

Supplementary materials S1-S4

Suppression of pyruvate dehydrogenase kinase by dichloroacetate in cancer and skeletal muscle cells is isoform-specific and partially independent of HIF-1 α

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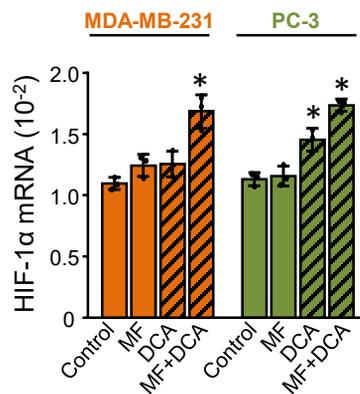
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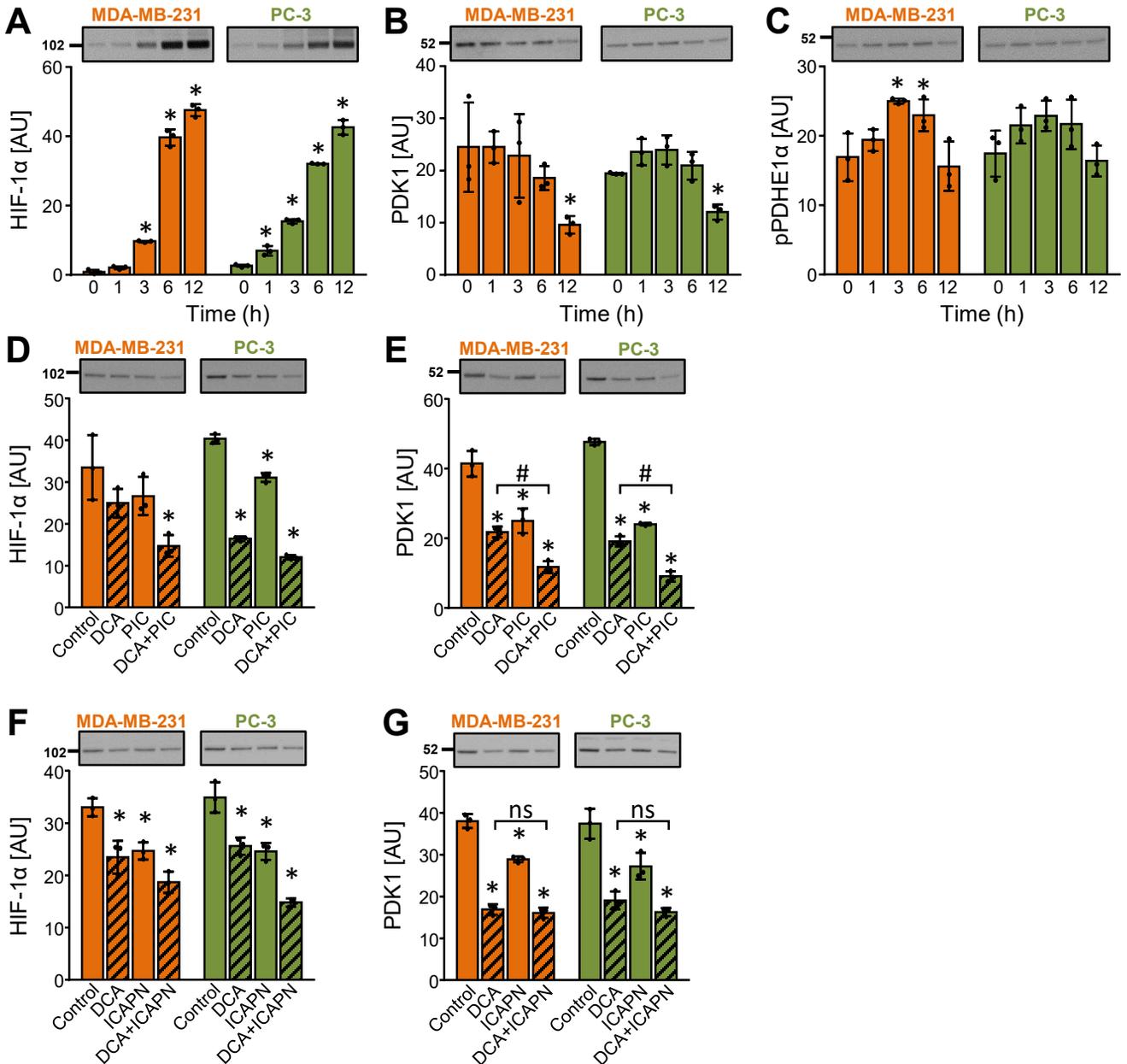
Supplementary figures S1-S3

Supplementary Figure S1



Supplementary figure S1 (supplementary figure to Figure 5): Inhibition of the mTOR pathway does not contribute to the DCA-induced suppression of PDK1. MDA-MB-231 and PC-3 cells were treated with 5 mM metformin (MF), 10 mM DCA, or both in serum-free RPMI medium for 24 hours. The mRNA expression of HIF-1 α was measured with qPCR (endogenous control: cyclophilin). Results are means \pm SD of one experiment performed in triplicate for each cell line ($n = 3$). * $p < 0.05$ vs. Control.

