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## Publications (15)

Megat S, Mora N, Sanogo J, Roman O, Catanese A, Alami N, Freischmidt A, Mingaj X, De Calbiac H, Muratet F, Dirrig-Grosch S, Dieterle S, Van Bakel N, Müller K, Sieverding K, Weishaupt J, Andersen P, Weber M, Neuwirth C, Margelisch M, Sommacal A, van Eijk K, Veldink J, PROJECT MINE ALS SEQUENCING CONSORTIUM, Lautrette G, Couratier P, Camuzat A, Le Ber I, Grassano M, Chio A, Boeckers T, Ludolph A, Roselli F, Yilmazer-Hanke D, Millecamps S, Kabashi E, Storkebaum E, Sellier C, Dupuis L. Author Correction: Integrative genetic analysis illuminates ALS heritability and identifies risk genes. *Nat Commun* 2023; 14:8026.

Van Daele S, Moisse M, Farei-Campagna J, Zwamborn R, van der Spek R, van Rheenen W, van Eijk K, Kenna K, Corcia P, Vourc'h P, Couratier P, Hardiman O, McLaughlin R, Gotkine M, Drory V, Ticozzi N, Silani V, Ratti A, de Carvalho M, Mora J, Povedano M, Andersen P, Weber M, Başak N, Shaw C, Openshaw P, Morrison K, Landers J, Glass J, van Es M, van den Berg L, Al-Chalabi A, Veldink J, Van Damme P. Genetic variability in sporadic amyotrophic lateral sclerosis. *Brain* 2023; 146:3760-3769.

Megat S, Mora N, Sanogo J, Roman O, Catanese A, Alami N, Freischmidt A, Mingaj X, De Calbiac H, Muratet F, Dirrig-Grosch S, Dieterle S, Van Bakel N, Müller K, Sieverding K, Weishaupt J, Andersen P, Weber M, Neuwirth C, Margelisch M, Sommacal A, van Eijk K, Veldink J, PROJECT MINE ALS SEQUENCING CONSORTIUM, Lautrette G, Couratier P, Camuzat A, Le Ber I, Grassano M, Chio A, Boeckers T, Ludolph A, Roselli F, Yilmazer-Hanke D, Millecamps S, Kabashi E, Storkebaum E, Sellier C, Dupuis L. Integrative genetic analysis illuminates ALS heritability and identifies risk genes. *Nat Commun* 2023; 14:342.

Al Khleifat A, Iacoangeli A, Jones A, Van Vugt J, Moisse M, Shatunov A, Zwamborn R, van der Spek R, Cooper-Knock J, Topp S, van Rheenen W, Kenna B, van Eijk K, Kenna K, Byrne R, López V, Opie-Martin S, Vural A, Campos Y, Weber M, Smith B, Fogh I, Silani V, Morrison K, Dobson R, van Es M, McLaughlin R, Vourc'h P, Chio A, Corcia P, de Carvalho M, Gotkine M, Panades M, Mora J, Shaw P, Landers J, Glass J, Shaw C, Başak N, Hardiman O, Robberecht W, Van Damme P, van den Berg L, Veldink J, Al-Chalabi A. Telomere length analysis in amyotrophic lateral sclerosis using large-scale whole genome sequence data. *Front Cell Neurosci* 2022; 16:1050596.

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## Projects (0)

No results found.

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