Supplementary information to:

Original article:

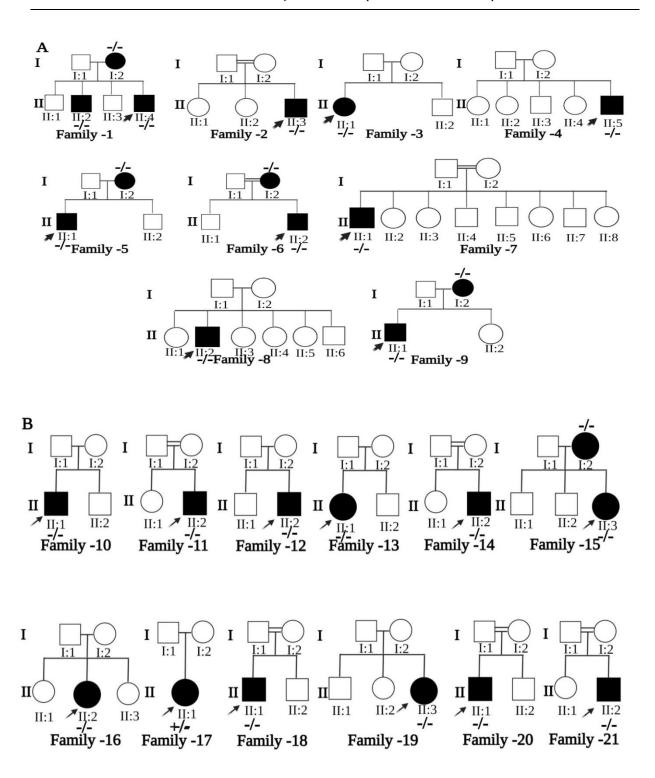
EXPLORING MITO-NUCLEAR GENETIC FACTORS IN LEBER'S HEREDITARY OPTIC NEUROPATHY: INSIGHTS FROM COMPREHENSIVE PROFILING OF UNIQUE CASES

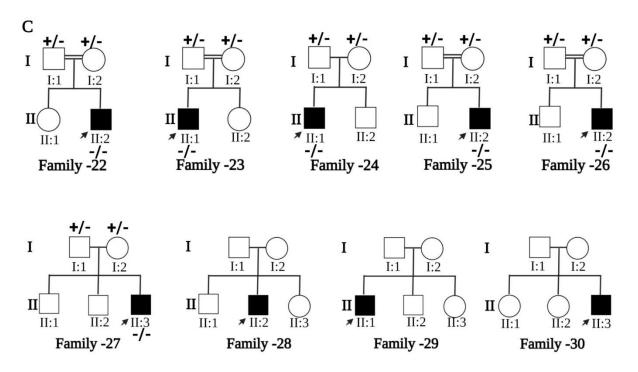
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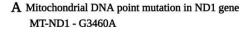
https://dx.doi.org/10.17179/excli2023-6297

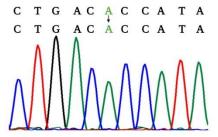
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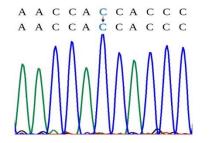


Supplementary Figure 1: Pedigree of thirty unrelated probands involved in this study. Solid symbols with an arrow represent proband. Consanguinity is represented by a double line. The genotype is denoted by (-/-) for homozygous variant carriers, (+/-) for heterozygous variant carriers. **A-** Group I; **B-** Group II and **C-** Group III

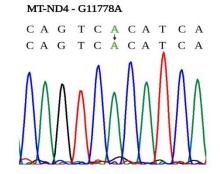




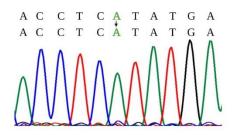
C Mitochondrial DNA point mutation in ND6 gene MT-ND6 - T14484C



 \boldsymbol{B} Mitochondrial DNA point mutation in ND4 gene

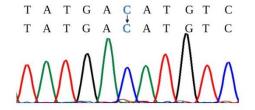


D Mitochondrial DNA point mutation in MT-ND1 gene MT-ND1- C4171A

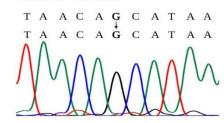


Supplementary Figure 2: Representative chromatogram of the probands in group I harboring primary mitochondrial DNA mutations (MT-ND1, MT-ND4 and MT-ND6 genes)

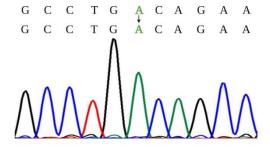
A Mitochondrial DNA point mutation in ND1 gene MT-ND1 - T4216C



Mitochondrial DNA point mutation in ND4 gene MT-ND4 - A12033G

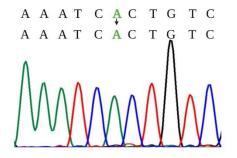


C Mitochondrial DNA point mutation in ND5 gene MT-ND5 - G13708A

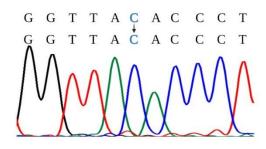


Supplementary Figure 2a: Sequencing chromatogram of the probands in group I harboring a co-occurrence secondary mitochondrial DNA mutations (MT-ND1, MT-ND4 and MT-ND5 genes)

