

Real-world clinical outcomes of intraventricular cerliponase alfa in CLN2 disease: Comparison with a historical cohort

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Conflict of interest disclosure

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CLN2 disease: A rare and rapidly progressive childhood neurodegenerative disorder



CLN2 disease is a rare, genetic, neurodegenerative disorder of childhood caused by deficient tripeptidyl peptidase 1 (TPP1) enzyme activity^{1–3}



Deficient TPP1 activity results in the intralysosomal accumulation of autofluorescent storage material and subsequent neuronal and retinal cell loss^{1,2}



Symptom onset of classic late-infantile phenotype is typically at 2–4 years of age, presenting with unprovoked seizures and history of language delay followed by psychomotor decline and vision loss^{1–3}

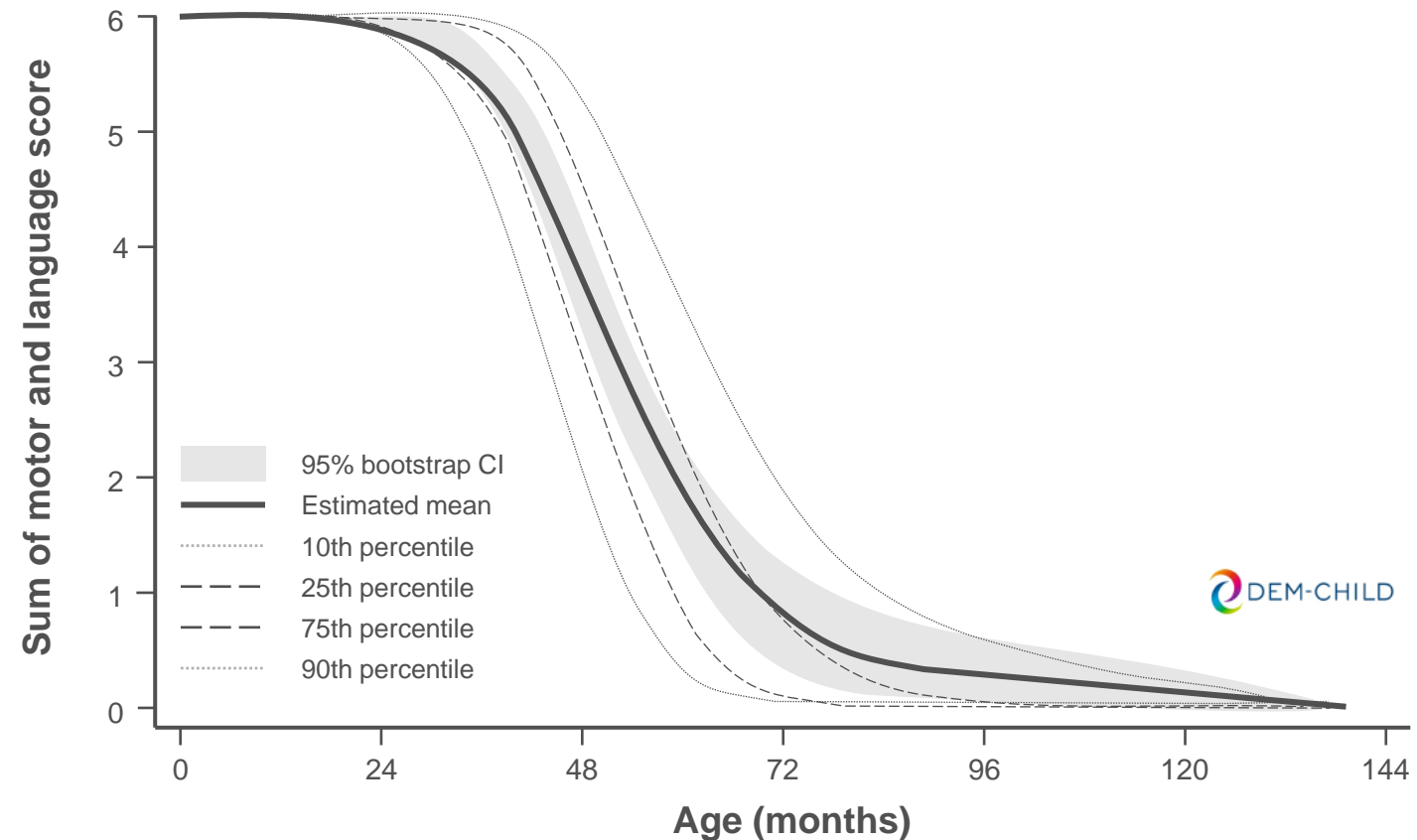


Neurodegeneration is rapidly progressive, leading to early death^{1–3}

CLN2 disease is associated with rapid decline in motor and language function

DEM-CHILD database, natural history cohort (2002–2016) (N = 41)

Motor and language domains are each scored from 3 to 0, with lower scores indicating more extensive impairment of function



DEM-CHILD

Cerliponase alfa

- Cerliponase alfa is a recombinant human form of TPP1 enzyme and was approved for the treatment of CLN2 disease in the US and EU in 2017^{1–4}
 - In clinical trials, intracerebroventricular administration of cerliponase alfa was shown to slow decline in motor and language function compared to natural history^{1,2}

Study objective:

To assess the real-world effectiveness and safety of cerliponase alfa in children with CLN2 disease who initiated treatment outside of the clinical trial setting using data from the independent DEM-CHILD database (international NCL patient registry)

Analysis methods

Patients

- Enrolled in the DEM-CHILD database (international NCL patient registry) (NCT04613089) with confirmed diagnosis of CLN2 disease by genetic and enzyme testing