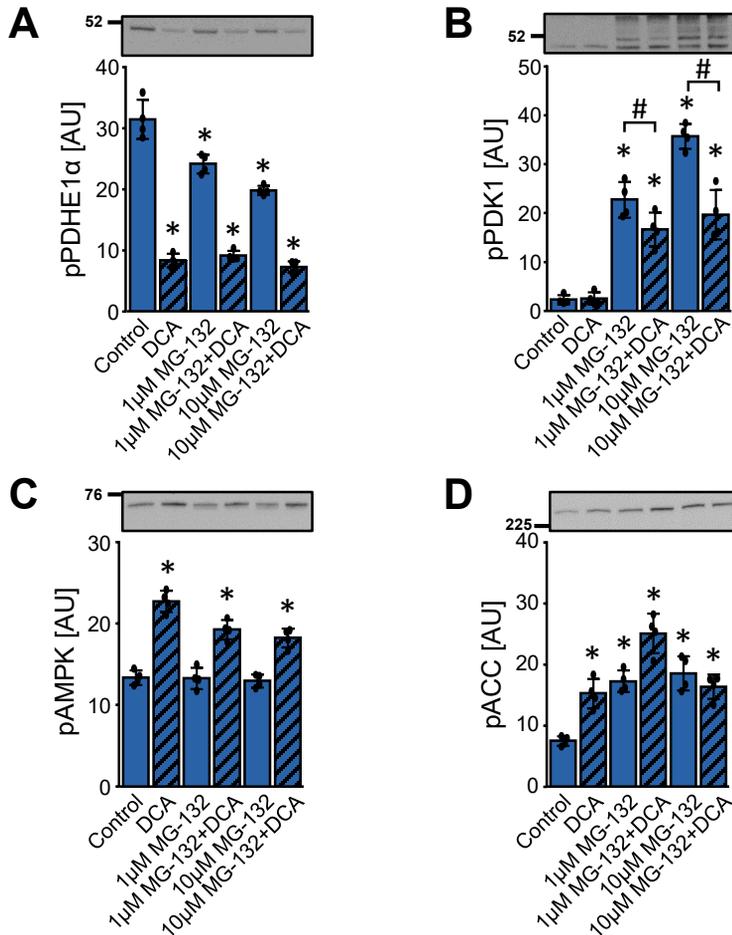
Supplementary Figure S2



Supplementary figure S2 (supplementary figure to Figure 6): Inhibition of the proteasome or mitochondrial proteases does not prevent the DCA-induced suppression of PDK1 in cancer cells.

(A–C) Cells were treated with MG-132 (10 μ M) for 12 hours in serum-free RPMI medium. (D,E) Cells were treated with DCA (10 mM) and/or the protease inhibitor cocktail (PIC, 1:600) in the RPMI medium without serum (PC-3) or with 10% FBS (MDA-MB-231) for 24 hours. (F,G) Cells were treated with 9 μ M (MDA-MB-231) or 45 μ M (PC-3) calpain inhibitor III (ICAPN) in serum-free RPMI medium for 24 hours. DCA was added during the final 14 hours of treatment. The protein levels of HIF-1 α , PDK1, and phospho-PDHE1 α ^{Ser293} (pPDHE1 α) were determined with immunoblotting. Numbers next to the blots indicate molecular weight markers in kDa. Results are means \pm SD of one experiment, performed in triplicate, for each cell line and each condition ($n = 3$). * $p < 0.05$ vs. 0 h or Control and # $p < 0.05$ between selected samples (as indicated).

Supplementary Figure S3

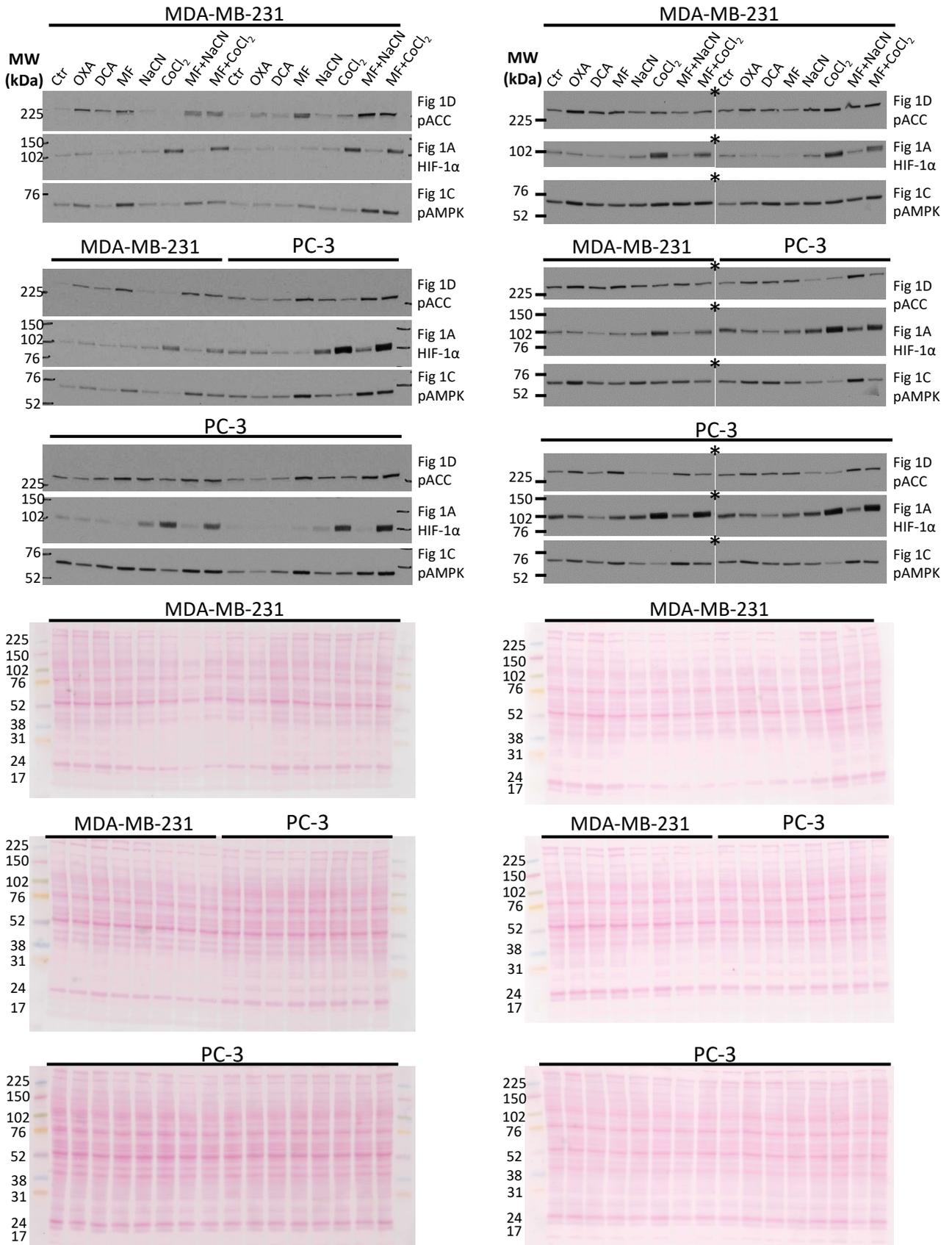


Supplementary figure S3 (supplementary figure to Figure 7): DCA reduces the abundance of PDK1 and PDK2 in L6 myotubes despite upregulation of HIF-1 α and inhibition of the proteasome. The differentiated L6 cells (myotubes) were treated with MG-132 (1 or 10 μ M) for 1 hour, which was followed by a 24-hour treatment with DCA (10 mM) and/or MG-132 (1 or 10 μ M). Immunoblotting was used to estimate the abundance of (A) phospho-PDHE1 α ^{Ser293} (pPDHE1 α), (B) phospho-PDK1^{Thr338} (pPDK1), (C) phospho-AMPK α ^{Thr172} (pAMPK), and (D) phospho-ACC^{Ser79} (pACC). Numbers next to the blots indicate molecular weight markers in kDa. Results are means \pm SD of one experiment in four replicates ($n = 4$). * $p < 0.05$ vs. Control and # $p < 0.05$ between selected samples (as indicated).

