

1 **In vitro characterization of mitochondrial function and structure in rat and human cells with**
2 **a deficiency of the NADH:ubiquinone oxidoreductase Ndufc2 subunit.**

3

4 Salvatore Raffa^{1,4}, Cristina Scrofani¹, Sabatino Valente¹, Andrea Micaloni¹, Maurizio Forte², Franca
5 Bianchi², Roberta Coluccia², Aron M. Geurts³, Sebastiano Sciarretta⁵, Massimo Volpe^{1,2}, Maria
6 Rosaria Torrisci^{1,4}, Speranza Rubattu*^{1,2}

7

8

- 9 1. Department of Clinical and Molecular Medicine, School of Medicine and Psychology,
10 Sapienza University of Rome, Rome, Italy;
- 11 2. I.R.C.C.S. Neuromed, Pozzilli (Isernia), Italy;
- 12 3. Department of Physiology and Human and Molecular Genetics Center, Medical College of
13 Wisconsin, Milwaukee, WI, USA.
- 14 4. Cellular Diagnostics Unit, Azienda Sant'Andrea University Hospital, Rome, Italy.
- 15 5. Department of Medical-Surgical Sciences and Biotechnologies, Sapienza University of
16 Rome, Latina, Italy.

17

18 *Corresponding author:

19 Speranza Rubattu

20 Dept. of Clinical and Molecular Medicine

21 School of Medicine and Psychology. Sapienza University, Rome, Italy

22 IRCCS Neuromed, Pozzilli, Italy

23 Tel. 0039 06 33775979; Fax 0039 06 33775061

24 e-mail: rubattu.speranza@neuromed.it

25

33 **a. Performing the characterization of fibroblast primary cultures from the *Ndufc2* knockout**
34 **rat model**

35 To analyze the biological characteristics of primary cells, we used all cultures at the second passage
36 (P2) to avoid in vitro differentiation (Leone L. et al; doi: 10.1371/journal.pone.0146365).

37 The mesenchymal immunophenotype of the primary cultures was verified by immunofluorescence
38 assessment of the expression of vimentin, a component of the intermediate filaments widely used as
39 a fibroblast marker. The analysis showed that all cultures were characterized by a strong staining for
40 vimentin, displaying a signal compatible with the structure and localization of perinuclear
41 cytoplasmic bundles of filaments (Supplementary Fig. 1A). To confirm the purity of cultures, we
42 ruled out the presence of epithelial cells by assessment of the cytokeratins expression, components
43 of the epithelial intermediate filaments; no signal was detected in all isolated primary cultures
44 (Supplementary Fig. 1A).

45 To evaluate the degree of fibroblasts differentiation, we used a morphological characterization to
46 distinguish in vitro the differentiation stages based on distinct morphological features and we
47 calculated the L:E differentiation index, as defined in Materials and Methods. The quantitative
48 analysis revealed that the L:E ratio was homogeneous, with no appreciable differences among all
49 fibroblast cultures (median of L:E index: 0.83 vs 0.85 for SHRSR and SHRSR_KO *Ndufc2*,
50 respectively; p=NS; Supplementary Fig. 1B).

51 To better evaluate the differentiated phenotype of all fibroblast cultures, we analyzed by
52 quantitative immunofluorescence the percentage of positive cells for the expression of α -smooth
53 muscle actin, the most commonly used marker for activated and differentiated fibroblasts. Both
54 fibroblast cultures from KO and WT rats displayed a percentage of positive staining for the α -SMA
55 close to 25%, with fluorescent signal localized in the peripheral areas of the cytoplasm,
56 representative of an higher content of actin bundles (median of α -SMA positive staining: 23.2% vs
57 26.9% for SHRSR and SHRSR_KO *Ndufc2*, respectively; p=NS; Supplementary Fig. 1C).

58 To verify the typical ultrastructural features of primary fibroblasts, all cultures were analyzed by
59 transmission electron microscopy (TEM). The cells showed a typical spindle-like shape and a
60 cytoplasm with a few rough endoplasmic reticulum cisternae and intracellular filaments; in
61 addition, some fibroblasts were characterized by a cytoplasm with very abundant organelles,
62 numerous cisternae of rough endoplasmic reticulum and evident intracellular filaments, also
63 organized in bundles, localized near the plasma membrane, and consistent with differentiated
64 fibroblasts (Supplementary Figs. 1E-G)

65 Taken together, these data showed that no appreciable morphological, immunophenotypic and
66 ultrastructural findings were observed between fibroblast cultures from SHRSR_WT and
67 SHRSR_KO *Ndufc2* rats.

