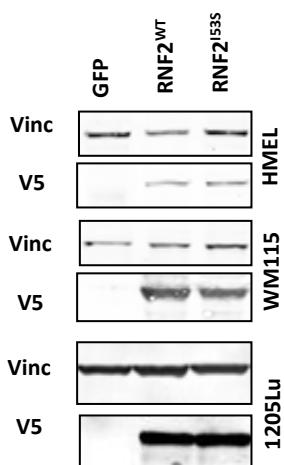
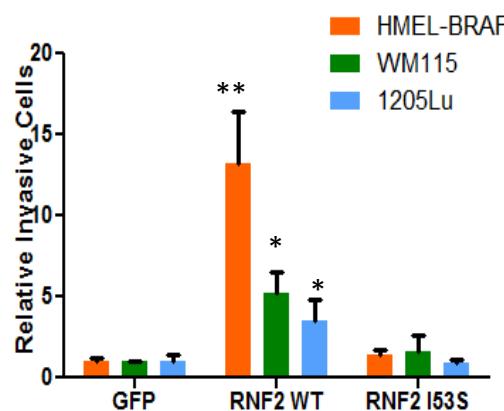
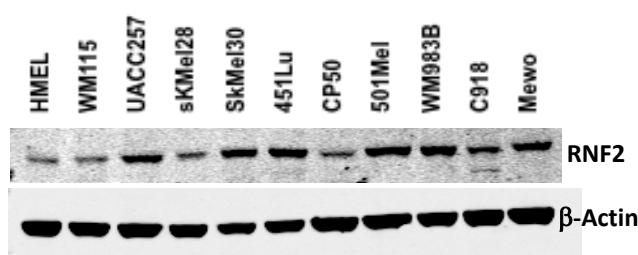
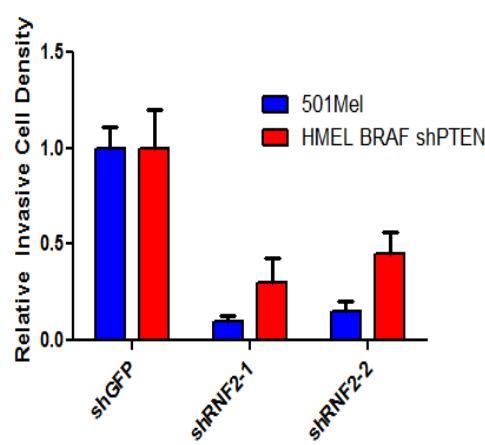
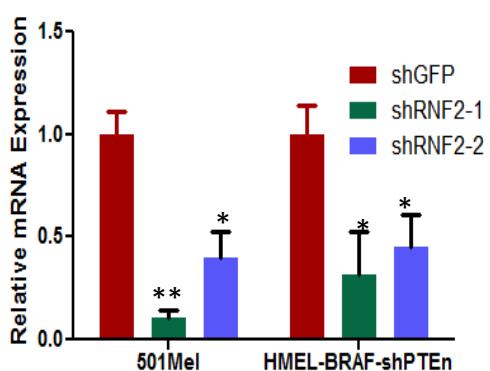
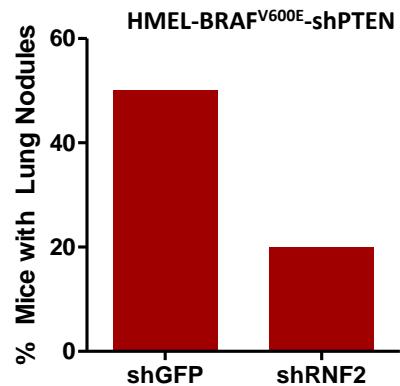
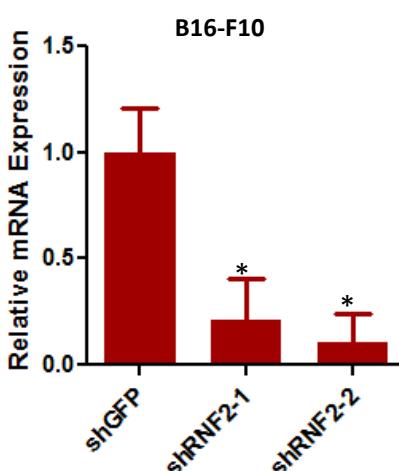
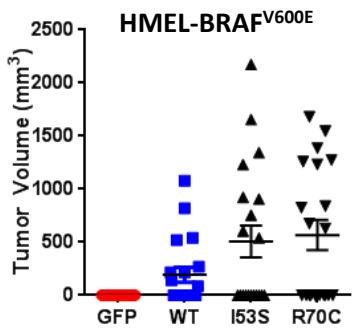