Supplement S4: Raw images

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.

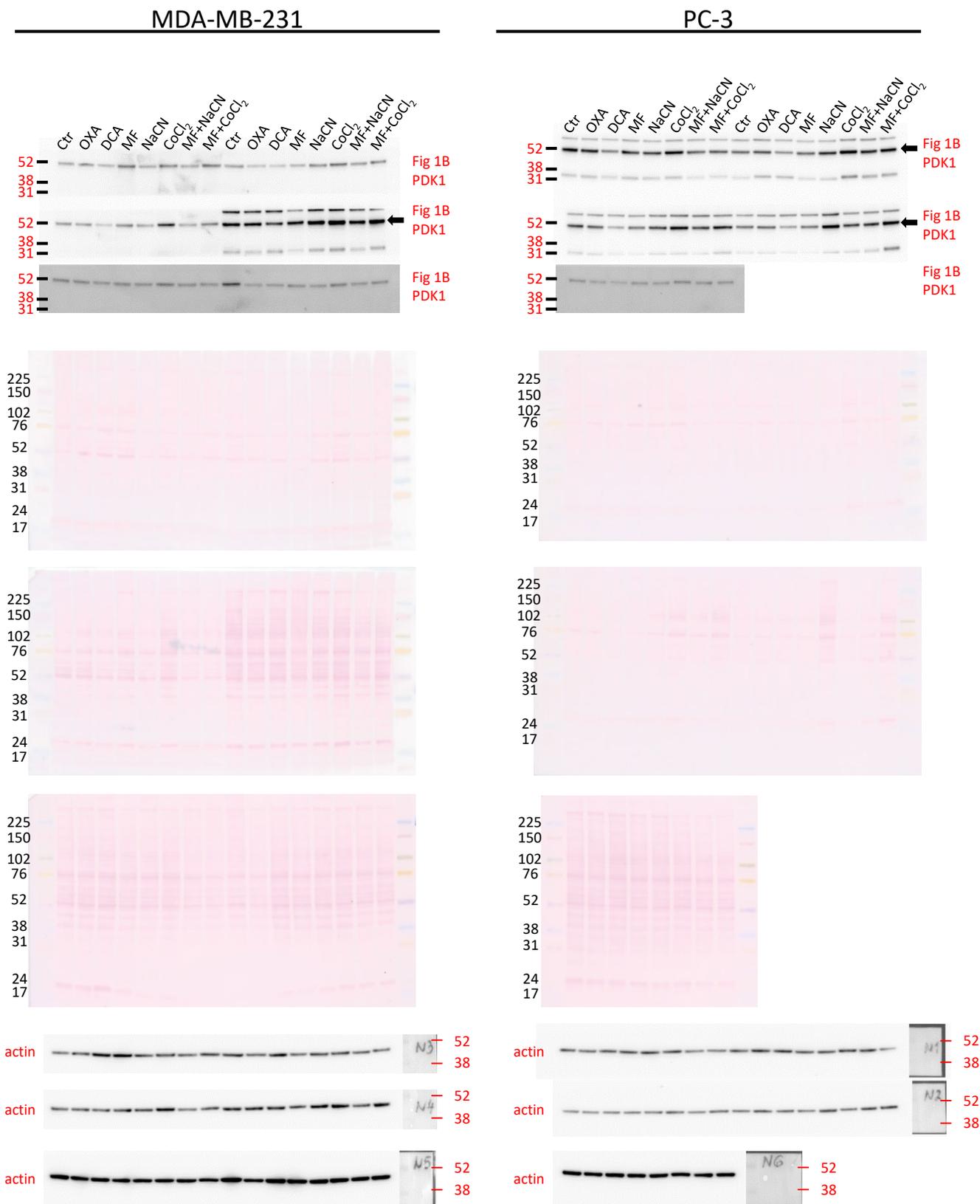


* Blots shown as originally taken on a Densitometer

Original uncropped blots presented in Figure 1 (1/2)

Immunoreactive bands were visualized with Fusion FX (Vilber) using enhanced chemiluminescence and quantified using Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

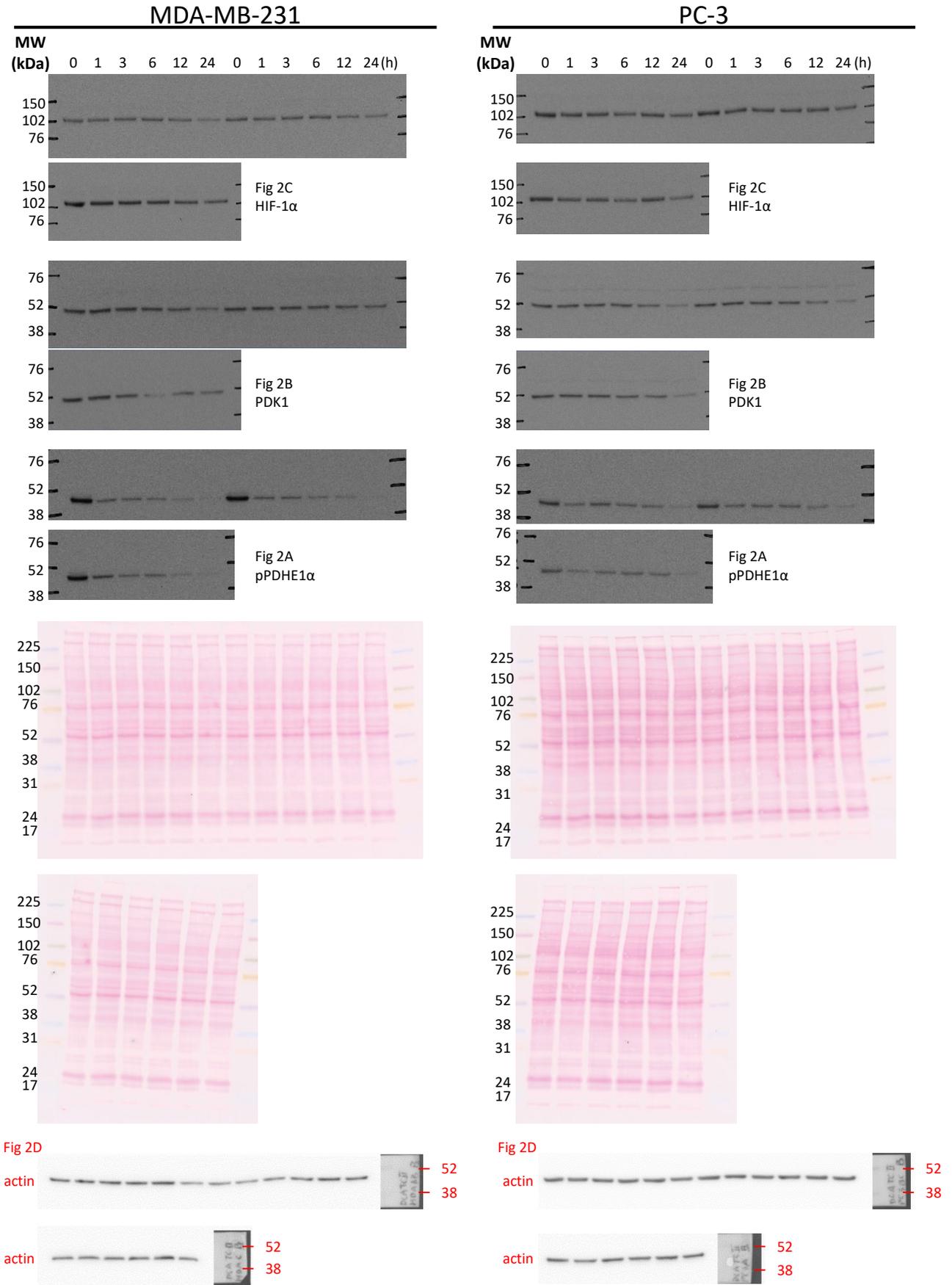
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 1 (2/2)

