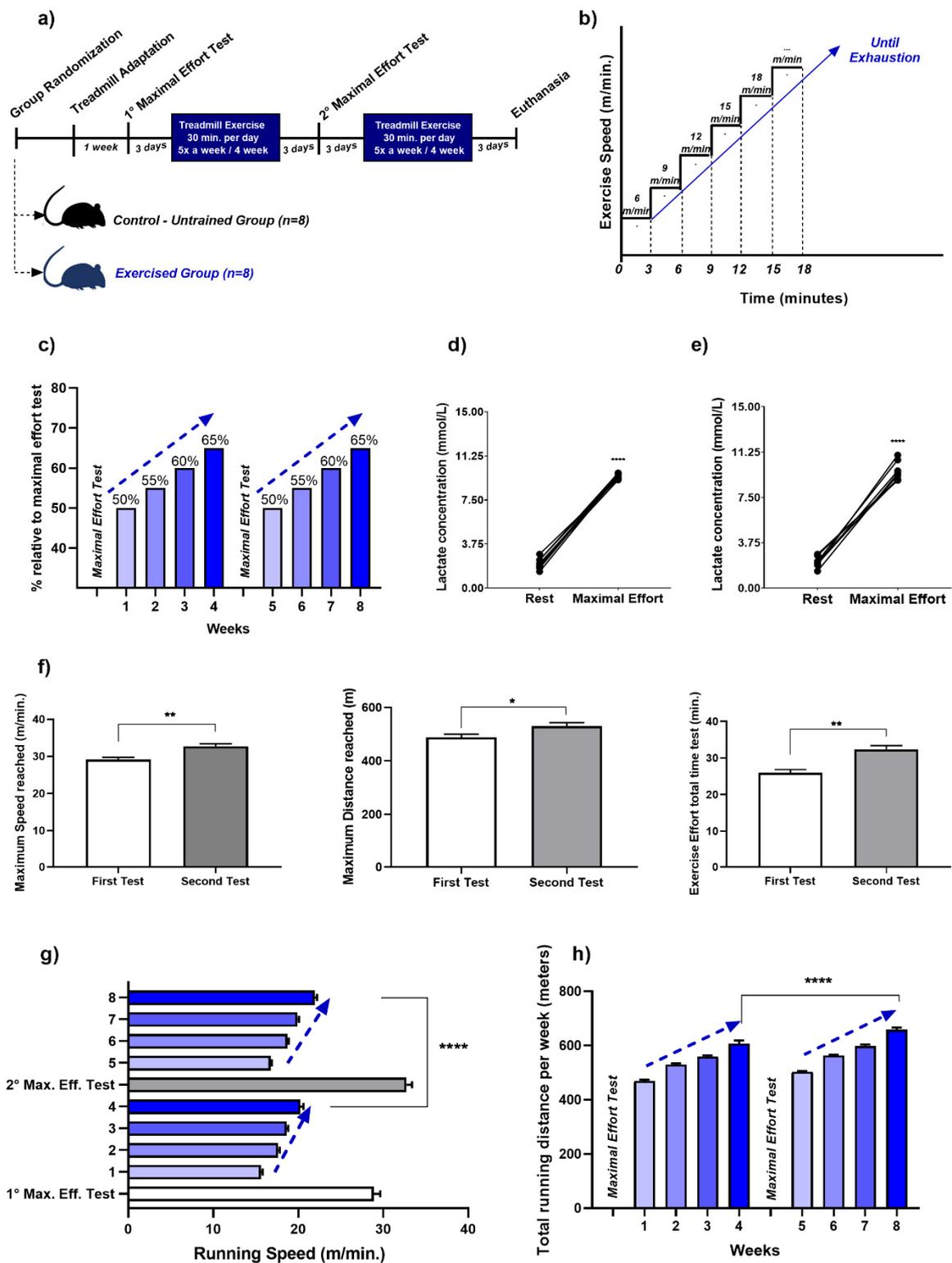


SUPPLEMENTARY MATERIAL

Supplementary Table S1. *Forward and reverse sequences of primers.*

GENE	Accession Number	SPECIES	FORWARD	REVERSE
<i>Nox2</i>	NM_023965.1	<i>Rattus norvegicus</i>	<i>CCATTACACCATTGCACATC</i>	<i>CGAGTCACAGCCACATAC G</i>
<i>Nox4</i>	NM_053524.1		<i>TCCATCAAGCCAAGATTCTGAG</i>	<i>GGTTTCCAGTCATCCAGTAGAG</i>
<i>Cat</i>	NM_012520.2		<i>CAAGCTGGTTAATGCGAATGG</i>	<i>TTGAAAAGATCTCGGAGGCC</i>
<i>Gpx1</i>	NM_030826.4		<i>AATCAGTTCGGACATCAGGAG</i>	<i>GAAGGTAAAGAGCGGGTGAG</i>
<i>Gpx2</i>	NM_183403.2		<i>GACACGAGGAAACCGAAGCA</i>	<i>GGCCCTTACAACGTCT</i>
<i>Gpx3</i>	NM_022525.4		<i>CAGCTACTGAGGTCTGACAG</i>	<i>ACTAGGCAGGATCTCCGAG</i>
<i>Sod1</i>	NM_017050.1		<i>TGTGTCCATTGAAGATCGTGTG</i>	<i>CTTCCAGCATTTCAGTCTTTG</i>
<i>Sod2</i>	NM_017051.2		<i>GGACAAACCTGAGCCCTAAG</i>	<i>CAAAAGACCCAAAGTCACGC</i>
<i>Sod3</i>	NM_012880.2		<i>GACCTGGAGATCTGGATGGA</i>	<i>GTGGTTGGAGGTGTTCTGCT</i>
<i>Nef2l2</i>	NM_031789.3		<i>TTGTAGATGACCATGAGTCGC</i>	<i>TGCCTGCTGTATGCTGCTT</i>
<i>Hmox1</i>	NM_012580.2		<i>ATCGTGCTCGCATGAACACT</i>	<i>CAGCTCCTCAAAACAGCTCAATG</i>
<i>Gclm</i>	NM_017305.2		<i>CAGTGGGCACAGGTAACC</i>	<i>AATGCAGTCAAATCTGGTGGC</i>
<i>Gclc1</i>	NM_017305.2		<i>GGTGACGAGGTGGAGTACAT</i>	<i>AACATCGCCGCCATTAGTA</i>
<i>Ppargc1a</i>	NM_031347.1		<i>CACCGCAATTCTCCCTTGTA</i>	<i>TGCGGTATTCATCCCTCTTG</i>
<i>Ppargc1b</i>	NM_176075.3		<i>TGCCACAACCCAACAGTCTCA</i>	<i>AGCAGTCTCCAGCAGCCCAAAG</i>
<i>Vegfa</i>	NM_031836.3		<i>ACCACAGTCCATGCCATCAC</i>	<i>TCCACCACCTGTTGCTGTA</i>
