

In vitro and in vivo characterization of MCT1 inhibitor AZD3965 confirms preclinical safety compatible with breast cancer treatment

--- Supplementary data ---

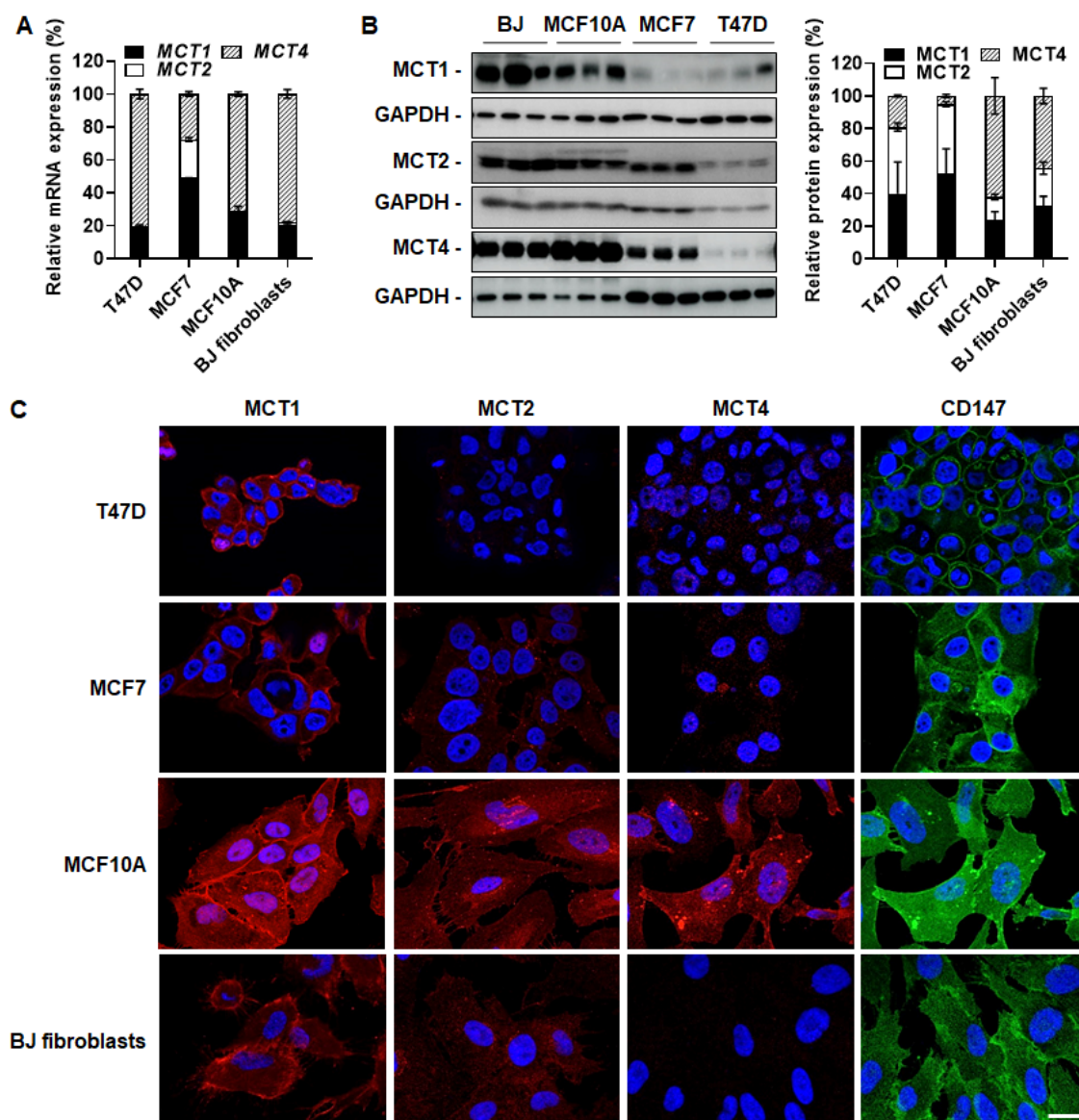


Figure S1. MCTs and CD147/basigin expression in breast-associated cells. (A-C) T47D, MCF7, MCF10A cells and BJ fibroblasts were assayed in medium containing 25 mmol/L glucose with 10 mmol/L GlutaMAX, 10% FBS and without added lactate. **(A)** Relative basal mRNA expression of *MCT1*, *MCT2* and *MCT4*. Cumulated expression served for normalization (100%) ($n = 3$). **(B)** Relative basal protein expression of *MCT1*, *MCT2* and *MCT4*. Representative western blots are shown with GAPDH as a loading control. Cumulated expression served for normalization (100%) ($n = 3$). **(C)** Representative pictures of immunocytochemical staining of *MCT1* (red), *MCT2* (red), *MCT4* (red) and CD147/basigin (green) on T47D, MCF7, MCF10A cells and BJ fibroblasts. Cell nuclei are stained in blue with DAPI. Bar = 20 μ m. All data are show as means $\pm$ SEM.

