

Neuroprotective pentapeptide CN-105 is associated with reduced sterile inflammation and improved functional outcomes in a traumatic brain injury mouse model

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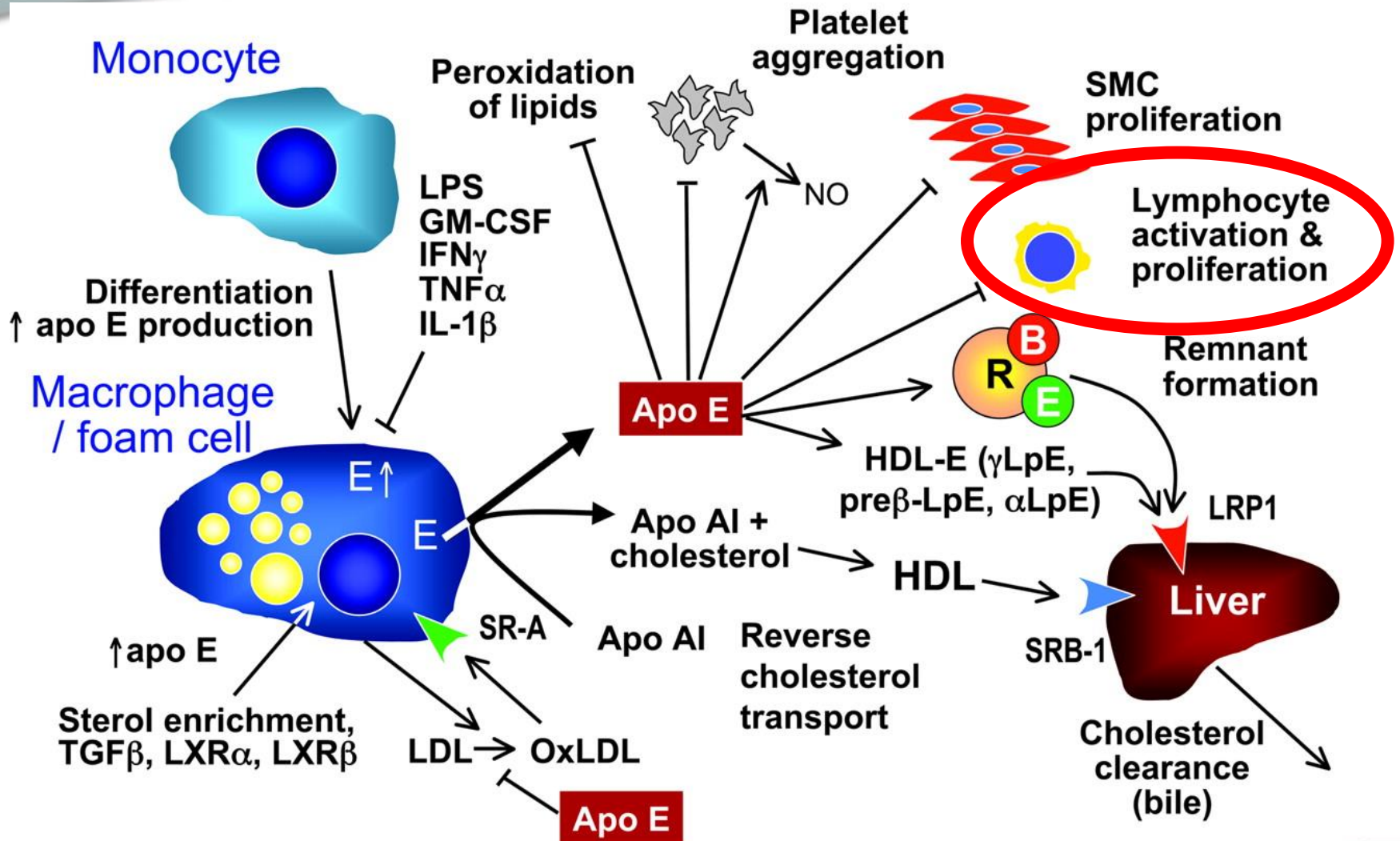
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- Aim of the study
- Material and Methods
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Apo E



Apo E

- Apolipoprotein E
 - Lipoprotein: transport of triglyceride acids and cholesterol in blood
- Polymorphism on Chr. 19
 - apoE2: reduced binding affinity to LDL-receptor, atherosclerosis,
 - apoE3: the „neutral“ Apo E genotype
 - apoE4: associated with atherosclerosis, alzheimers disease, cerebrovascular disease
- Function:
 - found to reduce glial activation and inflammatory cytokine release in vitro
 - → extend to mouse model,
 - but: Apo E holoprotein does not cross the blood brain barrier and would not be suitable for peripheral administration
- CN-105:
 - smaller peptide
 - Designed modeling the polar receptor binding face of the helical apoE receptor binding region (Ac-VSRRR-amide)

Aim of the study

- → to investigate the therapeutic potential of CN-105 in a murine model of closed head injury
- Hypothesis: intravenous administration of CN-105 dampens neuroinflammatory responses and thus possibly improves the functional outcomes

Material and Methods

- **Closed head injury model**
 - 12-14 weeks old male mice
 - anesthesia induction, tracheal intubation and lungs were mechanically ventilated
 - secured in a stereotactic device on acrylic cast to allow 3mm of space below the head for acceleration/deceleration; no ear bars to avoid basilar skull fracture
 - metallic disc was adhered at the skull, immediately caudal to bregma
 - 2.0-mm diameter pneumatic impactor → single midline impact
 - sham mice were treated identically except for absence of impact
- **Drug administration**
 - Animals were placed in restrainer
 - A) single i.v. dose of 100 μ L drug was administered by tail vein
 - B) vehicle treated animals received 100 μ L of normal saline

Material and Methods

- **Immunohistochemistry**

- to assess effect of CN-105 on inflammation, neuronal injury etc.
- immunohistochemical staining performed using the F4/80 antibody and Fluoro-Jade B stain on days 10 and 1, respectively, after TBI
- IHC was performed on separate cohorts of mice from those used in neurobehavioral tests
- for histological assessment: secondary antibody, biotinylated goat anti-mouse IgG (1:3000), ABC, DAB (all from vector laboratories)

- **Cell quantification and image analysis**

- A) F4/80 quantification:
 - brains of 5 TBI treated and 6 TBI vehicle treated mice were counted
 - sections of dorsal hippocampus was analyzed by stereoinvestigator software
 - immunopositive microglia identified with 20x objective and total number estimated by optical fractionator method

Material and Methods

- B) Fluoro-Jade B:
- Brain sections of 6 TBI treated and 6 TBI vehicle treated mice were counted
- dorsal hippocampus was examined for degenerating neurons using epifluorescence microscope

