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The mediating role of reading-related endophenotypes in the association between genes and reading skills.

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KEYWORDS: candidate genes, reading skills, endophenotypes, mediation, academic achievement.

ABSTRACT:

Reading is a fundamental skill for academic achievement. However, up to 20% of school-aged children have reading difficulties that affect their school functioning (Écalle et Magnan, 2015). Reading skills have already been associated with candidate genes and reading-related endophenotypes (Lervåg et Hulme, 2009; Mascheretti et al., 2018). Nonetheless, processes explaining the relationship between candidate genes and reading remain poorly studied (Mascheretti, 2018). It has been suggested that endophenotypes could be mediators between genes and reading skills (Mascheretti et al., 2020). The aim of the study was to examine the mediating role of reading-related endophenotypes in the association between genes and reading skills, in a population-based twin sample (Quebec Newborn Twin Study). Reading skills (decoding and comprehension), endophenotypes and DNA were measured when children were 8 years old ($n = 253$). Four endophenotypes (i.e., phonological awareness (PA), rapid automatized naming (RAN), rapid bimodal temporal processing (RTP) and rapid auditory processing (RAP)), as well as 14 markers spanning five reading-related candidate genes (i.e., DYX1C1, DCDC2, KIAA0319, ROBO1 and GRIN2B) using a structural equation modeling were tested. Results showed that PA and RAN were mediators in the pathway from DYX1C-rs3743205 and GRIN2B-rs2192973 to reading, respectively. Specifically, the minor allele (T/T) of DYX1C-rs3743205 predicted higher scores in PA, which, in turn, predicted higher decoding skills; and the minor allele (A/A) of GRIN2B-rs2192973 predicted higher score in RAN, which, in turn, predicted poorer comprehension skills. Our results support the use of endophenotypes in the explanation of the complex pathways between genes and reading.

References

- Écalle, J. et Magnan, A. (2015). *L'apprentissage de la lecture et ses difficultés* (2^e éd.). Dunod.
- Lervåg, A. et Hulme, C. (2009). Rapid automatized naming (RAN) taps a mechanism that places constraints on the development of early reading fluency. *Psychological Science*, 20(8), 1040–1048. <https://doi-org.acces.bibl.ulaval.ca/10.1111/j.1467-9280.2009.02405.x>
- Mascheretti, S., Andreola, C., Scaini, S., & Sulpizio, S. (2018). Beyond genes: A systematic review of environmental risk factors in specific reading disorder. *Research in Developmental Disabilities*, 82, 147–152. <https://doi.org/10.1016/j.ridd.2018.03.005>
- Mascheretti, S., Riva, V., Feng, B., Trezzi, V., Andreola, C., Giorda, R., ... & Facoetti, A. (2020). The Mediation Role of Dynamic Multisensory Processing Using Molecular Genetic Data in Dyslexia. *Brain sciences*, 10(12), 993.

GRANT SUPPORT: The Quebec Newborn Twin Study is supported by grants from the Canadian Institutes of Health, the Social and Humanities Research Council of Canada and the multiple Quebec Research Funds. Specifically, this research is supported by the Canadian Institutes of the Social and Humanities Research Council and the Quebec Research Fund- Society and Culture.