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**Robert J. Robbins**



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genetic component and more than 6 000 conditions have been linked to a known molecular cause In the last 13 years the adoption of human genomic sequencing has enabled the more efficient and accurate diagnosis and research of rare genetic disorders Genomic sequencing has become a first tier diagnostic test for many patients with congenital syndromes and suspected genetic disorders in high income countries as well as an effective method for the study of undiagnosed and novel genetic disorders in the research arena The implementation of genomic sequencing has dramatically changed the diagnosis and research of rare diseases in high income countries In contrast the reality in low and middle income countries LMICs is strikingly different where disparities on accessibility to these technologies exist The high cost of genomic sequencing and other molecular technologies remains a limiting factor in the common implementation of these methods for diagnosis and research of rare diseases in resource limited settings The study of rare genetic diseases in LMICs may be underestimated when compared to large scale genomic studies performed in developed countries despite being performed under much strained circumstances Consequently research publications on the genetics of RDs within LMICs might be underrepresented in the literature limiting the understanding of genetic and phenotypic variability across populations and contributing to the lack of representation of non European individuals in genetic studies This Research Topic aims to provide an opportunity for researchers and clinicians from LMICs who are dedicated to the study of rare genetic disorders to share their findings with the global genetics scientific community as well as their challenges and perspectives on the implementation of modern technologies and approaches for the diagnosis and study of rare genetic disorders

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