Mitochondrial maturation drives germline stem cell differentiation

in Caenorhabditis elegans

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SUPPORTING INFORMATION

MmPOLRMT HsPOLRMT DmMTRNAPOL CeRPOM-1 ScRPO41	-RPLGPRGLDWLKIHLINLTGLKKGDSLRMRLAFADEVMEEILDSADNPL -RPLGPHGLDWLKIHLVNLTGLKKKEPLRKKLAFAEEVMDDILDSADQPL -QPLGVDGLSWLKH.CHNLTGLKKRSVVAERLLYAEEIMPDILDSADNPL -QKLGEKGFDWLKLHCINLTGTMKRSSVADRAVVAAERLIPLMDSARDPL GKKLGPSGLKWLKIHLSNLFGFDKLP-LKDRVAFTESHLQDIKDSAENPL :***::***	JIGRKWMMEAD EPWQILACCMEVJ JIGRKWMMGAE EPWQILACCMEVJ JIGRMWMSKD EPWQILACCMEUJ NGQKWMMSD EPWQILAACVEII JIGDRWWITAD KPWQALAICFELM	AHAVRS-PDPAAY: ANAVRA-SDPAAY ANVHRC-PDPAAYI ENATRYGSDVALFI NEVMKM-DNPEEF: : : :	ISHLPVHQDGSCNG VSHLPVHQDGSCNG LSRFPIHQDGSCNG PSQLPIHQDGSCNG ISHQPVHQDGICNG	LQHYAALGRDSVGAASVNLIP LQHYAALGRDSVGAASVNLEP LQHYAALGRDBEGGRSVNLAP LQHYAALGRDNEGGVQVNLIQ LQHYAALGGDVEGAIQVNLVP
Mm.POLRMT HsPOLRMT DmMTRNAPOL CeRBOM-1 ScRPO41	SDLPQDVYREVAIQVEEFRQQDAKEGLRVAQVLEGFISRKVVK SDVPQDVYSGVAAQVEVFRRQDAQRGMRVAQVLEGFIIRKVVK SAIPQDVYSAVAALVEKSRKADAQNGLHVAEALAGFVRRKVIK SDIPNDVYSDVAQRVEQKRQQDEQSNGEDCDVARKLREALPQNVFRKVIK SDKPQDVYAHVARLVQKRLEIAAEKGDENAKILKDKIIRKVVK	QTVMTVVYGV TRYGGRLQIEKR QTVMTVVYGV TRYGGRLQIEKR QTVMTTVYGV TRYGARLQIARQ QTVMTTVYGV TRYGAVLQIKRQ QTVMTNVYGV TYVGATFQIARQ	LRELSDFPQEFVW LRELSDFPQEFVW LKDIDDFPKDWVW LKAMD-IPGEDAA LSPIFDDRKE-SL	EASHYLVRQVFKSI EASHYLVRQVFKSI PASTYLTIKTFESI IFARYLARKTFASI DFSKYLTKHVFSAI : **. :.* ::	QEMFTSTRAIQHWLTESANLI QEMFSGTRAIQHWLTESARLI REMFTSTREIQDWFTECARLI NDAFTSSMALKDWFRLIAKGS RELFHSAHLIQDWLGESAKRI *.
Mm.POLRMT HsPOLRMT DmMTRNAPOL CCRPOM-1 ScRPO41	SHAGWEVEWVTPLGIPIIQPYHRESKVQVKGG SHMGSVVEWVTPLGVEVIQPYRLDSKVKQIGG SGVCSQNVEWVTPLGLEVVQPYNRQ-EMKHSPF SDLMKTVEWITPLGLEVVQPYCKL	-LQSIILISSVDESQKPNILKQ GIQSIIYIINGDISRKPNIRKQ RSGFKVSANMPMDLYERPNILKQ VERKGKLILAPVPMKQ LQTVFISDPFAVNPVNARRQ :*	ONGFPPNFIHSLD: ONGFPPNFIHSLD: ONAFPPNFIHSLD: /DAFPPNFVHSLD: CAGLPPNFIHSLD: .:****:***	SSHMMLTALHCYRK SSHMMLTALHCYRK SSHMMLTSLHCERQ STHMMLTSLNCAQR ASHMLLSAAECGKQ	GLIFVSVHDCFWIHAADIFIM GLIFVSVHDCYWIHAADVSVM GIIFVSVHDCFWIHANTVPEL GIIFAAVHDCFWIHANSVDQM GLDFASVHDSYWIHASDIDIM
Mm.POLRMT NEVCREQFVRLHSQPILEDLAKFLKKRFCSVSS					
			Identities	Conservative substitutions	
MmPOLRMT HsPOLRMT DmMTRNAPOL CCRPOM-1 ScRPO41	VIKLQEILQSLPKTGIFDLGQVIRSTYFFS ASQLKEILQAVPKPGAFDLEQVKRSTYFFS KRQLNRILKQMPQKGDFDLENVLDSVYFFS FQKYSEIFTANIEHGDLDIEKVKDSVYFFS SVLLPLRLPEIPPKGDFDVTVLRNSQYFFS	ScRPO41	39%	56%	
		DmMTRNAPOL	53%	65%	
		MmPOLRMT	48%	63%	
		HsPOLRMT	47%	62%	