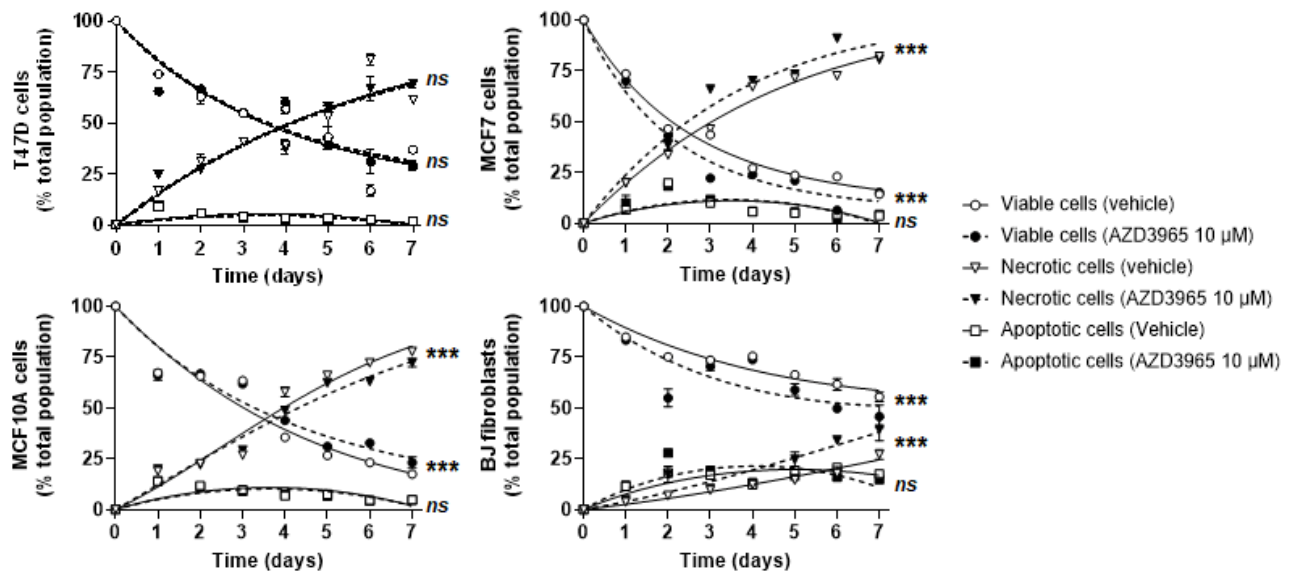


Figure S2. Long-term culture with lactate as only exogenous resource induces breast-associated cell necrosis with limited impact of additional MCT1 inhibition by AZD3965. T47D, MCF7, MCF10A cells and BJ fibroblast density was assayed in medium containing 10 mmol/L sodium *L*-lactate, no glucose, no glutamine, and 1 % FBS. On day 0, cells were treated $\pm$ 10 μ mol/L of AZD3965. Graphs show the percentage of viable cells, necrotic cells and apoptotic cells over time determined using flow cytometry after Annexin V and propidium iodide labeling ($n = 4-6$). All data are show as means $\pm$ SEM. *** $P < 0.05$, ns $P > 0.05$ comparing whole curves; by two-way ANOVA.

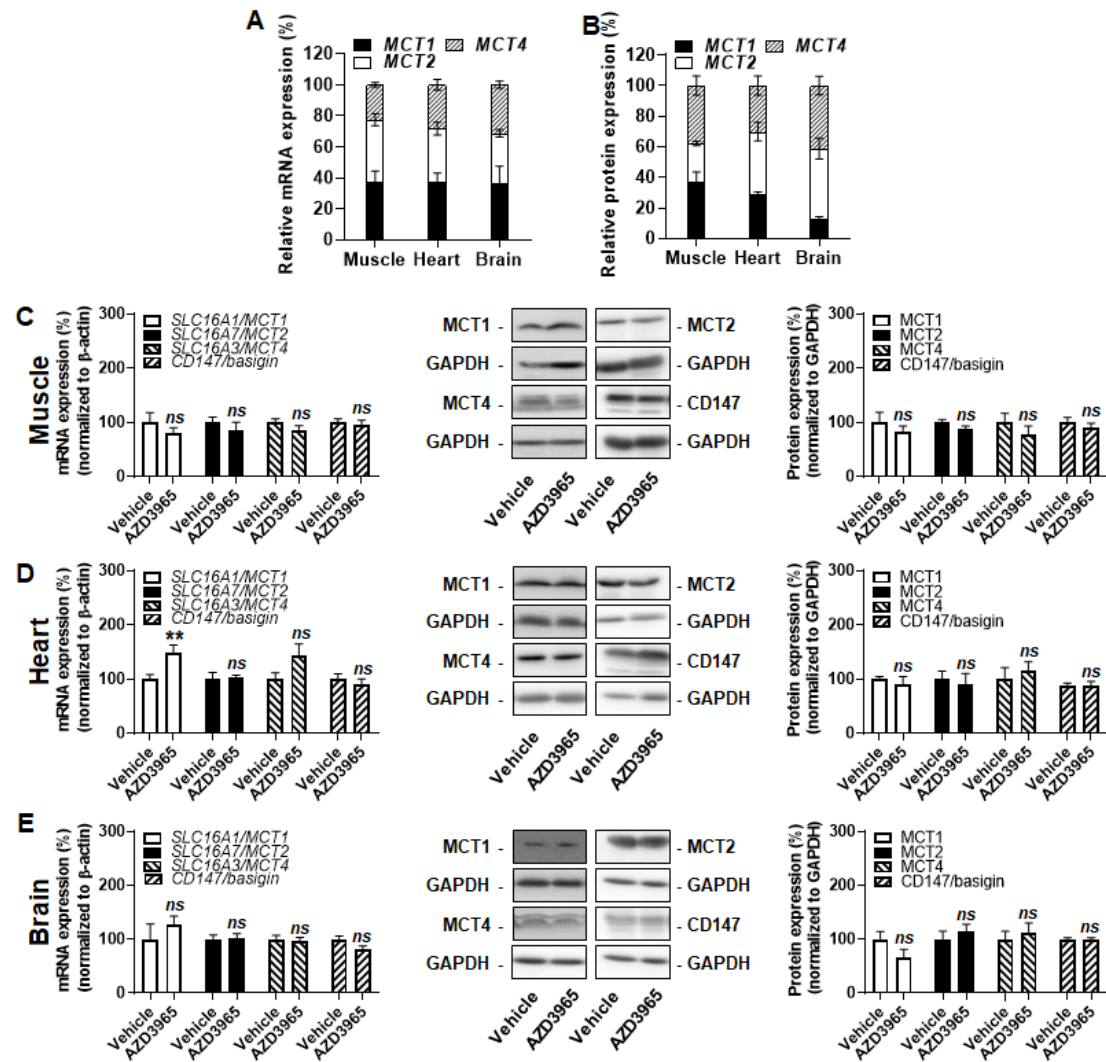
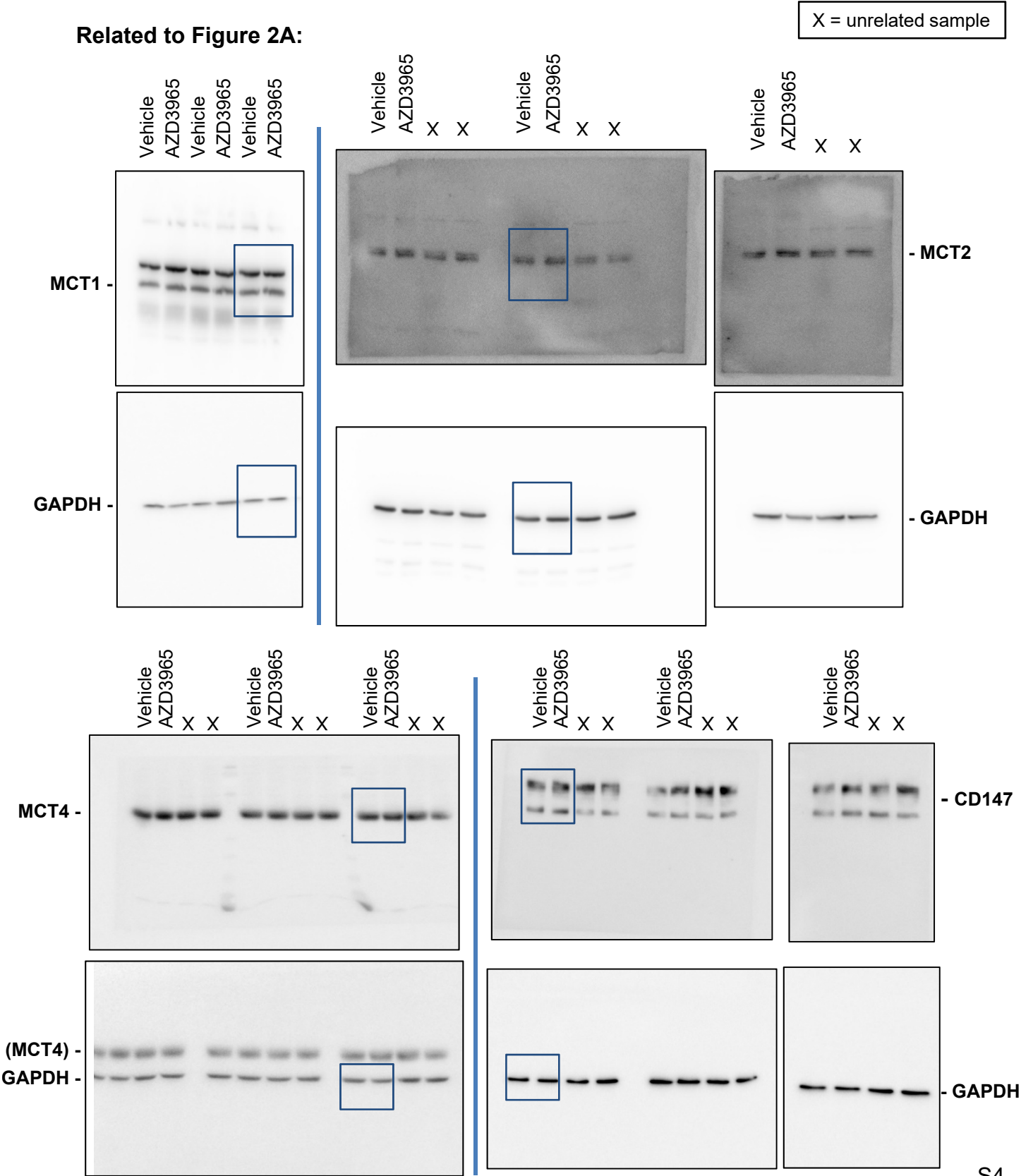


