



1-3 SEPTEMBER 2021 • VIRTUAL



Interaction of Aging and Stroke on Cellular and Molecular Level – Implications for Stroke Treatment

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Faculty Disclosure



X No, nothing to disclose

Yes, please specify:

7th European Stroke Organisation Conference

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Ischemic stroke

- Disease of aging
 - Most strokes in people > 65 years
 - Elderly shows higher mortality and poorer quality of life after stroke compared to young
- Sexually dimorphic disease (age modifies the influence of sex)
- These factors are often ignored in preclinical research
 - Most preclinical studies have used only young, male animals

• Complex factors underlying worsened stroke outcome in the elderly thus remain poorly understood, particularly in females





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Experimental design

- Female young (3m) and aged (18m) mice
- Permanent middle cerebral artery occlusion (MCAO) to model ischemic stroke (STAIR recommendation)
- Brains (penumbral cortex) collected at 3 days after MCAO and analyzed by RNA-Seq





Hypotheses

- Two components underlie aggravated stroke outcome in aged animals:
 - 1) Changes during normal aging making brain more susceptible to ischemic injury
 - 2) Difference in the response between young and aged brain leading to secondary injury and impaired regeneration





Aging is accompanied with increased neuroinflammation involving primarily glial cells



Aged control vs young control

- Minor changes detected in aged control brain
 - 52 genes UP, 13 genes DOWN

3 month

18 month

physiological aging

- Altered processes associated with increased inflammation and axonal degeneration
- Metanalysis identified involvement of pro-inflamatory astrocytes and microglia



Aging alters the magnitude of the transcriptional response to ischemic stroke

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Different response to ischemic impact



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Stroke does not activate neuroprotective pathways in young compared to aged mice







Combination of aging and stroke leads to massive activation of type I interferon signaling and aggravated inflammatory response



Combination of aging and stroke leads to massive activation of type I interferon signaling and aggravated inflammatory response





Age-dependent activation of type I IFN regulatory modules after stroke





Age-dependent activation of type I IFN regulatory modules after stroke



Cell

Volume 164, Issue 3, 28 January 2016, Pages 564-578

Resource

Parsing the Interferon Transcriptional Network and Its Disease Associations

Sara Mostafavi ^{1, 2, 8}, Hideyuki Yoshida ^{1, 8}, Devapregasan Moodley ¹, Hugo LeBoité ¹, Katherine Rothamel ¹, Towfique Raj ^{3, 5}, Chun Jimmie Ye ³, Nicolas Chevrier ⁴, Shen-Ying Zhang ⁶, Ting Feng ¹, Mark Lee ³, Jean-Laurent Casanova ⁶, James D. Clark ⁷, Martin Hegen ⁷, Jean-Baptiste Telliez ⁷, Nir Hacohen ³, Philip L. De Jager ^{3, 5}, Aviv Regev ³ ... Christophe Benoist ¹ ⁸ ¹⁰





CellPres



Temporal and cell-specific analysis of IFN-I signaling in young and aged mice after stroke





Summary

- Brain aging is accompanied by increasing inflammation involving primarily glial cells
- Transcriptional response to stroke is highly similar in young and aged brain and differs primarily in magnitude
- Differential stroke outcome is associated with overactivation of pro-inflammatory pathways in aged mice, rather than activation of neuroprotective program in young mice
- Increased activation of IFN-I signaling represents a key difference in response to stroke between young and aged mice
- Microglia and oligodendrocytes, but not astrocytes and endothelial cells massively upregulate IFN-I signature following stroke

Cell Reports

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Decoding the Transcriptional Response to Ischemic Stroke in Young and Aged Mouse Brain

Graphical Abstract



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In Brief

Cerebral stroke is a leading cause of mortality affecting mainly aged populations. Androvic et al. use RNA-seq to analyze aging, stroke, and their interaction in mouse brain. They identify pathways associated with agedependent vulnerability to stroke, including overactivation of type I interferon signaling and downregulation of the synaptic maintenance program.

Highlights

RNA-seq analysis of aging, ischemic stroke, and their interaction in female mice

• Response to stroke in young and aged brain is similar, but differs in magnitude

- Aged ischemic brain is characterized by upregulation of typel interferon signaling
- Aged mice downregulate axonal and synaptic maintenance
 program after stroke

Androvic et al., 2020, Cell Reports 31, 107777 June 16, 2020 © 2020 The Author(s). https://doi.org/10.1016/j.celrep.2020.107777

CellPress



Implication for stroke treatment

• Targeting IFN-I signaling



Moses Zhang ^a, Catherine E. Downes ^a, Connie H.Y. Wong ^b, Kate M. Brody ^a, Pedro L. Guio-Agulair ^a, Jodee Gould ^c, Robert Ates ^a, Paul J. Hertzog ^c, Juliet M. Taylor ^a, Peter J. Crack ^a $\stackrel{\otimes}{\sim}$ \boxtimes

• Targeting inflammatory response



Neurobiology of Disease Volume 151, April 2021, 105257



Inhibition of TGFβ-activated kinase 1 promotes inflammation-resolving microglial/macrophage responses and recovery after stroke in ovariectomized female mice

Yaan Liu ¹, Sicheng Li ¹, Rongrong Wang, Hongjian Pu, Yongfang Zhao, Qing Ye, Yejie Shi ^A

- Targeting cell populations responsible for IFN-I signaling
 - In progress...



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