The American Journal of Human Genetics, Volume 104

Supplemental Data

ELP1 Splicing Correction Reverses

Proprioceptive Sensory Loss in Familial Dysautonomia

Elisabetta Morini, Dadi Gao, Connor M. Montgomery, Monica Salani, Chiara Mazzasette, Tobias A. Krussig, Brooke Swain, Paula Dietrich, Jana Narasimhan, Vijayalakshmi Gabbeta, Amal Dakka, Jean Hedrick, Xin Zhao, Marla Weetall, Nikolai A. Naryshkin, Gregory G. Wojtkiewicz, Chien-Ping Ko, Michael E. Talkowski, Ioannis Dragatsis, and Susan A. Slaugenhaupt

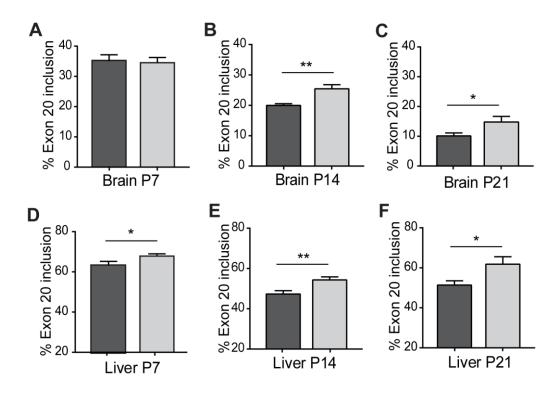


Figure S1. Kinetin treatment improves *ELP1* splicing in nursing pups. At the day of delivery the dams were randomly assigned to vehicle diet or kinetin diet, and, continued to be fed these diets until the time of weaning. *ELP1* splicing was analyzed in nursing pups carrying the human transgene with the major FD splicing mutation, TgFD9. (A-F) Percent of exon 20 inclusion in vehicle-assigned (dark grey) and kinetin-assigned (light grey) TgFD9 pups. (A) In brain at P7 no significant differences were detected in percent of exon 20 inclusion (P = 0.73) between vehicle-assigned (n=5) and kinetin-assigned (n=7) TgFD9 pups, two-tailed unpaired Student's *t*-test. (B) In brain at P14 a significant difference of **P< 0.01 was detected in percent of exon 20 inclusion between vehicle-assigned (n=5) and kinetin-assigned (n=8) TgFD9 pups, two-tailed unpaired Student's *t*-test. (C) In brain at P21, weaning time, a significant difference of *P< 0.05 was detected in percent of exon 20 inclusion between vehicle-assigned (n=5) and kinetin-assigned (n=6) TgFD9 pups, two-tailed unpaired Student's *t*-test. (D) In liver at P7 a significant

difference of *P< 0.05 was detected in percent of exon 20 inclusion between vehicle-assigned (n=5) and kinetin-assigned (n=7) TgFD9 pups, two-tailed unpaired Student's t-test. (E) In liver at P14 a significant difference of **P< 0.01 was detected in percent of exon 20 inclusion between vehicle-assigned (n=5) and kinetin-assigned (n=8) TgFD9 pups, two-tailed unpaired Student's t-test. (F) In liver at P21, weaning time, a significant difference of *P< 0.05 was detected in percent of exon 20 inclusion between vehicle-assigned (n=5) and kinetin-assigned (n=6) TgFD9 pups, two-tailed unpaired Student's t-test. Means and s.e.m. are shown.

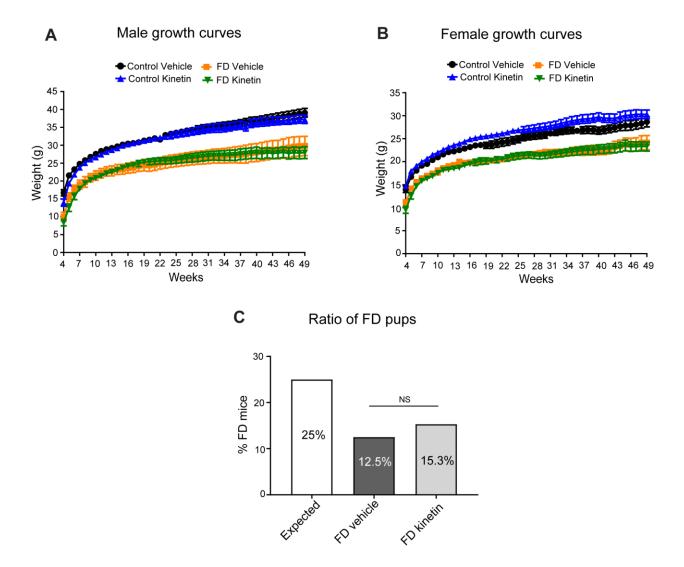


Figure S2. Kinetin treatment did not affect growth or ratio of the FD mice. (A) Postnatal growth curves for vehicle-treated (n=12, black), kinetin-treated (n=16, blue) control male mice and vehicle-treated (n=7, orange), kinetin-treated (n=11, green) FD male mice. (B) Postnatal growth curves for vehicle-treated (n=13, black), kinetin-treated (n=18, blue) control female mice and vehicle-treated (n=12, orange), kinetin-treated (n=15, green) FD female mice. Means and s.e.m. are shown. (C) The expected Mendelian ratio of TgFD9; $Ikbkap^{A20/flox}$ mice obtained by crossing TgFD9; $Ikbkap^{flox/flox}$ mice × $Ikbkap^{A20/flox}$ mice is 25% (white bar). The observed ratio was 12.5% (38/305) for the vehicle-treated TgFD9; $Ikbkap^{A20/flox}$ mice (dark grey bar) and 15.3% (52/340) for the kinetin-treated TgFD9; $Ikbkap^{A20/flox}$ mice (light grey bar). No significance

differences were observed in the actual ratio between vehicle-treated and kinetin-treated TgFD9; $Ikbkap^{\Delta 20/flox} \ \text{mice, (P=0.1539)} \ \chi^2 - \text{Test.}$

Kinetin tissue distribution (im/bit o b/bit o

Figure S3. Kinetin distribution in different tissues. The levels of compound were measured in cortex, trigeminal ganglia, lung, liver, heart, kidney and plasma from kinetin-treated FD mice (n = 13) using mass spectrometry. Means and s.e.m. are shown.

Table S1. Fibroblast cell lines from FD patients used in the RNAseq experiment to assess kinetin specificity.

Coriell #	Genotype	Sex	Age	Race
0850	Homozygous for <i>ELP1</i> T>C splice mutation	Male	26	White
2341	Homozygous for <i>ELP1</i> T>C splice mutation	Male	17	White
4589	Homozygous for <i>ELP1</i> T>C splice mutation	Male	16	White
2343	Homozygous for <i>ELP1</i> T>C splice mutation	Female	24	White
4663	Homozygous for <i>ELP1</i> T>C splice mutation	Female	2	White
4899	Homozygous for <i>ELP1</i> T>C splice mutation	Female	12	White