# Utility of next generation sequencing in paediatric neurological disorders: experience from South Africa

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## INTRODUCTION

Next generation sequencing (NGS)-based tests have become routine first-line investigative modalities in paediatric neurology clinics in many high-income countries. These tests are both cost-effective and reliable in diagnosing many complex childhood neurological diseases. However, NGS-based testing in low-and middle-income countries is limited due to affordability constraints. The primary objective of this study was to evaluate the diagnostic yield and impact of targeted gene panel sequencing in a selected paediatric cohort attending a tertiary paediatric neurology clinic in the Western Cape Province of South Africa.

### **OBJECTIVES**

The objective of this study was to evaluate the diagnostic yield of targeted gene panel sequencing in a paediatric cohort attending a tertiary neurology clinic in the Western Cape Province of South Africa.

### **METHODS**

Retrospective study of 124 consecutive paediatric patients with neurological disease.

Patient's age: 6 weeks to 17 years

Test: NGS-based multi-gene panels

Questionnaire: given to caregivers where

pathological variant found

Duration: 41-month period

Results: Pathological variants phenotypical

matching

RESULTS			
Did the result help explain the reason for your child illness?	Results		
Yes	25 (86%)		
No	3 (10%)		
Unknown	1 (3%)		
Did the result help you to take better care for your child?			
Yes	14 (48%)		
No	13 (44%)		
Some/little	2 (7%)		
Should prenatal testing become available, would you make use of it?			
Yes	28 (97%)		
Not applicable	1 (3%)		
Did knowing the result bring you and			
your family any closure?			
Yes	28 (97%)		
Some/little	1 (3%)		

DISORDER	Pathogenic	Patho-	phenotype
GROUPS	gene	genic	correlation
Neuromuscular	RYR1	3	3
Disorders			
	SMN1	1	1
	TTN	1	0
	CAPN3	1	0
	GBE1	1	1
	STAC3	2	2
	DMD	4	4
	SCN9A	2	2
	SMN1,	1	1
	SMNcopies2		
	COL6A1	1	1
TOTAL		17(68%)	15(52%)

DISORDER GROUPS	Pathogenic	Patho-	phenotype
	gene	genic	correlation
Epilepsies	SCN1A	7	7
	COG5	1	(
	DYRK1A	1	1
	GABRB3	1	1
	UBE3A	1	(
	STXBP1	1	1
	KCNMA1	1	1
	KANSL1	1	1
	SCN2A	1	1
	TPP1	1	1
	CDKL5	4	۷
	KCNQ2	3	3
	GNAO1	1	1
	SLC6A5	1	1
	NPRL3	1	1
	PCDH19	1	1
	CDKL5	1	1
	SMC1A	1	1
TOTAL		28 (44%)	26(41%)
Heredodegenerative	ARSA	2	2
disorders			
	ACAD5	1	(
	UGT1A1	2	(
	ABCD1	1	1
	PLP1	1	
TOTAL		7(77%)	4(44%)
<b>CP</b> spectrum disorders	SCN2A	1	1
	ADAR	2	1
	SLC16A2	1	1
	BTD	2	(
	KCNA2	1	1
	QDPR	1	1
	ATM	1	(
	NPHP1	1	(
TOTAL		10(63%)	5 (31%)

#### RESULTS

The overall study diagnostic yield (DY) was 45%. Neuromuscular disorders 52% (13/25), cerebral palsy spectrum disorders 31% (5/16) and early-onset epilepsies 41% (26/63). With variants of uncertain significance in 38% (47/124).

The majority of caregivers (97%) viewed NGS-based testing as a positive experience.

#### CONCLUSION

The study enforces that NGS testing is achievable in resource-constrained settings. The high diagnostic yield in this study suggests that it is feasible to recommend NGS as a first-tier testing approach for children with neurological disorders.

#### REFERENCES

van Niekerk et al, 2025 In Press at the European Journal of Human Genetics. (accepted awaiting publication)

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