- **Testing for functional deficits**
 - Automated Rotarod to assess vestibulomotor function
 - one day before TBI: clinical training trial at accelerating rotational speed for 200 seconds and then three additional test trials (n=11-12 per group)
 - → average time to fall from cylinder was recorded as baseline latency
 - Mice were tested on consecutive days post-injury and received three consecutive daily trials with accelerating rotational speed

Material and Methods

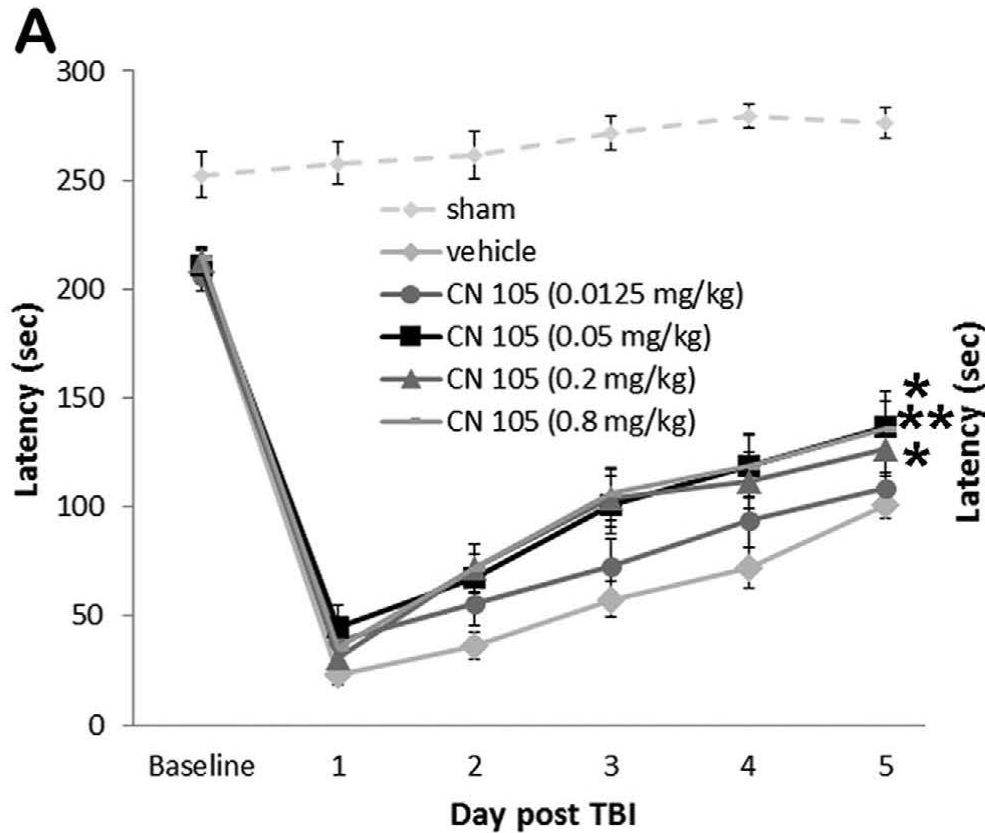
- Morris Water Maze to assess spatial learning and memory
- submerged platform placed in a pool with 105cm diameter, opaque water
- Four trials /day for 4 consecutive days: days 28-31 after TBI (n=11-12 per group)
- Mice were introduced to varying quadrants of the pool
- Probe trial on day 4 (=last day) of the experiment: platform was removed and mice were allowed to swim freely for 60 sec → percent of time the mice spent in the platform quadrant was quantified
- **RNA extraction and RT-PCR**
 - brain tissue was processed for RNA extraction from a separate cohort of treated and untreated mice on day 1 post injury (CN-105, n=4; vehicle, n=3; control (sham-operated), n=3)
 - gene expression was measured using the Mouse Inflammatory Response and Autoimmunity PCR Array (profiles the expression of 84 inflammation and autoimmunity related genes)

Material and Methods

- **Gene expression data analysis:**
 - inflammatory and autoimmunity related genes (84 genes)
- **Pharmacokinetic analysis:**
 - to assess CNS penetration: quantitative whole body autoradiography analysis (QWBA) was performed and areas of interest in blood and brain were compared

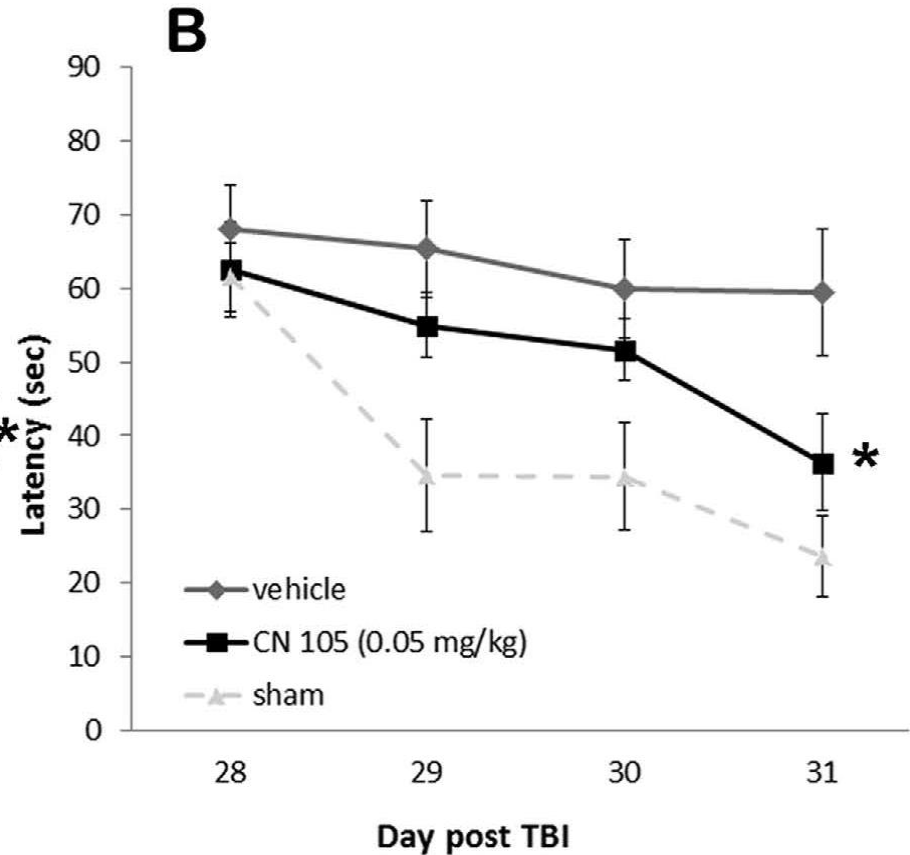
Results

Assessment by Rotarod (CN-105: 2h post TBI)



