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Employment

Research Fellow

Research Fellow
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1 Aug 2016 → present

Research outputs

MRSD: a quantitative approach for assessing suitability of RNA-seq in the investigation of mis-splicing in Mendelian disease
Rowlands, C., Taylor, A., Rice, G., Whiffin, N., Hall, H. N., Newman, W., Black, G., kConFab Investigators, O'Keefe, R., Hubbard, S., Douglas, A. G. L., Baralle, D. & Briggs, T., 12 Dec 2021, (Accepted/In press) In: American Journal of Human Genetics.

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Lepelley, A., Mina, E. D., Nieuwenhove, E. V., Waumans, L., Fraitag, S., Rice, G. I., Dhir, A., Frémond, M-L., Rodero, M. P., Seabra, L., Carter, E., Bodemer, C., Buhas, D., Callewaert, B., Lonlay, P. D., Somer, L. D., Dyment, D. A., Faes, F., Grove, L., Holden, S. & 10 others, Hully, M., Kurian, M. A., McMillan, H. J., Suetens, K., Tyynismaa, H., Chhun, S., Wai, T. , Wouters, C., Bader-Meunier, B. & Crow, Y. J., 13 Aug 2021, In: Journal of Experimental Medicine. 218, 10, e20201560.
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