Immunoreactive bands were visualized on X-ray films or with Fusion FX (Vilber) using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

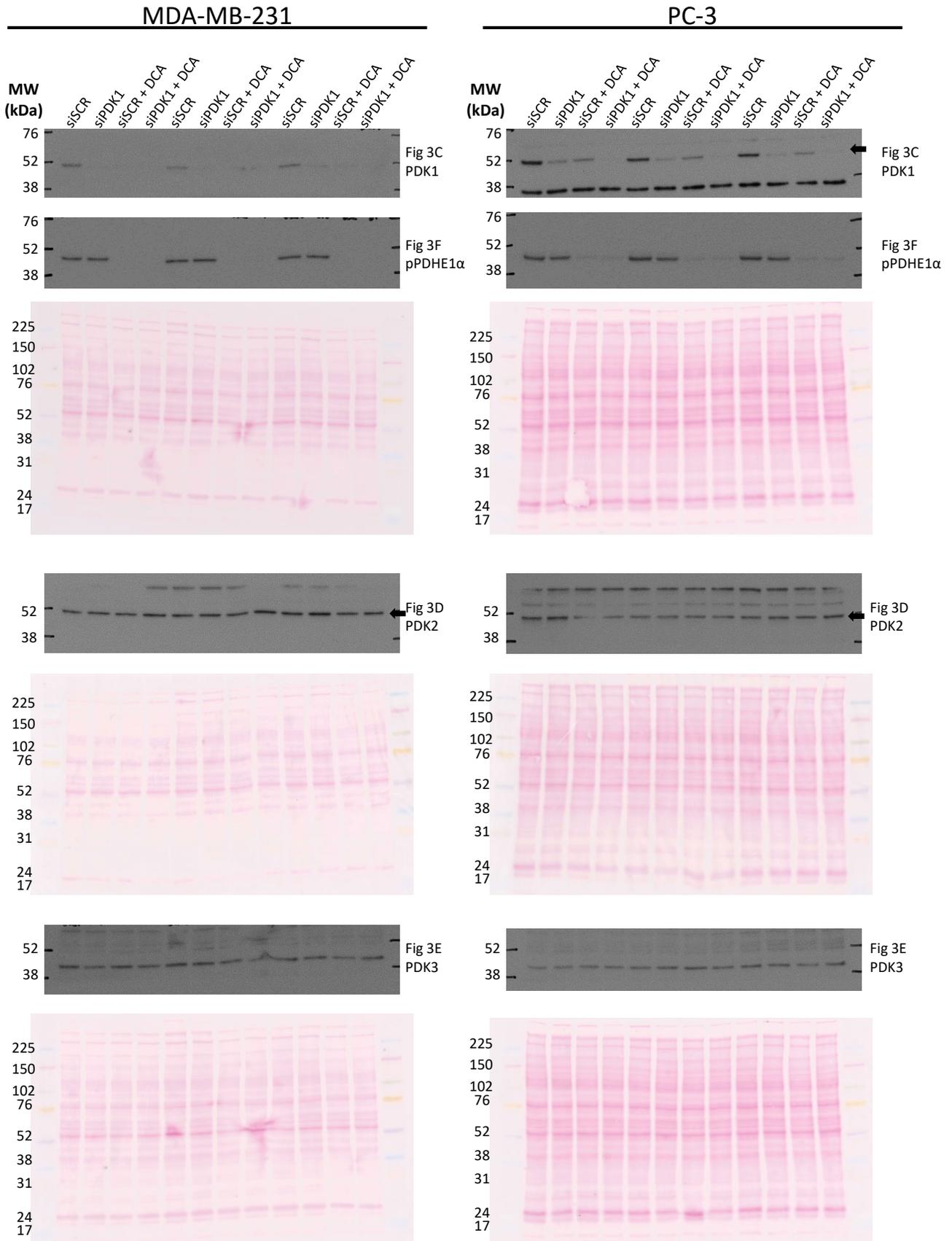
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 2