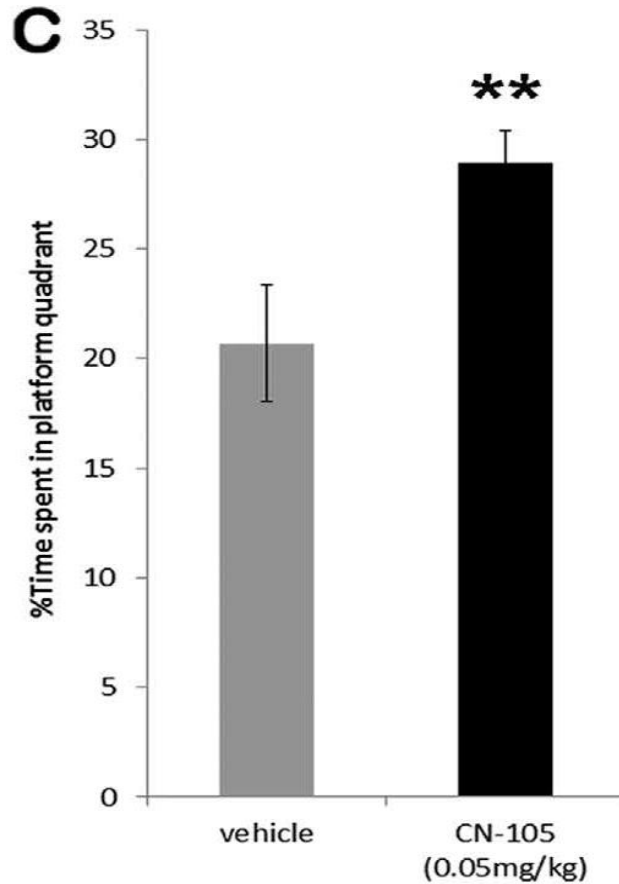
* $p < 0.05$ and ** $p < 0.01$

Assessment by MWM (CN-105: 2h post TBI)



