfolding and aggregations of synuclein protein is a feature of many neurodegenerative disorders, particularly Parkinson's disease (PD) and Lewy Body Dementia (LBD). Neuroinflammation, characterized by reactive microglia and astrocytes are also a central marker of PD pathology. The inflammatory hypothesis suggests that dysregulated pathways in glial cells contribute to degeneration. Using a public dataset from Broad Institute, containing snRNA seq data from post-mortem midbrain tissue, we attempt to characterize functional alterations in several glial cell populations in Parkinson's disease. We specifically focus on functions that are associated with inflammation and damage response.

METHODS

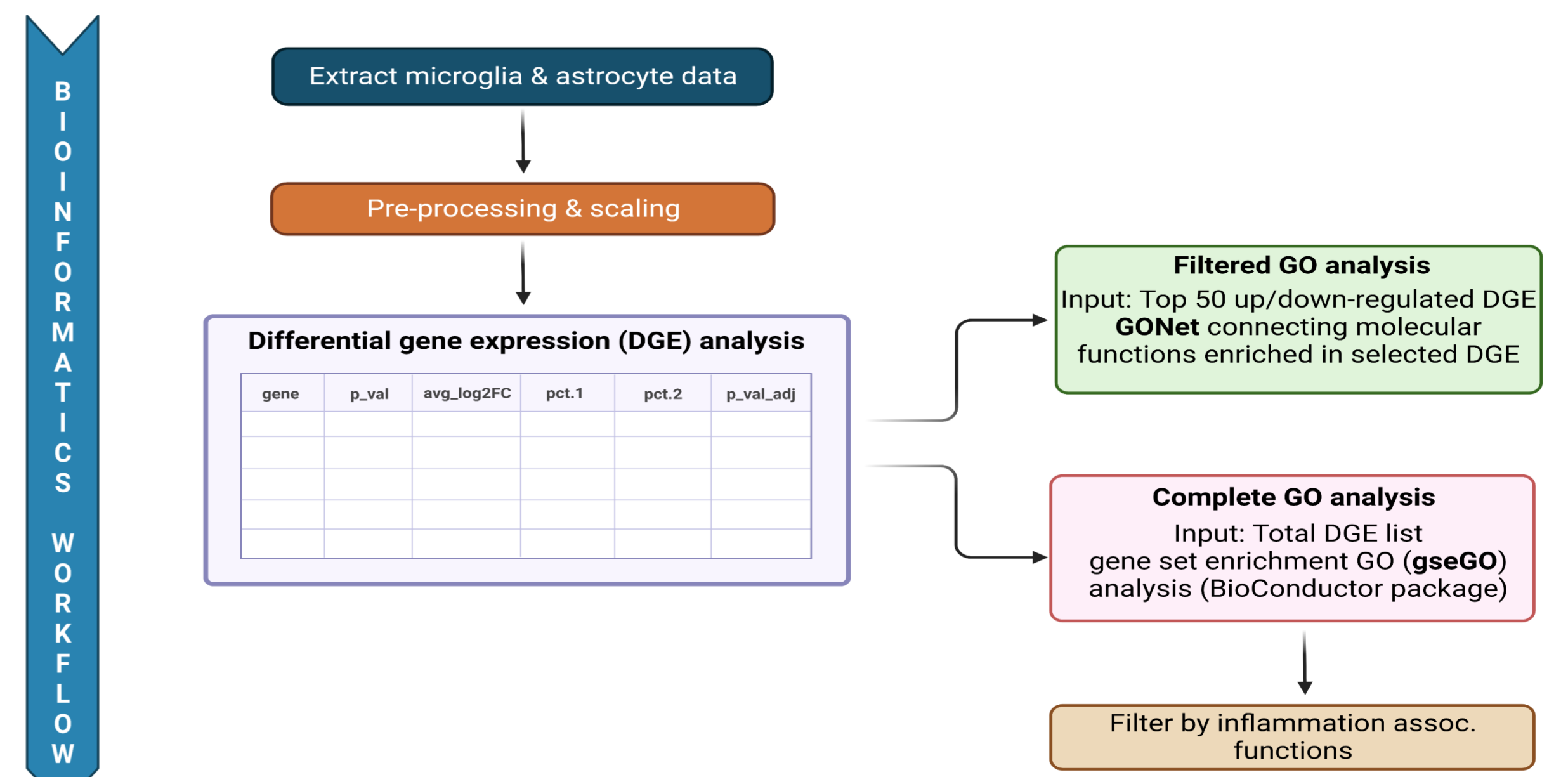


8 ctrl 6 PD 4 LBD

Data source: Single-cell genomic profile of human dopamine neurons identifies a population that selectively degenerates in Parkinson's disease (Kamath et al., 2022)

