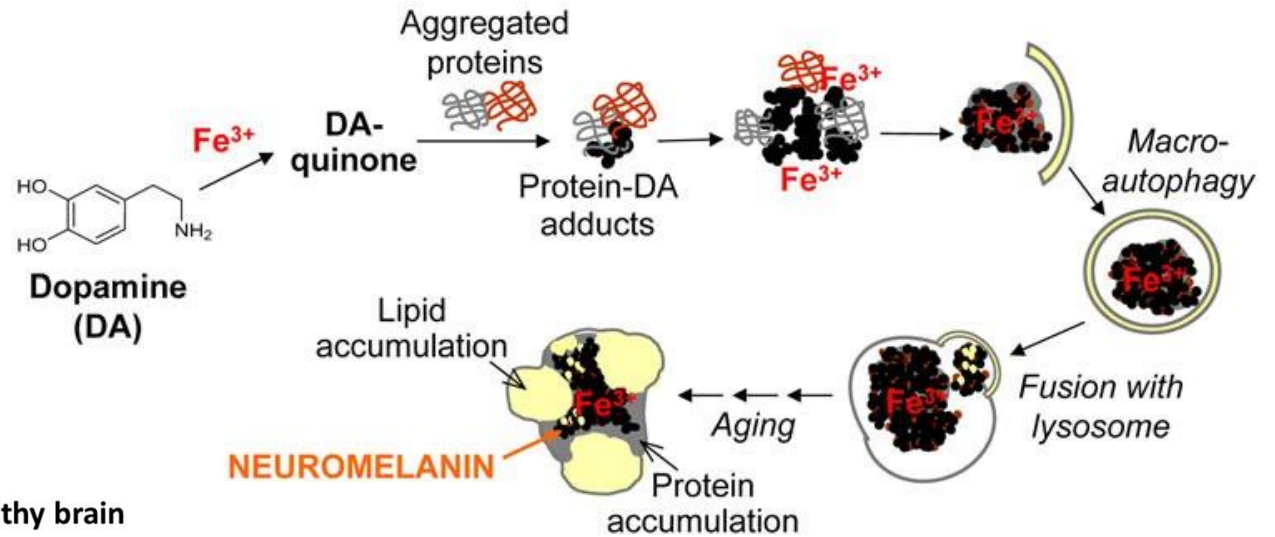
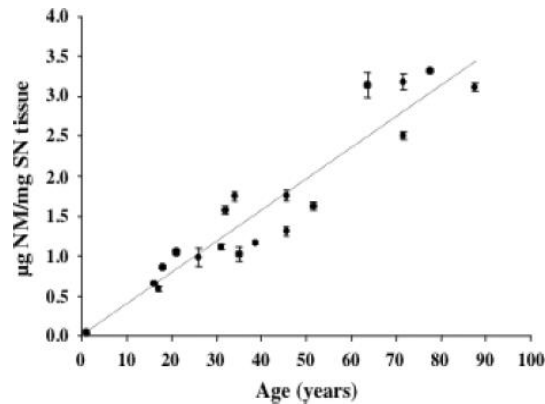
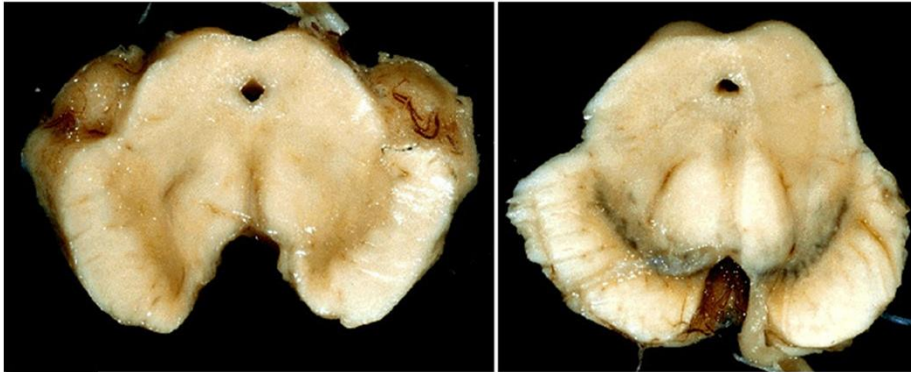