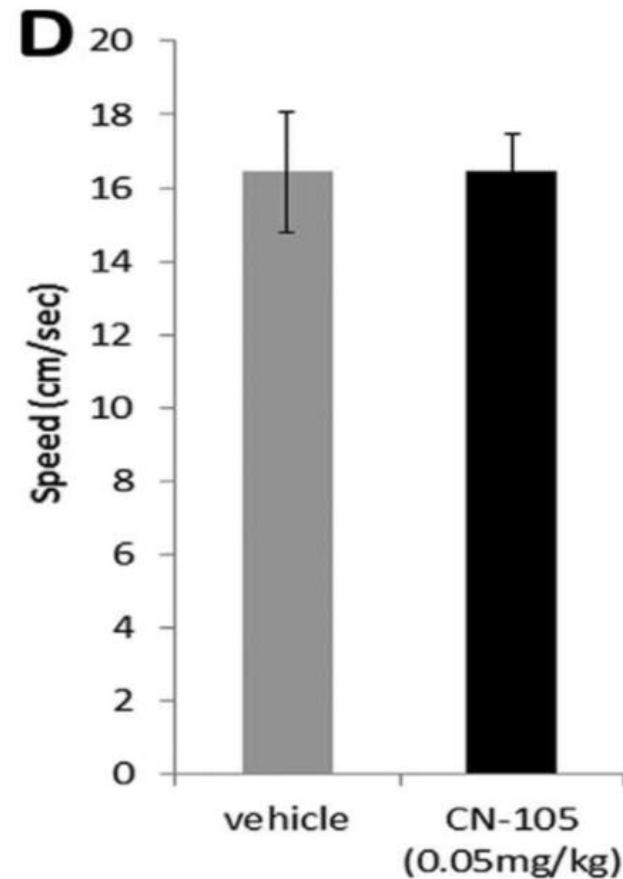
Results

Probe trial on day 4
of MWM experiment



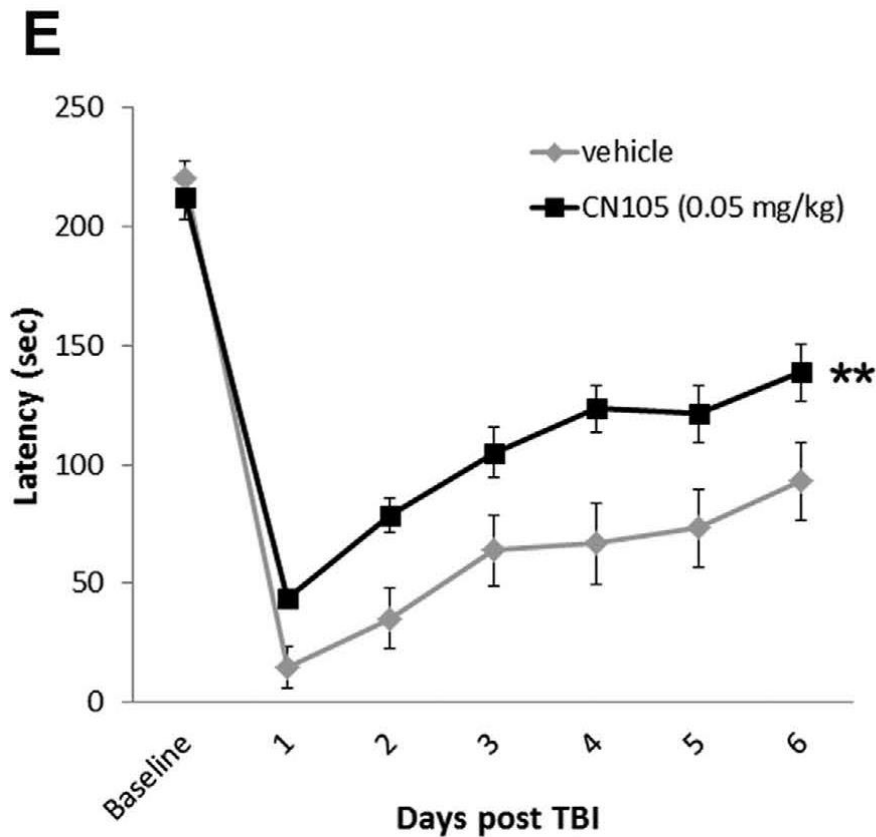
* $p < 0.05$ and ** $p < 0.01$

Swim speed during
whole MWM experiment



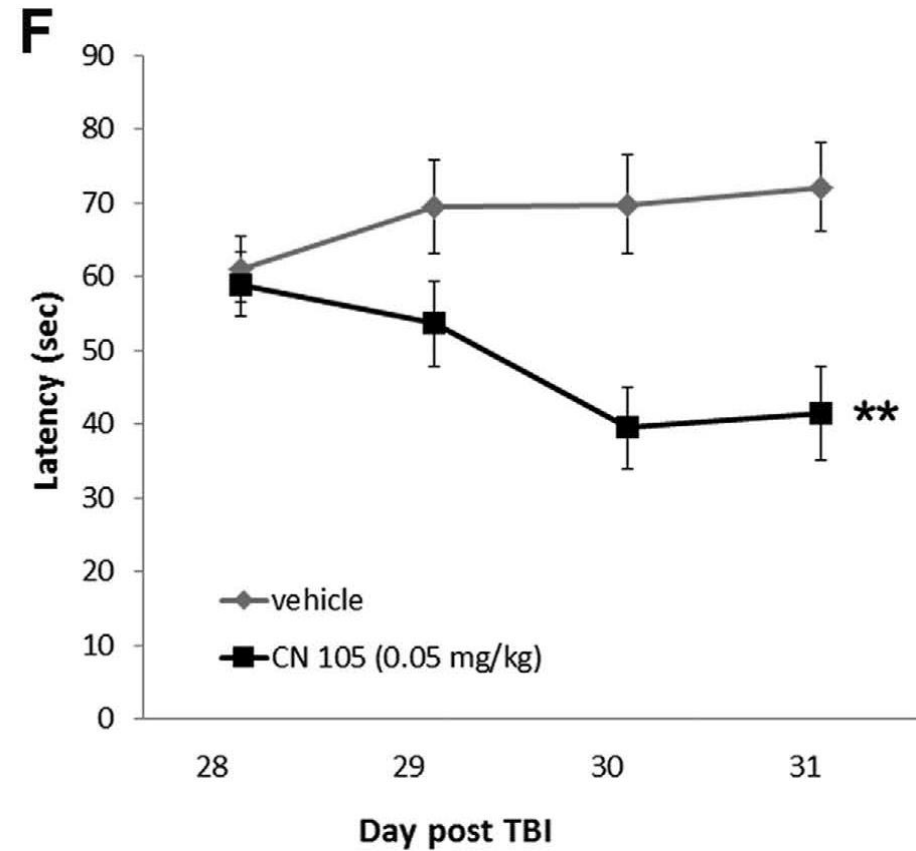
Results

Assessment by Rotarod (CN-105: 4h post TBI)



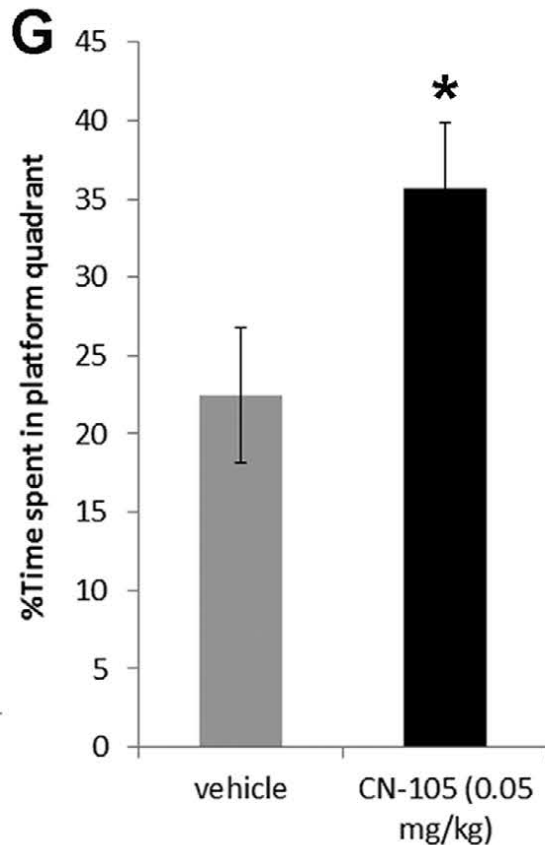
* $p < 0.05$ and ** $p < 0.01$

Assessment by MWM (CN-105: 4h post TBI)

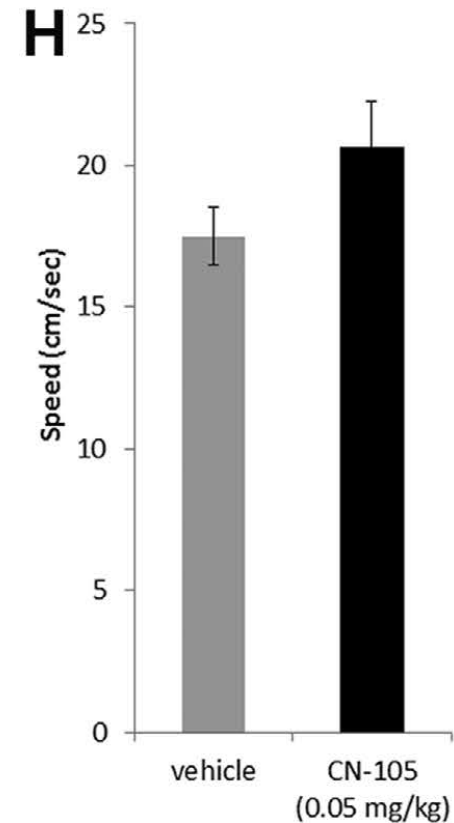


Results

Probe trial on day 4 of MWM experiment
(CN-105: 4h post TBI)



Swim speed during whole MWM experiment
(CN-105: 4h post TBI)



* $p < 0.05$ and ** $p < 0.01$

Results

Comparison of activated F4/80
immunostained microglia in hippocampus
(10 days post TBI)

