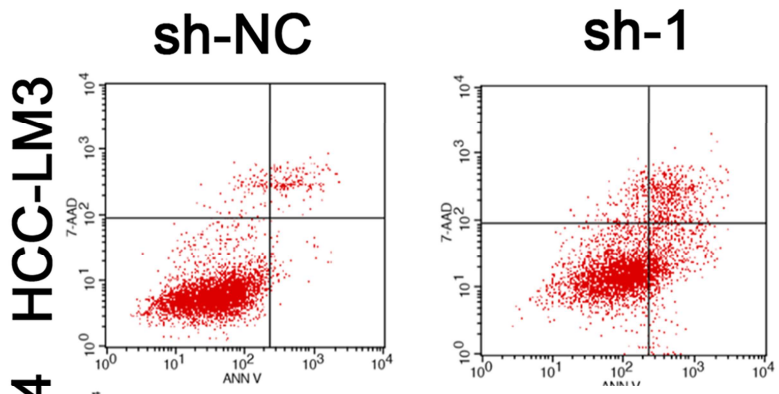
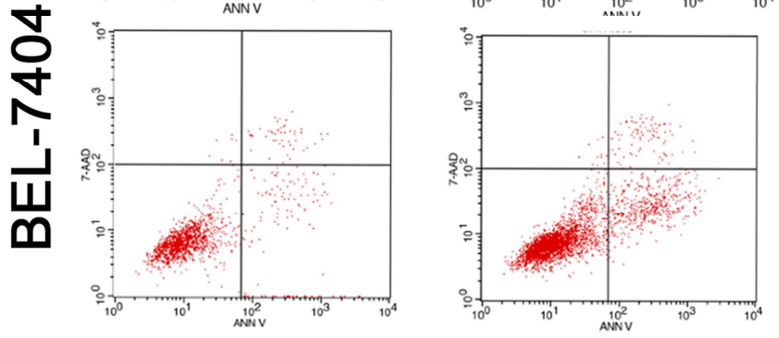
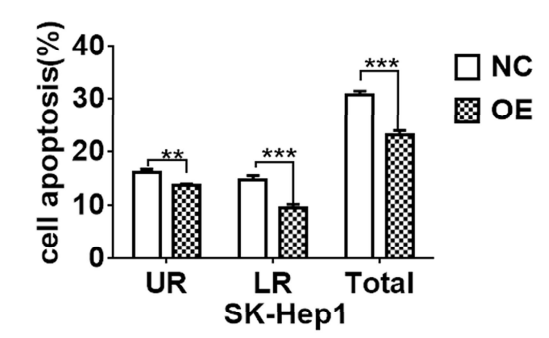
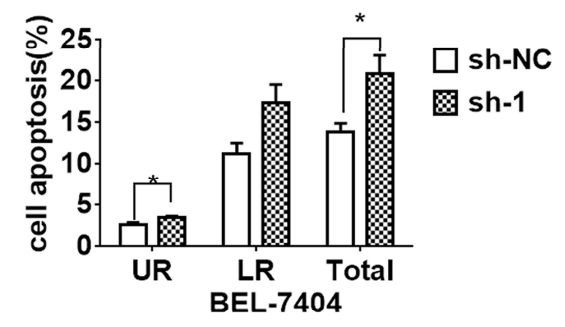
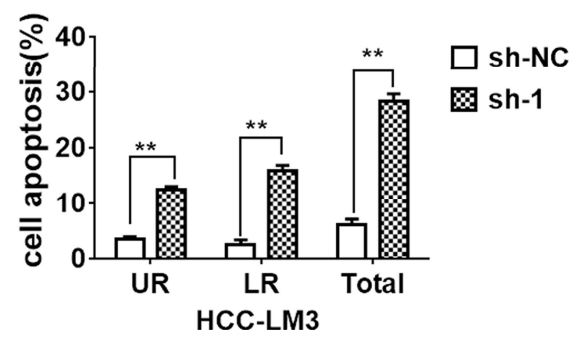
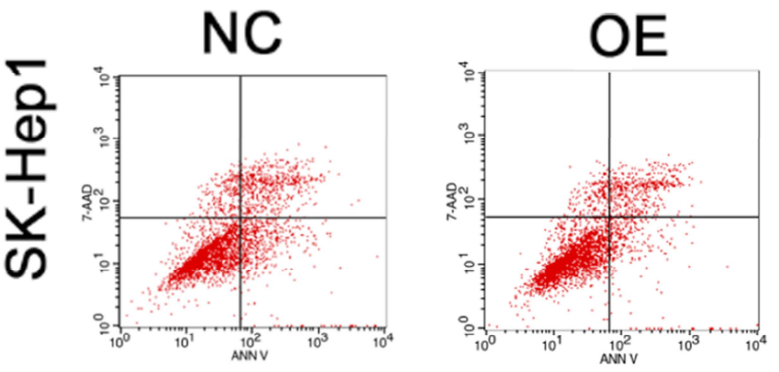
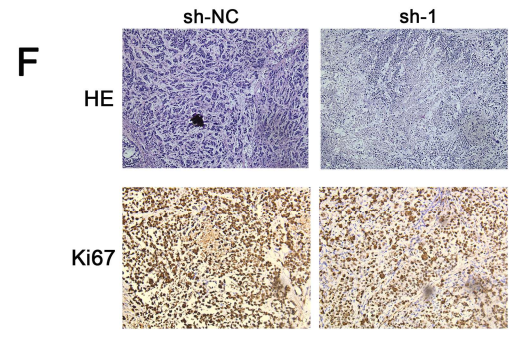
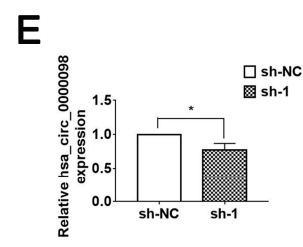
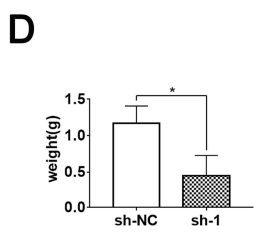
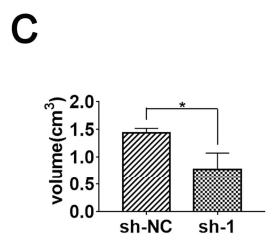
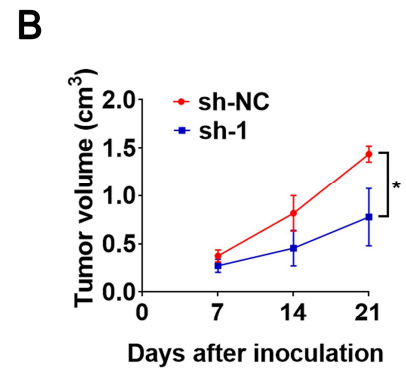