<i>Srebf1</i>	NM_001276707.1		<i>GGAGCCATGGATTGCACATT</i>	<i>GCTTCCAGAGAGGAGCCCAG</i>
<i>Ppargc1a</i>	NM_031347.1		<i>GGTGAAACTCTGGGAGATCCTCC</i>	<i>AGCAACCATTGGGTCAGCTCT</i>
<i>Il18</i>	NM_031512.2		<i>GCAATGGTCGGACATAGTT</i>	<i>AGACCTGACTTGGCAGAGGA</i>
<i>Il6</i>	NM_012589.2		<i>TCTCTCCGAAGAGACTTCCA</i>	<i>ATACTGGTCTGTTGTGGG</i>
<i>Tnf</i>	NM_012589.2	<i>ACCACGCTCTTCTGTCTACTG</i>	<i>CTTGGTGGTTTGCTACGAC</i>	
<i>Ccl2</i>	NM_031530.1	<i>GTGCTGACCCCAATAAGGAA</i>	<i>TGAGGTGGTTGTGGAAAAGA</i>	
<i>Cdkn2a</i>	NM_031550.2	<i>TCCGAGAGGAAGGCGAACTC</i>	<i>GCTGCCCTGGCTAGTCTATCTG</i>	
<i>Cdkn2d</i>	NM_130812.4	<i>ACCCCAAGTGAGGGTTTCT</i>	<i>GATCCTCTCTGGCCTCAACA</i>	
<i>Actb</i>	NM_031144.3	<i>CAC TTTCTACAATGAGCTGCG</i>	<i>CTGGATGGCTACGTACATGG</i>	



Supplementary Figure S1. Experimental design to aerobic exercise capacity evaluation and periodization of the EX-group. a) Timeline of interventions; b) Graphic representing the maximum speed test protocol; c) Progressive periodization model of aerobic training over the weeks; d) Lactate plasma concentration at rest and maximum post-first and second; (e) maximal speed tests measured in plasma by spectrophotometry using the BioClin® Kit; f) Comparison between the physical performance of animals in the first and last test; g) Maximal speed reached in the 65% of the maximal capacity of AE program; h) Total running distance reached in the 65% of the maximal capacity of AE program. Data were expressed as the mean \pm standard error of the mean (n = 8/group). * $p < 0.05$; ** $p < 0.01$; **** $p < 0.0001$.