A, C, E, G → treated by vehicle

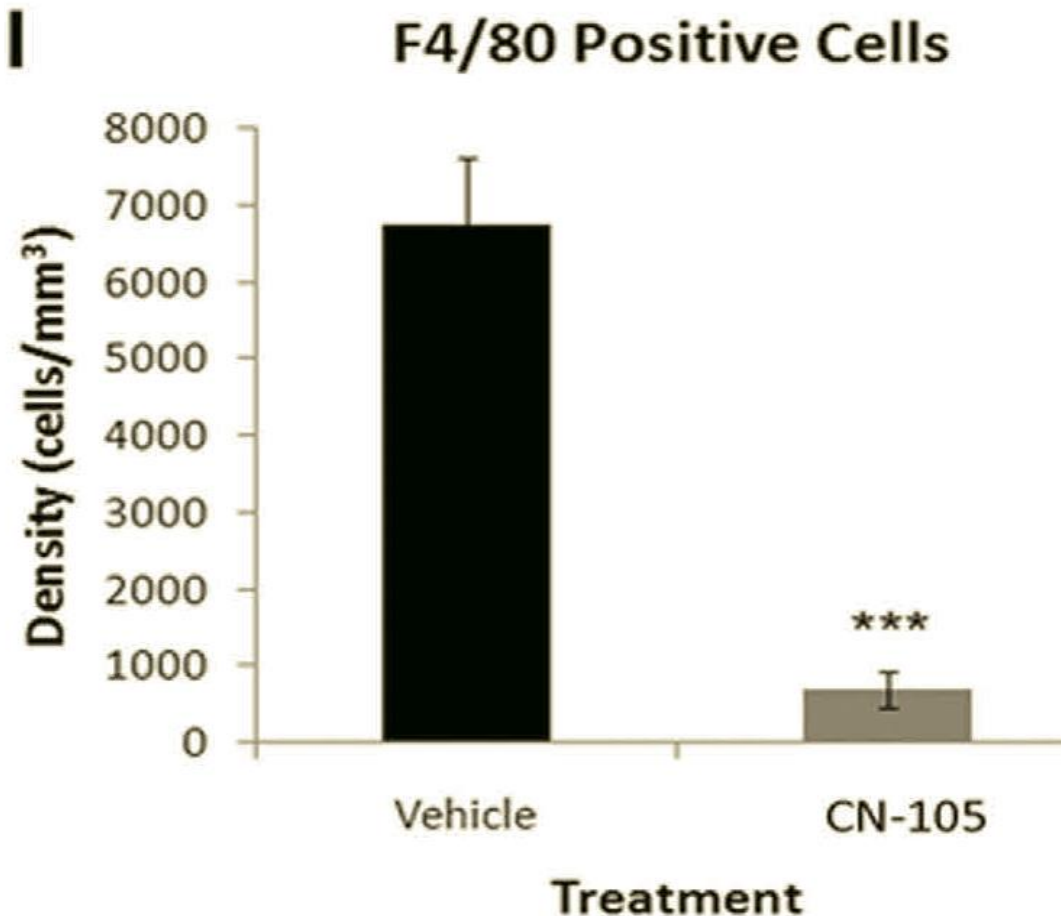
B, D, F, H → treated by CN-105

C+D: CA3 region (cornu ammonis
mit Pyramidenzellen)

E+F: polymorphic region (with
marked microglia cells)

G+H: periventricular region
(corpus callosum and fimbria)

Results



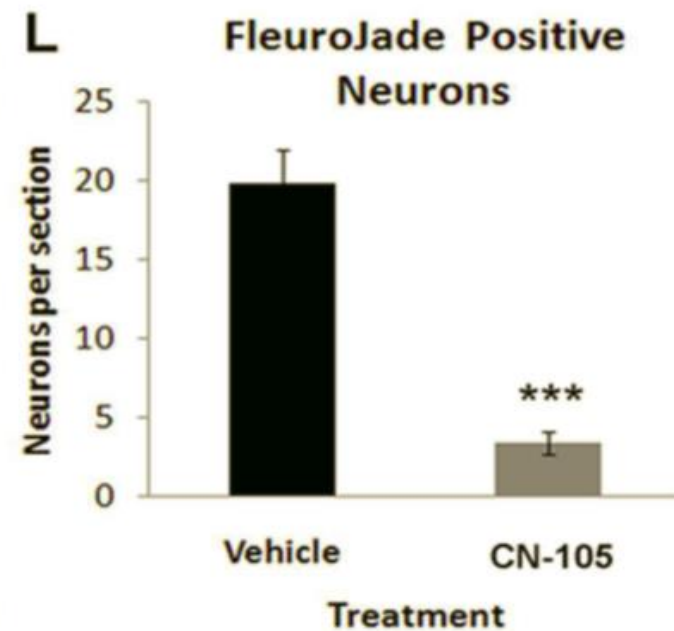
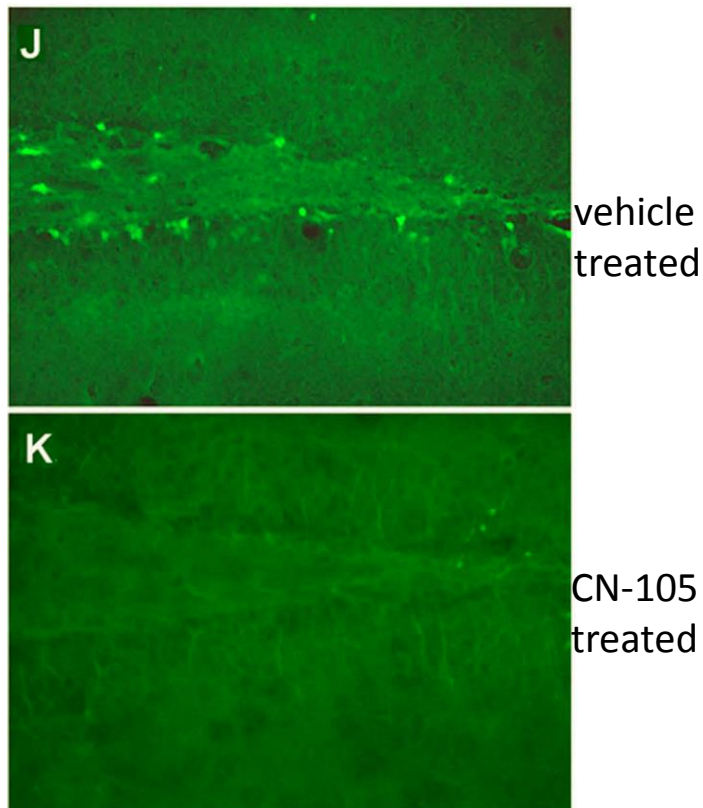
**Comparison of activated
F4/80 immunostained
microglia in hippocampus
(10 days post TBI)**