Brain regions: Caudate nucleus & substantia nigra

Disease ontology	# of cells	Cell type	# of cells
normal	231530	dopaminergic neurons	22048
Parkinson's disease	135344	astrocytes	33506
Lewy body dementia	67466	microglia	33041
		non-dopamine neurons	91479
		oligodendrocytes	178815
		oligo-progenitor cells	13691
		endothelial cells	14903



RESULTS

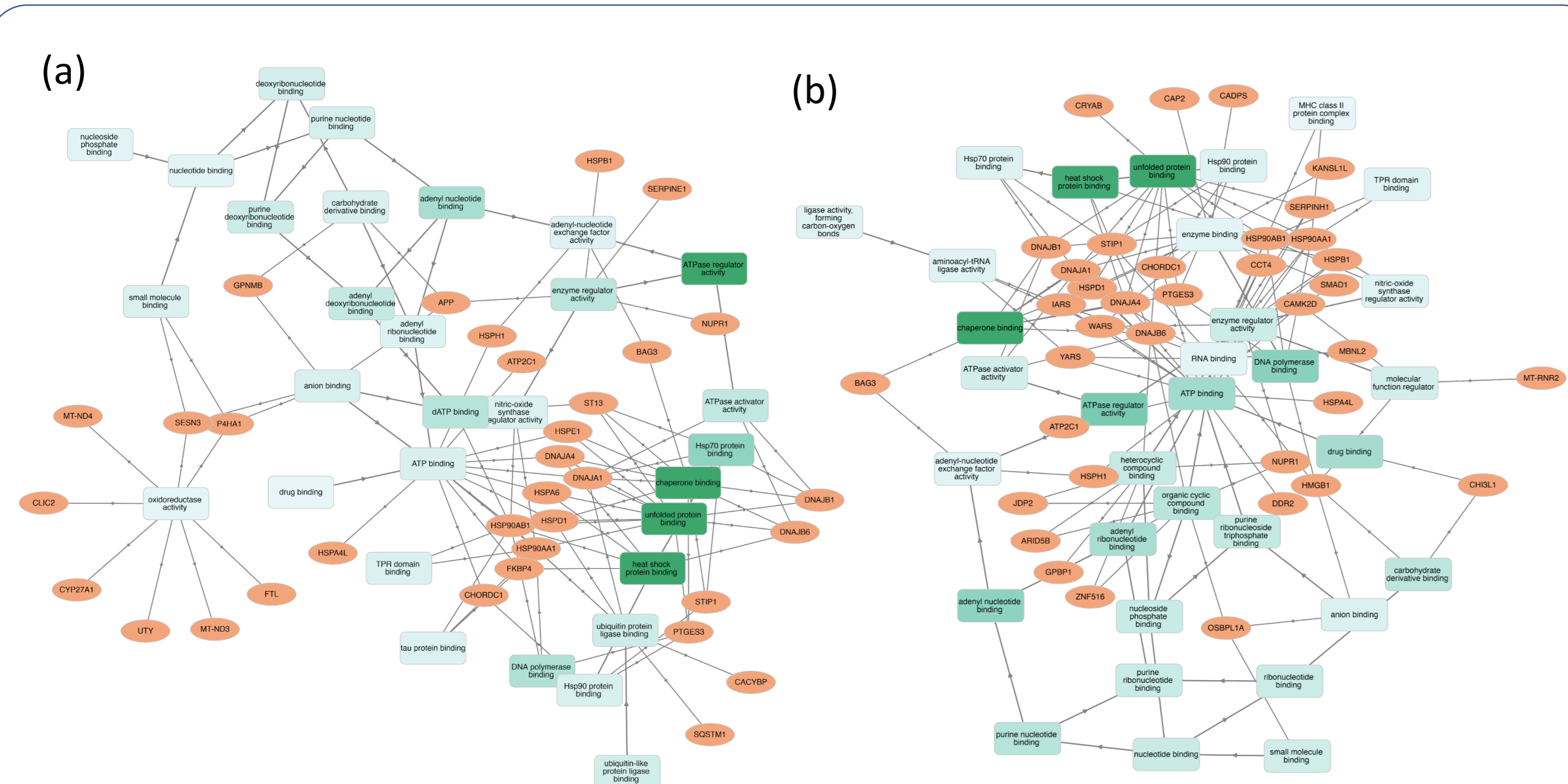


Figure 1: GONet enrichment plot of molecular functions associated with the 50 most upregulated genes in PD microglia (a) and PD astrocytes (b).

