

## Supplementary Materials for

# MITF regulates IDH1, NNT, and a transcriptional program protecting melanoma from reactive oxygen species

### Authors

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### This PDF file includes:

Fig. S1. MITF regulates genes involved in oxidative-reductive processes in melanoma

Fig. S2. Features of zebrafish melanoma model

Fig. S3. MALDI-TOF mass spectra

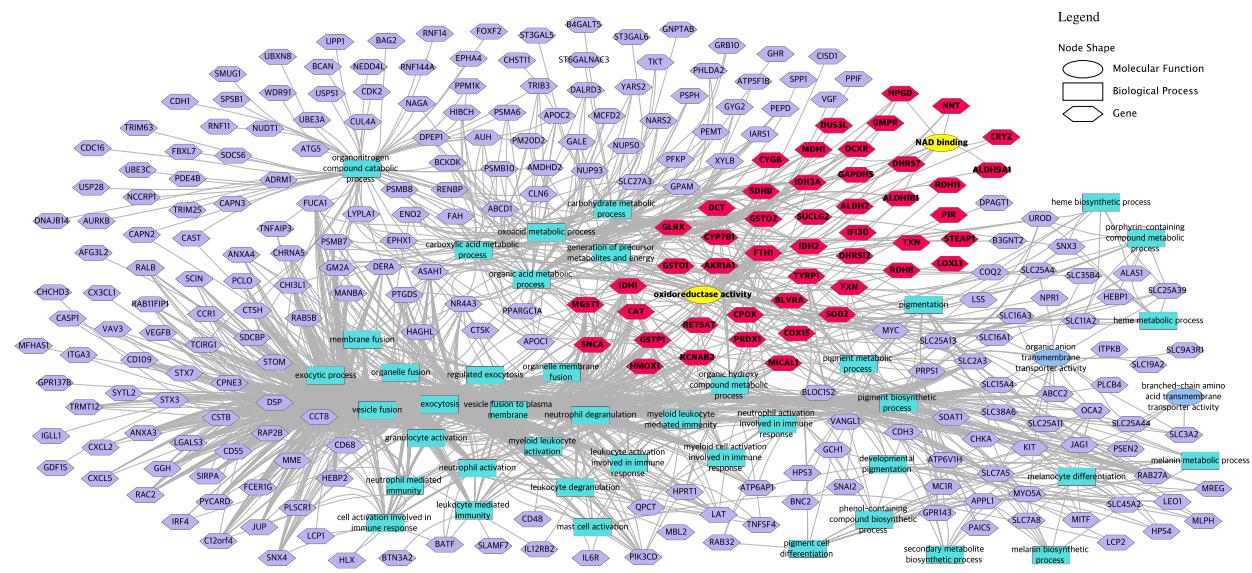
Fig. S4. Relative MITF mRNA levels after MITF silencing or overexpression

