## *Arntl* deficiency in myeloid cells reduces neutrophil recruitment and delays skeletal muscle repair

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Supplementary Figure S1. *Arntl* deletion from myeloid cells does not significantly change skeletal muscle phenotype pre-injury.

a; Bodyweight of untreated Ctrl (n = 8) and cKO (n = 8) mice. b; TA muscle mass per body weight of untreated Ctrl (n = 14) and cKO (n = 13) muscles. c,d; Hematoxylin/eosin staining of untreated Ctrl and cKO TA muscles. Representative images from n = 4 Ctrl and cKO mice. e; Mean fiber diameter of untreated Ctrl (n = 6) and cKO (n = 8) TA muscles. f; Fiber diameter distributions of Ctrl (n = 6) and cKO (n = 8) untreated TA muscles. Data are expressed as the means  $\pm$  SEM. \* p < 0.05 with unpaired two-tailed Student's *t*-test for (a, b, e) or two-way ANOVA with Bonferroni's multiple comparisons test (f). Scale bar = 100  $\mu$ m.



## Supplementary Figure S2. *Arntl* deletion from myeloid cells does not significantly change the number of regenerating fibers.

The number of regenerating fibers in Ctrl (n = 6) and cKO (n = 5) TA muscles on day 7 postinjury. Data are expressed as the means  $\pm$  SEM. not significant (n.s.); *p* > 0.05 with unpaired two-tailed Student's *t*-test.



Supplementary Figure S3. *Arntl* deletion from macrophages increases the expression of *Il6*. a; RT-PCR assessment of interleukin 6 (*Il6*) expression in untreated and treated Ctrl (n = 3) and cKO (n = 3) peritoneal macrophages with Kdo2-lipid A (KLA) for 6h. b; Expression level of IL6 protein in untreated and treated Ctrl (n = 3) and cKO (n = 3) peritoneal macrophages with KLA for 24h. c; RT-PCR assessment of *Il6* expression in sorted myeloid cells (CD45<sup>+</sup>CD11b<sup>+</sup>) from Ctrl (n=3) and cKO (n=4) muscles on day 3 after injury. Data are expressed as the means  $\pm$  SEM. \* *p* < 0.05 with unpaired two-tailed Student's *t*-test for (b, c) or two-way ANOVA with Bonferroni's multiple comparisons test for (a).



Supplementary Figure S4. Culture medium conditioned by *Arntl*-depleted macrophages significantly inhibited differentiation of muscle stem cells.

a; Immunostaining of skeletal muscle myosin (green) and Hoechst staining of nuclei (blue) from wild-type skeletal muscle stem cells cultured in conditioned medium (CM) from Ctrl and cKO peritoneal macrophages. Representative images from n = 3 Ctrl and cKO wells. b; Quantification of the fusion index from the image in (a). Ctrl (n = 3) and cKO (n = 3) wells. Data are expressed as the mean  $\pm$  SEM. \* p < 0.05 with unpaired two-tailed Student's *t*-test. Scale bar = 200 µm.



Supplementary Figure S5. *Arntl* deletion from myeloid cells does not significantly change the number of circulating neutrophils and the migration-related gene expressions in monocytes.

a; Quantification of neutrophils (CD45<sup>+</sup> CD11b<sup>+</sup> Ly6G<sup>+</sup>) in bone marrow cells from Ctrl (n = 5) and cKO (n = 5) mice left untreated. b; Quantification of neutrophils (CD45<sup>+</sup> CD11b<sup>+</sup> Ly6G<sup>+</sup>) in blood from Ctrl (n = 5) and cKO (n = 5) mice left untreated. c; RT-PCR assessment of *Arntl* expression in neutrophils (CD45<sup>+</sup> CD11b<sup>+</sup> Ly6G<sup>+</sup>) sorted from untreated bone marrow cells from

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Ctrl (n = 5) and cKO (n = 4) mice. d; RT-PCR assessment of chemokine (C-C motif) receptor 2 (*Ccr2*), chemokine (C-X3-C motif) receptor 1 (*Cx3cr1*), and *Arntl* expression in monocyte (CD45<sup>+</sup> CD11b<sup>+</sup> Ly6G<sup>-</sup> Siglec-F<sup>-</sup> Ly6C<sup>hi</sup>) sorted from bone marrow cells from Ctrl (n = 5) and cKO (n = 4) mice left untreated. Data are expressed as the means  $\pm$  SEM. \* *p* < 0.05 with unpaired two-tailed Student's *t*-test.



Supplementary Figure S6. *Arntl* deletion from myeloid cells does not significantly change the expression of chemokines related to neutrophil and monocyte migration in skeletal muscle.

a; RT-PCR assessment of chemokine (C-X-C motif) ligand 1 (*Cxcl1*) and chemokine (C-X-C motif) ligand 2 (*Cxcl2*) expressions in the TA muscles from Ctrl (n = 7) and cKO (n = 6) muscles on day one after muscle injury. b; RT-PCR assessment of (C-C motif) receptor 2 (*Ccl2*) and chemokine (C-X3-C motif) ligand 1 (*Cx3cl1*) expressions in the TA muscles from Ctrl (n = 7)

and cKO (n = 6) muscles on day 2 after muscle injury. Data are expressed as the means  $\pm$  SEM. \* p < 0.05 with unpaired two-tailed Student's *t*-test.



1 : Arntl<sup>flox/flox</sup> 2 : Lyz2Cre<sup>+/-</sup> Arntl<sup>flox/flox</sup>

## Supplementary Figure S7. PCR confirms the Arntl deletion in bone marrow-derived

macrophages (BMDMs) from myeloid cell-specific Arntl knockout mice.

The image of the original blots of the experiment corresponds to Figure 1b.



Supplementary Figure S8. Western blots confirm the *Arntl* deletion in BMDM and peritoneal exudate cells (PECs) from myeloid cell-specific *Arntl* knockout mice.

a; Image of the original blots of the first experiment corresponds to Figure 1c. The membrane hybridized with ARNTL antibody was reprobed with tubulin beta (TUBB) antibody. b; Image of the original blots of the second experiment corresponds to Figure 1c.