68 **b. KCTD21 and NDUFC2 expression study in a larger group of healthy subjects**

69 For this purpose we recruited an overall number of 34 subjects (n=34) homozygous for the major
70 allele (TT/rs11237379; n=9), heterozygous (TC/rs11237379; n=16), or homozygous for the minor
71 allele (CC/rs11237379; n=9). Genotype assessment for the NDUFC2 variant was performed as
72 previously reported (Rubattu S. et al.; doi: 10.1161/JAHA.115.002701). Analysis of NDUFC2
73 expression was performed by RTPCR of cDNA using specific oligonucleotides following
74 previously published procedures (Rubattu S. et al.; doi: 10.1161/JAHA.115.002701). Primers used
75 for KCTD21 were the following: KCTD21 forward GAG GGC AGG AGG ACT ACT T; KCTD21
76 reverse GTC GCC AGT GAG GTT GTA TAG. The amount of cDNA target was calculated using
77 GAPDH (primer for: CAAGGCTGTGGGCAAGGT; primer rev: GGAAGGCCATGCCAGTGA)
78 as housekeeping gene.

79

80

81

82

83

84 **Legend to Supplementary Figure 1**

85 **Characterization of fibroblast primary cultures from the *Ndufc2* knockout rat model**

86 (A) Morphological evaluation (DIC) and immunofluorescence analysis of expression of vimentin
87 and cytokeratins on a representative sample of fibroblast cultures isolated from the mesenchymal
88 tissue of SHRSR and SHRSR_KO *Ndufc2*. In all cultures, the cells showed a strong staining for the
89 mesenchymal marker vimentin (red), which appears as perinuclear cytoplasmic bundles of
90 filaments. No fluorescent signal was detected for cytokeratins (green), ruling out the presence of
91 epithelial cells in the cultures. Nuclei are stained with DAPI (blue).

92 (B) Morphological evaluation of the differentiation degree of fibroblasts cultures by differential
93 interference contrast microscopy. The quantitative analysis revealed that the L:E differentiation
94 index was homogeneous among both cultures. The scatter dot-plot showed the percent of L:E ratio
95 values obtained from three independent experiments for each culture; the median values and the
96 interquartile ranges of the groups are indicated (Kruskal-Wallis test: $\wedge p=NS$ versus SHRSR_WT).

97 (C) Evaluation of α -smooth muscle actin expression. The immunofluorescence analysis of the
98 expression of the differentiation marker α -SMA showed a signal localized in intracellular filaments,
99 also organized in bundles, situated in the peripheral cytoplasmic areas of the cells. The quantitative
100 analysis displayed a similar percentage of positive staining in both fibroblast cultures (scatter dot-
101 plot with median values and interquartile ranges of α -SMA staining; Kruskal-Wallis test: $\wedge p=NS$
102 versus SHRSR_WT).

103 (D-G) Ultrastructural features of the cell cultures. Both the fibroblasts isolated from skin of
104 SHRSR_WT (D-E) and SHRSR_KO *Ndufc2* (F-G) showed a typical spindle-like shape with
105 cytoplasm characterized by high cytoplasmatic complexity, with abundant organelles and
106 intracellular filaments localized in the peripheral areas of the cell, near the plasma membrane,
107 consistent with the intracellular localization of α -smooth muscle actin. (TEM micrographs, uranyl
108 acetate/lead citrate; Legend: Nu, nucleus; NM, nuclear membrane, PM, plasma membrane; Mt,
109 mitochondrion, IFb, intermediate filament bundles; rER, rough endoplasmic reticulum).

110 **Legend to Supplementary Figure 2**

111 **NDUFC2 expression study in a larger group of healthy subjects and GTEx analysis**

112 (A) NDUFC2 expression, assessed by RTPCR, in PBMCs of the subjects used for all *in vitro*
113 studies described in the main text (n=6 for each rs11237379/NDUFC2 genotype).

114 **P<0.001 for both CT and TT versus CC genotype. Values are expressed as means ±SEM.