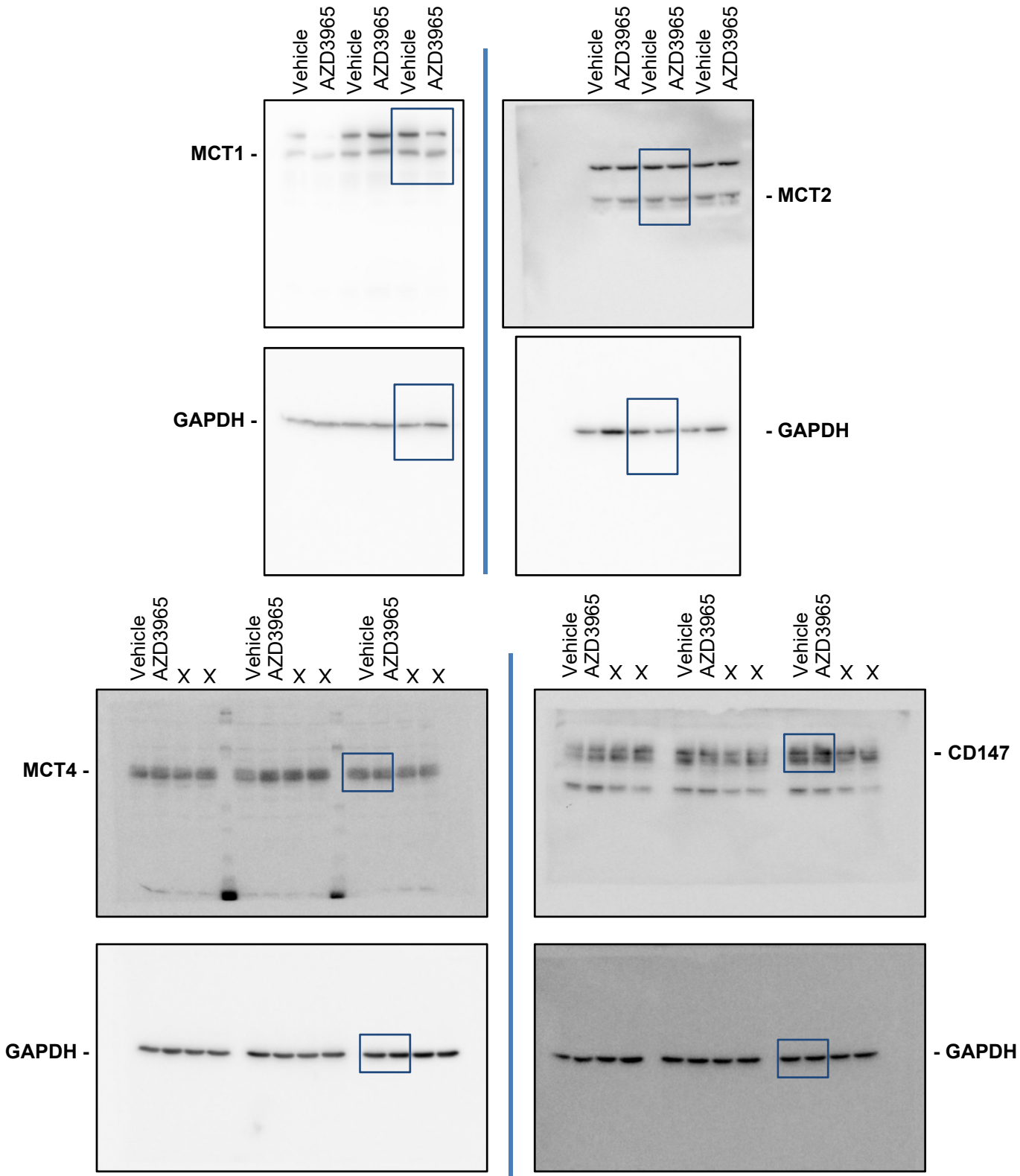
Figure S3. A chronic treatment with AZD3965 does not alter the expression of MCTs and CD147/basigin in mouse skeletal muscles, heart and brain. (A-E) Mouse tissues were collected on the day of sacrifice of Group 2 depicted in **Figure 7A**. (A) Relative mRNA expression of MCT1, MCT2 and MCT4 in the muscles, heart and brain of vehicle-treated mice. Cumulated expression served for normalization (100%) ($n = 4-6$). (B) Relative protein expression of MCT1, MCT2 and MCT4 in the muscles, heart and brain of vehicle-treated mice. Representative western blots are shown with GAPDH as a loading control. Cumulated expression served for normalization (100%) ($n = 4-6$). (C) mRNA (left panel) and protein (middle and right panels) expression of MCT1, MCT2, MCT4 and CD147/basigin in the gastrocnemius muscles of mice treated $\pm$ 100 mg/Kg AZD3965 ($n = 4-24$ for RT-qPCR, $n = 5-6$ for WB). (D) As in (C) but in whole mouse hearts ($n = 3-6$ for RT-qPCR, $n = 5-6$ for WB). (E) As in (C) but in whole mouse brains ($n = 5-6$ for RT-qPCR, $n = 5-6$ for WB). All data are shown as means $\pm$ SEM. ** $P < 0.01$, *ns* $P > 0.05$ compared to corresponding tissues from vehicle-treated animals; by Student's *t* test (C-E).

