

Table S1. Analysis of variance for mitochondrial and nuclear effects on organismal traits

Dependent variable ¹	Factor	Df	F value	P value
Development time (mean emergence time) N=10-20 replicate vials	mtDNA	2	152.2	<2.2e-16
	Nuclear	1	216.5	<2.2e-16
	Sex	1	6.631	0.0109
	mtDNA*nuclear	2	169.5	<2.2e-16
	mtDNA*sex	2	0.041	0.9601
	Nuclear*sex	1	6.309	0.0130
	mtDNA*nuclear*sex	2	0.072	0.9307
	Residuals	164		
Development time (day of first emergence) N=10-20 replicate vials	mtDNA	2	185.7	<2.2e-16
	Nuclear	1	465.2	<2.2e-16
	Sex	1	7.608	0.006
	mtDNA*nuclear	2	105.8	<2.2e-16
	mtDNA*sex	2	0.761	0.469
	Nuclear*sex	1	0.579	0.448
	mtDNA*nuclear*sex	2	0.025	0.976
	Residuals	164		
Fecundity (total eggs over ten days) N=7-14 replicate females	mtDNA	2	19.38	3.22e-07
	Nuclear	1	299.7	<2.2e-16
	mtDNA*nuclear	2	7.325	0.0014
	Residuals	60		
Bristle length (mm) N=10-14 individuals per sex per genotype	mtDNA	2	1372	<2.2e-16
	Nuclear	1	501.3	<2.2e-16
	Sex	1	37.03	1.30e-08
	mtDNA*nuclear	2	250.3	<2.2e-16
	mtDNA*sex	2	0.470	0.6264
	Nuclear*sex	1	0.474	0.4925
	mtDNA*nuclear*sex	2	0.264	0.7680
	Residuals	126		

¹ Data include the following genotypes: (*ore*);*OreR*, (*ore*);*AutW132*, (*simw*⁵⁰¹);*OreR*, (*simw*⁵⁰¹);*AutW132*, (*sm21*);*OreR*, (*sm21*);*AutW132*.