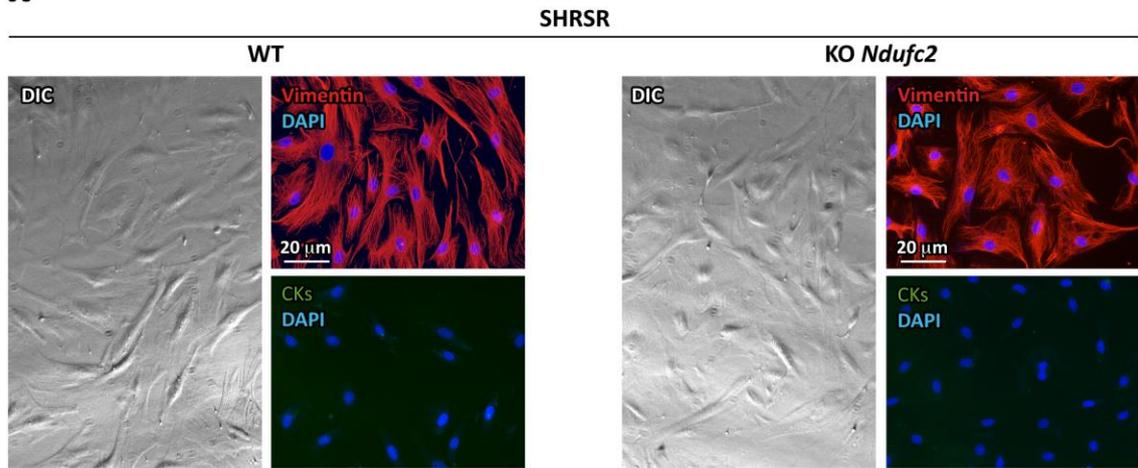
115 (B) Box plots from Genotype-Tissue Expression GTEx for brain NDUFC2 and KCTD21. The
116 GTEx Project was supported by the Common Fund of the Office of the Director of the National
117 Institutes of Health, and by NCI, NHGRI, NHLBI, NIDA, NIMH, and NINDS. The data used for
118 the analyses described in this manuscript were obtained from: the GTEx Portal on 06/20/17

119 (C) NDUFC2 and KCTD21 expression assessed by RTPCR in PBMCs of a larger group of healthy
120 subjects carrying the three genotypes for rs11237379/NDUFC2.

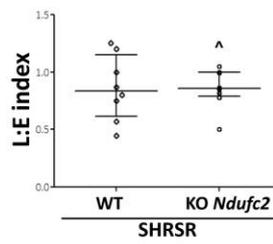
121 **P<0.001 for both CT and TT versus CC genotype. Values are expressed as means ±SEM.

122

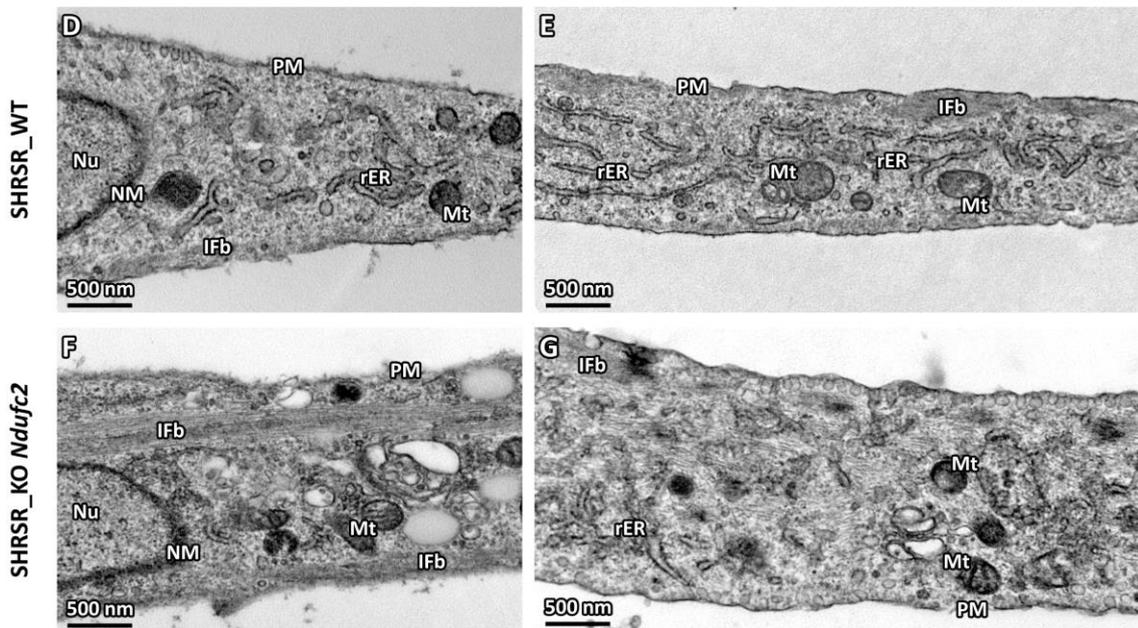
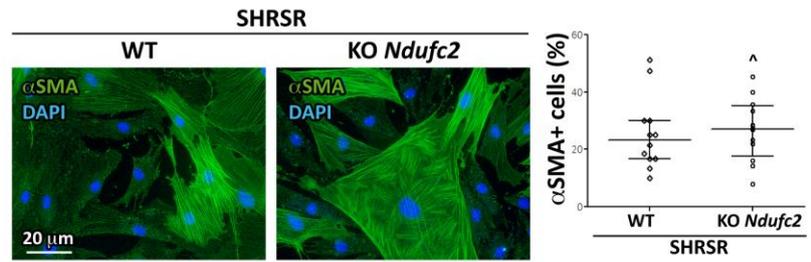
A