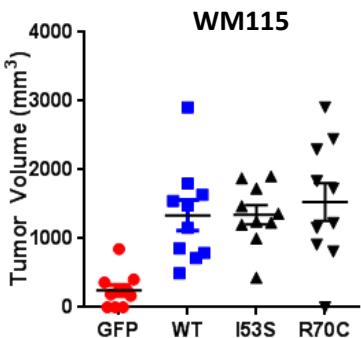
A**B****C****D****E****F****G**

Supplemental Figure 1, Rai et al

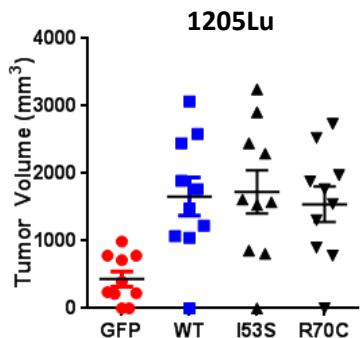
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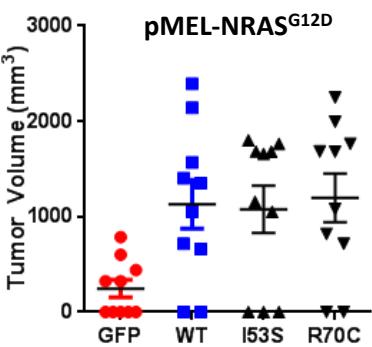
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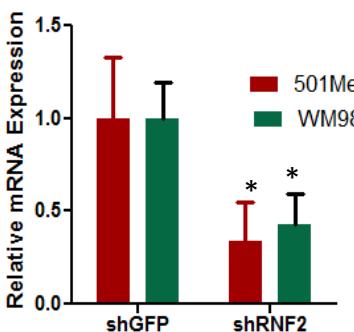
