Therapeutic potential of NIrp1 inflammasome, Caspase-1, or Caspase-6 against Alzheimer disease cognitive impairment

Joseph Flores¹, Anastasia Noël¹, Marie-Lyne Fillion¹, Andréa C. LeBlanc^{1,2*}

 ¹Lady Davis Institute for Medical Research, Jewish General Hospital, 3755 Chemin Côte Ste Catherine, Montreal, Quebec, Canada H3T 1E2
 ²Department of Neurology and Neurosurgery, McGill University, 3755 Rue University, Montreal, Quebec, Canada, H3T 1E2

*Corresponding author: Andréa LeBlanc PhD, Lady Davis Institute for Medical Research, Jewish General Hospital, 3755 Chemin Côte Ste Catherine, Montreal, QC, Canada H3T 1E2. Tel.: +1 (514) 340 8222 ext 25303; e-mail: andrea.leblanc@mcgill.ca

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Table S1. Behavioral data comparing J20 versus genetically ablated J20, WT versus genetically ablated WT, and J20 and genetically ablated J20 versus WT mice.

Nirp1, Casp ablated J20	1, Casp6 genetica vs. J20 mice	lly	J20	J20/NIrp1-	J20/Casp1 ^{./-}	J20/Casp6 ^{./-}		J20/NIrp1+/-	J20/Casp1+/-	J20/Casp6*/-
	Mice (n)		15	10	13	11		14	11	11
NOR	NOR Discrimination in			▲ (p<0.0001)	▲ (p<0.0001)	▲ (p<0.0001)		▲ (p<0.0001)	▲ (p<0.0001)	▲ (p<0.0001)
								1		
Barnes Maze	Mice (n)		13	10	13	9		12	11	10
Learning	Primary erro	rs		▼ (p=0.0017)	▼ (p=0.02)	▼ (p=0.03)		▼ (p=0.01)	Ш	=
acquisition	Primary laten	су		=	=	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	=			
	Primary erro	rs		=	▼ (p=0.0016)	▼ (p=0.03)		=	=	=
Probe test	Primary laten	су		=	=	=		=	=	=
	Target preference			no difference	no difference	no difference		no difference	no difference	no difference
Nirp1, Casp ablated WT	1, Casp6 genetica vs. WT mice	lly	WT	WT/Nirp1 ^{-/-}	WT/Casp1 ^{./-}	WT/Casp6 ^{-/-}		WT/NIrp1 ^{+/-}	WT/Casp1*'-	WT/Casp6 ^{+/-}
NOD	Mice (n)		16	13	12	12		12	13	11
NUR	Discrimination i	ndex		=	=	=		=	=	=
	1	1		1	1	1		I		I
Barnes Maze	Mice (n)		16	13	12	12		12	12	11
Learning	Primary erro	rs		=	=	=		=	=	=
acquisition	Primary laten	су		=	=	=	12 11 \bigvee = $(p=0.01)$ = $=$ = $=$ = $=$ = $=$ = no difference no difference NUT/NIrp1*/- WT/Casp1*/- 12 13 $=$ = 12 13 $=$ = <	=		
	Primary erro	rs		=	=	=		=	=	=
Probe test	Primary latency			=	=	=		=	=	=
	Target prefere	nce		no difference	no difference	no difference		no difference	no difference	no difference
J20 and ger J20 vs. WT	netically ablated mice	wт	J20	J20/NIrp1 ^{-/-}	J20/Casp1	J20/Casp6 ^{-/-}		J20/NIrp1*/-	J20/Casp1+/-	J20/Casp6 ^{+/-}
							1			

J20 and ger J20 vs. WT	netically ablated mice	wт	J20	J20/NIrp1 ^{./-}	J20/Casp1 ^{-/-}	J20/Casp6 ^{-/-}		J20/NIrp1+/-	J20/Casp1*'-	J20/Casp6 ^{+/-}
NOR	Mice (n)	16	15	10	13	11		14	11	11
	Discrimination index		▼ (p=0.0001)	=	=	=		=	=	=
Barnes Maze	Mice (n)	16	13	10	13	9		12	11	10
Learning	Primary errors		▼ (p=0.03)	=	=	=		=	=	=
acquisition	Primary latency		=	=	=	=		=	=	=
	Primary errors		NS p=0.06	=	=	=		=	=	=
Probe test	Primary latency		=	=	=	=		=	=	=
	Target preference		impaired	no difference	no difference	no difference		no difference	no difference	no difference

Episodic memory was measured with novel object recognition (NOR) while spatial memory was measured with the Barnes maze. Mice numbers are indicated in each group of mice. Upward arrow (▲) represents increased value,

downward arrow ($\mathbf{\nabla}$) represents decreased value, equal sign (=) represents unchanged value. Significant (p<0.05) or non-significant (NS) but close (0.1>p>0.05) statistical p values are indicated below or next to arrows.