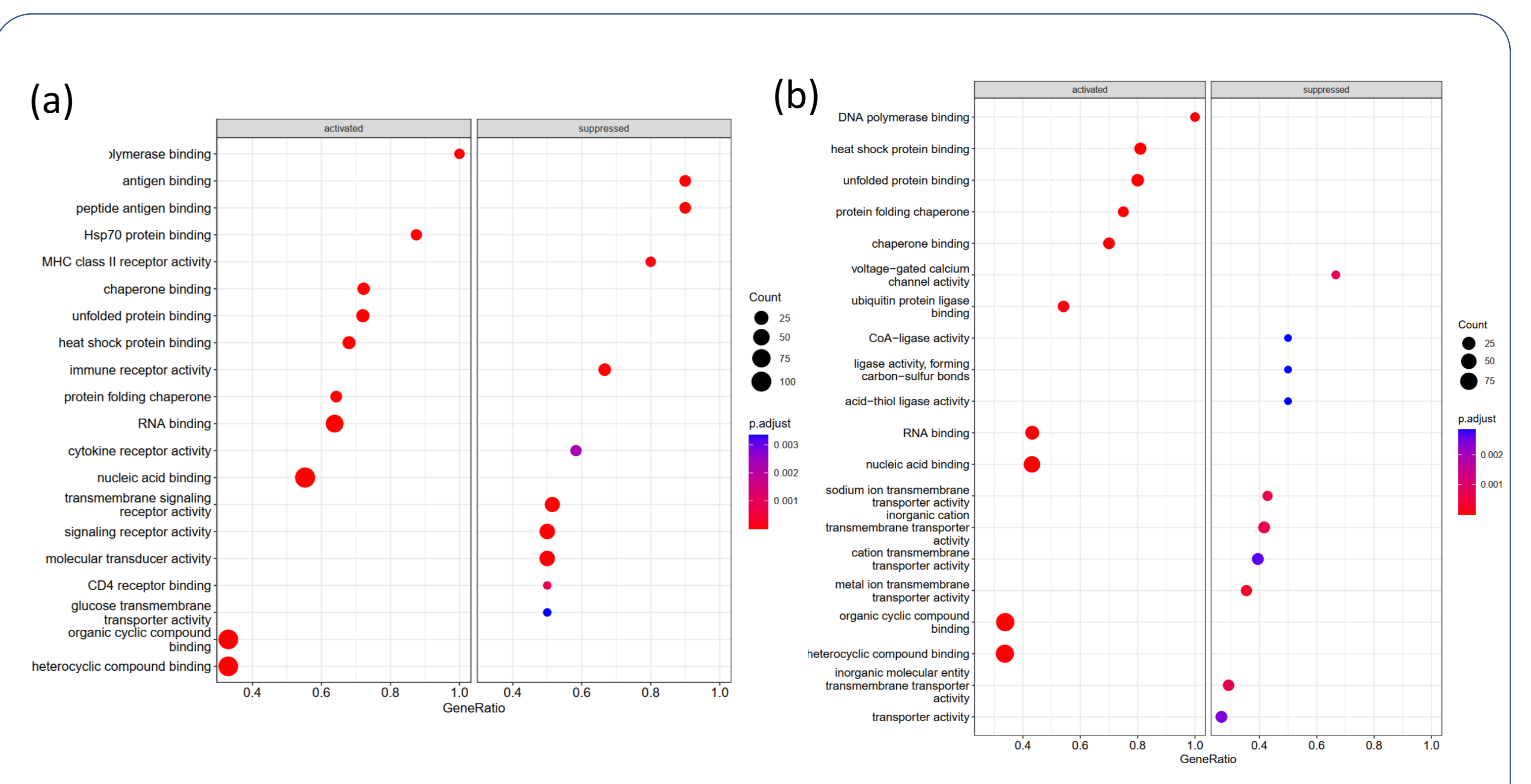


Figure 2: Dot plots of gseGO results showing the 10 most up and downregulated molecular functions in PD microglia (a) and PD astrocytes (b).

