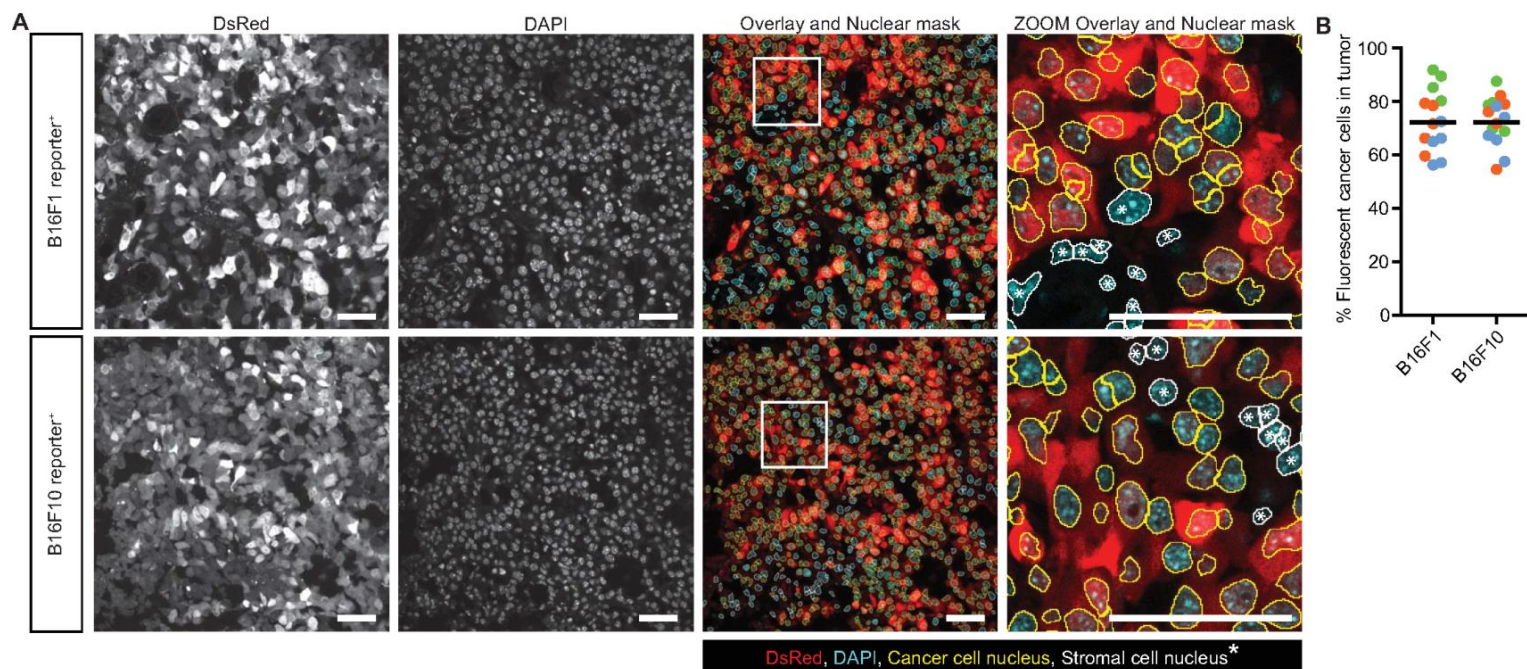


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Appendix Figure S1. Estimation of percentage of cancer cells within B16F1 and B16F10 tumors. (a) Representative images of sections of B16F1 (top) and B16F10 (bottom) reporter⁺ tumors for DsRed and DAPI staining. Nuclear regions were assessed for expression of DsRed and all nuclei were scored positive (cancer cell nucleus, yellow edge) or scored negative (stromal cell nucleus, white edge and marked with an asterisk) for DsRed. Scale bars are 50µm. (b) Quantification of 4-5 tumor regions per tumor for 3 mice per model, colors indicate tumor regions per mouse.



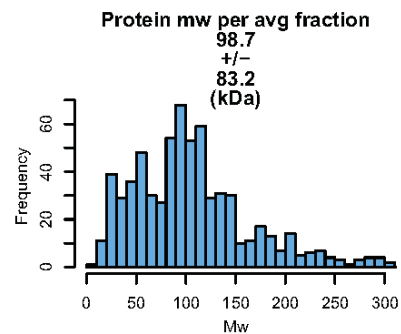
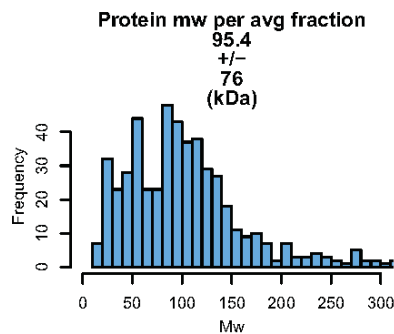
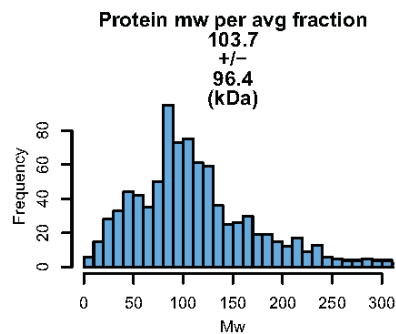
Appendix Figure S2. Predicted molecular weight of detected proteins per gel band. For each gel block per sample, the frequency of the predicted molecular weight of all identified proteins was plotted. Average molecular weight (in kDa, +/- SD) per fraction is displayed above each graph.