Table S2. Iba1⁺-microglial and GFAP⁺-astrocyte immunostaining density comparing J20 and genetically ablated J20 versus WT, or genetically ablated J20 versus J20 in the hippocampus and cortex.

	vs. WT mice	WT	J20	J20/NIrp1- ⁻	J20/Casp1	J20/Casp6 ^{-/-}	J20/NIrp1+/-	J20/Casp1*/-	J20/Casp6+/-
lba1	Mice (n)	8	6	5	6	4	8	6	6
vs WT	lba1 ⁺ cell density		▲ (p<0.0001)	▲ (p=0.0002)	=	=	▲ (p=0.002)	=	▲ (p=0.035)
GFAP	Mice (n)	6	6	5	6	4	6	6	6
vs WT	GFAP⁺ density		=	=	=	=	=	=	=
	vs. J20 mice								
lba1	Mice (n)	8	6	5	6	4	8	6	6
vs J20	lba1 ⁺ cell density	▼ (p<0.0001)		▼ (p=0.0006)	▼ (p<0.0001)	▼ (p<0.0001)	▼ (p<0.0001)	▼ (p<0.0001)	▼ (p<0.0001)
GFAP	Mice (n)	6	6	5	6	4	6	6	6
vs J20	GFAP⁺ density	=		=	=	=	=	=	=

Hippocampus

	Cortex										
	vs. WT mice	WT	J20	J20/NIrp1 ^{-/-}	J20/Casp1 ^{,,}	J20/Casp6 ^{-/-}		J20/NIrp1+/-	J20/Casp1+/-	J20/Casp6 ^{+/-}	
lba1	Mice (n)	8	6	5	6	4		8	6	6	
vs WT	lba1 ⁺ cell density		▲ (p<0.0001)	▲ (p=0.0001)	=	=		▲ (p<0.0001)	=	▲ (p=0.01)	
GFAP	Mice (n)	6	6	5	6	4		6	6	6	
vs WT	GFAP⁺ density		=	=	▲ (p=0.03)	=		=	=	=	
	vs. J20 mice										
lba1	Mice (n)	8	6	5	6	4		8	6	6	
vs J20	lba1⁺ cell density	▼ (p<0.0001)		=	▼ (p<0.0001)	▼ (p=0.002)		=	▼ (p<0.0001)	▼ (p<0.0001)	
GFAP	Mice (n)	6	6	5	6	4		6	6	6	
vs J20	GFAP⁺ density	=		=	=	=		=	=	=	

Immunohistochemical staining density was measured as described in methods. Upward arrow (\blacktriangle) represents increased value, downward arrow (\blacktriangledown) represents decreased value, equal sign (=) represents unchanged value. Significant (p<0.05) statistical p values are indicated.

Table S3. Immunohistochemical A β staining density and ELISA-measured A β in the hippocampus and cortex of genetically ablated J20 compared to J20 mice.

