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Publications (2)

Tazelaar G, Hop P, Seelen M, Van Vugt J, van Rheenen W, Kool L, van Eijk K, Gijzen M, Dooijes D, Moisse M, Calvo A, Moglia C, Brunetti M, Canosa A, Nordin A, Pardina J, Ravits J, Al-Chalabi A, Chio A, McLaughlin R, Hardiman O, Van Damme P, de Carvalho M, Neuwirth C, Weber M, Andersen P, van den Berg L, Veldink J, van Es M. Whole genome sequencing analysis reveals post-zygotic mutation variability in monozygotic twins discordant for amyotrophic lateral sclerosis. *Neurobiol Aging* 2022; 122:76-87.

Tazelaar G, Boeynaems S, De Decker M, Farei-Campagna J, Kool L, Goedee H, McLaughlin R, Sproviero W, Iacoangeli A, Moisse M, Jacquemyn M, Daelemans D, Dekker A, Van Der Spek R, Westeneng H, Kenna K, Assialioui A, Da Silva N, PROJECT MINE ALS SEQUENCING CONSORTIUM, Povedano M, Mora J, Hardiman O, Salachas F, Millicamps S, Vourc'h P, Corcia P, Couratier P, Morrison K, Openshaw P, Shaw C, Pasterkamp R, Landers J, Van Den Bosch L, Robberecht W, Al-Chalabi A, van den Berg L, Van Damme P, Veldink J, van Es M. repeat expansions confer risk for amyotrophic lateral sclerosis and contribute to TDP-43 mislocalization. *Brain Commun* 2020; 2:fcaa064.

Projects (0)

No results found.

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