

Supplementary Raw WB data

A clinically relevant SLC2A1-associated malignant epithelial cell state predicts prognosis and immunotherapy response in lung adenocarcinoma

Linqian Song^{1,2}, Shiyun Xing^{1,2}, Peijie Li^{1,2}, Yunliang Cao^{1,2}, Hu Ma^{1,2}, *

¹Department of Oncology, The Second Affiliated Hospital of Zunyi Medical University, Zunyi, Guizhou 563000, P. R. China;

²Key Laboratory for Cancer Prevention and Treatment of Guizhou Province, Zunyi, Guizhou 563000, P. R. China.

* Corresponding author

Correspondence to: Hu Ma, Email: mahuab@163.com; ORCID: 0000-0003-1654-1576.

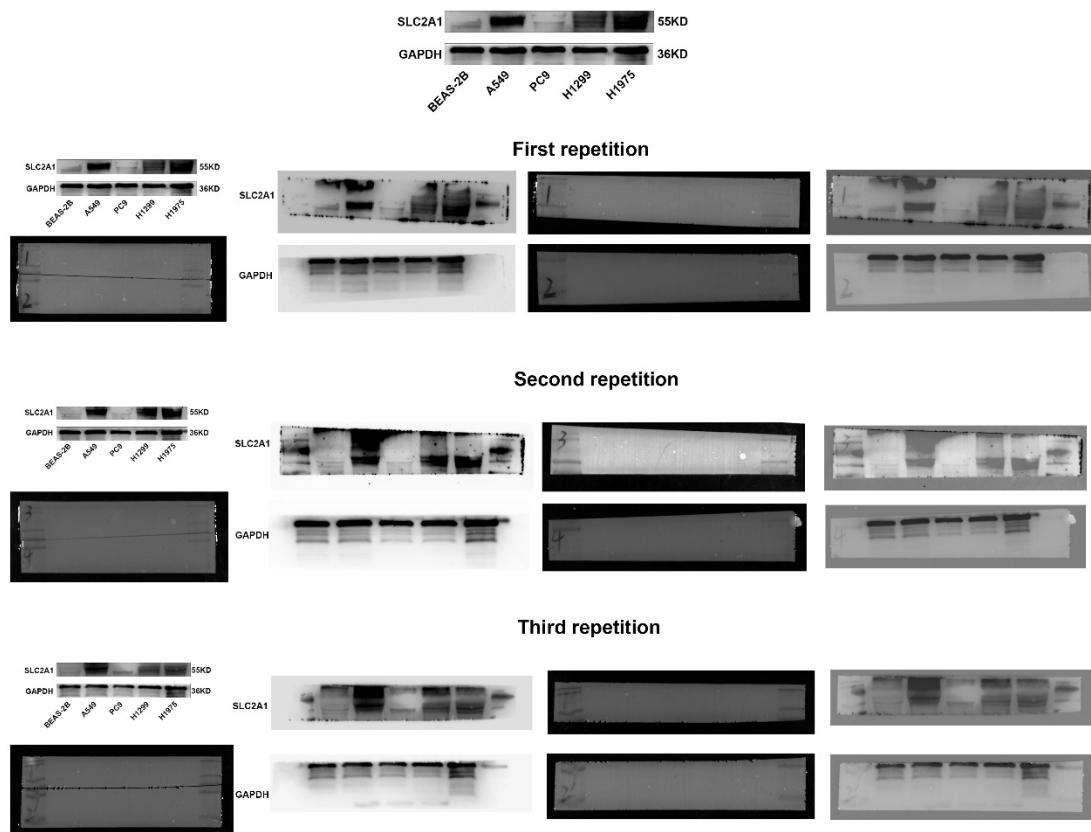
Supplementary Raw WB Images

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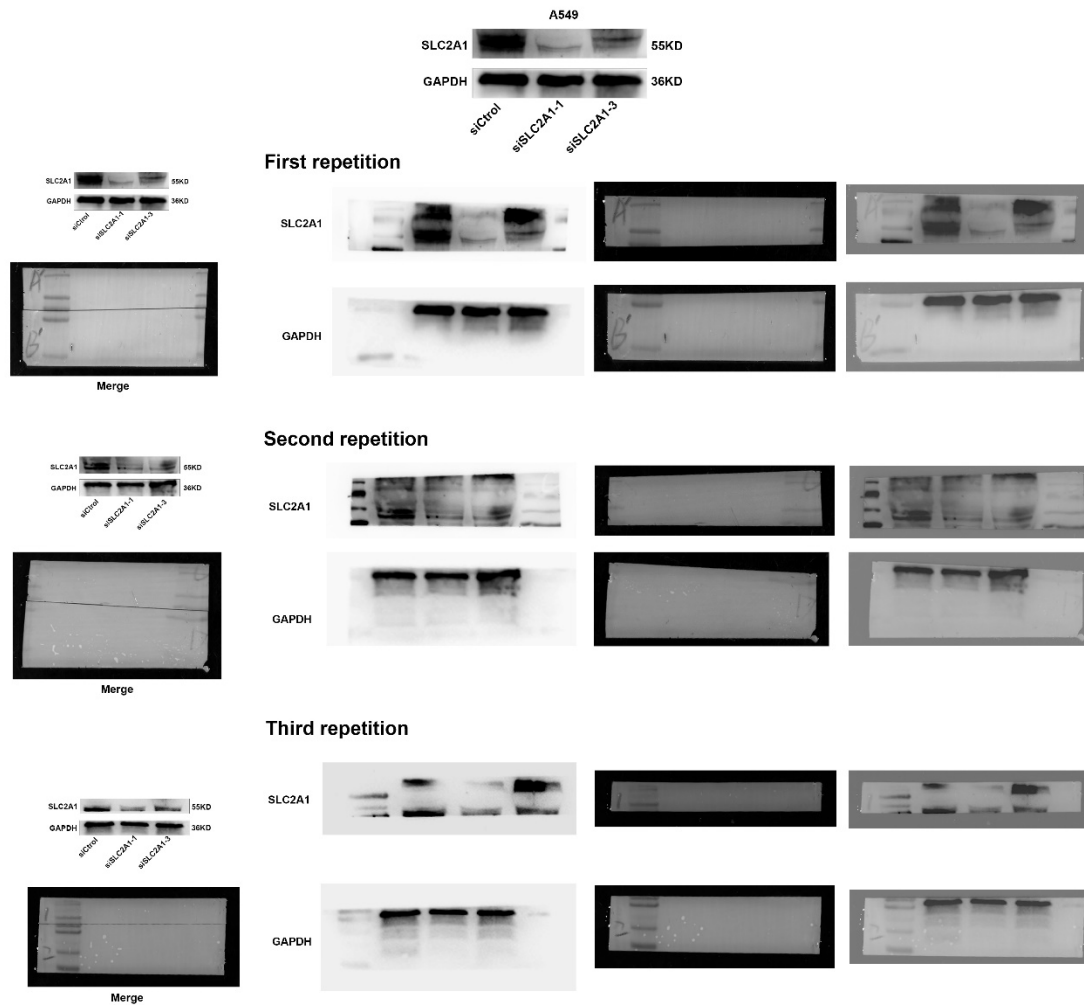
Raw WB Image 2. Original uncropped Western blot images of SLC2A1 knockdown in A549 cells.

Raw WB Image 3. Original uncropped Western blot images of SLC2A1 knockdown in H1975 cells.