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

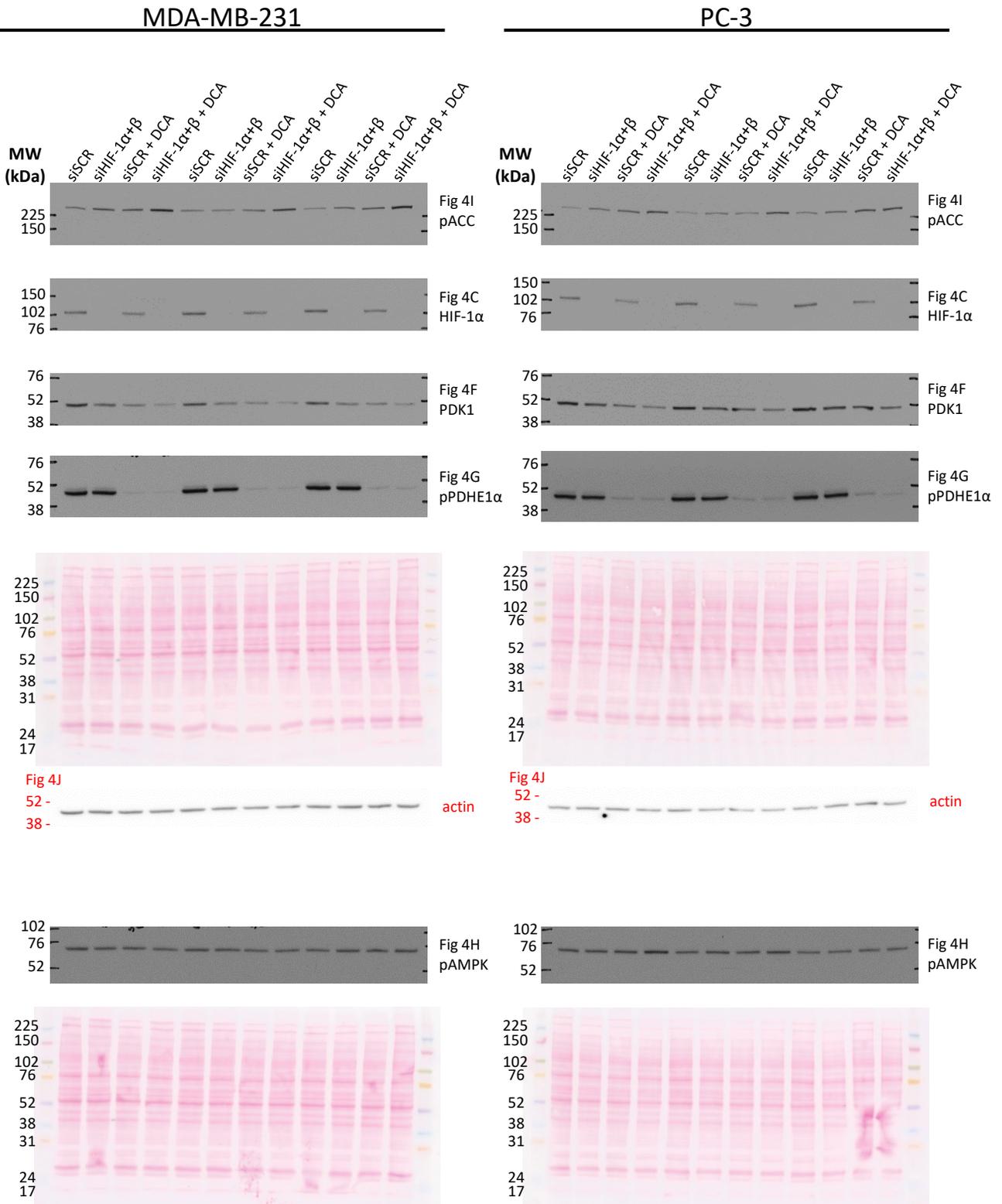
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 3

Immunoreactive bands were visualized on X-ray films or with Fusion FX (Vilber) using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.

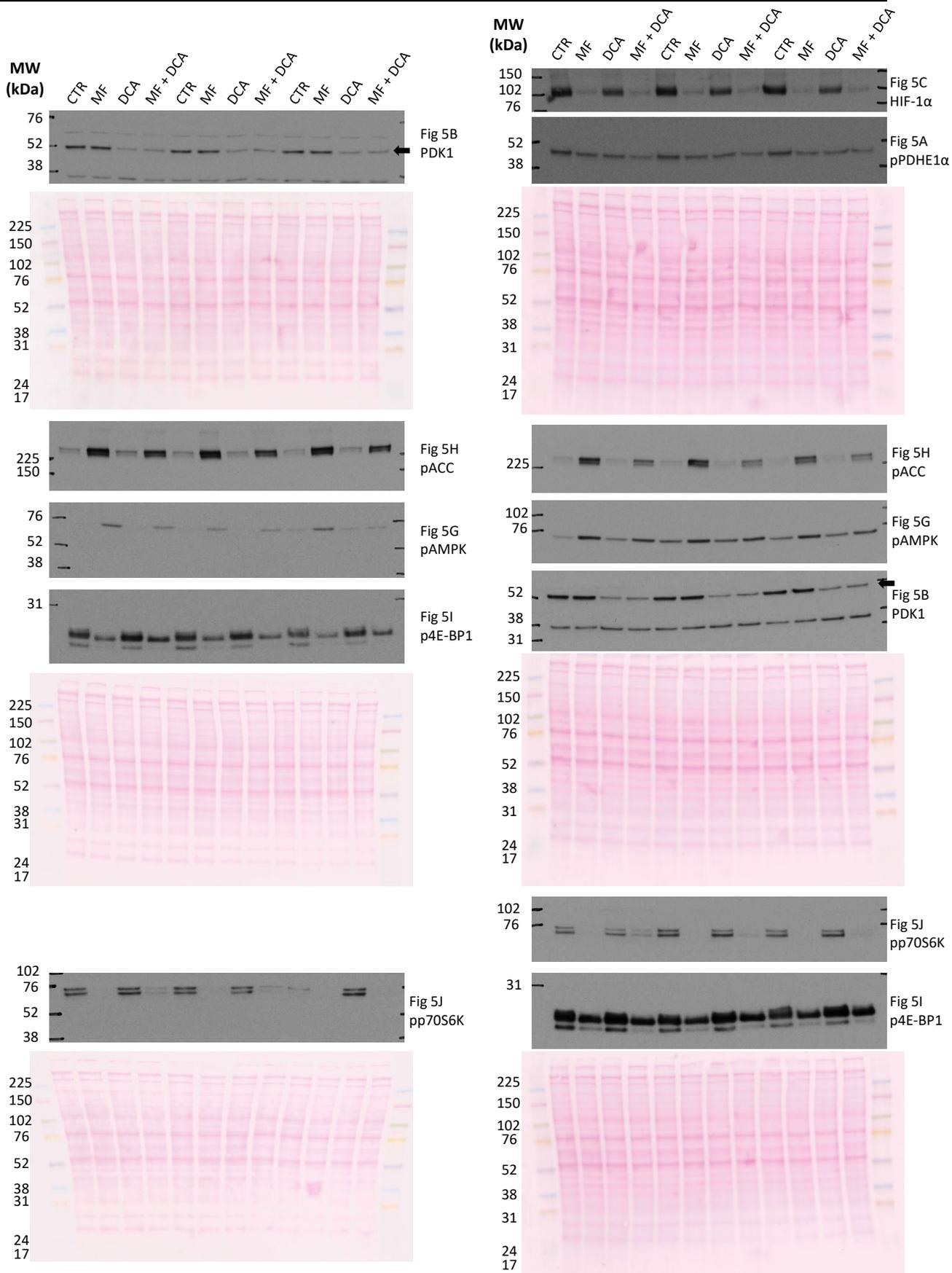


Original uncropped blots presented in Figure 4

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.