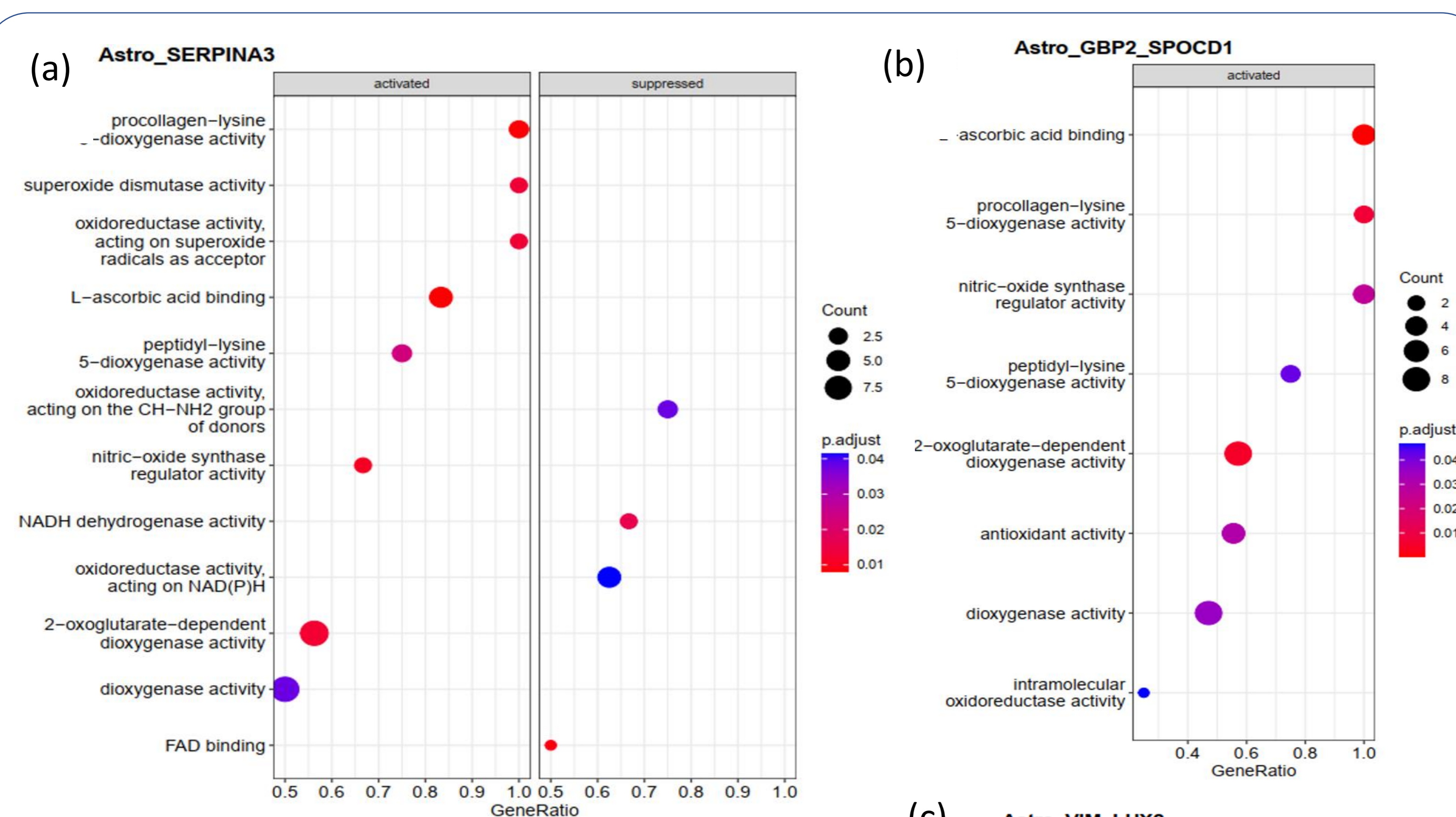


Figure 3: GseGO results for astrocyte subtypes filtered to show oxidoreductase activity related molecular functions in the three astrocyte subtypes that show increased expression in PD compared to control.