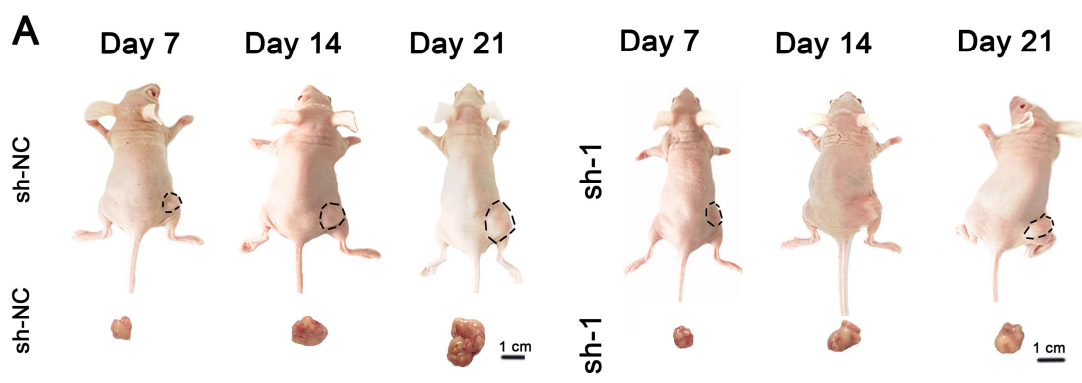
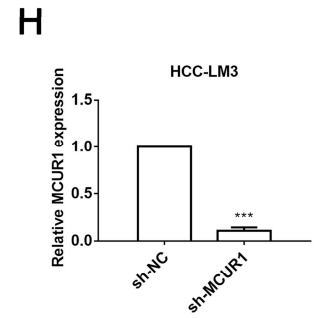
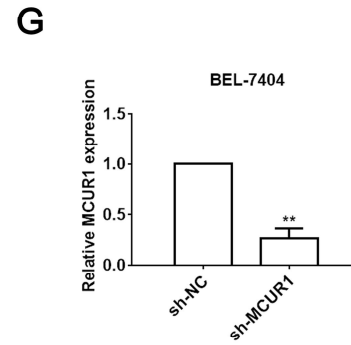
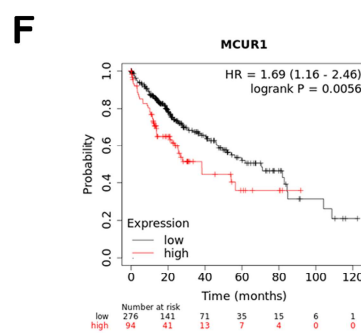
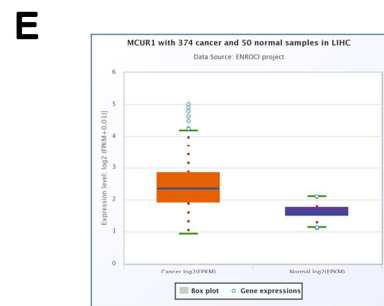
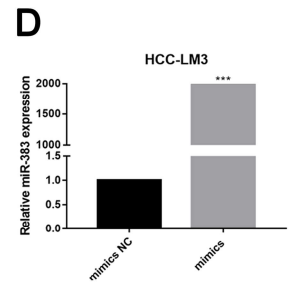
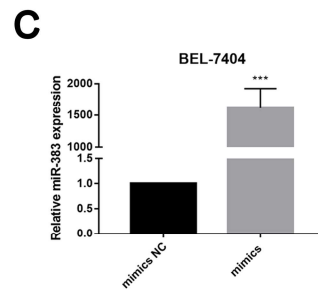
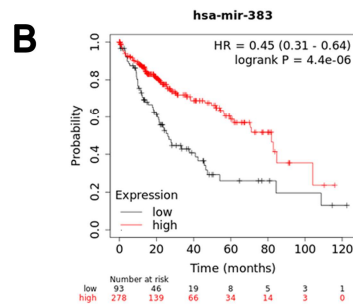
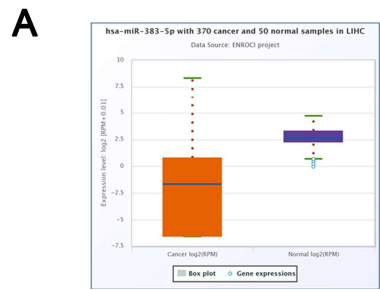


A**B****C**



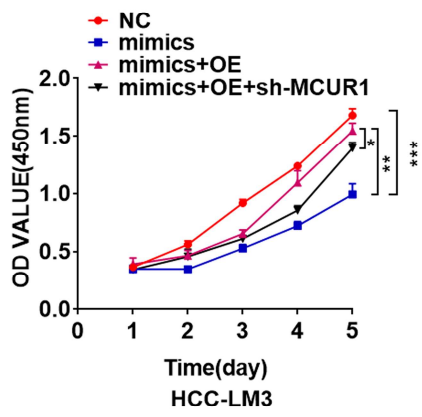