B

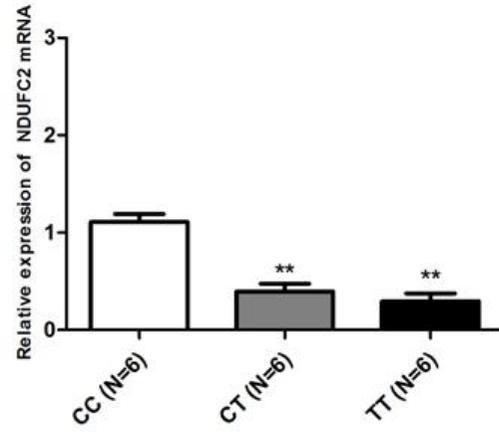


C

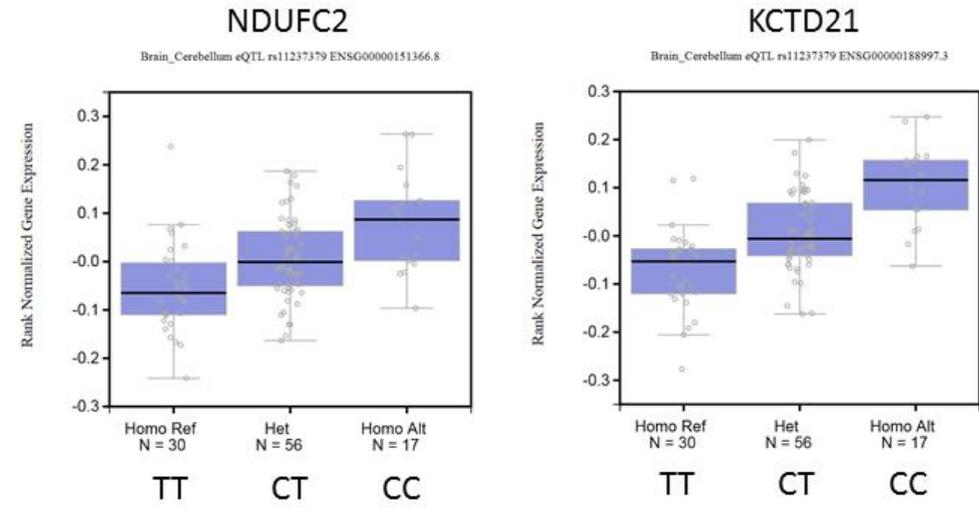


Supplementary Fig. 1

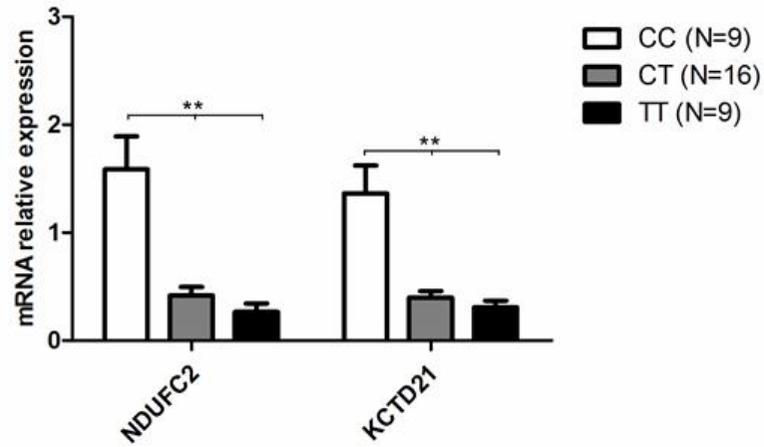
A



B



C



Supplementary Fig. 2