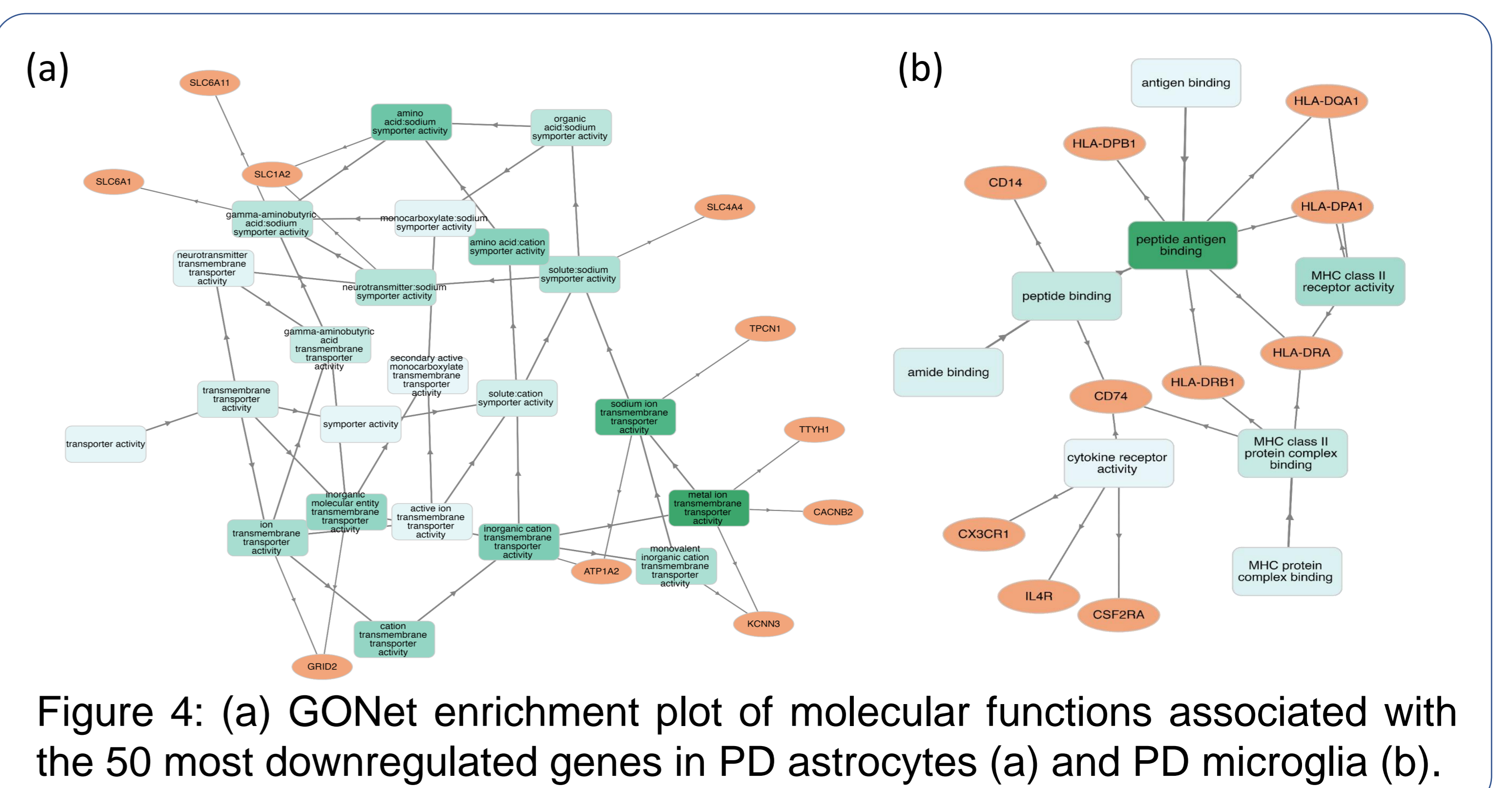


Figure 4: (a) GONet enrichment plot of molecular functions associated with the 50 most downregulated genes in PD astrocytes (a) and PD microglia (b).

CONCLUSIONS

- Proinflammatory pathways are downregulated in microglia, suggesting alternate reactive states of microglia at different stages of PD.
- Upregulated oxidoreductase activity in both astrocytes & microglia suggest involvement in responding to oxidative damage.
- Chaperone binding and protein folding activity are strongly upregulated in PD for both microglia and astrocytes, possibly in an attempt to clear misfolded proteins.

ACKNOWLEDGEMENTS