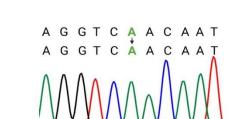
A Mitochondrial DNA point mutation in ATP6 gene B MT-ATP6 - G9139A



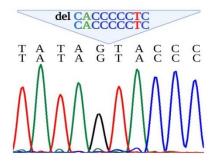
B Mitochondrial DNA point mutation in MT-TM gene MT-TM - T4454C



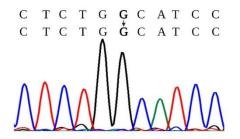
C Mitochondrial DNA point mutation in MT-CO2 gene MT-CO2- G7859A



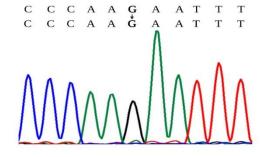
D Mitochondrial DNA deletion mutation in MT-NC7 MT-NC7 - (del CACCCCTC)



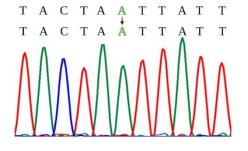
E Mitochondrial DNA point mutation in ND2 gene MT-ND2 - A4842G



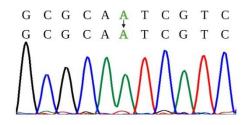
F Mitochondrial DNA point mutation in TL2 gene MT-TL2 - A12308G



G Mitochondrial DNA point mutation in ATP6 gene MT-ATP6 - G8950A

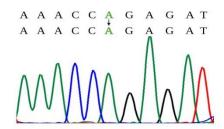


H Mitochondrial DNA point mutation in ND4 gene MT-ND4 - G11696A

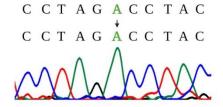


- I Mitochondrial DNA point mutation in MT-CO1 gene MT-CO1- G7444A

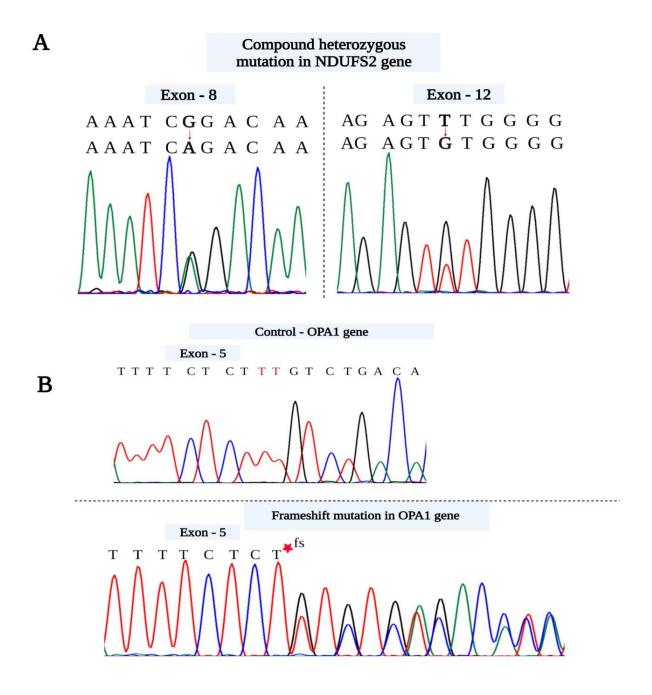
Mitochondrial DNA point mutation in MT-TT gene MT-TT - G15927A



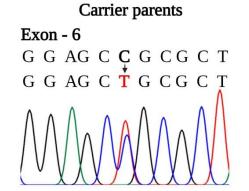
K Mitochondrial DNA point mutation in MT-ATP6 gene MT-ATP6- G8573A

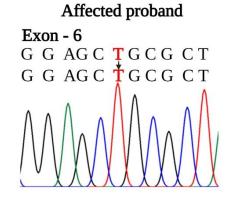


Supplementary Figure 3: Sequencing chromatogram from **A-K** represents the probands from group II harboring secondary mitochondrial DNA mutations.