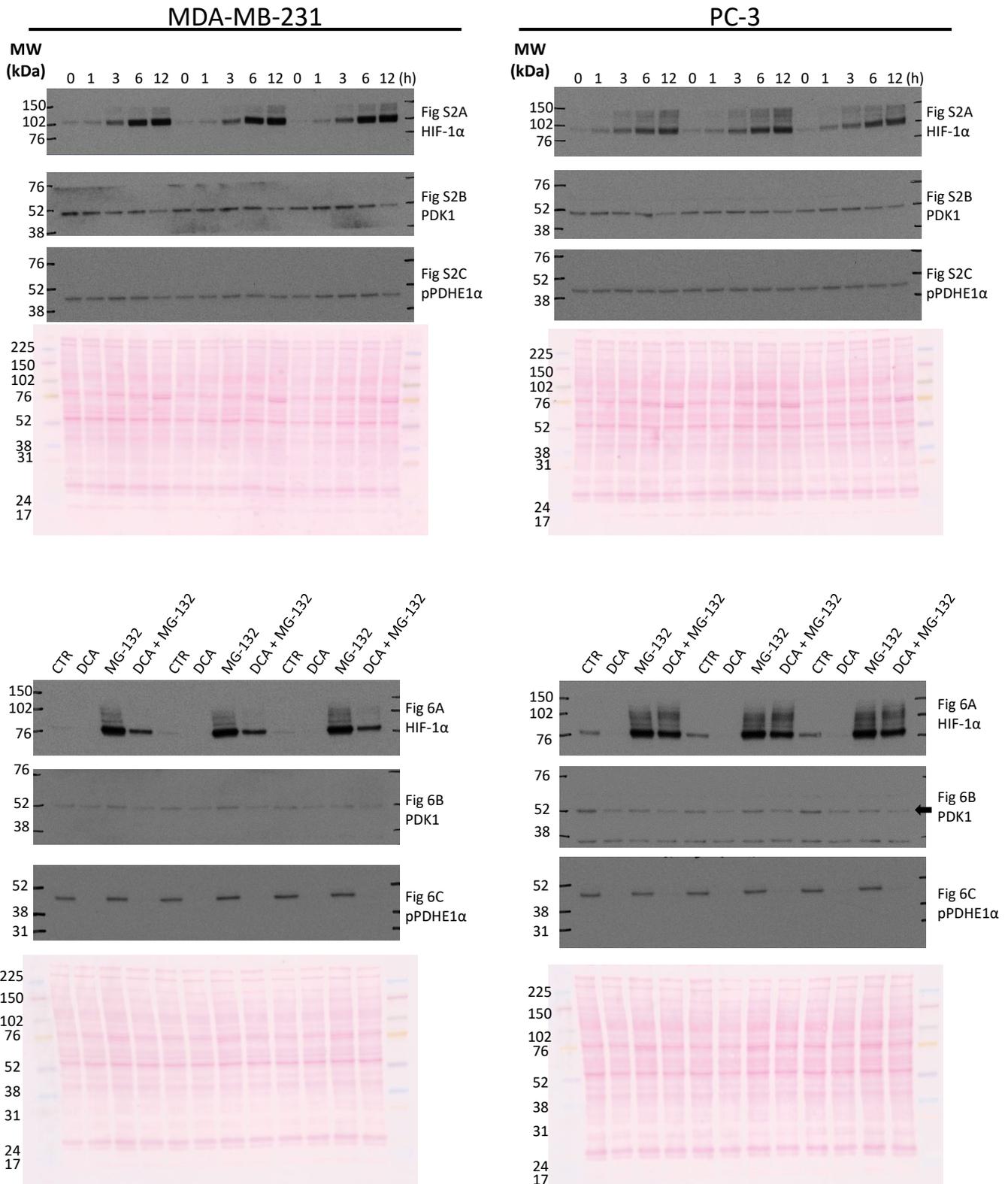
PC-3



Original uncropped blots presented in Figure 5 (2/2)

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

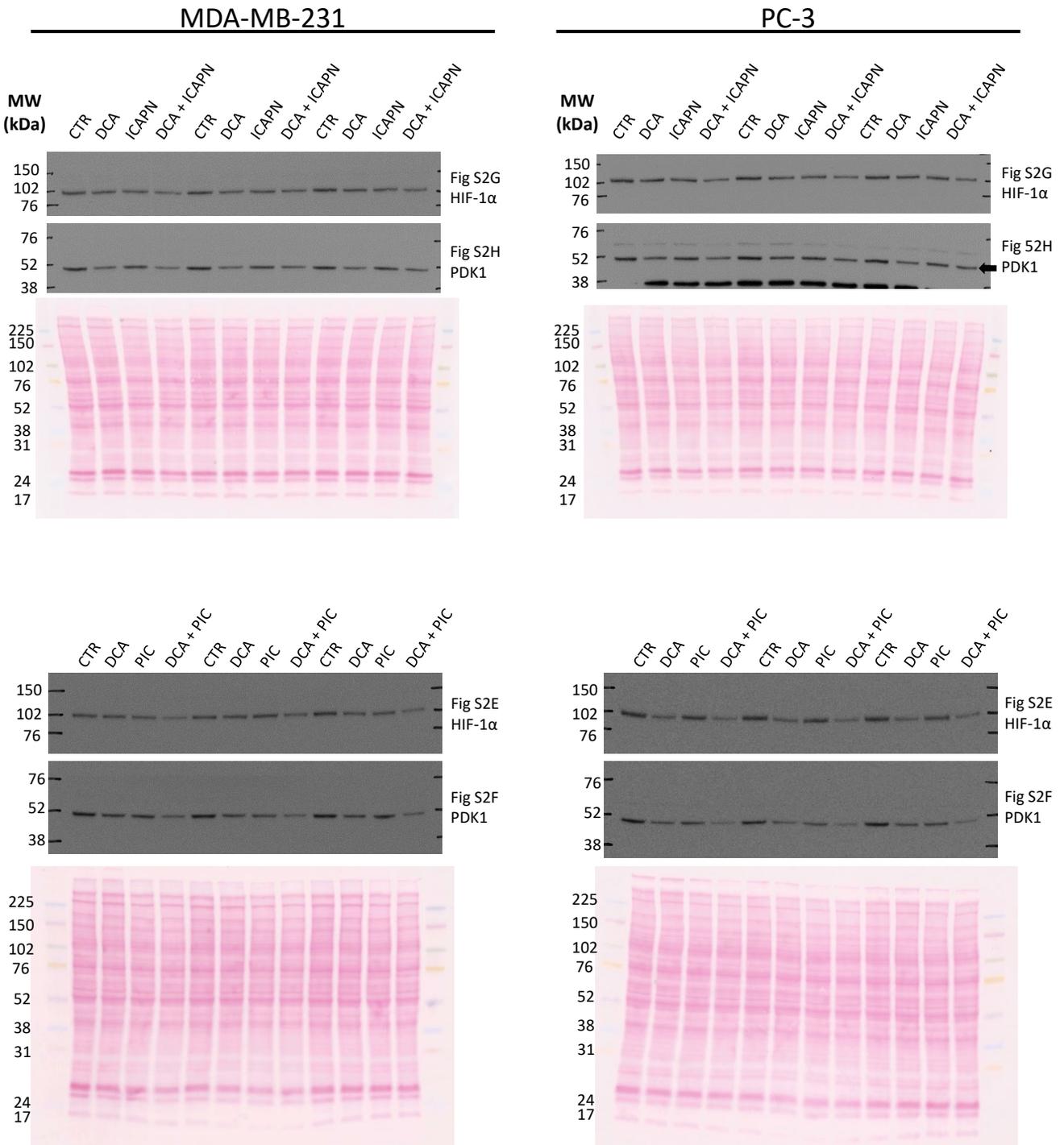
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 6 and S2 (1/3)