Treated patients: initiated cerliponase alfa outside of clinical trials and had at least 6 months of follow-up on treatment

Natural history patients: never received cerliponase alfa

Matching

- Treated patients were matched 1:1 with natural history patients on:
 - Baseline^a age (± 12 months) AND
 - Baseline score on motor and language (ML) domains (exact match)

Follow-up

- Data were considered from baseline until end of follow-up or data cut-off (December 30, 2020) for treated patients, and for period equivalent to that of the matched treated patient for natural history patients

Outcomes of interest

- Rate of decline in ML score
- Time to unreversed 2-point decline or ML score of 0
- Treatment-related adverse events

^aBaseline was defined as the last observation prior to the first dose (treated patients), or assessment at the age of matching (natural history patients).

CLN2, neuronal ceroid lipofuscinosis type 2; DEM-CHILD, A Treatment-Oriented Research Project of NCL Diseases as a Major Cause of Dementia in Childhood;

ML, motor-language; NCL, neuronal ceroid lipofuscinosis.

Patient characteristics

Patients matched on baseline age and ML score

		Natural history N = 21	Cerliponase alfa-treated N = 21
Sex	n (%)		
Female		5 (24)	11 (52)
Male		16 (76)	10 (48)
Age at baseline, years	Mean (SD) Median [min, max]	4.7 (1.9) 4.5 [0.7, 9.9]	4.7 (1.9) 4.5 [0.7, 9.9]
Baseline ML score	Mean (SD) Median [min, max]	3.9 (1.6) 4 [1, 6]	3.9 (1.6) 4 [1, 6]
Baseline ML score category	n (%)		
1		1 (5)	1 (5)
2		4 (19)	4 (19)
3		3 (14)	3 (14)
4		6 (29)	6 (29)
5		2 (10)	2 (10)
6		5 (24)	5 (24)
Genotype	n (%)		
2 common alleles ^a		13 (62)	14 (67)
1 common allele		5 (24)	4 (19)
0 common alleles		3 (14)	3 (14)
Age at disease onset, years	Mean (SD) Median [min, max]	3.0 (0.8) 3.0 [1.3, 4.4]	n = 20 3.4 (0.8) 3.3 [2.1, 6.0]
Age at diagnosis, years	Mean (SD) Median [min, max]	n = 17 5.2 (1.6) 4.8 [2.9, 9.8]	4.2 (2.0) 3.9 [0.2, 8.8]

^aThe common alleles are c.622C>T and c.509-1G>C.

max, maximum; min, minimum; ML, motor-language; n, number of valid observations; N, number of patients; SD, standard deviation.