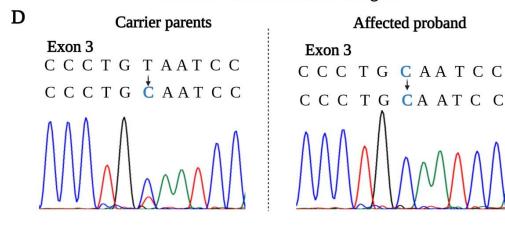
\mathbf{C} Missense mutation in NDUFS7 gene

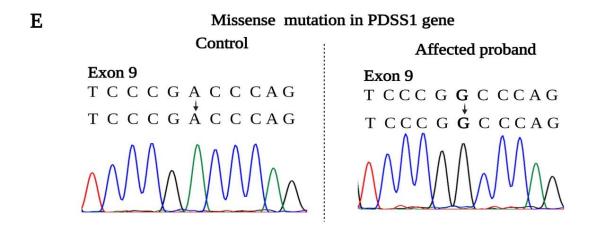


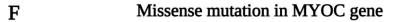


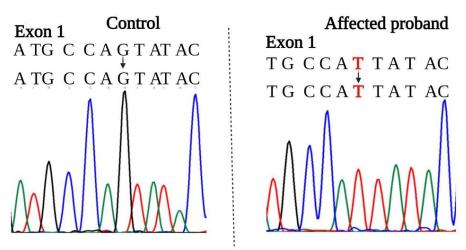
Affected proband

Missense mutation in MTFMT gene





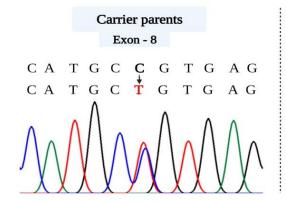


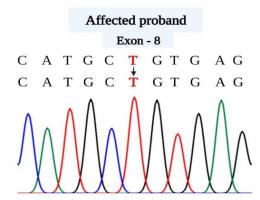


Supplementary Figure 4: Sequencing chromatogram from **A-F** represents the probands from group II harboring nuclear DNA mutations (NDUFS2, OPA1, NDUFS7, MTFMT, PDSS1 and MYOC gene)



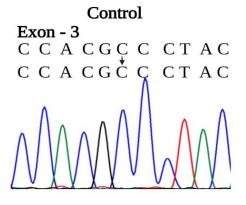
Missense mutation in NDUFV1 gene

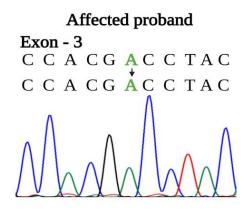




B

Missense mutation in NDUFA3 gene



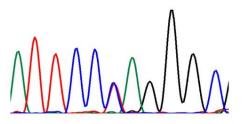


\mathbf{C}

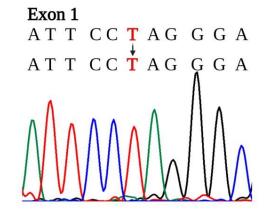
Non-sense mutation in DNAJC30 gene

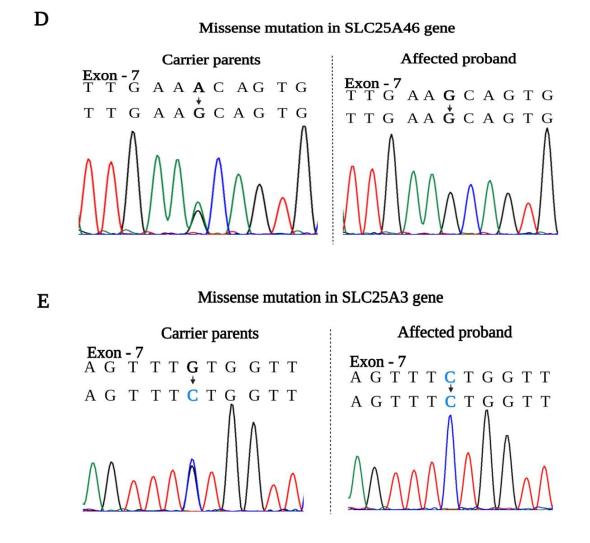
Carrier Parents

Exon 1 AT T C C C A G G G A AT T C C T A G G G A



Affected Proband





Supplementary Figure 5: Sequencing chromatogram from **A-E** represents the probands from group III harboring nuclear DNA mutations (NDUFV1, NDUFA3, DNAJC30, SLC25A46 and SLC25A3 gene

Supplementary Table 1: Illustrates the key component of the sample group, family history and male to female ratio

Group	Probands	Total (Familial/ Sporadic)	Male (Familial/ Sporadic)	Female (Familial/ Sporadic)	Ratio (Male: Female)
	Total screened (n=30)	30 (5/25)	24 (4/20)	6 (1/5)	4:01
Group I (1)	LHON primary mutation (n=9)	9 (4/5)	8 (4/4)	1 (-/1)	8:01
Group II (2)	Secondary mitochondrial DNA mutation (n=12)	12 (1/11)	7 (-/7)	5 (1/4)	1.4:1
	*Secondary mitochondrial DNA mutation alone (n=6)	6(1/5)	4 (-/4)	2 (1/1)	2:01
	#Secondary mitochondrial DNA mutation + Nuclear gene mutation (n=6)	6 (-/6)	3 (-/3)	3 (-/3)	1:01
Group III (no mtDNA mutation)	Nuclear gene mutation (n=6)	6 (-/6)	6 (-/6)	0 (-/-)	NA
	No mutation (n=3)	3 (-/3)	3 (-/3)	0 (-/-)	NA

^{*} indicates patients harbor only secondary mtDNA mutation # indicates patients harbor both secondary mtDNA mutation and nuclear gene mutation