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

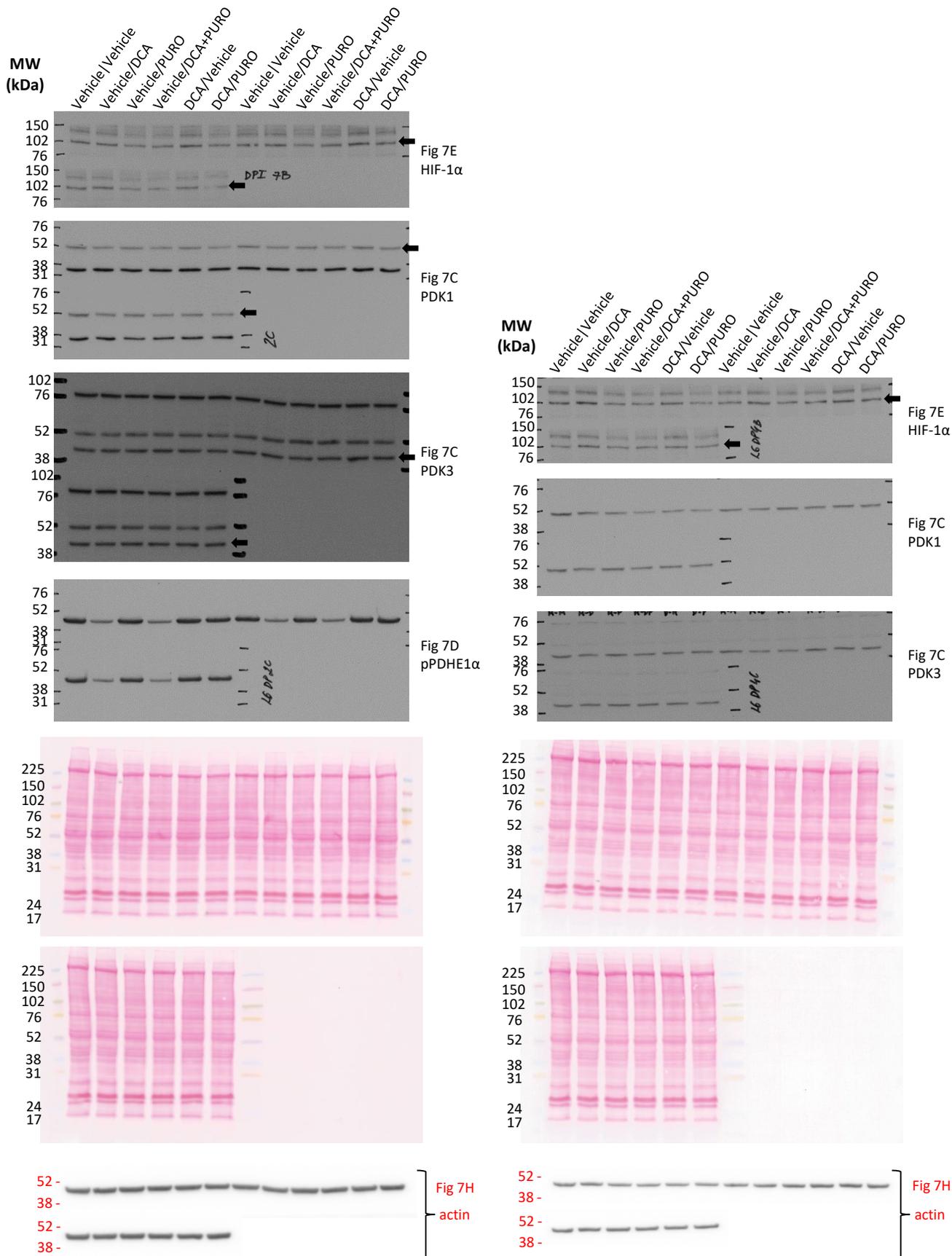
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 6 and S2 (2/3)

Immunoreactive bands were visualized on X-ray films or with Fusion FX (Vilber) using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