### Other Supplementary Material for this manuscript includes the following:

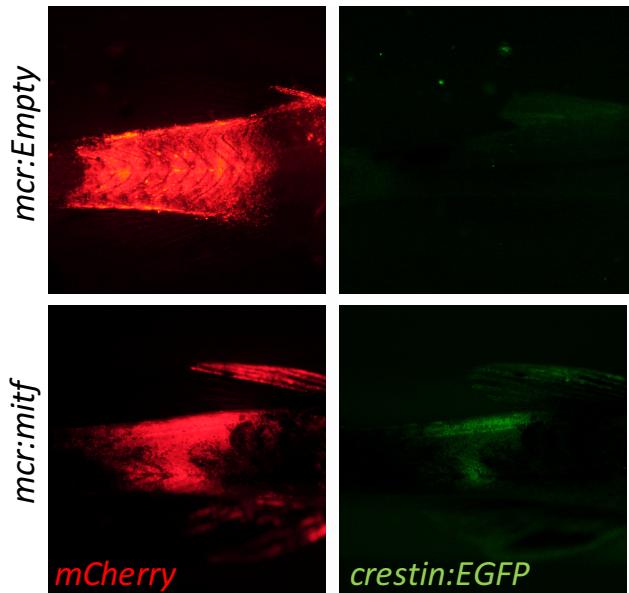
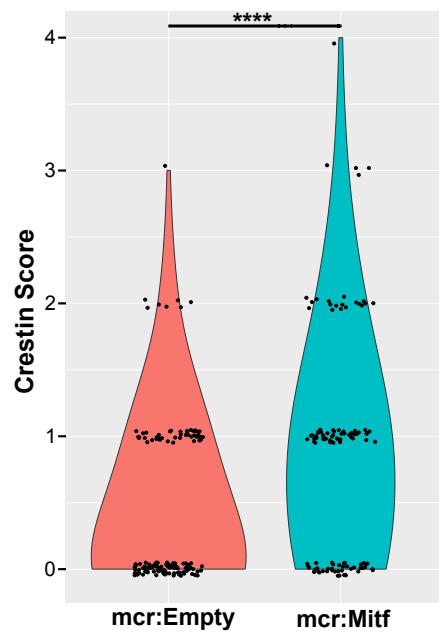
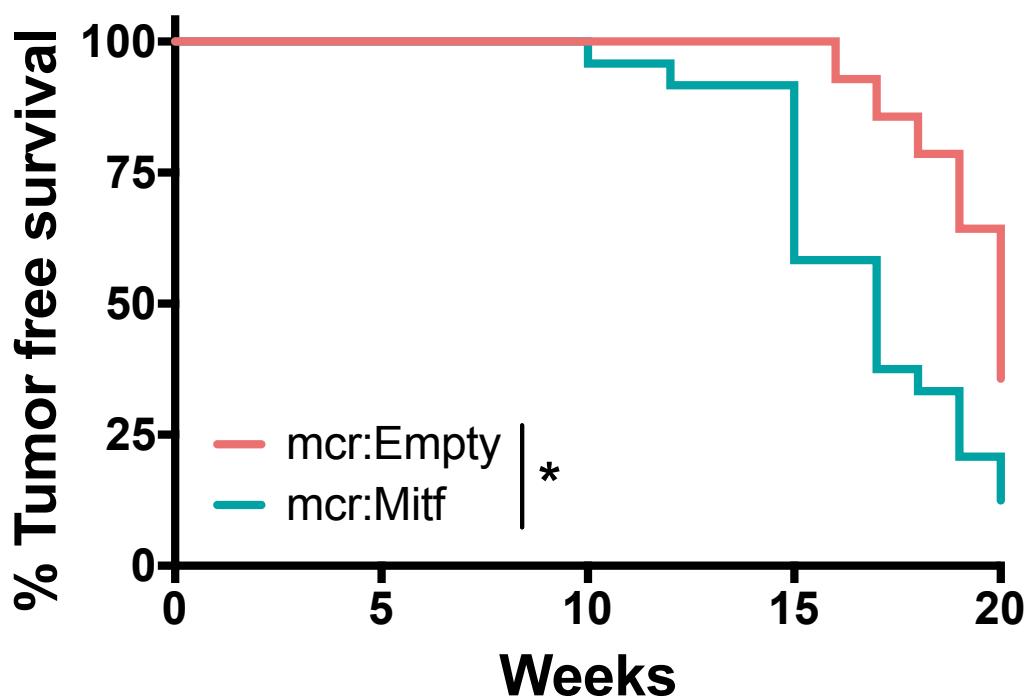
Table S1. Supplementary Table 1. Gene sets enriched in genes downregulated after MITF KD in MALME-3M cells by DAVID. Yellow highlight indicates gene sets associated with ROS that were used to define MITF-driven redox program score in Table S3.

Table S2. Supplementary Table 2. Gene sets enriched in genes downregulated after MITF KD in MALME-3M cells by DAVID after redundant gene set removal by REVIGO.

Table S3. Supplementary Table 3. Genes and gene sets to define the MITF-driven redox program score based on the genes annotated to redox processes whose expression in MALME-3M cells changes significantly after *MITF* knockdown.