→ significant reduction
in microgliosis

*** $p=0.0002$

Results

**Fluoro-Jade B stained brain slices 24h post-TBI:
degenerating neurons in dorsal dentate gyrus**



→ significant reduction
of neuron degeneration

Gene	S-v	S-CN-105	TBI-v	TBI-CN-105	S-CN-105/S-v	TBI-v/S-v	TBI-CN-105/S-v
Cxcl3	3.0E-05	2.3E-05	6.7E-01	1.6E+00	0.8	22171.7	54032.0
Ccl11	7.8E-05	2.5E-04	6.4E-01	1.8E-01	3.2	8199.7	2253.1
Ccr4	3.9E-06	6.0E-06	2.6E-02	3.1E-02	1.5	6537.5	7945.9
Ccl20	2.6E-05	1.1E-04	1.2E-01	7.9E-02	4.4	4574.0	3086.8
Tlr6	4.7E-04	6.2E-04	1.2E+00	7.0E-03	1.3	2528.8	14.9
Ccl24	3.2E-05	9.7E-05	7.6E-02	5.1E-03	3.0	2356.9	159.1
Myd88	7.8E-04	1.1E-03	1.8E+00	6.6E-03	1.5	2327.8	8.5
Cxcl9	6.8E-05	1.0E-04	1.5E-01	2.2E-03	1.5	2265.1	32.2
Lta	1.2E-04	2.6E-04	1.3E-01	1.1E-01	2.3	1133.5	917.0
Tnfsf14	2.5E-05	1.3E-04	2.5E-02	9.0E-03	5.2	1027.7	365.9
Il9	1.2E-05	*	1.2E-02	1.4E-02	**	957.7	1155.5
Ifng	7.3E-05	5.8E-05	5.6E-02	4.3E-03	0.8	767.1	59.7
Il10	1.9E-05	1.1E-04	1.5E-02	1.6E-03	5.7	766.1	82.2
Il1b	1.1E-03	2.0E-03	7.5E-01	4.6E-01	1.8	683.8	416.6
Il8rb	8.3E-06	6.1E-05	5.2E-03	5.4E-03	7.3	623.4	658.4
Il6	1.4E-04	1.1E-04	5.4E-02	1.8E-03	-1.3	395.1	13.2
Il1a	1.4E-03	1.9E-03	5.5E-01	5.4E-02	1.4	382.3	37.7
Cd40	5.4E-04	8.3E-04	1.6E-01	2.1E-02	1.5	301.4	38.8
Ccr2	2.6E-04	5.5E-04	7.1E-02	1.5E-03	2.1	271.6	5.6
Cxcl1	6.0E-05	1.6E-04	1.5E-02	1.8E-03	2.7	244.8	29.4
Ccr7	1.2E-04	2.1E-04	2.7E-02	3.4E-02	1.8	228.3	283.2
Il18rap	1.1E-04	4.5E-05	2.4E-02	6.6E-03	0.4	211.3	58.6
Tnf	2.8E-05	3.5E-05	5.5E-03	3.8E-02	1.3	197.3	1367.7
C4b	3.6E-03	7.0E-03	6.3E-01	5.9E-01	2.0	176.5	163.3
Il8ra	1.3E-05	*	2.2E-03	2.6E-03	**	168.5	196.8
Cxcr4	4.3E-03	6.6E-03	6.6E-01	9.0E-02	1.5	153.8	21.0
Cxcl11	2.4E-05	9.0E-05	3.4E-03	8.8E-03	3.8	141.3	372.3
Il6ra	3.4E-03	4.9E-03	4.8E-01	1.6E-01	1.5	140.3	47.9
Il22	4.3E-05	4.7E-04	5.6E-03	3.4E-03	10.8	129.4	77.8
Il1rn	2.3E-05	*	2.8E-03	1.7E-03	**	121.0	74.2
Ccl2	5.8E-04	8.3E-04	6.5E-02	2.5E-02	1.4	110.8	42.3
C3ar1	3.5E-03	4.1E-03	3.7E-01	1.2E-01	1.2	108.2	34.9

Assessment of gene expression

(24h post sham or TBI injury)

→ using a pathway array specific for inflammatory and immune responses (84 genes)

4 cohorts:

- Sham + vehicle (S-v)
- Sham + CN-105 (S-CN-105)
- TBI + vehicle (TBI-v)
- TBI + CN-105 (TBI-CN-105)

→ 57 genes were upregulated, 12 were downregulated, 10 were unchanged

*mRNA expression was not detectable

**relative values could not be calculated because one of the components was not detectable

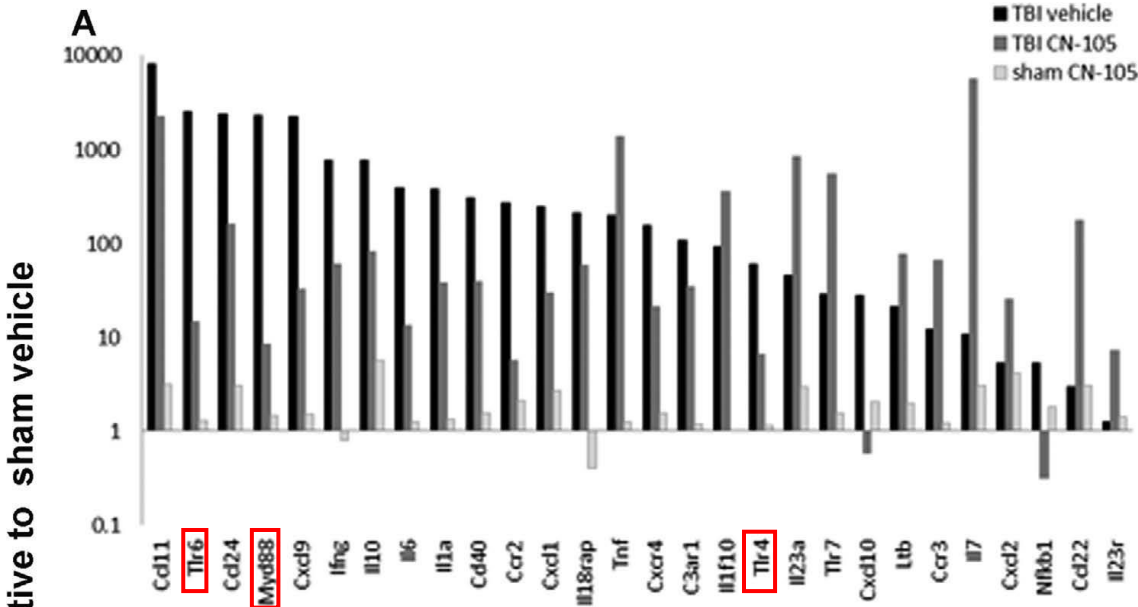
Gene	Fold Change	Gene	Fold Change	Gene	Fold Change
<i>Il22ra2</i>	10895	<i>Tlr3</i>	1.3	<i>Fasl</i>	-2.0
<i>Nfatc3</i>	10044	<i>Ccr7</i>	1.2	<i>Csf1</i>	-2.0
<i>Cxcl5</i>	6257	<i>Ccr4</i>	1.2	<i>Il1rap</i>	-2.2
<i>Bcl6</i>	921	<i>Il9</i>	1.2	<i>Ccl4</i>	-2.6
<i>Il7</i>	515	<i>Crp</i>	1.2	<i>Ccl2</i>	-2.6
<i>Cd12</i>	64.2	<i>Il8ra</i>	1.2	<i>Tnfsf14</i>	-2.8
<i>Ly96</i>	61.2	<i>Il8rb</i>	1.1	<i>Tollip</i>	-2.9
<i>Cd22</i>	60.2	<i>Tlr2</i>	1.0	<i>Il6ra</i>	-2.9
<i>Cd8</i>	31.2	<i>Rtpk2</i>	-1.1	<i>C3ar1</i>	-3.1
<i>Tlr7</i>	19.0	<i>Kng1</i>	-1.1	<i>Il18rap</i>	-3.6
<i>Il23a</i>	18.5	<i>Hdac4</i>	-1.1	<i>Cd11</i>	-3.6
<i>Nr3c1</i>	13.7	<i>C4b</i>	-1.1	<i>Cd19</i>	-5.2
<i>Il18</i>	7.2	<i>Fos</i>	-1.2	<i>Cxcr4</i>	-7.3
<i>Tnf</i>	6.9	<i>Cd40lg</i>	-1.2	<i>Cd40</i>	-7.8
<i>Il23r</i>	5.8	<i>Ccl7</i>	-1.2	<i>Cxcl1</i>	-8.3
<i>Ccr3</i>	5.4	<i>Cd25</i>	-1.2	<i>Tlr4</i>	-9.0
<i>Cxcl2</i>	4.8	<i>Lta</i>	-1.2	<i>Il10</i>	-9.3
<i>Il1f10</i>	3.8	<i>Flt3l</i>	-1.4	<i>Il1a</i>	-10.1
<i>Ltb</i>	3.6	<i>Tlr1</i>	-1.4	<i>Ifng</i>	-12.9
<i>Il1r1</i>	3.5	<i>Cd20</i>	-1.5	<i>Ccl24</i>	-14.8
<i>Cxcl11</i>	2.6	<i>Cebpb</i>	-1.5	<i>Nfkb1</i>	-16.6
<i>Il10rb</i>	2.6	<i>Tirap</i>	-1.5	<i>Il6</i>	-29.8
<i>Tlr5</i>	2.5	<i>Itgb2</i>	-1.5	<i>Cxcl10</i>	-47.9
<i>Cxcl3</i>	2.4	<i>Il1rn</i>	-1.6	<i>Ccr2</i>	-48.3
<i>Ccr1</i>	2.3	<i>Ccl5</i>	-1.6	<i>Cxcl9</i>	-70.3
<i>C3</i>	1.5	<i>Il1b</i>	-1.6	<i>Tlr6</i>	-170
<i>Ccl3</i>	1.5	<i>Il22</i>	-1.7	<i>Myd88</i>	-273
<i>Ccl1</i>	1.5	<i>Ccl17</i>	-1.7	<i>Nos2</i>	*

