

## Supplementary Materials:

**Table S1.** Clinical, histological, and molecular features of the nine patients.

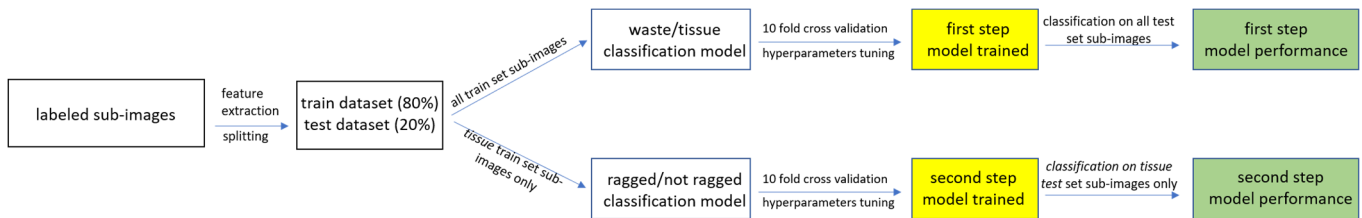
	Patient 1	Patient 2	Patient 3	Patient 4	Patient 5	Patient 6	Patient 7	Patient 8	Patient 9
Sex	M	M	F	F	M	M	F	M	F
Age at muscle biopsy	5 yrs	11 yrs	41 yrs	8 yrs	60 yrs	3 months	46 yrs	67 yrs	58 yrs
Clinical features	Cardiomyopathy, myopathy	Retinitis pigmentosa, ptosis, ataxia, hearing loss	Cardiomyopathy, hearing loss, diabetes	Retinitis pigmentosa, hearing loss, diabetes, psychomotor regression	Ophthalmoplegia	Hypotonia, severe metabolic acidosis	Ptosis, decreased visual acuity	Ophthalmoplegia	Ophthalmoplegia
Histology	RRFs; subsarcolemmal rims, lipid storage	RRFs, COX-negative SDH-positive fibers, lipid storage	RRFs, COX-negative SDH-positive fibers	RRFs, COX-negative SDH-positive fibers, lipid storage	Variation in fiber size, RRFs, COX-negative SDH-positive fibers	RRFs, diffuse reduced COX staining, SDH-positive fibers	RRFs, COX-negative SDH-positive fibers	RRFs, COX-negative SDH-positive fibers	RRFs, COX-negative SDH-positive fibers, lipid storage
Number of analysed images	2	4	6	5	5	2	2	6	5
Mutation	ACAD9 c.170C>T (p.Pro57Leu)+ c.1240C>T (p.Arg414Cys)	8 Kb mtDNA single deletion	m.5522G>A MT-TW (tRNA <sup>Trp</sup> )	7 Kb mtDNA single deletion	mtDNA common deletion	Muscular mtDNA depletion	OPA1 c.124C>T (p.H42Y)	mtDNA common deletion	5 Kb mtDNA single deletion
Diagnosis	Mitochondrial Cardiomyopathy	Kearns-Sayre syndrome	Encephalomyopathy	Kearns-Sayre syndrome	CPEO	Mitochondrial DNA depletion syndrome	DOA, dominant optic atrophy plus syndrome	CPEO	CPEO

Abbreviation: M, man; F, woman; CPEO, chronic progressive external ophthalmoplegia; RRFs, ragged red fibers; COX, cytochrome c oxidase.; SDH, succinate dehydrogenase, mtDNA, mitochondrial DNA.

**Table S2.** Hyperparameters values for each model after the tuning operation.

	Waste-tissue classification	Ragged – not ragged classification
RF	min size of nodes: 15 number of trees: 625 number of variables: 9 priors of the classes: 0.487, 0.316	min size of nodes: 19 number of trees: 1466 number of variables: 4 priors of the classes: 0.322, 0.792
GBM	distribution: adaboost	distribution: adaboost

	<b>number of trees:</b> 300 <b>shrinkage:</b> 0.483373 <b>fraction selected:</b> 0.630 <b>fraction for train:</b> 0.946	<b>number of trees:</b> 173 <b>shrinkage:</b> 0.0540611 <b>fraction selected:</b> 0.514 <b>fraction for train:</b> 0.988
SVM	<b>kernel:</b> radial <b>degree:</b> - <b>gamma:</b> 0.01572429 <b>coef0:</b> - <b>cost:</b> 5.31966	<b>kernel:</b> radial <b>degree:</b> - <b>gamma:</b> 0.01732987 <b>coef0:</b> - <b>cost:</b> 4.905418



**Figure S1:** protocol used to select the model with the best overall fit for both classification steps.