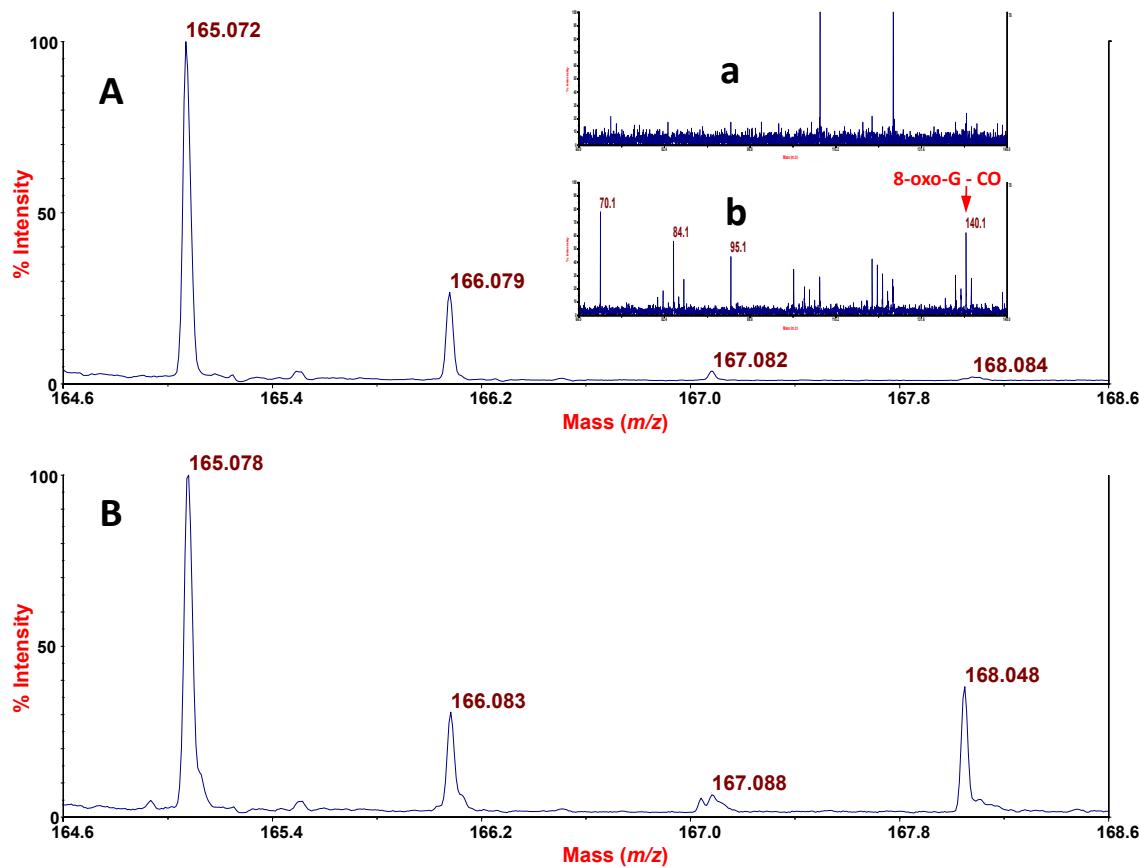


**Fig. S1. MITF regulates genes involved in oxidative-reductive processes in melanoma.** The oxidoreductase activity gene set is enriched in high MITF melanoma cell lines in short-term melanoma cultures. GSEA results were visualized by Cytoscape. Abbreviations: MITF: Microphthalmia-associated transcription factor

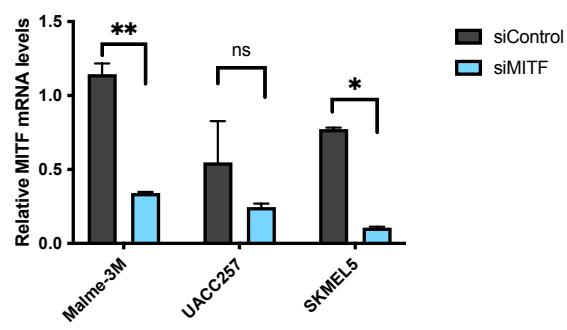
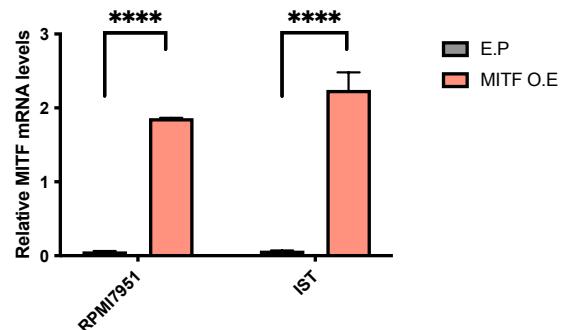
**A****B****C**

**Fig. S2. Features of zebrafish melanoma model.** Fluorescent images (A) and quantitation (B) show rescued melanocytes (*mitf:mCherry*+ cells) and neural crest reactivation in early melanoma patches (*crestin:EGFP*+ cells) in 6-week-old *MITF* overexpressing (*mcr:MITF*) fish compared with control

(*mcr:Empty*) fish (scored on a 0-5 scale, with 5 representing a tumor). **(C)** Kaplan-Meier curve depicting the percent of *mcr:MITF* fish with tumor-free survival compared to control *mcr:Empty* fish. The *mcr:Empty* vectors was used as the negative control. Abbreviations: MITF: Microphtalmia-associated transcription factor



**Fig. S3. MALDI-TOF mass spectra.** (A) CCA matrix; (B) DNA from Mitf1, showing a peak for 8-oxo-G at  $m/z$  168.048; a corresponding TOF/TOF spectrum of  $m/z$  168 gave a peak at  $m/z$  140.1 from loss of CO (insets a and b correspond to A and B, respectively). Abbreviations: 8- oxo-G: 8-Dihydro- 8-oxoguanine

**A****B**

**Fig. S4. Relative MITF mRNA levels after MITF silencing or overexpression.** (A) Comparison of MITF mRNA levels following MITF gene knockdown compared to MITF control. (B) Comparison of MITF mRNA levels after MITF gene overexpression compared to MITF control. Non-targeting siRNA (siControl) and an empty plasmid (E.P.) were used as negative controls. All mRNA levels were normalized to RPL11. Values represent the mean  $\pm$  SEM. \* $p < 0.05$  \*\* $p < 0.01$ , \*\*\*\* $p < 0.0001$ . Abbreviations: MITF: Microphthalmia-associated transcription factor, E.P.: empty plasmid, O.E.: overexpression