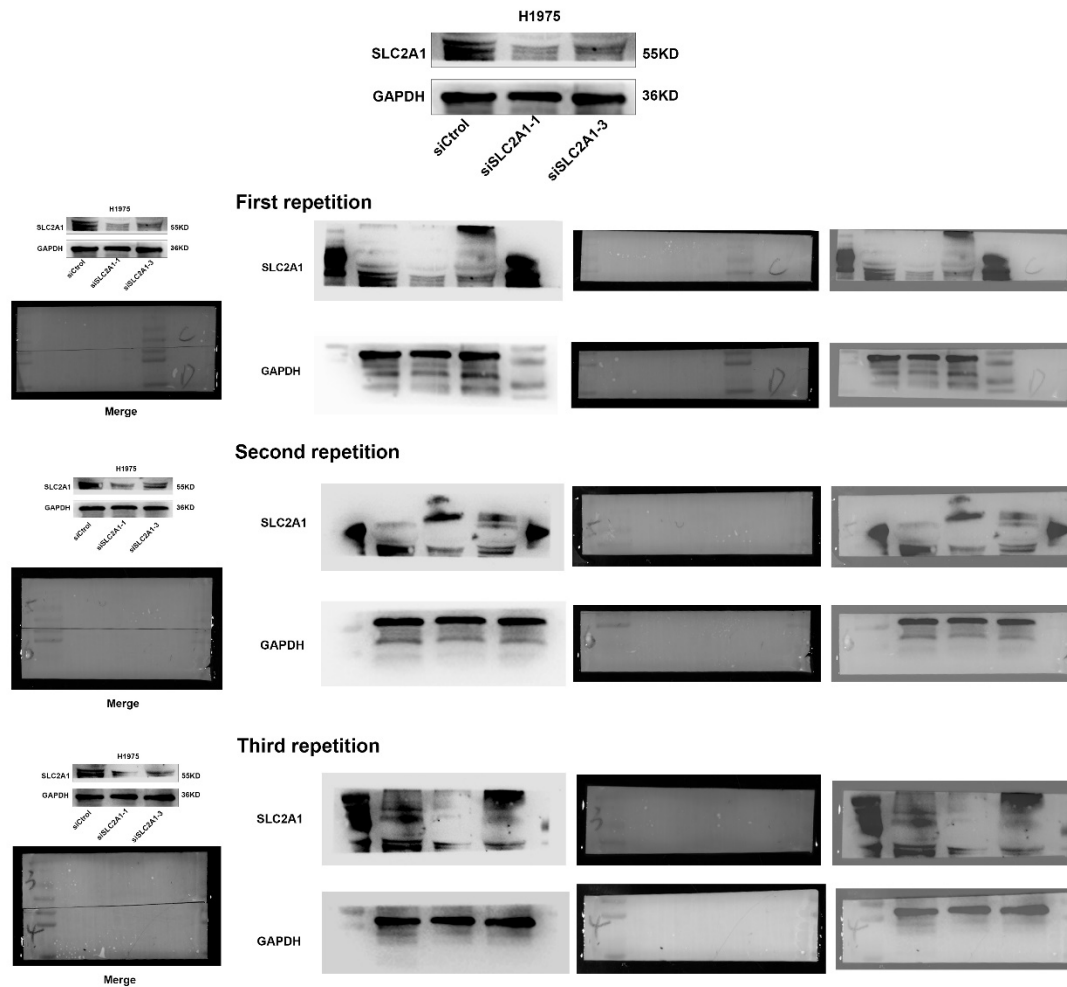
Raw WB Image 1 Original uncropped Western blot images of SLC2A1 expression across lung cell lines.

Uncropped and full-length Western blot images corresponding to Figure 10B are shown. Protein expression of SLC2A1 (55 kDa) and GAPDH (36 kDa) was detected in normal bronchial epithelial cells (BEAS-2B) and lung adenocarcinoma cell lines (A549, PC9, H1299, and H1975). Three independent experimental repetitions are presented.



Raw WB Image 2. Original uncropped Western blot images of SLC2A1 knockdown in A549 cells.

Uncropped and full-length Western blot images corresponding to Figure 10D are shown. A549 cells were transfected with control siRNA (siCtrl) or siRNAs targeting SLC2A1 (siSLC2A1-1 and siSLC2A1-3). Protein levels of SLC2A1 (55 kDa) and GAPDH (36 kDa) were detected. Three independent experimental repetitions are presented.



Raw WB Image 3. Original uncropped Western blot images of SLC2A1 knockdown in H1975 cells.

Uncropped and full-length Western blot images corresponding to Figure 10D are shown. H1975 cells were transfected with control siRNA (siCtrl) or siRNAs targeting SLC2A1 (siSLC2A1-1 and siSLC2A1-3). Protein levels of SLC2A1 (55 kDa) and GAPDH (36 kDa) were detected. Three independent experimental repetitions are presented.

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