# Vulnerability of dopaminergic neurons in Parkinson's disease: the role of neuromelanin



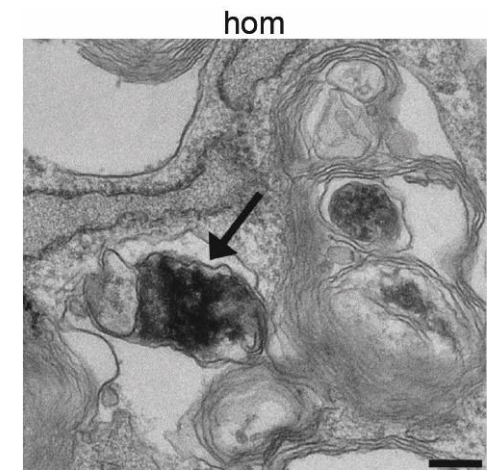
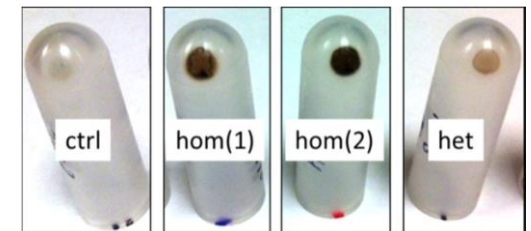
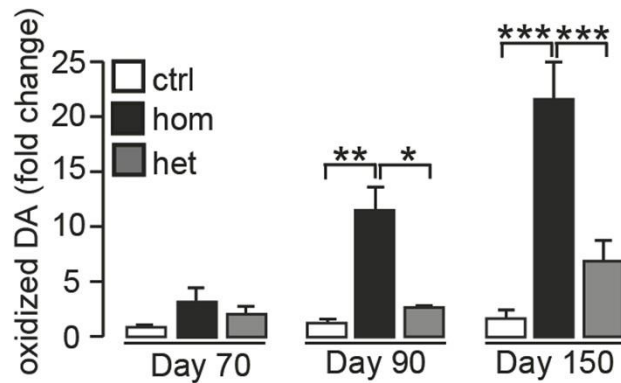
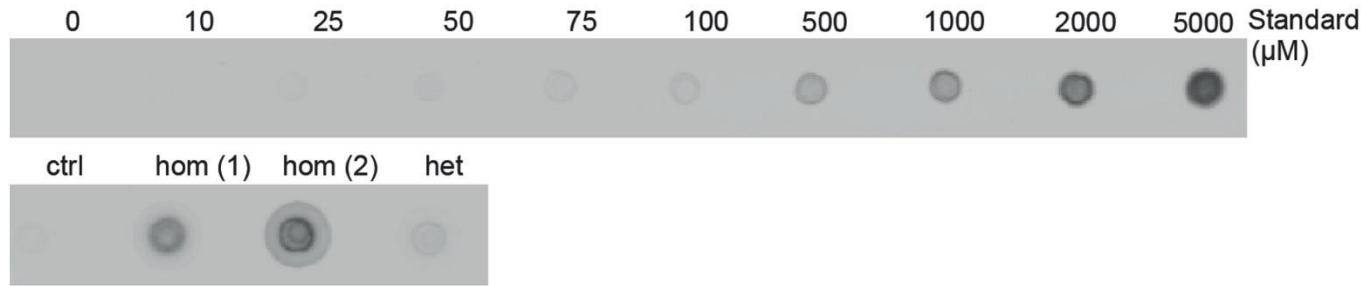
PD brain

Healthy brain



- Neuromelanin normally accumulates with age in nigral dopaminergic neurons
- Neuroprotective role - neuromelanin is chelator for metals, oxidized products
- Dopamine quinone-derived adducts not sequestered in neuromelanin considered toxic
- No neuromelanin detected in rodent substantia nigra

