

# CNV Mendelian Precision Report

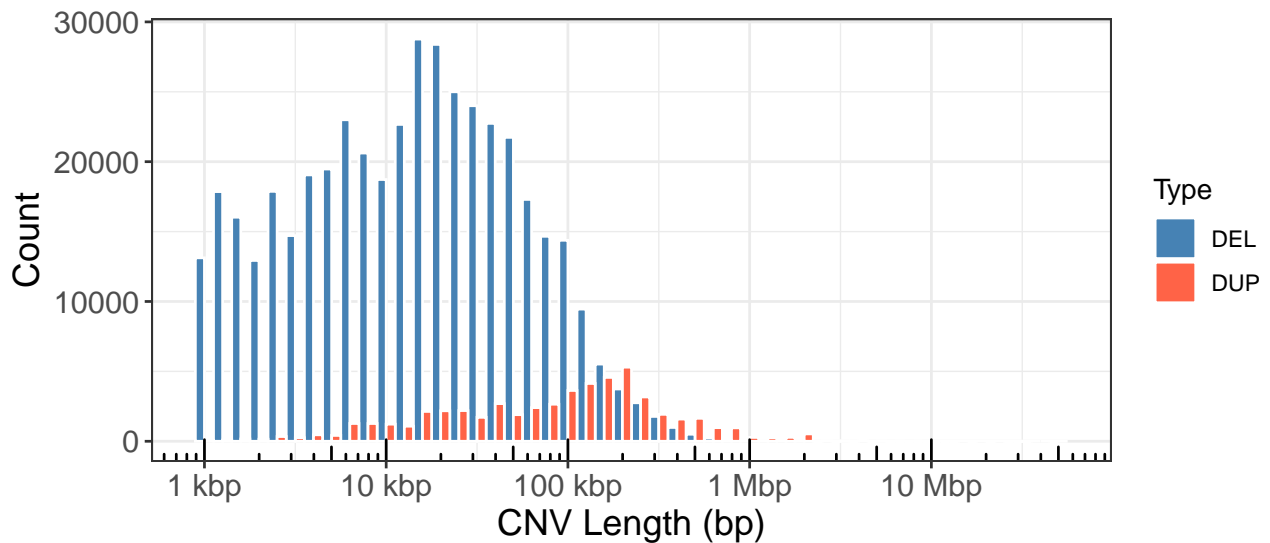
Jacquemont Lab

**Dataset: SPARK\_PennCNV\_raw**

**Date: 04/08/2025**

**Number of Trios or Children analyzed: 16421**

**CNV Length Distribution Across Samples**



##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1000	5026	16553	53641	47163	49397679

**Distribution of CNVs by Copy Number State**

##	0	0,1	1	1,3	2	3	3,4	4
##	1925	3	436550	25	315	53663	5	96

**Summary of CNV Counts per Individual**

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1	8	13	30	35	1183

Mendelian precision across CNV length bins: assessing dataset quality (proportion of child CNVs observed in parents with  $\geq 50\%$  reciprocal overlap) across multiple metrics.

*Note: bins with fewer than 1000 CNVs were excluded from the analysis.*