Cell Lysate

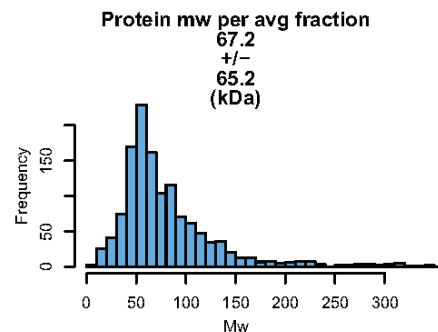
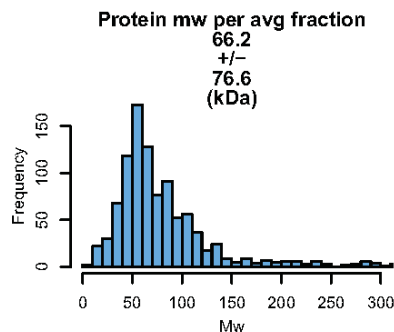
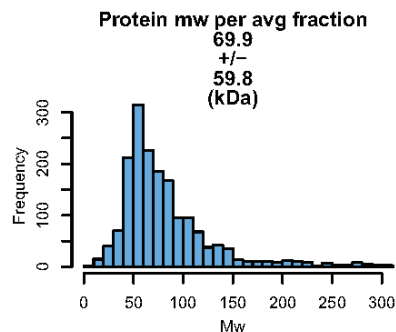
16.5K EV

100K EV

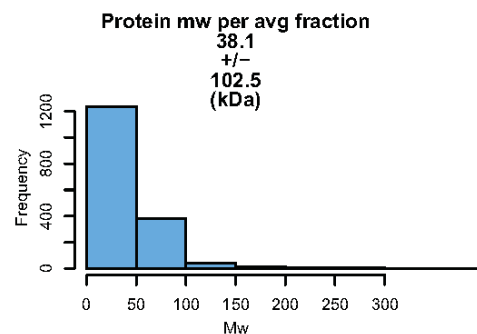
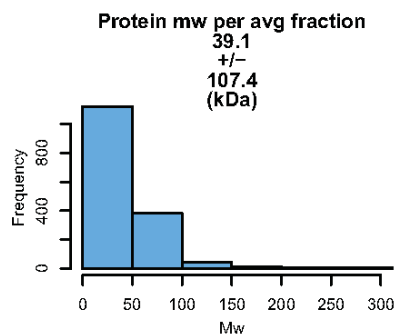
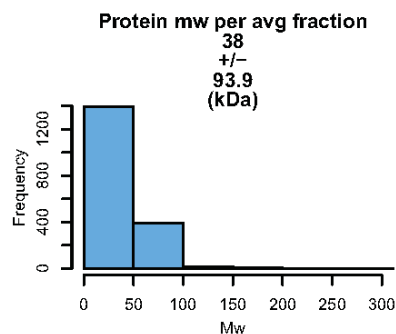
Gel band 1



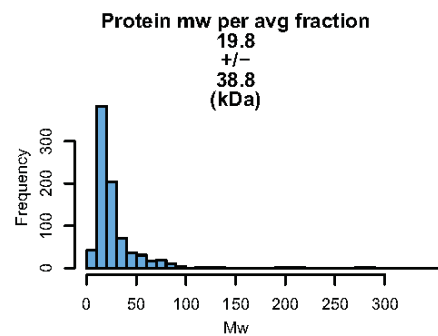
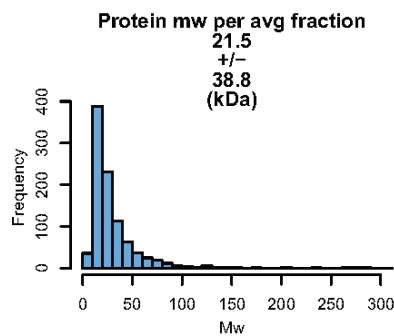
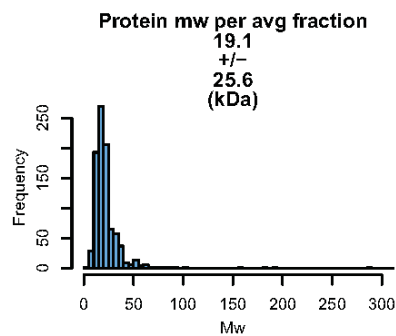
Gel band 2



Gel band 3



Gel band 4



Gel band 5