# Accumulation of oxidized dopamine and neuromelanin in iPS-derived DA neurons from patients with Parkinson's disease

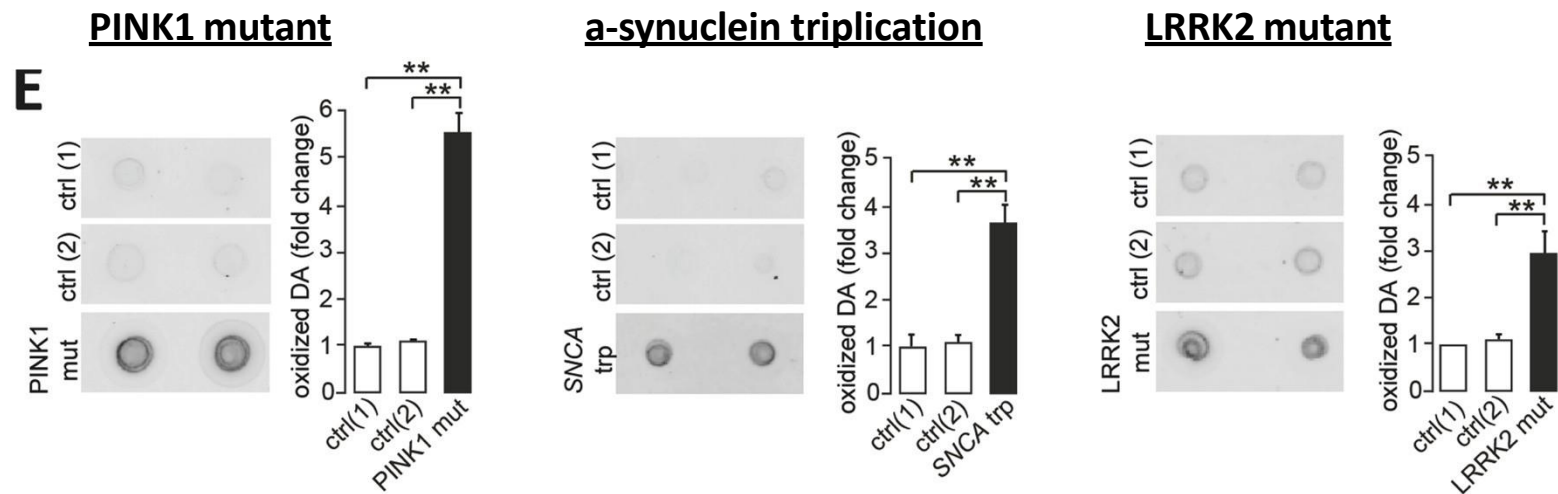


Lena Burbulla

Burbulla et al., *Science*, 2017

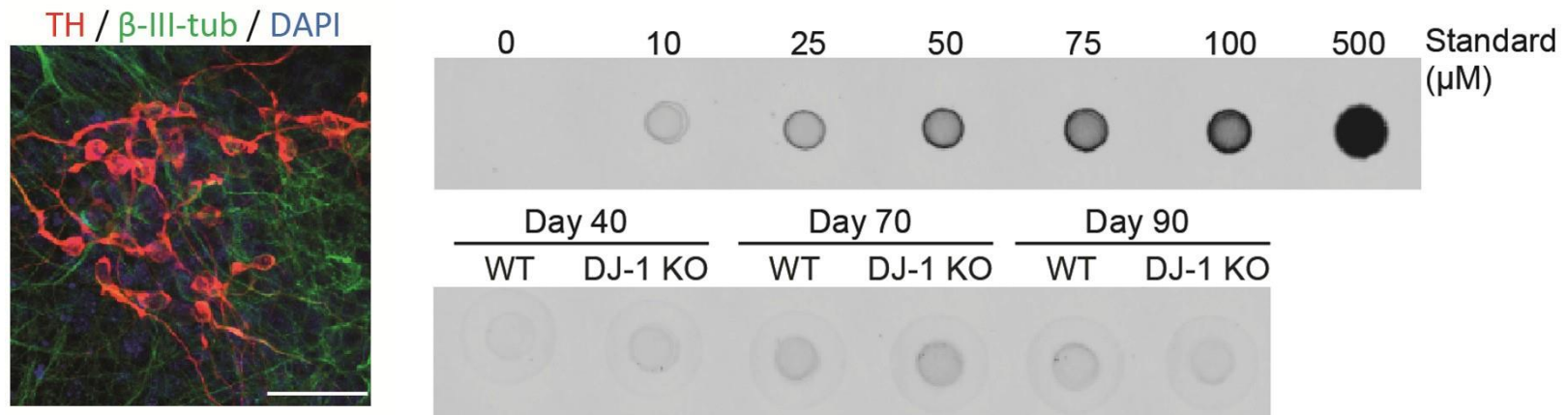
# Accumulation of oxidized dopamine in neurons from patients with genetic forms of PD