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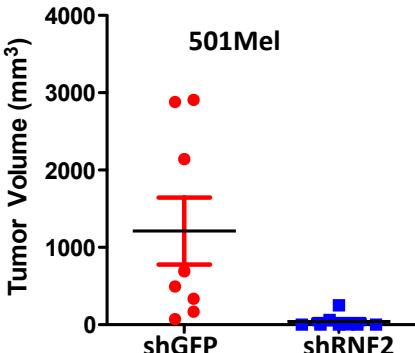
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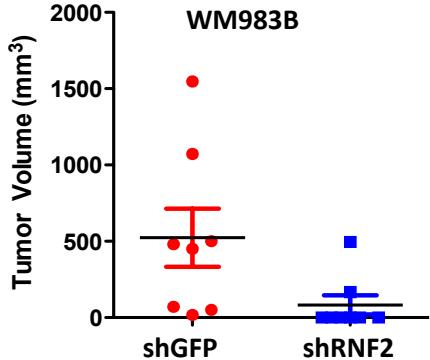
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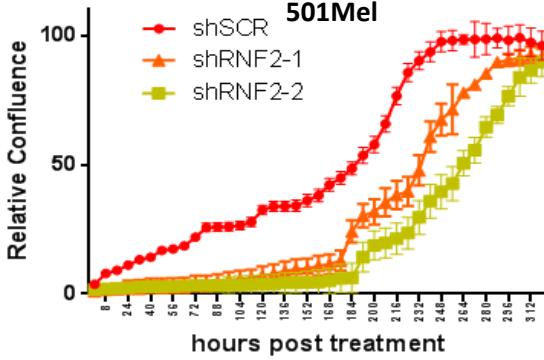
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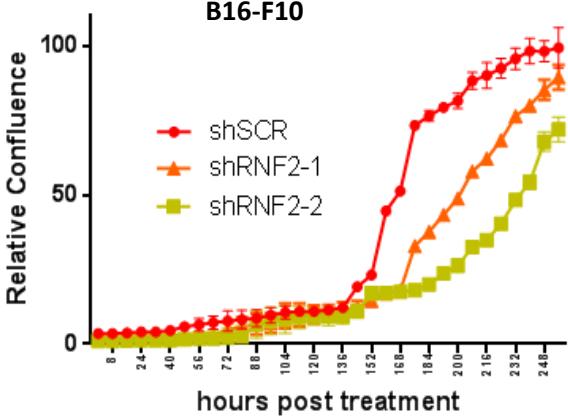