Rate of decline in ML score

Patients matched on baseline age and ML score

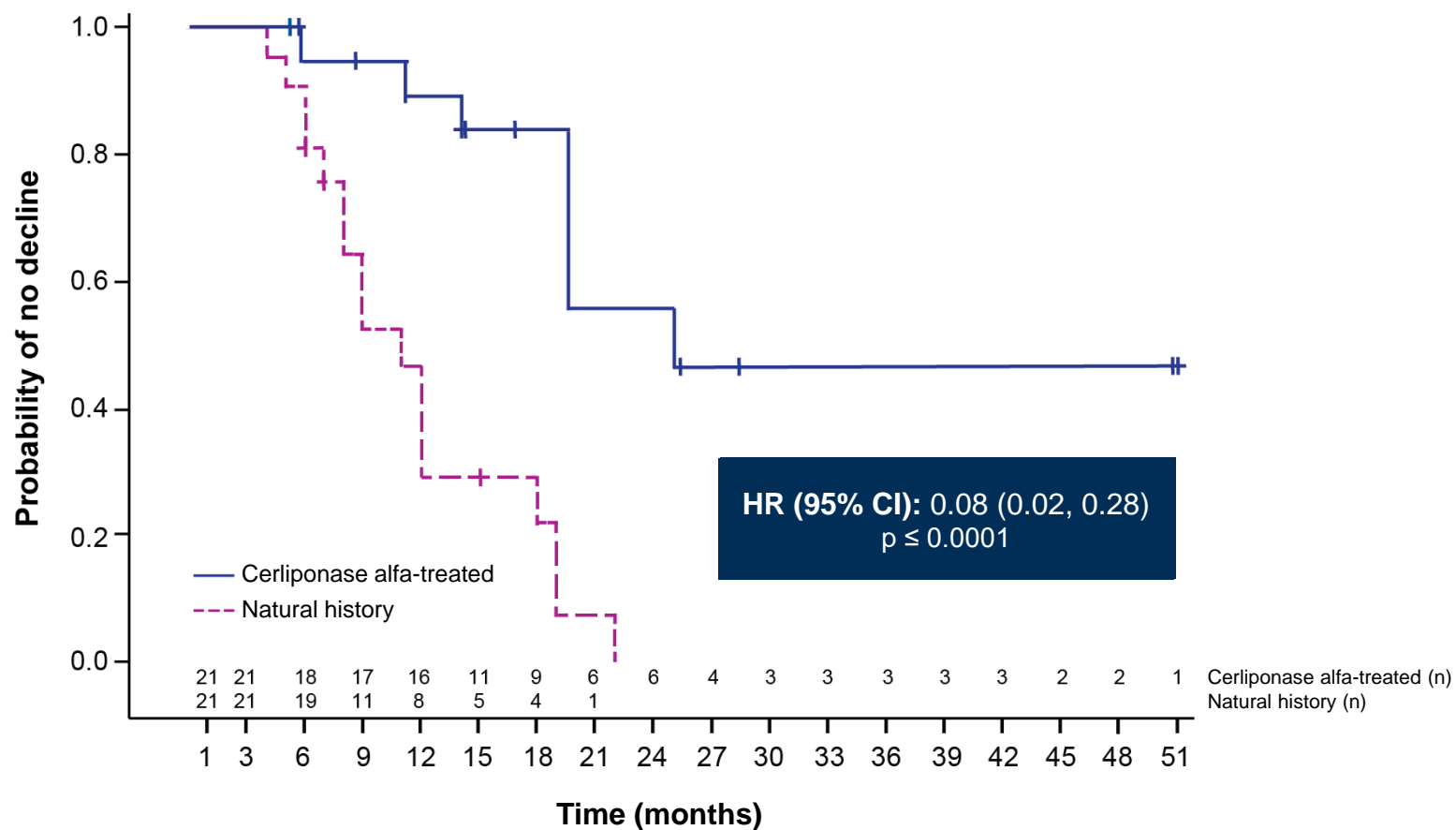
	Natural history N = 21	Cerliponase alfa-treated N = 21
Rate of decline in ML score, points per 48 weeks		
Mean (SD)	1.88 (1.45)	0.46 (0.43)
Median [min, max]	1.66 [0.00, 5.60]	0.44 [0.00, 1.33]
95% CI	1.22, 2.54	0.26, 0.64