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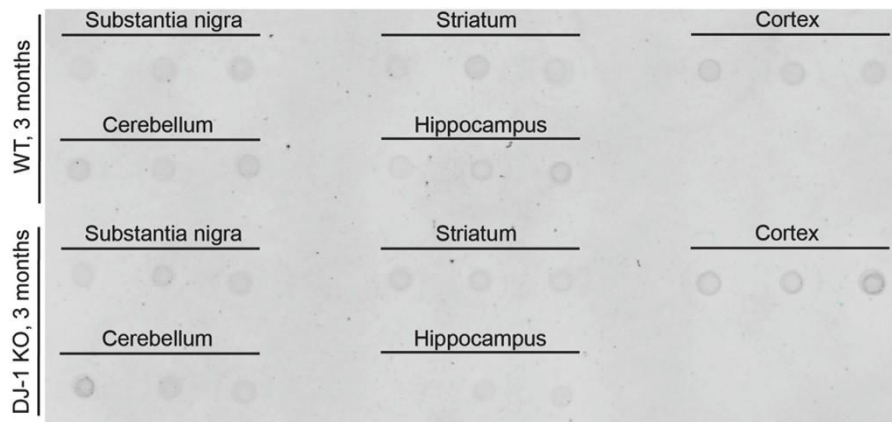
# No accumulation of oxidized dopamine in iPS-derived dopaminergic neurons from PD mouse models

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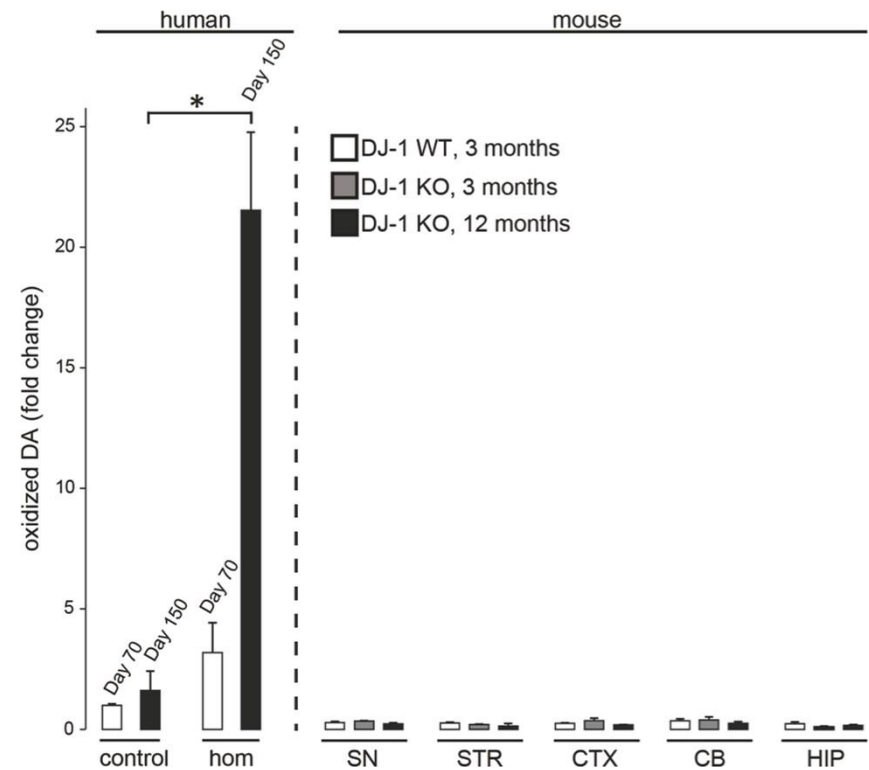
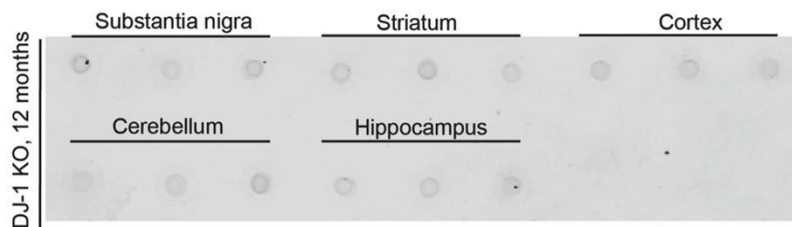