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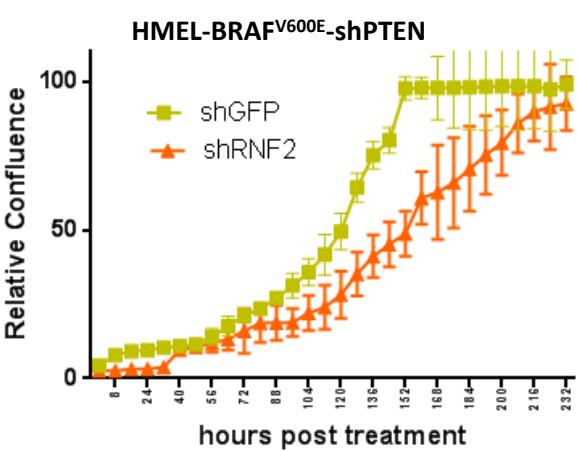
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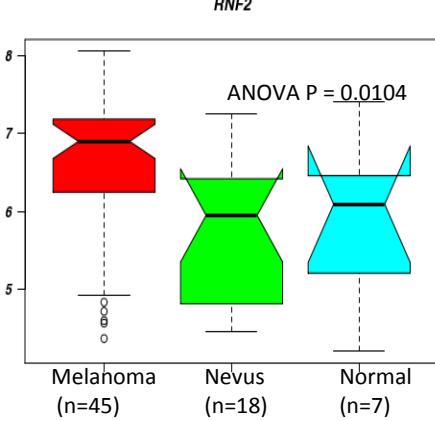
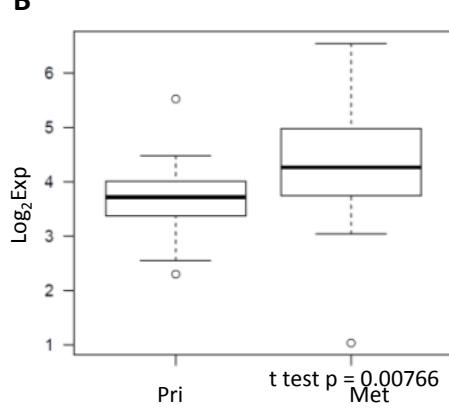
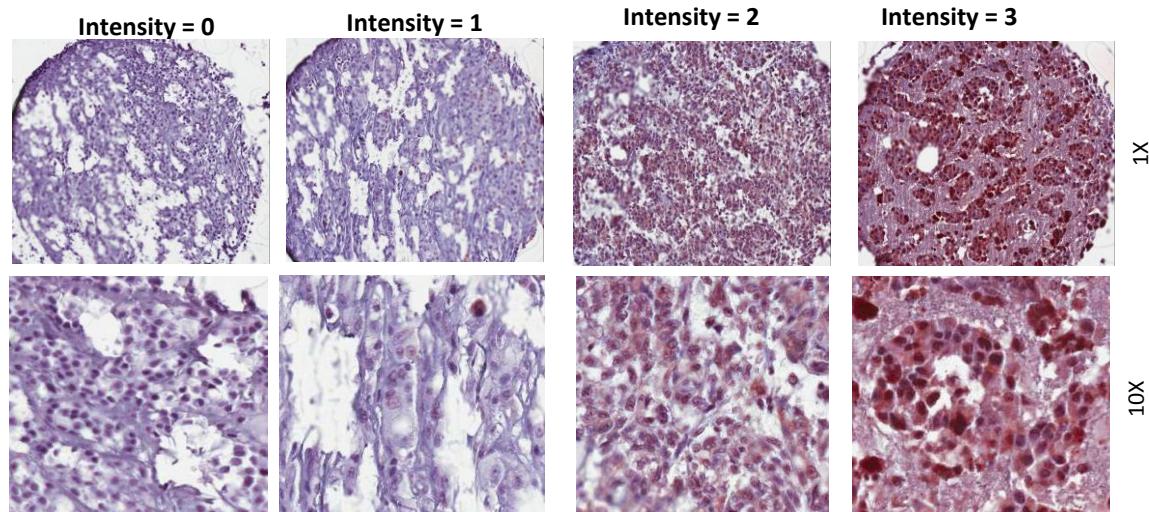
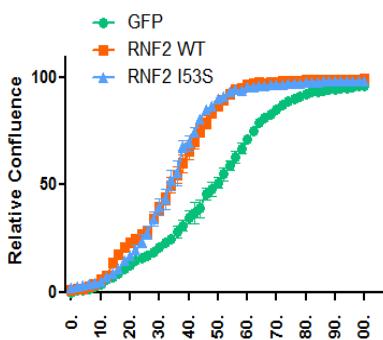
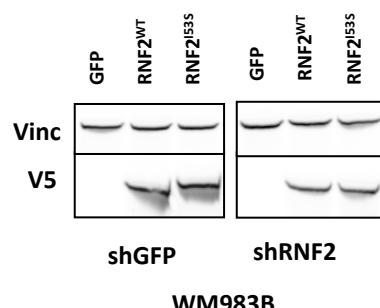
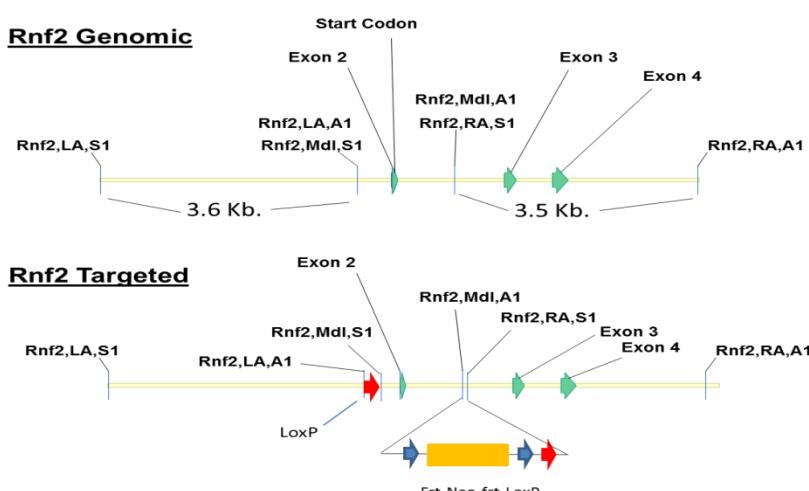