Mean (SE) difference: 1.42 (0.33)
95% CI: 0.74, 2.10; p = 0.0003

- At the end of follow-up, cerliponase alfa-treated patients had:
 - Higher ML score than matched natural history controls in 15/21 pairs (71%)
 - Equivalent ML score to matched natural history controls in 6/21 pairs (29%)

Time to unreversed 2-point decline or score 0 in ML domains

Patients matched on baseline age and ML score



Treatment-related adverse events

All cerliponase alfa-treated patients (N = 24)

	All treatment-related AEs			Treatment-related AEs resulting in hospitalization		
	N = 24			N = 24		
	Incidence n (%)	Events	Rate (events/100 PY)	Incidence n (%)	Events	Rate (events/100 PY)
Any treatment-related AE	16 (66.7)	75	13.4	7 (29.2)	19	3.4
Treatment-related AEs						
Pyrexia	12 (50.0)	43	7.7	4 (16.7)	5	0.9
Vomiting	8 (33.3)	10	1.8	5 (20.8)	6	1.1
Nausea	5 (20.8)	5	0.9	2 (8.3)	2	0.4
Arrhythmia	3 (12.5)	3	0.5	0	0	0
Device-related infection	3 (12.5)	3	0.5	3 (12.5)	3	0.5
Seizure	3 (12.5)	6	1.1	0	0	0
Device leakage	2 (8.3)	2	0.4	2 (8.3)	2	0.4
Flushing	1 (4.2)	2	0.4	0	0	0
Headache	1 (4.2)	1	0.2	1 (4.2)	1	0.2

There were no deaths in the cerliponase alfa-treated group; 6 (29%) patients in the natural history control group died during the equivalent length of follow-up (median time to death: 313 weeks)

Conclusions

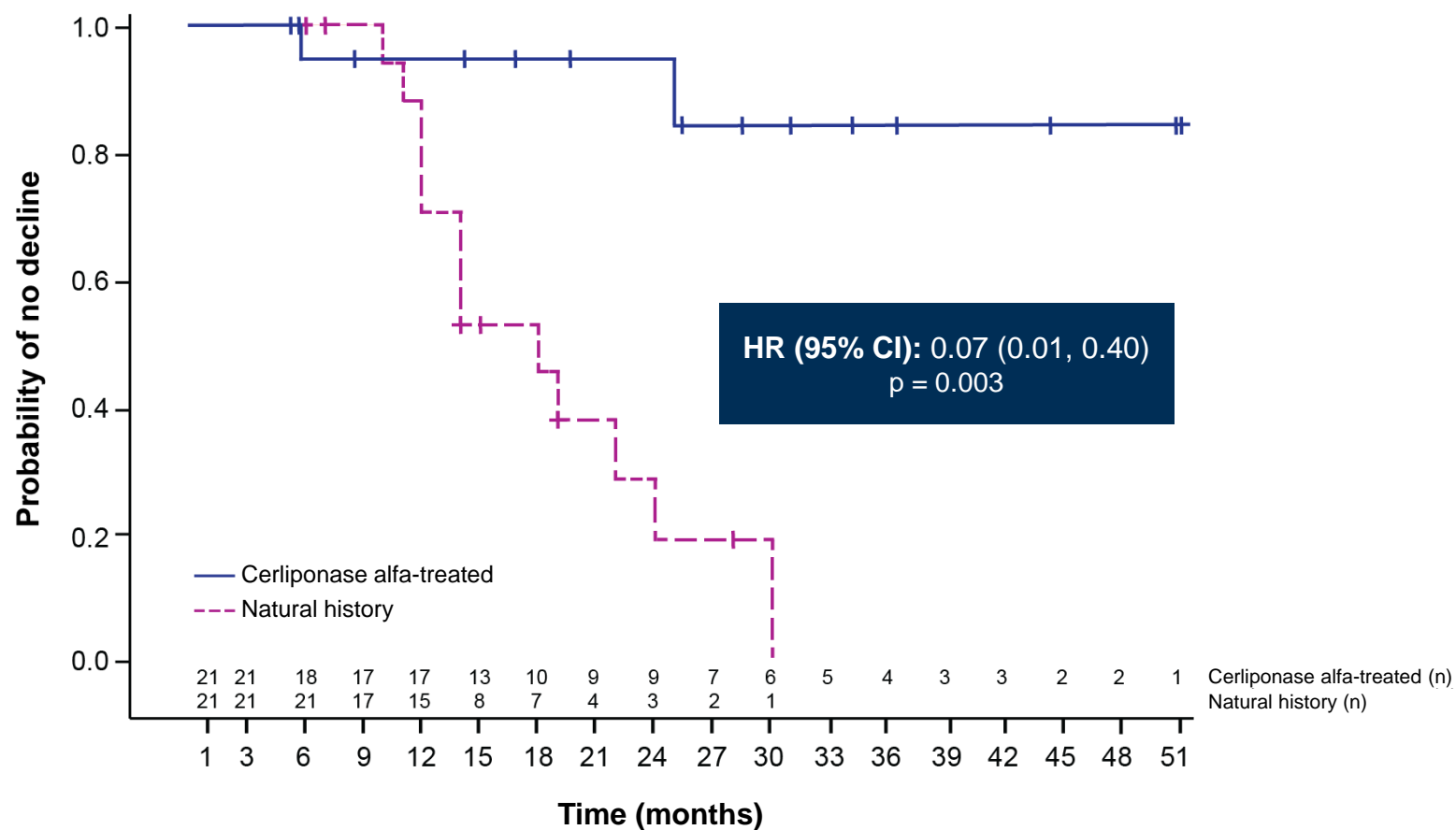
- Cerliponase alfa treatment slowed deterioration in motor and language function in children with CLN2 disease in a real-world setting
 - These results from the real-world setting support efficacy findings from cerliponase alfa clinical trials
- No new safety signals were identified
 - Treatment-related AEs observed were consistent with those seen in clinical trials
- Study limitations:
 - Historical controls
 - Only AEs suspected by the treating physician to be treatment (drug or device)-related were considered

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Time to unreversed score of 0 in ML domains

Patients matched on baseline age and ML score



Rate of decline in motor score

Patients matched on baseline age and ML score

	Natural history N = 21	Cerliponase alfa-treated N = 21
Rate of decline in motor score, points per 48 weeks		
	n = 20	n = 20
Mean (SD)	0.99 (0.75)	0.23 (0.28)
Median [min, max]	1.02 [0.00, 2.40]	0.11 [0.00, 0.79]
95% CI	0.64, 1.34	0.10, 0.36

Mean (SE) difference: 0.75 (0.18)
95% CI: 0.38, 1.12; p = 0.0003

Rate of decline in language score

Patients matched on baseline age and ML score

	Natural history N = 21	Cerliponase alfa-treated N = 21
Rate of decline in language score, points per 48 weeks		
	n = 19	n = 18
Mean (SD)	1.04 (0.96)	0.16 (0.28)
Median [min, max]	0.80 [0.00, 3.73]	0.00 [0.00, 0.72]
95% CI	0.58, 1.50	0.02, 0.30

Mean (SE) difference: 0.88 (0.23)
95% CI: 0.40, 1.36; p = 0.001

Sensitivity analysis: Rate of decline in ML score

Patients matched on baseline age, ML score, and genotype (3-criteria match)

	Natural history N = 18	Cerliponase alfa-treated N = 18
Rate of decline in ML score, points per 48 weeks		
Mean (SD)	2.07 (1.61)	0.49 (0.43)
Median [min, max]	2.18 [0.00, 5.60]	0.47 [0.00, 1.33]
95% CI	1.27, 2.87	0.28, 0.70

Mean (SE) difference: 1.58 (0.39)
95% CI: 0.76, 2.40; p = 0.0007