# No accumulation of oxidized dopamine in PD mouse models

## 3 months old mice

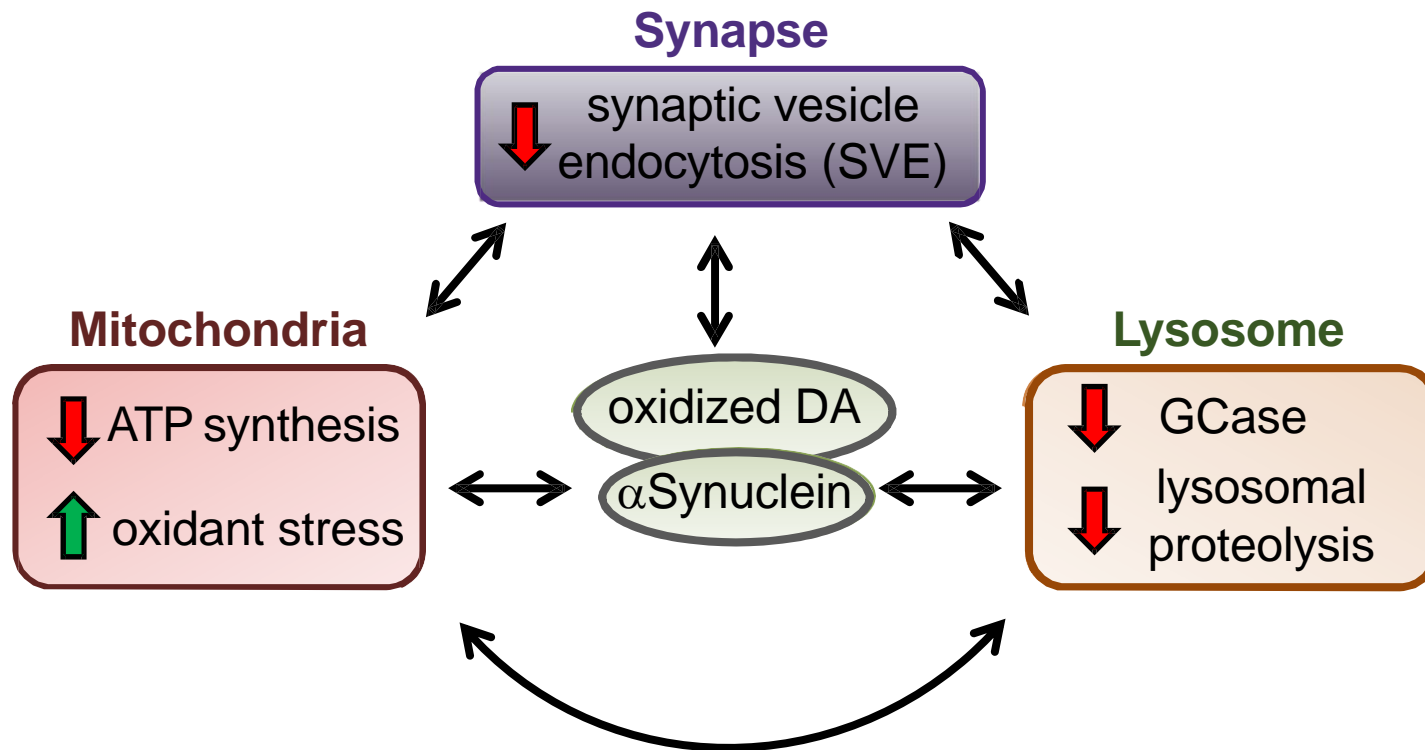


## 12 months old mice



# Mitochondrial, lysosomal and synaptic dysfunction in human dopaminergic neurons in PD

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Burbulla et al., *Science*, 2017

Wong et al. *Nature*, 2018

Nguyen & Krainc, *PNAS*, 2018

Burbulla et al., *Sci Transl Med*, 2019

Wong, Peng et al, *Dev Cell*, 2019

Kim et al, *Nature Commun* (2021)

Mazzulli et al., *Cell*, 2011

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## Session 3- Key Discussion Questions:

- What is the role of different animal and human model systems for discovery and validation?
- How can these approaches be used to gain a better understanding of different variants' impact on the disease state (e.g., variant to function), including phenotypic expression and differential vulnerability.
- How can greater ancestral diversity and the incorporation of environmental influences in genetic studies provide better insight into the phenotypic expression of neuropsychiatric disease states, and what are the implications for target validation?
- What scientific findings and lessons learned from rare variants and monogenetic diseases can be applied to more genetically complex common disorders?
- What criteria do different decision makers use when deciding how to validate biomarkers and select which targets to invest in and advance to clinic?

### Speakers

Helen Willsey, University of California, San Francisco

Fenna Krienen, Harvard Medical School

Paola Arlotta, Harvard University

Martin Kampmann, University of California, San Francisco

Daphne Koller, Insitro

Alice Zhang, Verge Genomics