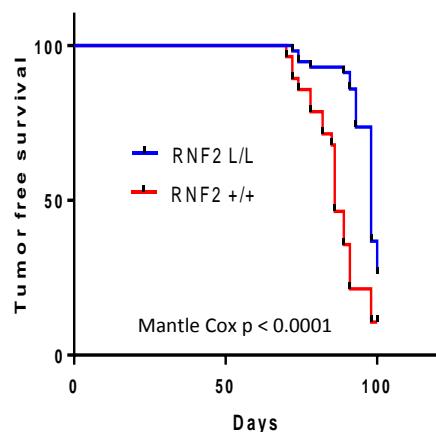
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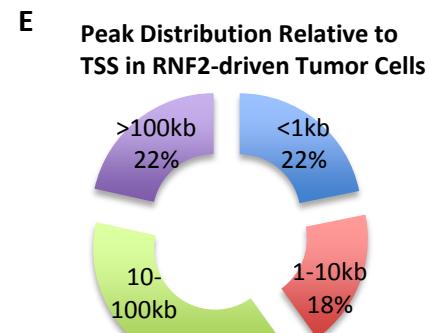
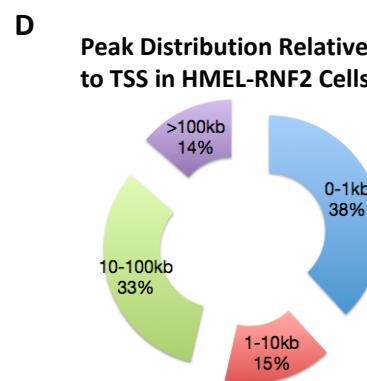
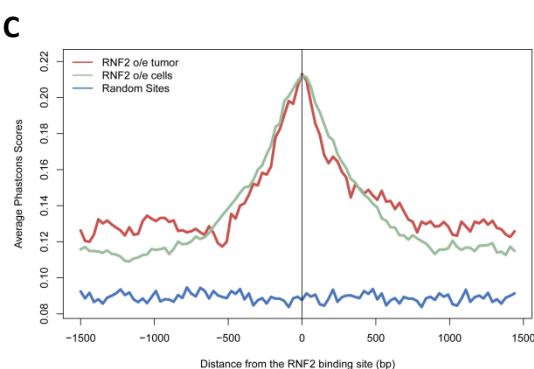
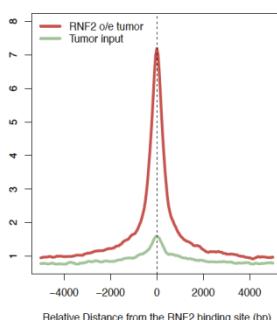
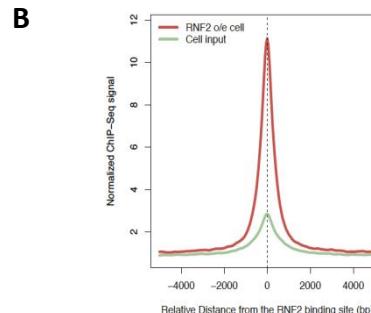
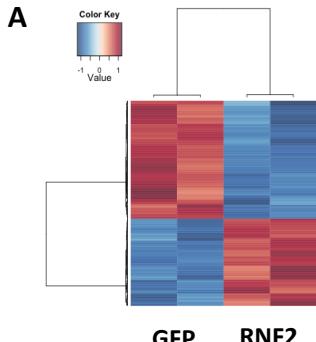
J