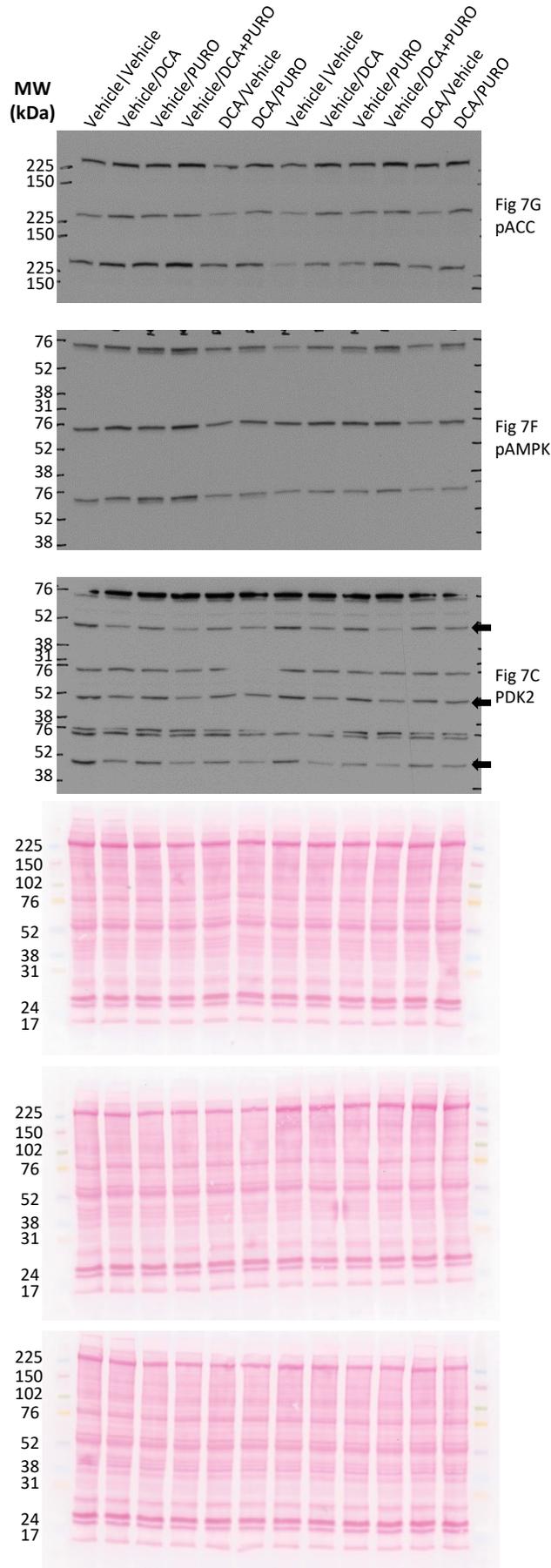
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 7 and S3 (1/4)

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

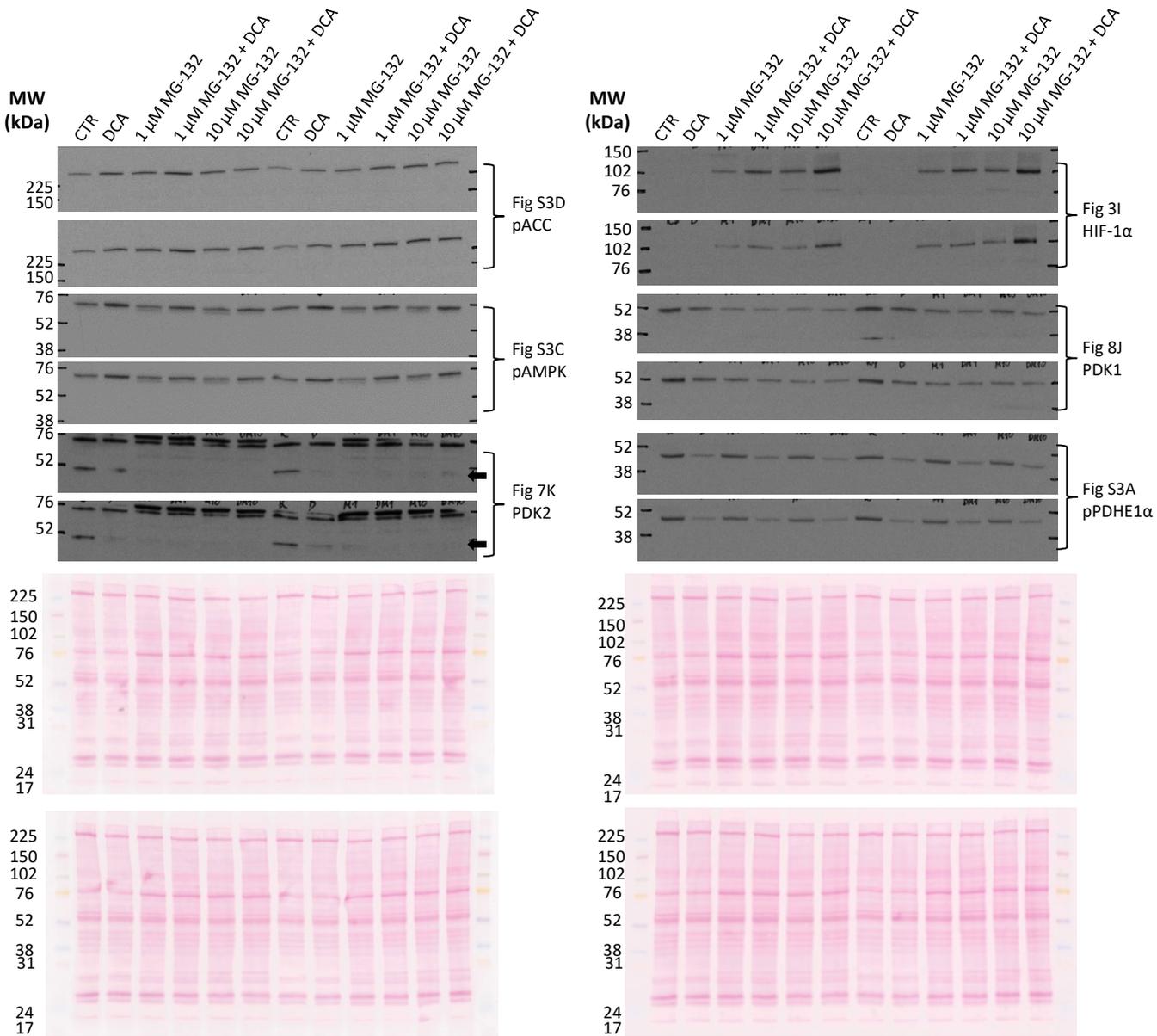
Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 7 and S3 (2/4)

Immunoreactive bands were visualized on X-ray films using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.



Original uncropped blots presented in Figure 7 and S3 (3/4)

Immunoreactive bands were visualized on X-ray films or with Fusion FX (Vilber) using enhanced chemiluminescence and quantified using GS-800 Densitometer and Quantity One 1-D Analysis Software 4.6.8. (Bio-Rad, Hercules, CA, U.S.)

Loading and transfer were evaluated by Ponceau S (0.1% (w/v) in 5% (v/v) acetic acid) staining.

