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An integrated modelling methodology for estimating the prevalence of centronuclear myopathy

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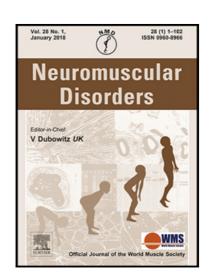
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Highlights

- There is a lack of comprehensive data on CNM prevalence by age and disease subtype
- There is a need for an integrated model enhancing current epidemiologic data
- The CNM incidence can be stratified by causative gene, severity and geographic region
- The CNM prevalence is estimated by causative gene, severity and geographic region
- The current data provide insight into CNM outcomes, e.g. severity and life years lost



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