Supplemental Figure 2, Rai et al

A**B****C****D****E****F****G**

Supplemental Figure 3, Rai et al

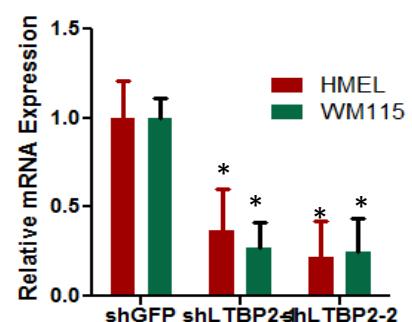
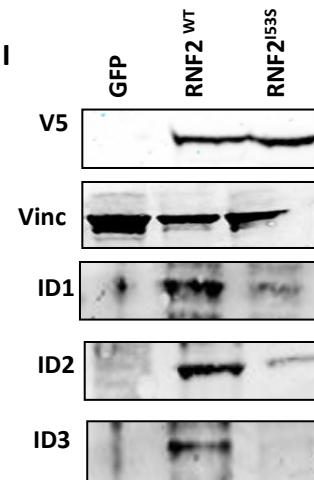
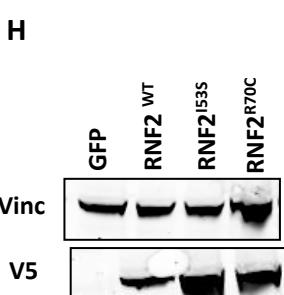


F

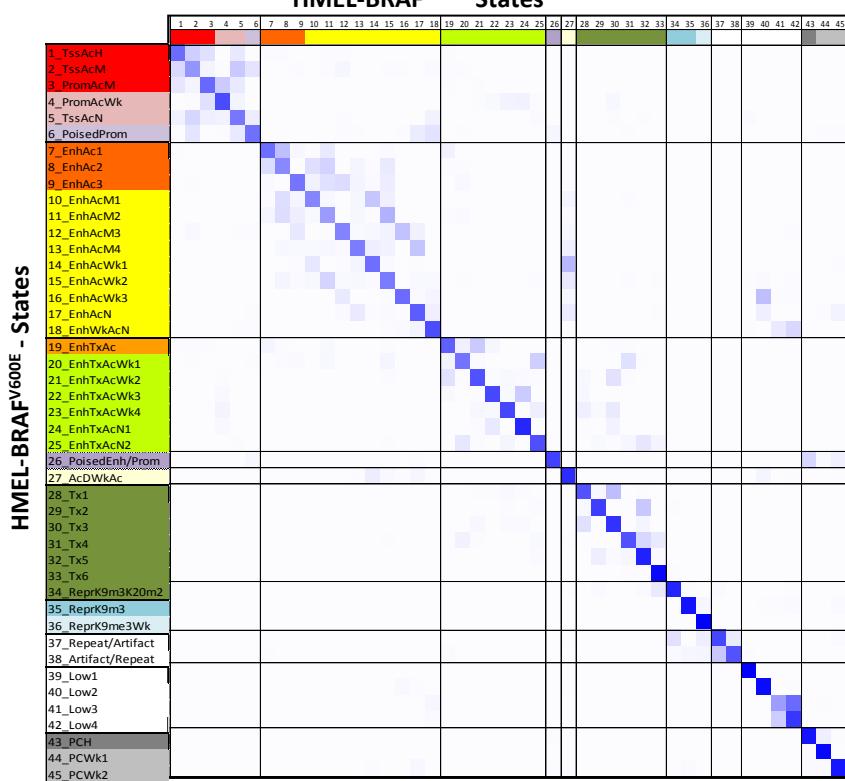
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GO:0050680~negative regulation of epithelial cell proliferation	4.02E-04
GO:0050678~regulation of epithelial cell proliferation	7.78E-04
GO:0045935~positive regulation of nucleobase, nucleoside, nucleotide and nucleic acid metabolic process	9.57E-04
GO:0001666~response to hypoxia	0.001109

G

GO Term	PValue
GO:0045449~regulation of transcription	0.00173241
GO:0016481~negative regulation of transcription	0.003218314
GO:0010629~negative regulation of gene expression	0.005640153
GO:0045934~negative regulation of nucleobase, nucleoside, nucleotide and nucleic acid metabolic process	0.006187558
GO:0051172~negative regulation of nitrogen compound metabolic process	0.00669907



Supplementary Figure 4, Rai et al

A**B**