Figure S1: The catalytic domain of mitochondrial RNA polymerases is highly conserved. CLUSTALW alignment of the catalytic domains from five model organisms (*S. cerevisiae*, *D. melanogaster*, *M. musculus*, *H. sapiens*) reveals high conservation of POLRMTs in evolutionarily diverse species. The table displays the percentage of identical residues as well as conservative substitutions in pair-wise comparisons of *C. elegans* mitochondrial RNA polymerase (RPOM-1) and its identified homologues.



Figure S2: Reduction of mitochondrial transcription compromises mitochondrial activity. A) Quantification of *hmg-5*, *tfbm-1* and *rpom-1* mRNA levels upon feeding with RNAi expressing bacteria. Mitochondrial function following genetic inhibition of *hmg-5*/TFAM, *rpom-1*/POLRMT and *tfbm-1*/TFB1M was assayed by staining with potential-based dyes, such as TMRE (B), Mitotracker ROS (D) and by measuring ATP production (C). E) RPOM-1 depletion reduces mitochondrial DNA content, at levels comparable to HMG-5 depletion. One-way ANOVA was used for multiple comparisons (n=40; ****P* < 0.001). Unpaired *t*-test was used for pairwise comparisons (n=40; ****P* < 0.001). Error bars, s.e.m.



Figure S3: The somatic gonad and sperm number are not affected by RPOM-1 depletion. The gonad sheath and the distal tip cell do not display any observable morphological defect following RPOM-1 depletion, as indicated by the p_{lim-7}GFP (A) and p_{lag-2}MYR::GFP (B) transcriptional reporters, respectively. C) Confocal images of the proximal gonad arm upon treatment with control and *rpom-1(RNAi)*. sp stands for sperm and -1 denotes the most proximal oocyte. D) Quantification of sperm nuclei per gonad in day 2 adult animals upon treatment with control and *rpom-1(RNAi)*. Images were acquired using a X40 objective lens. Scale bars, 20μm.



Figure S4: RPOM-1-depleted gonads are sensitive to mild heat stress. *rpom-1* knockdown in animals subjected to mild heat stress(25°C) is detrimental for gonads, as monitored with DAPI staining in fixed animals (A) and with a HIS-72::GFP nuclear reporter *in vivo* (B). The red stars highlight dead corpses in the proximal gonad arm. Images were acquired using a X40 objective lens. Scale bars, 20µm.



Figure S5: RPOM-1 is broadly expressed in *C. elegans* **somatic tissues and localizes in mitochondria.** RPOM-1 is expressed in various somatic tissues, such as the intestine (A), muscles (B), in neurons of the nerve ring (C), as well as the vulva (D). E) Confocal image of RPOM-1::GFP translational reporter animals stained with TMRE (Tetramethylrhodamine, ethyl ester, perchlorate). The expression pattern is reminiscent of proteins localized in the mitochondrial matrix. Images were acquired using a X40 objective lens. Scale bars, 20µm.









7



Figure S8: The Perceval reporter responds to changes in ATP production. Representative, lowmagnification images of transgenic animals expressing the Perceval ATP sensor at D3 of adulthood. Perceval is expressed in the oocytes located in proximal gonad arm, as well as in early embryos. Treatment for two days with 50nM Antimycin A, an inhibitor of ETC complex III and mitochondrial ATP production, reduces the fluorescence the Perceval sensor emits. Images were acquired using a X20 objective lens. -1 denotes the most proximal oocyte. Ds; distal, pr; proximal. Scale bars, 100µm.

8



Figure S9: RPOM-1 is required for mitochondrial maturation in the proximal gonad arm. DIOC6(3) staining of D1 adult hermaphrodites treated with control or *rpom-1(RNAi)* from hatching. Arrowheads highlight tubular mitochondria in the proximal gonad arm. The white dashed lines indicate dorsal oocyte membranes. -1 denotes the most proximal oocyte. Images were acquired using a X40 objective lens. Scale bars, 20µm.



Figure S10: POLRMT and MTCO1 expression increase during mammalian stem cell differentiation.

Staining of J1 mammalian cells with antibodies against POLRMT and MTCO1. Low expression of both proteins can be detected at the core of the stem cell colony (white circle) and higher in the periphery. Upon LIF removal, as the cells progress towards differentiation, an elevation of the expression of both proteins can be observed. In parallel, mitochondria with elongated shape (red rectangle, arrowheads) can be visualized. Images were acquired using a X40 objective lens. Scale bars, 20µm.