Figure S4. Uncropped western blots



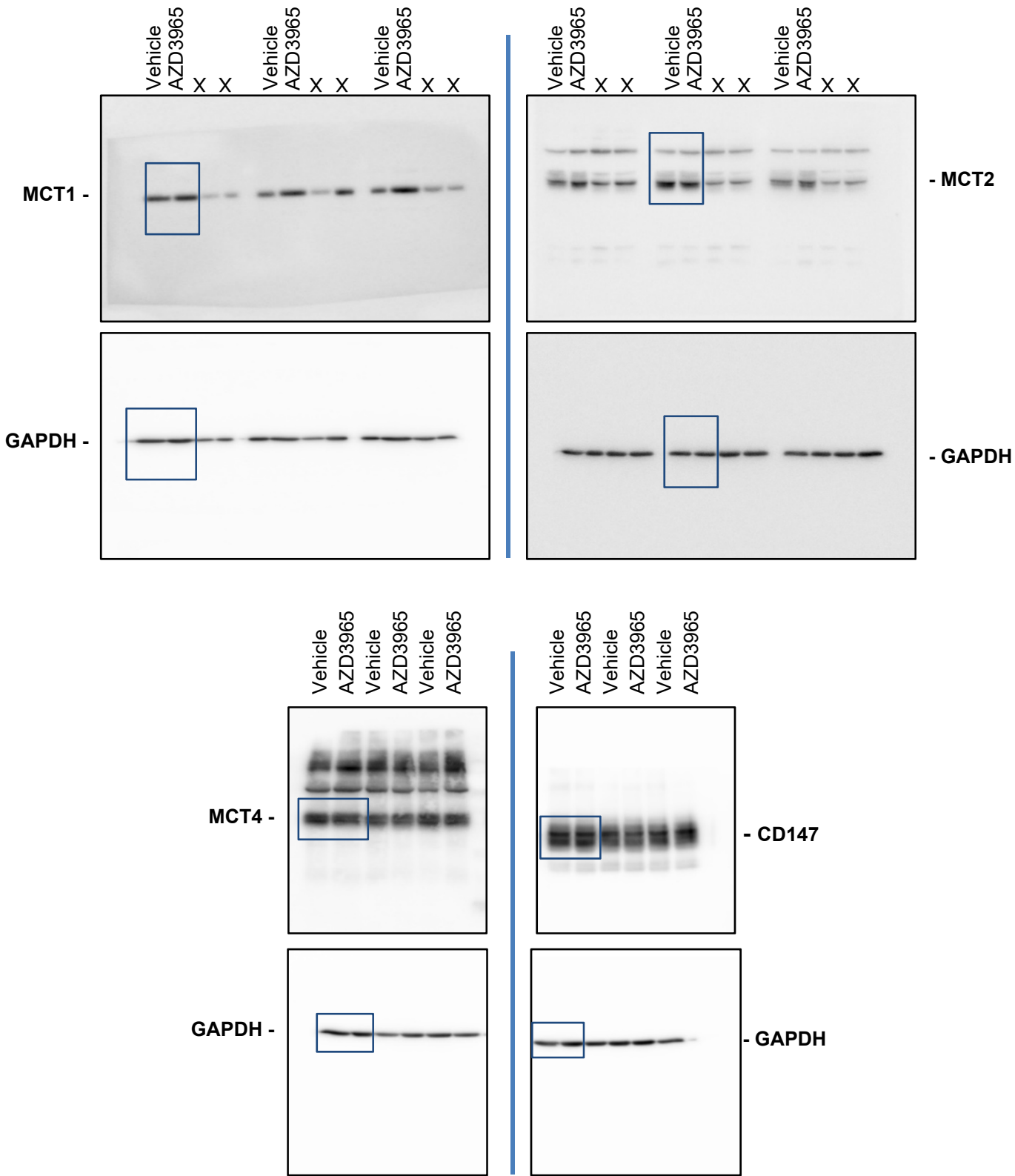
Related to Figure 2B:

X = unrelated sample



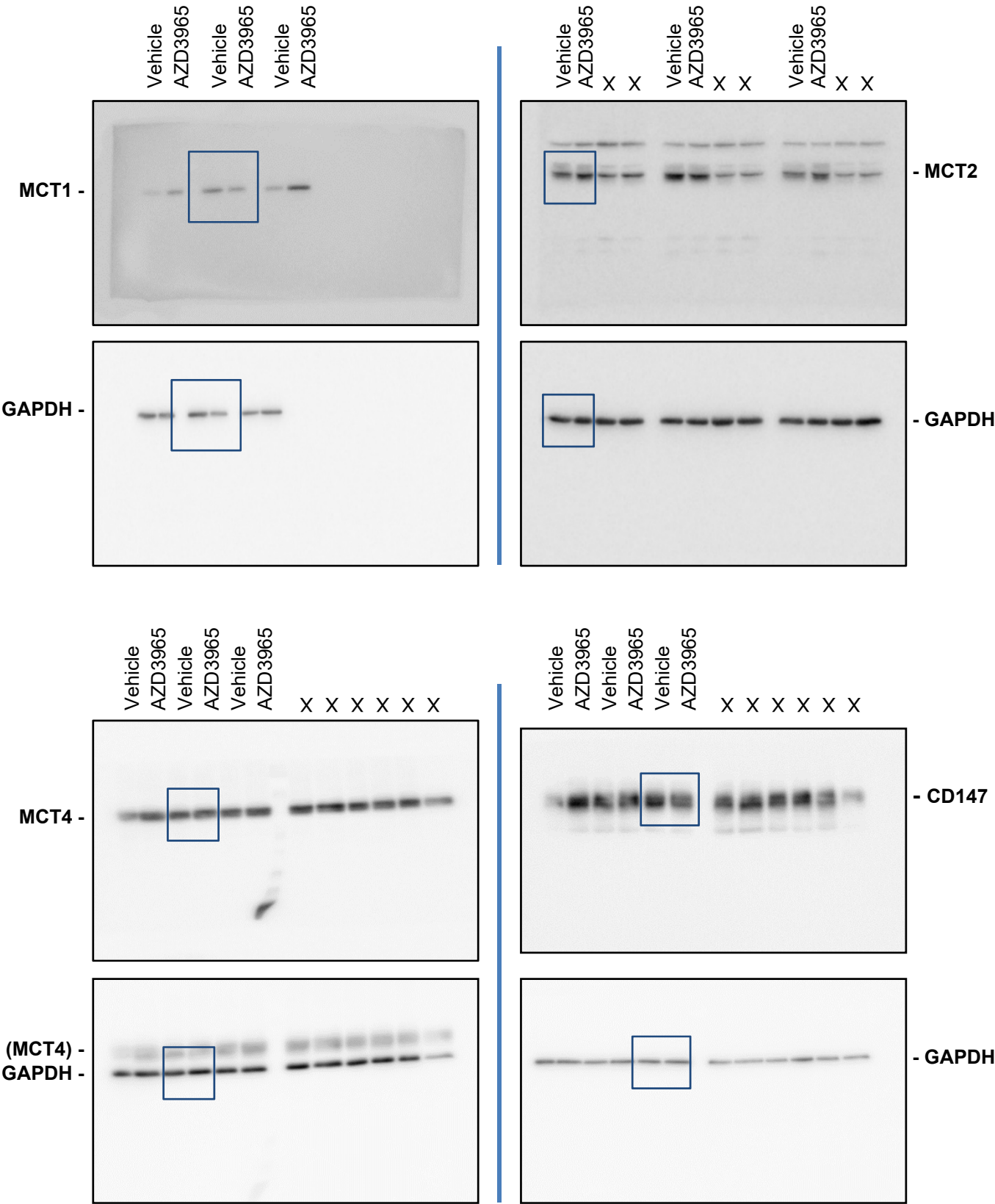
Related to Figure 2C:

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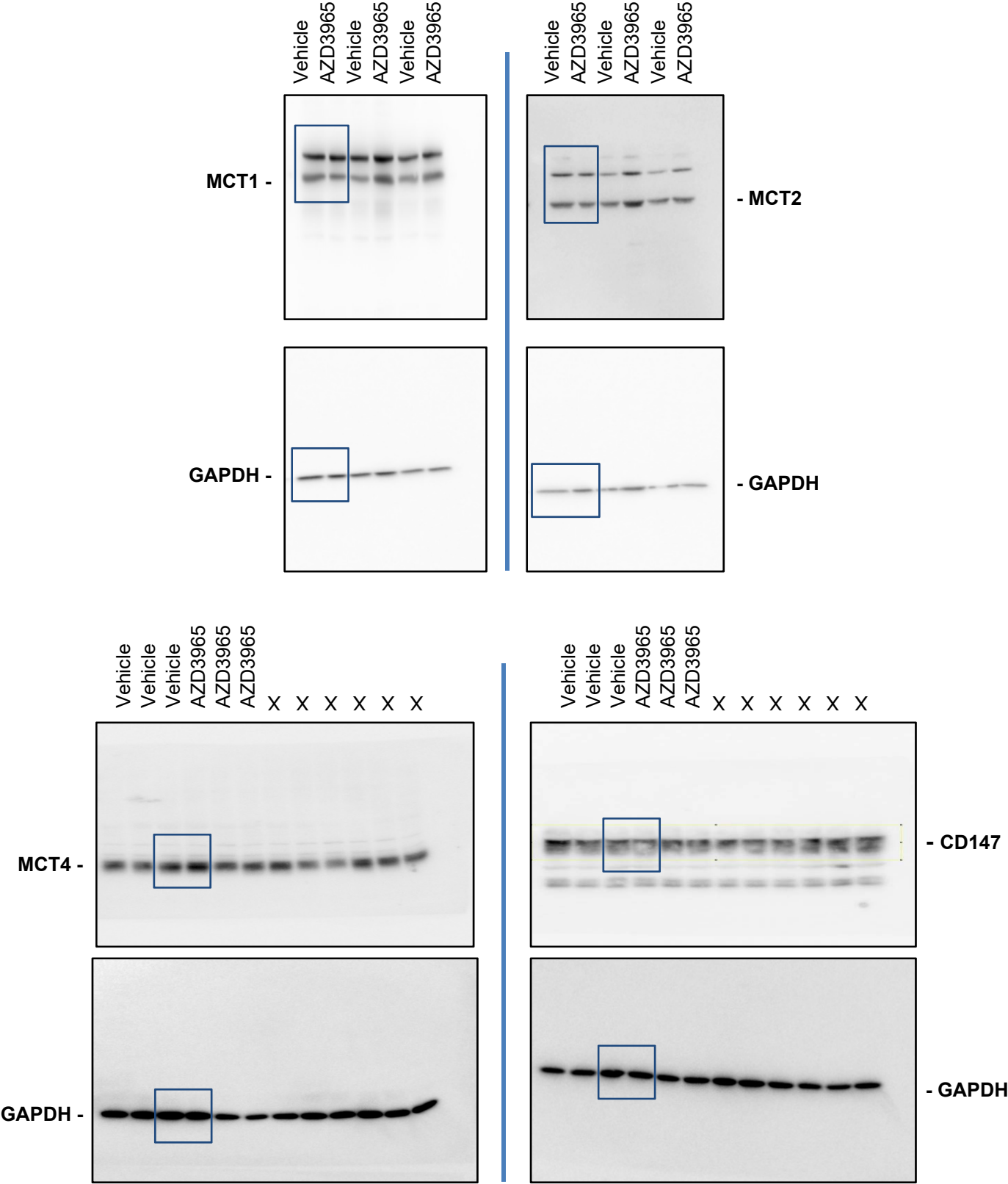
Related to Figure 2D:

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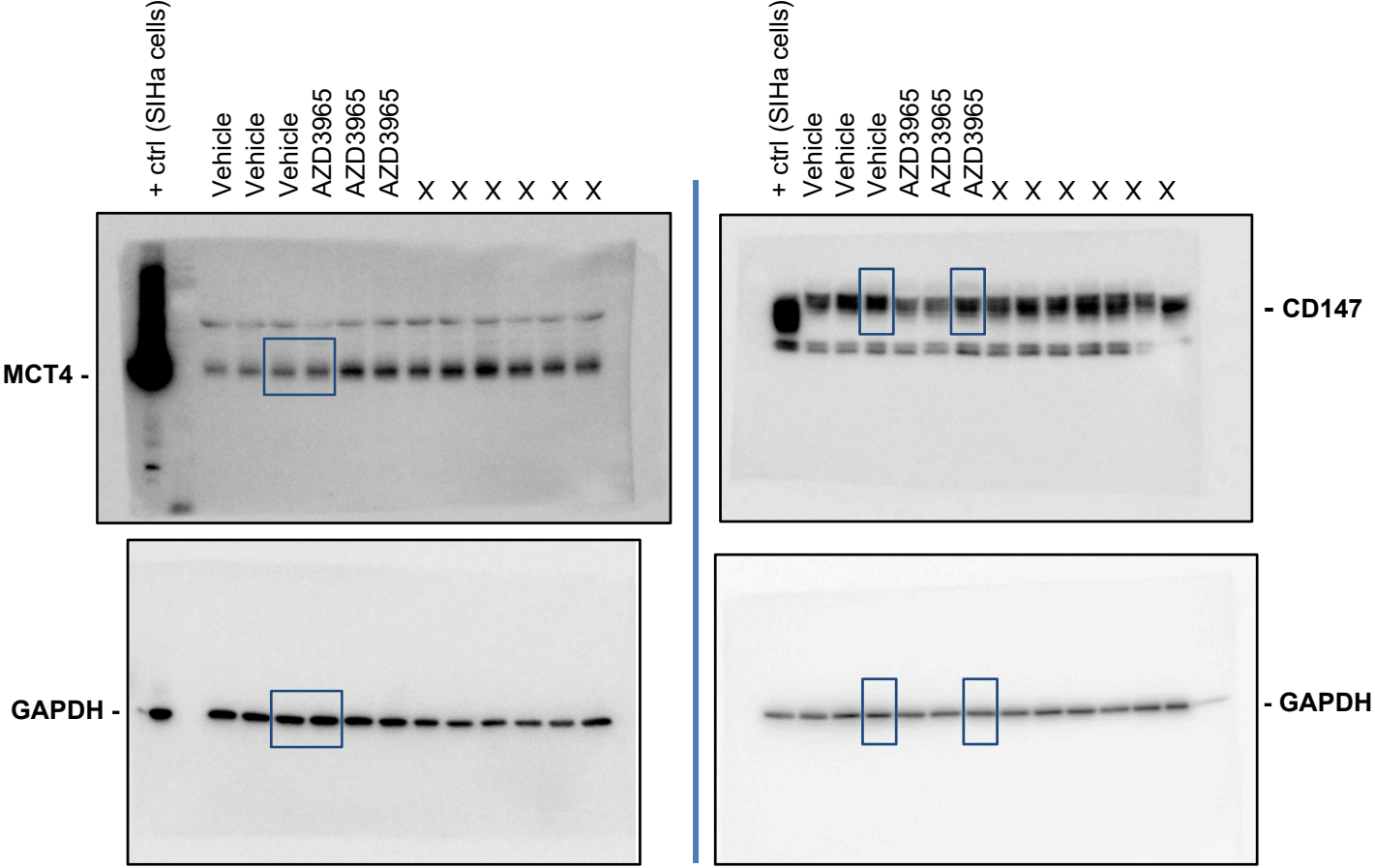
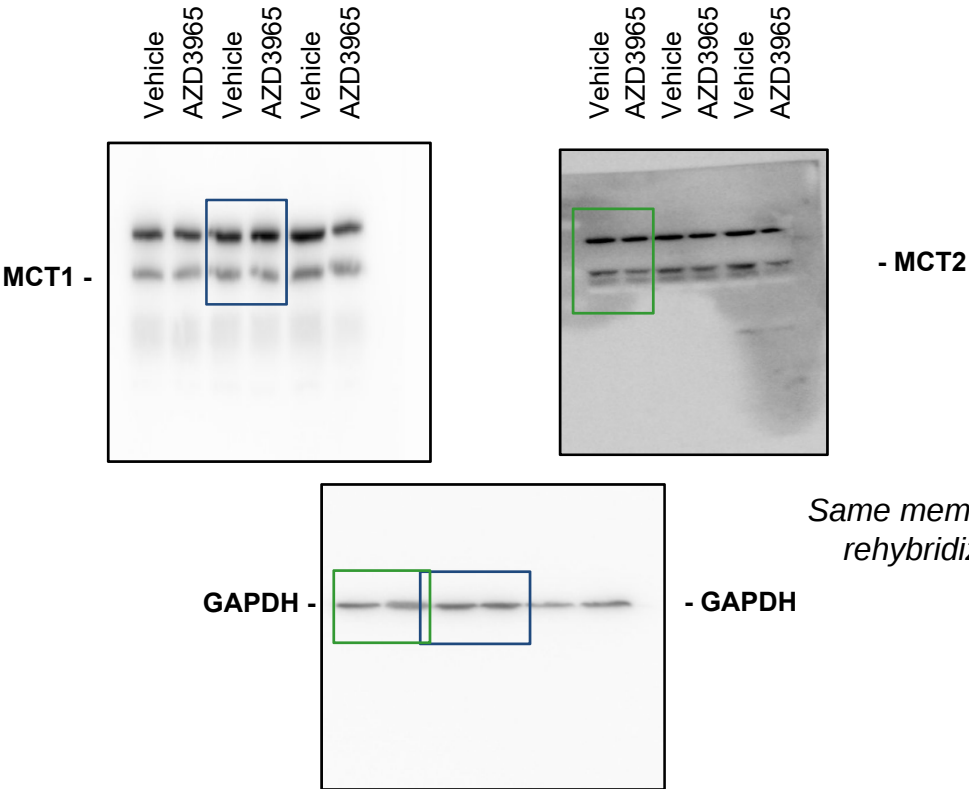
Related to Figure 5A:

X = unrelated sample



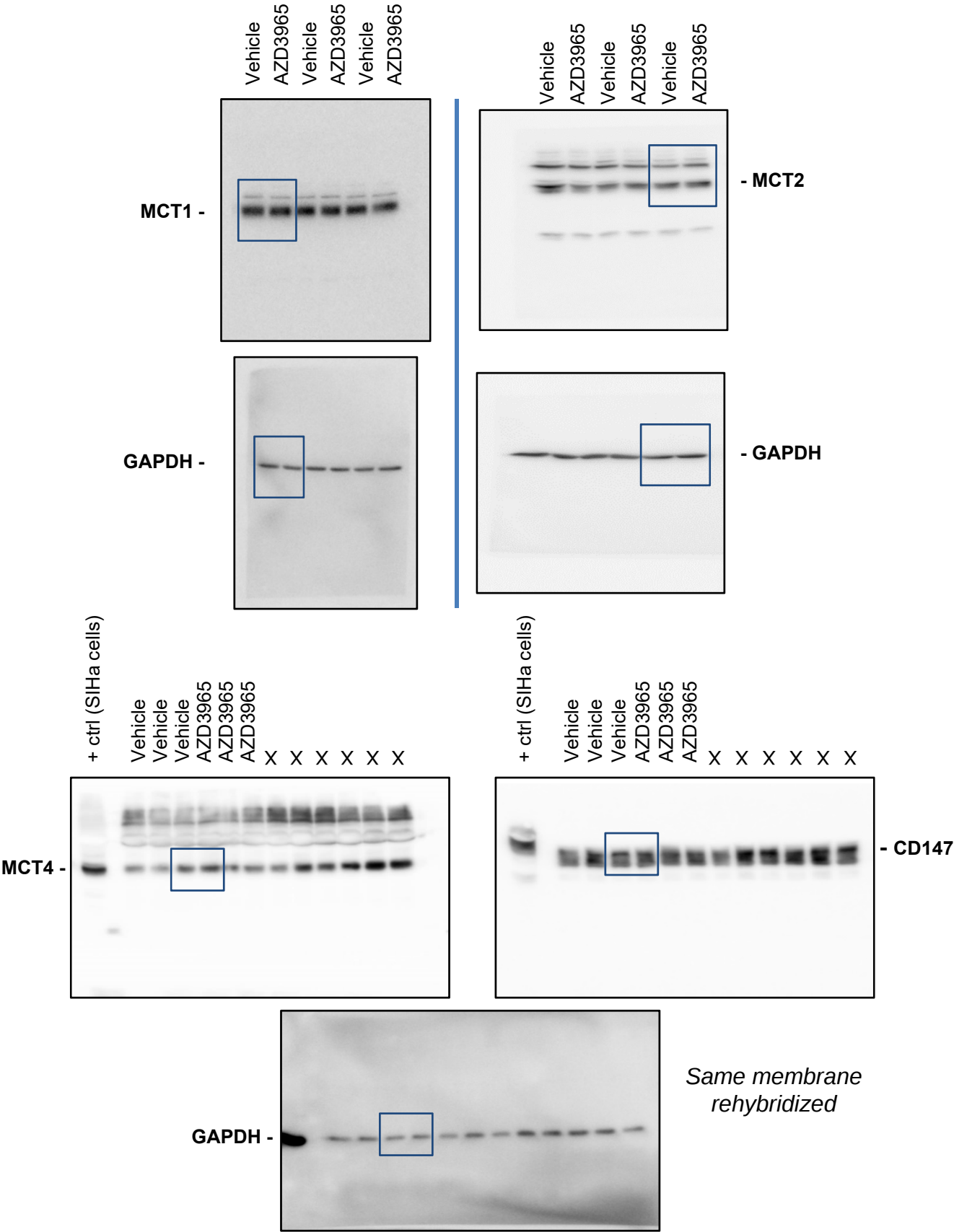
Related to Figure 5B:

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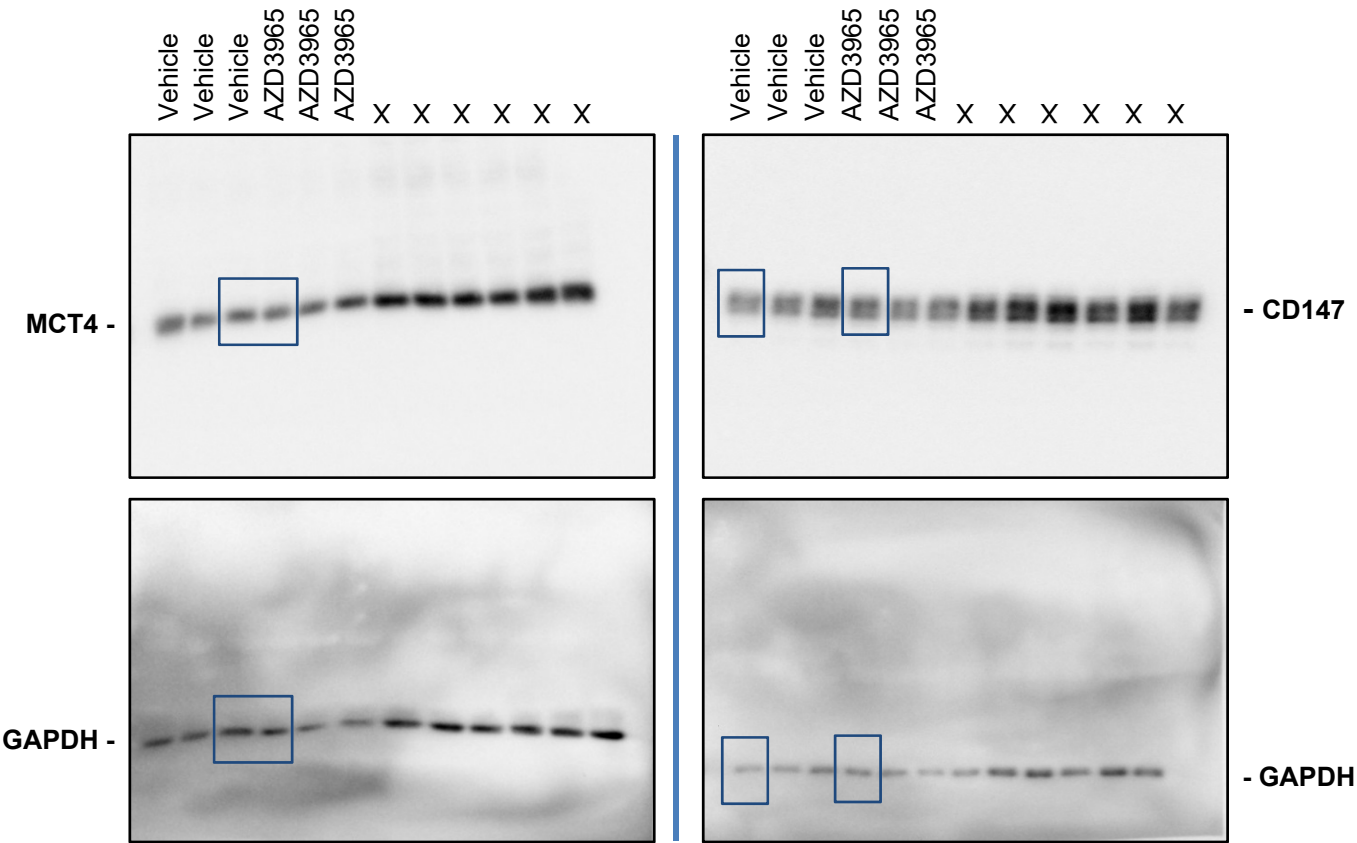
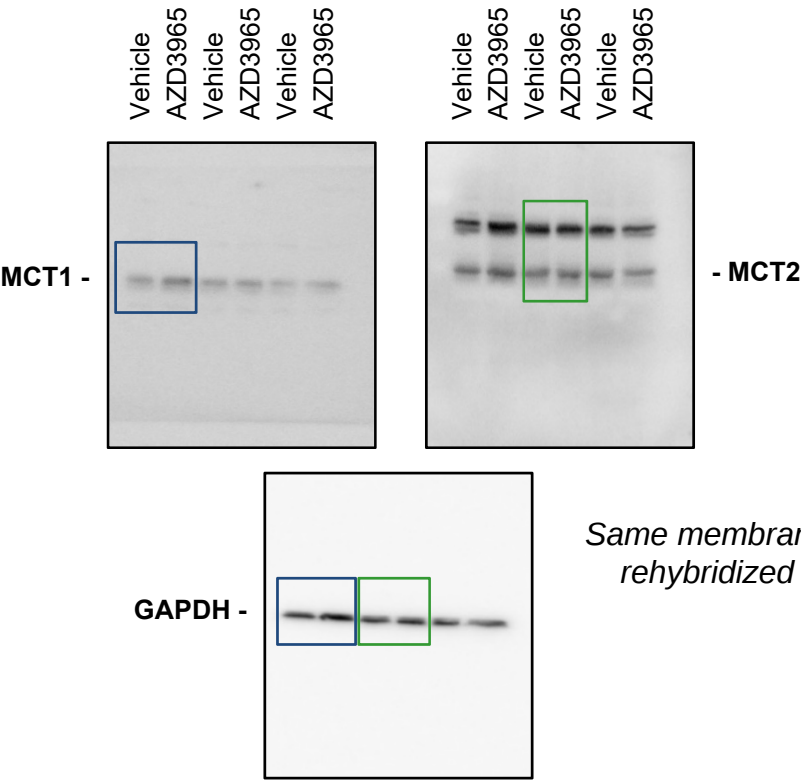
Related to Figure 5C:

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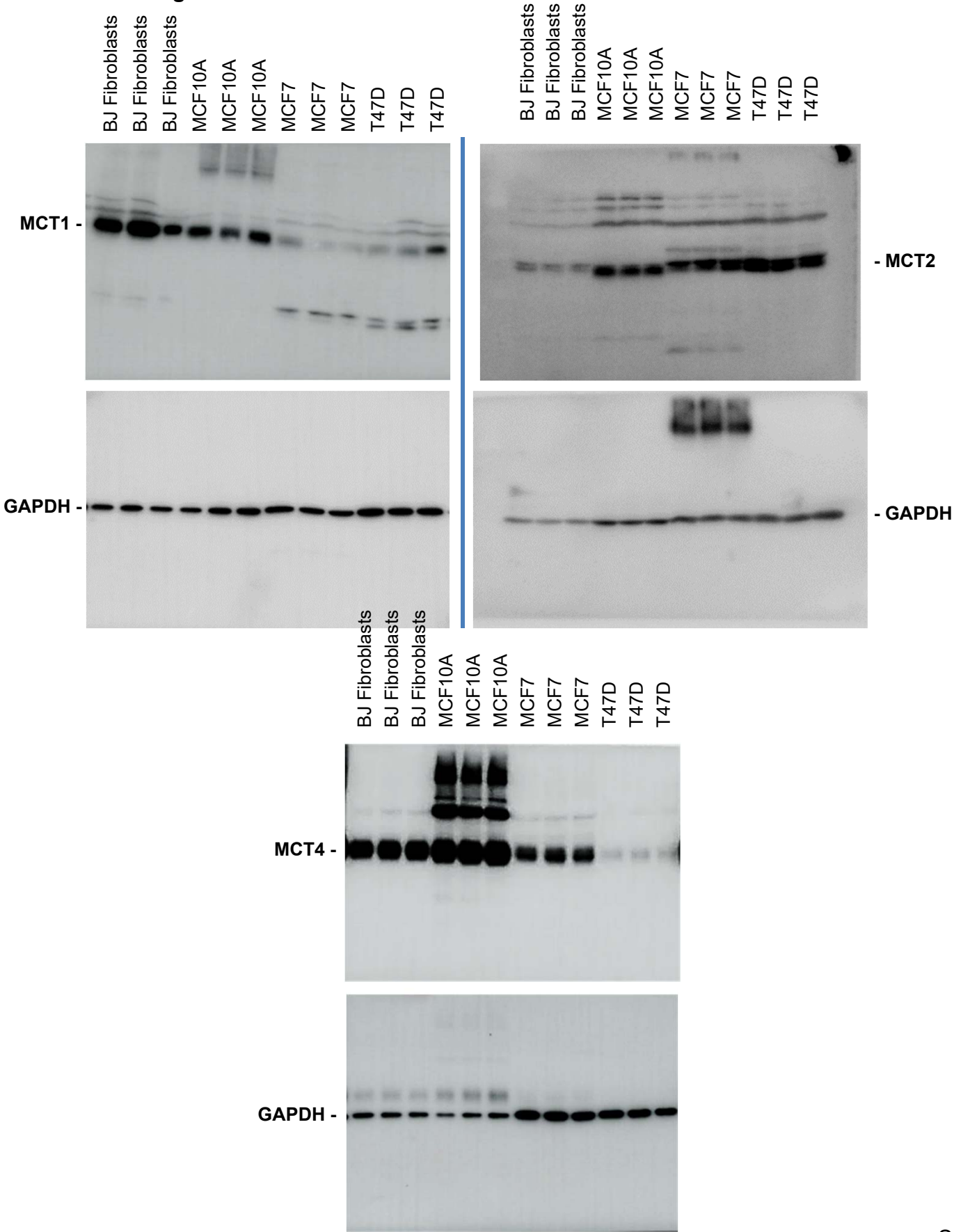


Related to Figure 5D:

X = unrelated sample



Related to Figure S1B:



Related to Figure S2C:

