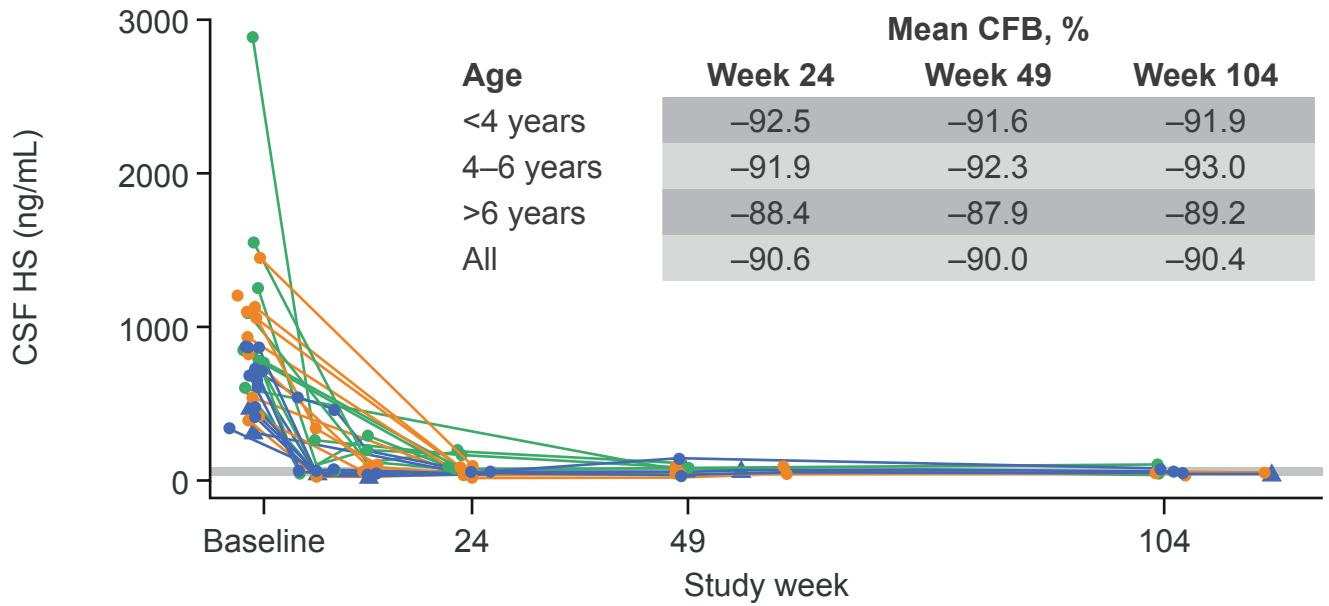


Supplemental Table 1. Baseline demographics and disease characteristics of participants

	Cohorts A–E (n=37)
Age	
Category, n (%)	
<4 years	10 (27.0)
4–6 years	12 (32.4)
>6 years	15 (40.5)
Median (min–max), years	5.1 (1.8–12.6)
MPS II phenotype, n (%)	
Neuronopathic	34 (91.9)
Non-neuronopathic	3 (8.1)
Variant type, n (%)	
Missense/synonymous	19 (51.4)
Large deletions/rearrangement	5 (13.5)
Other ^a	13 (35.1)
ERT status at enrollment	
SOC ERT-naive, ^b n (%)	10 (27.0)
SOC ERT-exposed, n (%)	27 (73.0)
Duration of SOC ERT, median (min–max), years ^c	2.2 (0.4–11.2)
Race, n (%)	
Asian	4 (10.8)
Black or African American	1 (2.7)
White	21 (56.8)
Not reported, unknown or other	11 (29.7)
Ethnicity, n (%)	
Hispanic or Latino	6 (16.2)
Not Hispanic or Latino	29 (78.4)
Not reported	2 (5.4)

Data are presented for the safety population. ^aIncludes nonsense and splice site/intron. ^bIncludes one participant who underwent allogeneic hematopoietic stem cell transplantation. ^cn=26. ERT, enzyme replacement therapy; max, maximum; min, minimum; MPS II, mucopolysaccharidosis type II; SOC, standard of care.

Supplemental Figure 1. CSF HS levels with tivenofusf alfa treatment^a



Age
 <4 years
 4–6 years
 >6 years
 All

Mean CFB, %		
Week 24	Week 49	Week 104
–92.5	–91.6	–91.9
–91.9	–92.3	–93.0
–88.4	–87.9	–89.2
–90.6	–90.0	–90.4

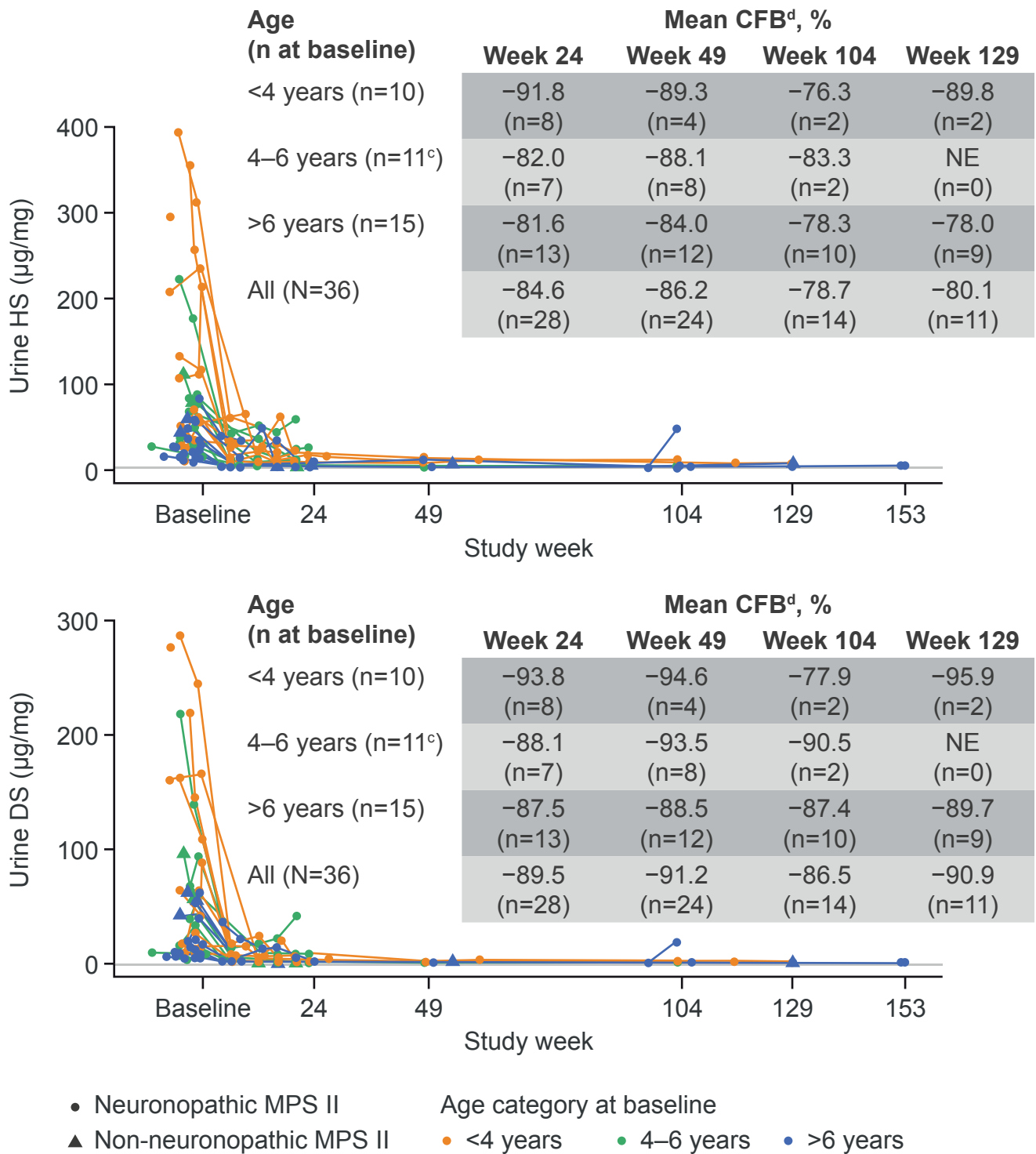
Number of participants, n

<4 years	10	9	6	3
4–6 years	11	9	8	3
>6 years	15	13	13	10
All	36	31	27	16

- Neuronopathic MPS II
 - ▲ Non-neuronopathic MPS II
- Age category at baseline
- <4 years
 - 4–6 years
 - >6 years

^aShaded gray area indicates the preliminary normal range of 39.1–92.5 ng/mL (10th to 90th percentiles) based on an analysis of CSF samples from 30 healthy adults (median [range] age, 52 [18–81] years). Total CSF GAG levels are similar in adults and children.¹ CFB, change from baseline; CSF, cerebrospinal fluid; GAG, glycosaminoglycan; HS, heparan sulfate; MPS II, mucopolysaccharidosis type II.