Assessment of gene expression

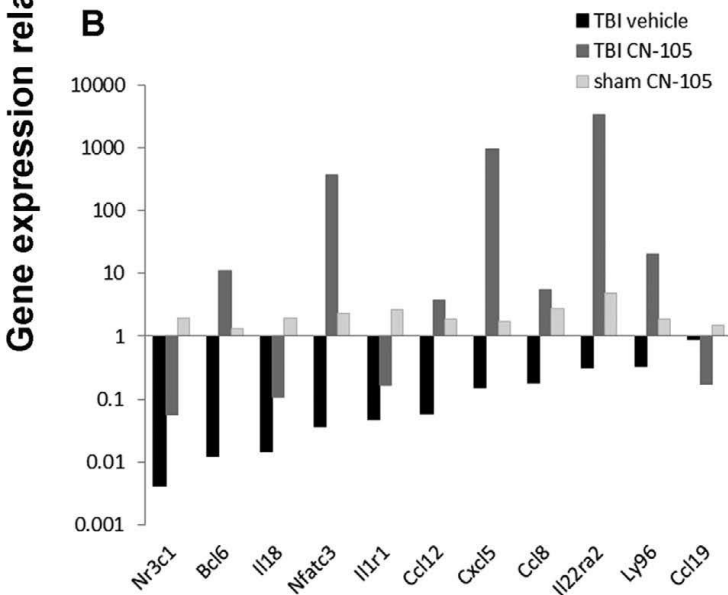
(24h post TBI with CN-105)

→ inflammatory gene expression of CN-105 relative to vehicle treated group

*mRNA expression was not detectable



CN-105 ameliorates changes in inflammatory gene expression (24h post TBI)