Hippocampus											
NIrp1, Casp1, Casp6 genetically ablated J20 vs. J20 mice			J20/ <i>NIrp1</i>	J20/Casp1	J20/Casp6 ^{-/-}		J20/NIrp1*'-	J20/Casp1 ^{+/-}	J20/Casp6 ^{+/-}		
Amyloid plaques	Mice (n)	6	5	6	4		6	6	6		
	A β staining density		▼ (p=0.001)	▼ (p=0.001)	▼ (p=0.007)		▼ (p=0.001)	▼ (p=0.001)	▼ (p=0.004)		
Amyloid protein levels (ELISA)	Mice (n)	6	6	6	5		6	6	6		
	Total Aβ		▼ (p<0.0001)	▼ (p=0.001)	▼ (p=0.03)		▼ (p=0.002)	▼ (p=0.0001)	▼ (p=0.0002)		
Formic acid soluble	Aβ38/total Aβ	otal Aβ otal Aβ		▼ (p=0.005)	=	=		=	▼ (p=0.007)	=	
amyloid levels	Aβ40/total Aβ		▲ (p<0.0001)	NS p=0.054	=		=	▲ (p=0.0003)	▲ (p=0.003)		
	Aβ42/total Aβ		▼ (p=0.0005)	NS p=0.062	=		\checkmark (p=0.001) \checkmark (p=0.001) \checkmark (p=0.004) 6 6 6 (p=0.002) (p=0.0001) (p=0.0002) = (p=0.007) = = (p=0.0003) (p=0.003) = (p=0.028) NS p=0.087 = = = = = = = = =				
	Total Aβ		▼ (p=0.01)	=	=		=	=	=		
RIPA soluble amyloid	A β 38/total A β		=	=	=		=	=	=		
levels	Aβ40/total Aβ		=	=	=		=	=	=		
	A β 42/total A β		=	=	=		=	J20/Casp1*/- J20/C 6 - $(p=0.001)$ \checkmark (p=0.001) $(p=0.0001)$ $(p=0.0001)$ $(p=0.0001)$ $(p=0.0001)$ $(p=0.0003)$ $(p=0.0003)$ $(p=0.028)$ NS p: = = = = = = = = = = = = = = = = = =	=		

Cortex										
Nirp1, Casp1, Casp6 genetically ablated J20 vs. J20 mice			J20/NIrp1 ^{-⊦-}	J20/Casp1	J20/Casp6 ^{-/-}		J20/NIrp1+'-	J20/Casp1 ^{+/-}	J20/Casp6*/-	
Amyloid plaques	Mice (n)	6	5	6	4		6	6	6	
	$A\beta$ staining density		=	=	=		=	=	=	
Amyloid protein levels (ELISA)	Mice (n)	6	6	6	5		6	6	6	
	Total Aβ		=	=	=		=	=	=	
Formic acid soluble	Aβ38/total Aβ		=	=	=		=	=	=	
amyloid levels	Aβ40/total Aβ		=	=	=		=	=	=	
	Aβ42/total Aβ		=	=	=		=	Irp1*/- J20/Casp1*/- J20/Casp6*/- 3 6 6 = = = 3 6 6 = = = 3 6 6 = = = = = = = = = = = = = = = = = = = = = = = = = = = = = = = = =		
	Total Aβ		=	=	=		=	=	=	
RIPA soluble amyloid	Aβ38/total Aβ		=	=	=		=	=	=	
levels	Aβ40/total Aβ		=	=	=		=	=	=	
	Aβ42/total Aβ		=	=	=		=	J20/Casp1*/- J20/ 6 = - = 6 = - = - = - = - = - = - = - = - = - = - = - = - = - = - = - = - = - =	=	

Immunohistochemical staining density and ELISA were done as described in methods. Upward arrow (\blacktriangle) represents increased value, downward arrow (\blacktriangledown) represents decreased value, equal sign (=) represents unchanged

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value. Significant (p<0.05) or non-significant (NS) but close (0.1>p>0.05), statistical p values are indicated. NS indicates not significant.



Fig. S1. *Nlrp1*, *Casp1*, or *Casp6* can be ablated from J20 and WT mice. Representative PCR of A 360 bp *hAPP^{Sw/Ind}* transgene amplicon in J20 mice, **B** 490 bp null and 792 bp WT allele for *Nlrp1*, **C** 300 bp null and 500 bp WT allele for *Casp1*, and **D** 340 bp null and 620 bp WT allele for *Casp6*.



Fig. S2. *Nlrp1*, *Casp1*, or *Casp6* genetic ablation does not alter locomotor behaviour. Open field **A** number of quadrant entries [F = 7.779, p < 0.0001], **B** percentage time moving [F = 4.745, p < 0.0001], and **C** percentage time in periphery, indicative of thigmotaxis [F = 4.863, p < 0.0001]. Bars represent mean \pm SEM of all mice per group; symbols denote performance of individual mice. *n* = 18 WT, 15 J20, 10 J20/*Nlrp1*^{-/-}, 13 J20/*Casp1*^{-/-}, 10 J20/*Casp6*^{-/-}, 13 WT/*Nlrp1*^{-/-}, 12 WT/*Casp1*^{-/-}, 12 WT/*Casp6*^{-/-}, 14 J20/*Nlrp1*^{+/-}, 12 J20/*Casp1*^{+/-}, 11 J20/*Casp6*^{+/-}, 12 WT/*Nlrp1*^{+/-}, 13 WT/*Casp1*^{+/-}, 11 WT/*Casp6*^{+/-} for (**A**-**C**). One-way ANOVA, Bonferroni's post-hoc compared to WT ([#]) or J20 (*). [#] or * p < 0.05, ^{##}p < 0.01, ^{####}p < 0.0001.