Supplemental Figure 2. Urine (A) HS^a and (B) DS^b levels with tividinofusp alfa treatment

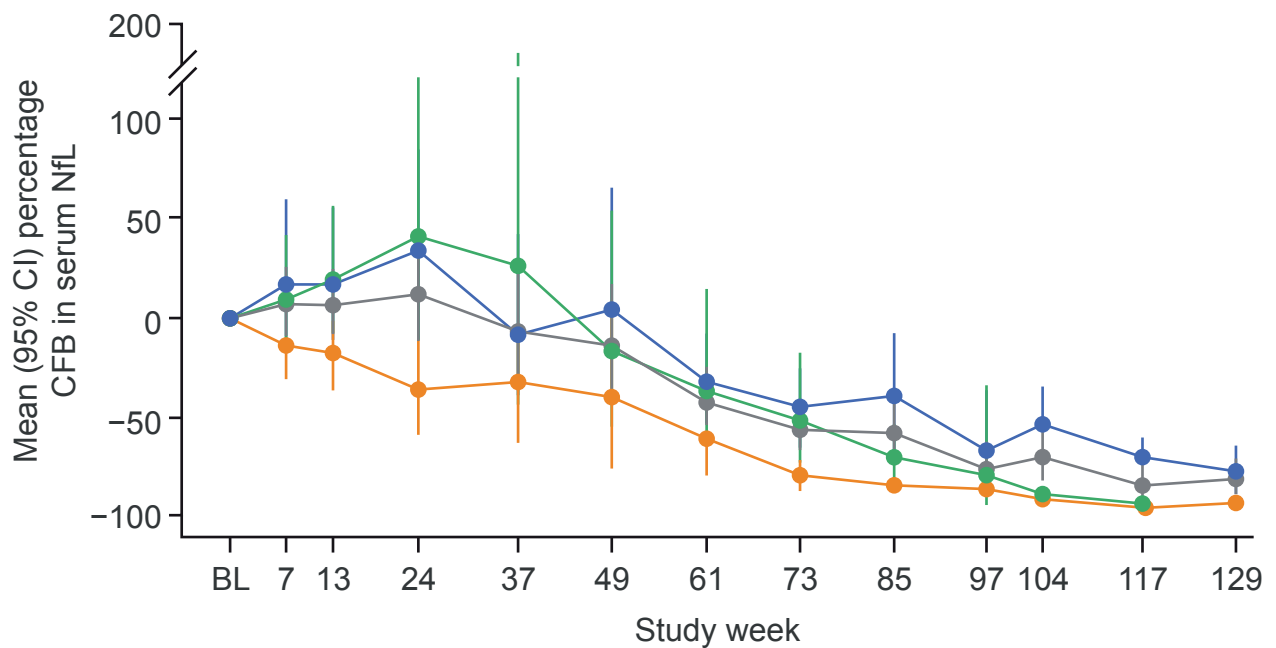


^aShaded gray area indicates the preliminary normal range of 1.61–3.6 µg/mg. ^bShaded gray area indicates the preliminary normal range of 0.34–0.83 µg/mg. ^cBaseline value missing for one patient. ^dData not shown for Week 153 because there were two participants in total. CFB, change from baseline; DS, dermatan sulfate; HS, heparan sulfate; MPS II, mucopolysaccharidosis type II; NE, not evaluable.