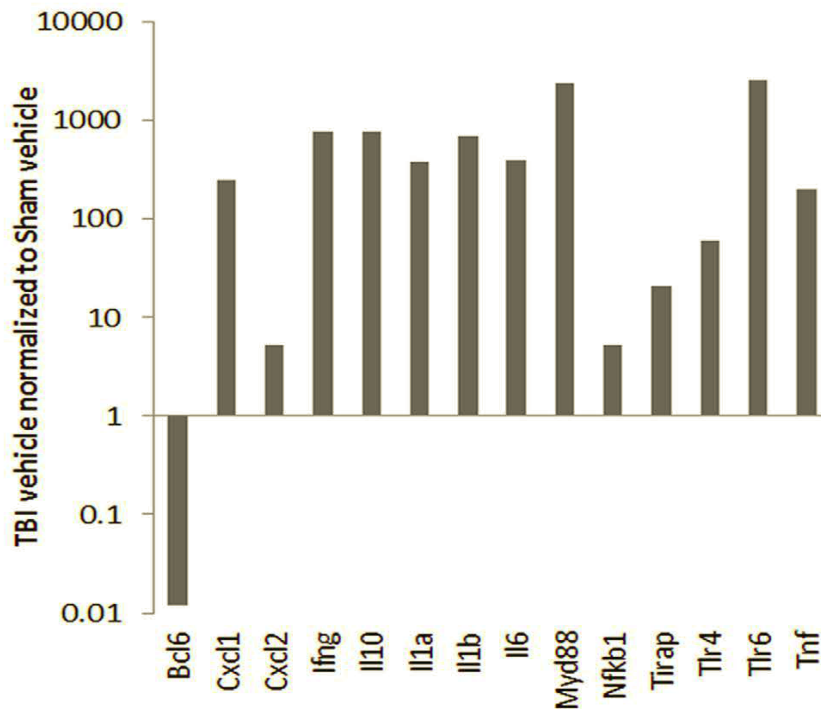


→ upregulated/downregulated expression of genes relative to sham vehicle

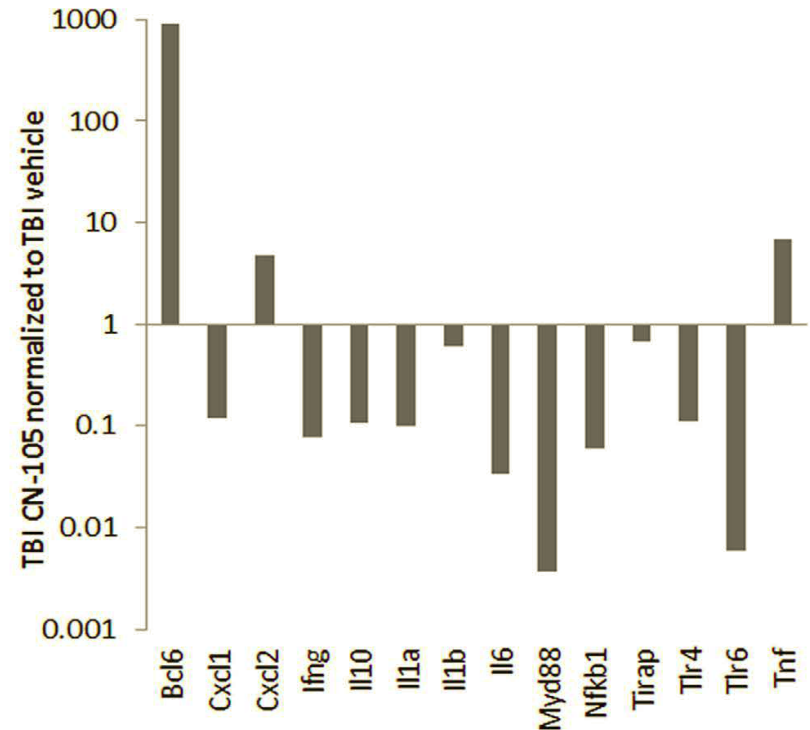
Results

**TLR signalling is
downregulated by CN-105
(24h post TBI)**

B

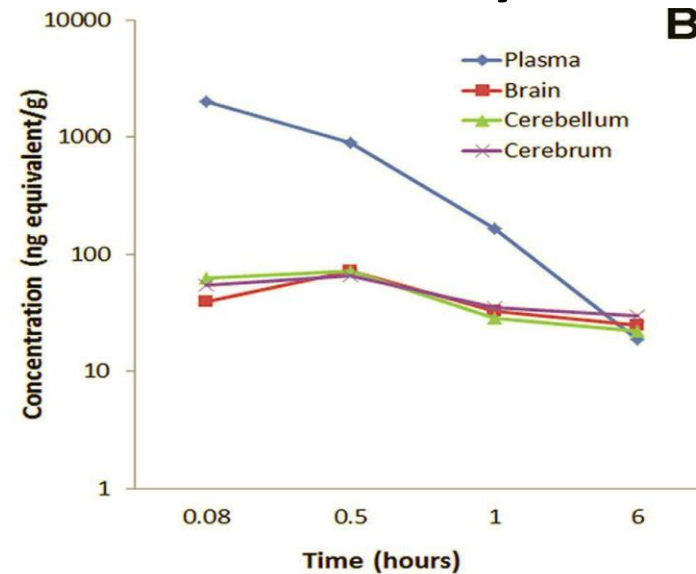
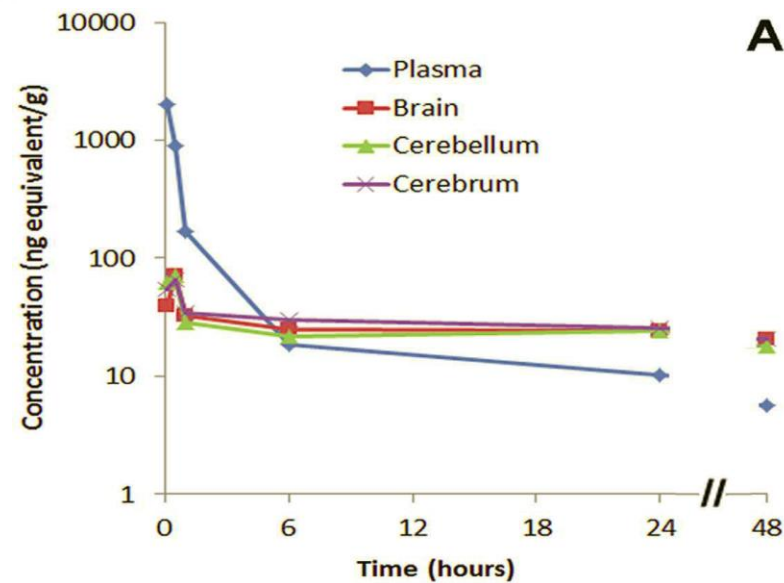


D



Results

Pharmacokinetic studies of CN-105 demonstrate CNS bioavailability



concentration of C14-radioactivity in plasma and CNS following intravenous dose of radiolabeled CN-105 peptide

	Radioactivity (ng equiv/g) in tissue following bolus (hours = h)					
	0.08h	0.5h	1h	6h	24h	48h
<i>Blood (Cardiac)</i>	1102	500	95	15	14.1	15.5
<i>Brain</i>	39.5	71.2	32.5	24.9	24.1	20.9
<i>% Brain/Blood</i>	3.58%	14%	34%	167%	170%	135%

Percent of radioactivity contributed to blood in the brain

Results

CN-105 is associated with:

- A) improved behavioral function
 - by Rotarod (better vestibulomotor performance)
 - by MWM (better preserved spatial learning memory)
 - no significant motor differences
 - pretreatment of mice 30min prior to TBI had no additional effects („data not shown“)
- B) reduced microgliosis and neuronal injury following TBI
 - esp. in CA3 und polymorphic region of hippocampus
 - degenerating neurons are significantly reduced
- C) changes in inflammatory gene expression patterns following TBI
 - 57 of 84 inflammatory genes were upregulated in TBI mice (esp. TLR-signalling pathway)
 - TBI-CN-105 lead to reduced inflammatory gene upregulation compared to TBI-v
- D) penetration into the CNS compartment
 - progressive increase of radioactivity in brain as compared to blood (3,6% at 5 minutes, 170% at 24h)

Discussion

- ApoE and ApoE-mimetic peptides decrease neuroinflammatory responses and secondary cell death
 - Already shown in several animal studies of acute brain injury
- mRNA levels of inflammatory cytokines and chemokines return to pre-injury levels by 24 hours or more?
- Difficulty to characterize endogenous microglia
 - Primitive macrophage entering embryonic brain vs. hematogenous macrophage that are recruited into brain following to injury
 - But number and activation status is decreased after CN-105 treatment
- Not all indicators are decreased by CN-105 treatment at 24 hours post injury
 - TNF alpha increases

Discussion

- MyD88 has recently shown to be significantly increased after TBI in several mouse model experiments
- Mechanisms by which CN-105 (ApoE) modulate inflammatory response is completely undefined
 - Via specific receptor interaction
 - Connection to LRP-1 receptor? NMDA-receptor via PSD-95?
- Gene expression suggests key role of BCL6 in inflammation process
 - Early changes of NF- κ B, chemokines and cytokines
 - Central key repressor in TLR signalling pathway?
- CN-105 may directly effect the blood brain barrier (tight junctions)

Discussion

- Administered up to 4 hours following injury
→ Temporal window should be further extended
 - Limitations of study:
 - ApoE polymorphism → modulate receptor binding via allosteric effects
 - Differential gene assay only focused on expression of inflammatory markers
 - Rodent models:
 - Brains have reduced ration of white: grey matter
→ Not ideal to model the diffuse axonal injury
 - Early changes of NF-kB, chemokines and cytokines
 - Central key repressor in TLR signalling pathway?
 - CN-105 may directly effect the blood brain barrier (tight junctions)
- promising therapeutic strategy in treatment of acute brain injury

Thank you for your attention!
Any questions?