Fig. S3. *Nlrp1*, *Casp1*, or *Casp6* genetic ablation prevents cognitive deficits in J20 mice. Barnes maze distribution of pokes to each hole during the probe, where (T) indicates target hole. Bars represent mean \pm SEM of all mice per group; symbols denote performance of individual mice.



Fig. S4. Synaptic density alterations after *Nlrp1*, *Casp1*, or *Casp6* genetic ablation in J20 mice. A Representative Golgi-Cox-stained dendritic spines in the stratum radiatum (SR) of the hippocampal CA1. Scale bar = 5 μ m. B Cell density in the pyramidal cell layer (PCL) of the hippocampal CA1. C Representative synaptophysin staining of the hippocampus. CA1 SR = stratum radiatum of the CA1, CA3 SLu = stratum lucidum of the CA3, DG-M = molecular layer of the dentate gyrus. Scale bar = 200 μ m.



Fig. S5. *Nlrp1*, *Casp1*, or *Casp6* genetic ablation reduces microglia activation in J20 mice. Representative micrographs of Iba1⁺ immunostained-microglia in the retrosplenial cortex. Scale bar = $50 \mu m$.



Fig. S6. *Nlrp1*, *Casp1*, or *Casp6* genetic ablation does not alter GFAP⁺ astrocytes in J20 mice. A Representative GFAP⁺ micrographs showing the stratum oriens (SO), pyramidal cell layer (PCL), SR, and stratum lacunosummoleculare (SLM) of the hippocampal CA1 (top) and retrosplenial and S1 cortex (bottom). Scale bar = 200 µm. **B** GFAP⁺ staining density from the SO to the SLM in the hippocampal CA1 [F = 4.317, p = 0.000032]. **C** GFAP⁺ staining density in the cortical retrosplenial and S1 area [F = 2.572, p = 0.0063]. Bars represent mean ± SEM of all mice per group; symbols denote individual results. *n* = 6 WT, 6 J20, 5 J20/*Nlrp1^{-/-}*, 6 J20/*Casp1^{-/-}*, 4 J20/*Casp6^{-/-}*, 6 WT/*Nlrp1^{-/-}*, 6 WT/*Casp6^{-/-}*, 6 J20/*Nlrp1^{+/-}*, 6 J20/*Casp1^{-/-}*, 6 WT/*Nlrp1^{+/-}*, 6 WT/*Casp6^{+/-}* for (**B**,**C**). One-way ANOVA, Bonferroni's post-hoc compared to WT ([#]) or J20 (*). [#] or * p < 0.05.



Fig. S7. *Nlrp1*, *Casp1*, or *Casp6* genetic ablation prevents Aβ deposition in J20 mice. A-B Representative Aβstained micrographs of the A hippocampus and B cortical retrosplenial and S1 area. Scale bars = 100 µm. C Hippocampal formic acid-soluble Aβ₃₈ [F = 6.346, p = 0.0001], Aβ₄₀ [F = 6.198, p = 0.0001], and Aβ₄₂ [F = 6.166, p = 0.0002] protein levels. D Hippocampal RIPA-soluble Aβ₃₈, Aβ₄₀ [F = 2.733, p = 0.02], and Aβ₄₂ protein levels. E Cortical formic acid-soluble Aβ₃₈, Aβ₄₀, and Aβ₄₂ protein levels. F Cortical RIPA-soluble total Aβ₃₈, Aβ₄₀, and Aβ₄₂ protein levels. Bars represent mean ± SEM of all mice per group; symbols denote performance of individual mice. *n* = 6 J20, 6 J20/*Nlrp1*^{-/-}, 6 J20/*Casp1*^{-/-}, 5 J20/*Casp6*^{-/-}, 6 J20/*Nlrp1*^{+/-}, 6 J20/*Casp6*^{+/-} for (C-F). One-way ANOVA, Dunnett's post-hoc compared to J20. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.