Supplemental Figure 3. Serum NfL mean percentage CFB

Age category at baseline

—●— All —●— <4 years —●— 4–6 years —●— >6 years



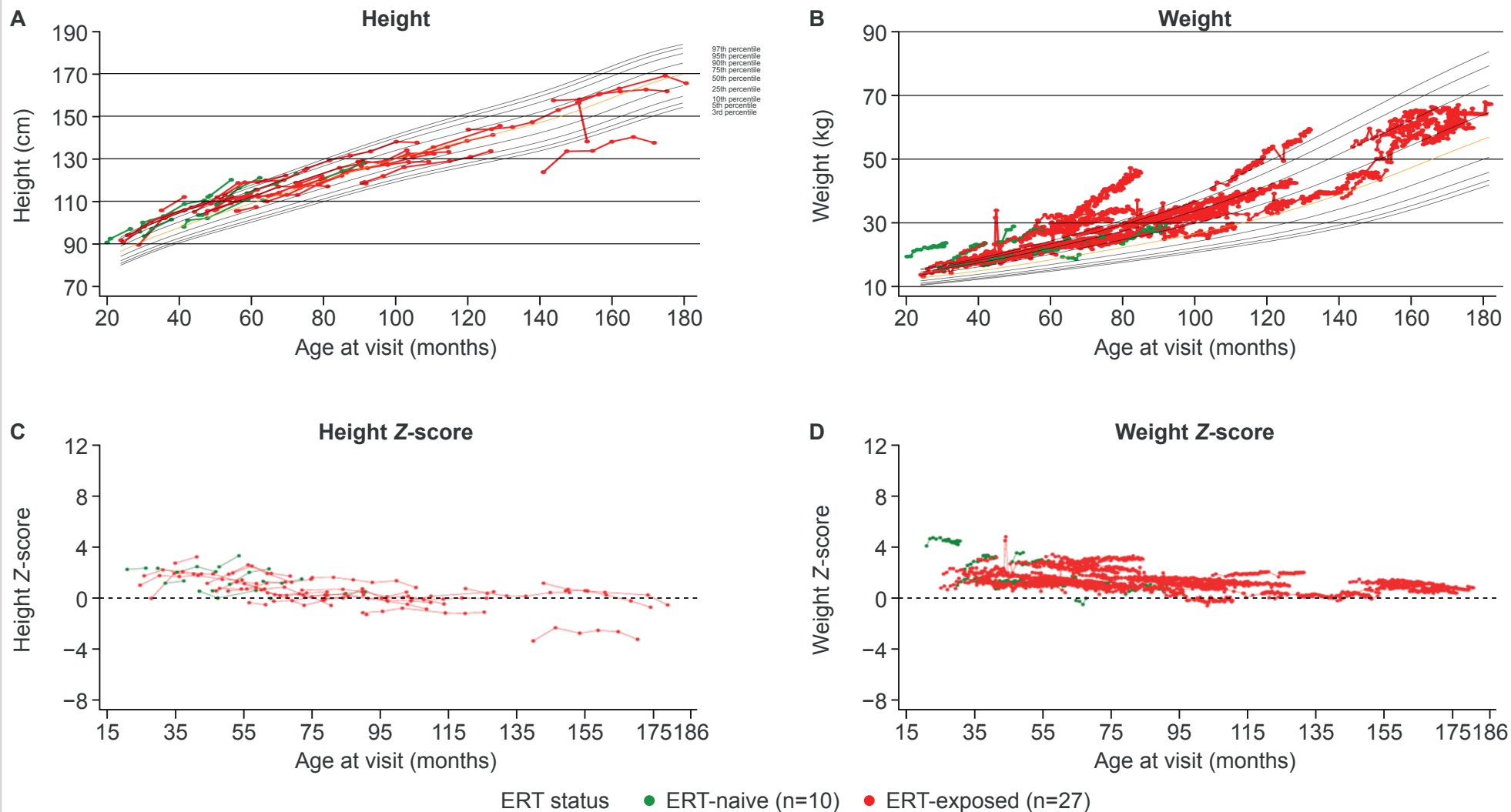
Number of participants, n

<4 years	7	6	7	7	6	5	6	5	3	3	2	3	2
4–6 years	8	8	8	8	7	7	7	7	5	4	3	2	0
>6 years	12	12	10	12	11	11	12	12	12	8	11	9	10
All	27	26	25	27	24	23	25	24	20	15	16	14	12

Age	Mean CFB, %					
	Week 24	Week 37	Week 49	Week 61	Week 104	Week 129
<4 years	-35.5	-31.7	-39.7	-59.8	-91.0	-92.7
4–6 years	40.4	26.4	-16.0	-36.4	-87.4	–
>6 years	32.8	-8.0	3.7	-31.8	-53.7*	-77.7*
All	11.9	-6.3	-13.5	-41.1*	-70.4*	-81.5*

* $P < 0.001$. Aggregate summaries by time point were provided for analysis visits that are common across all cohorts. The Week 7 analysis visit included observations closest to the target day (i.e. Day 43) from Weeks 5, 7 or 9. Mean CFB was computed from the geometric mean ratio relative to baseline. Corresponding 95% CI and P values were derived from the log ratio relative to baseline. Percentage CFB was derived as $100 \times x^{-1}$, in which x denotes the mean ratio, upper and lower limit for the mean ratio. BL, baseline; CFB, change from baseline; CI, confidence interval; NfL, neurofilament light chain.

Supplemental Figure 4. Individual participant (A) height, (B) weight, (C) height Z-score and (D) weight Z-score by age



ERT, enzyme replacement therapy.

REFERENCE

1. Hendriksz CJ *et al.* *Mol Genet Metab Rep* 2015;5:103–6.