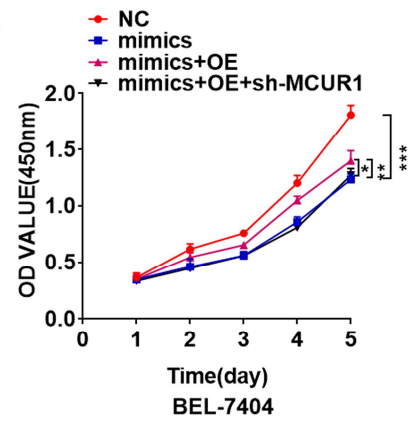


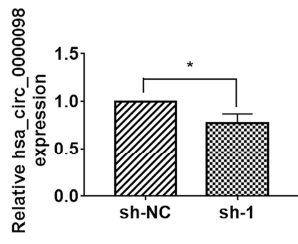
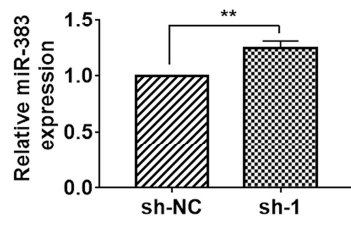
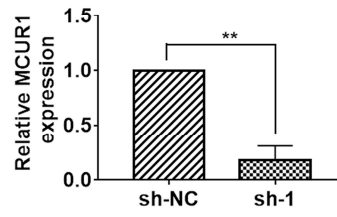
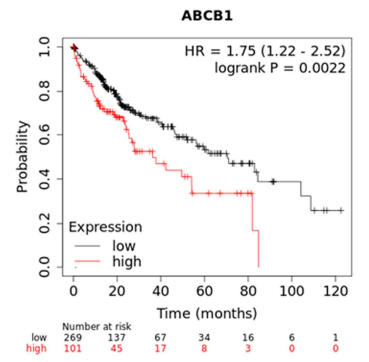
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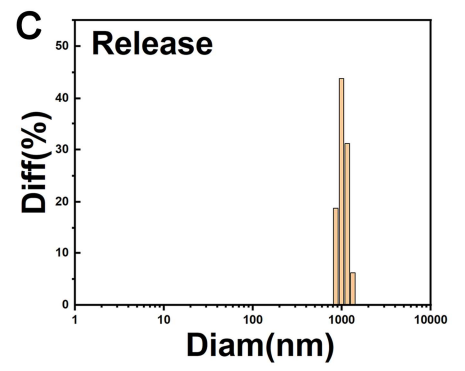
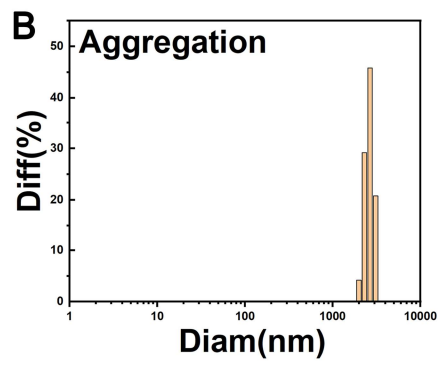
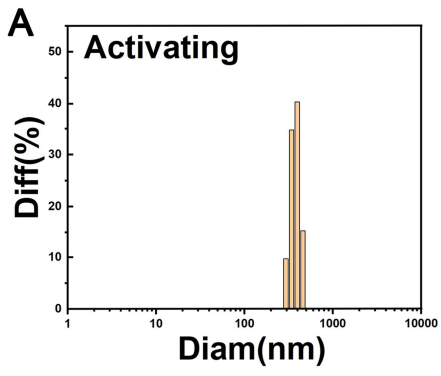
pathwayName	↕	log10(pval)
KEGG_Pathways_In_Cancer	↑	-5.555791
KEGG_Regulation_Of_Actin_Cytoskeleton		-3.55793
KEGG_Renal_Cell_Carcinoma		-3.61839
KEGG_P53_Signaling_Pathway		-3.69142
KEGG_Focal_Adhesion		-3.78243
KEGG_Vasopressin_Regulated_Water_Reabsorption		-3.40194
KEGG_Fc_Gamma_R_Mediated_Phagocytosis		-3.86731
KEGG_Pancreatic_Cancer		-3.09634
KEGG_Chronic_Myeloid_Leukemia		-2.90887
KEGG_Glioma		-2.91483
KEGG_Endocytosis	↑	-2.738511
KEGG_Gnrh_Signaling_Pathway		-2.74670
KEGG_Neurotrophin_Signaling_Pathway		-2.43671

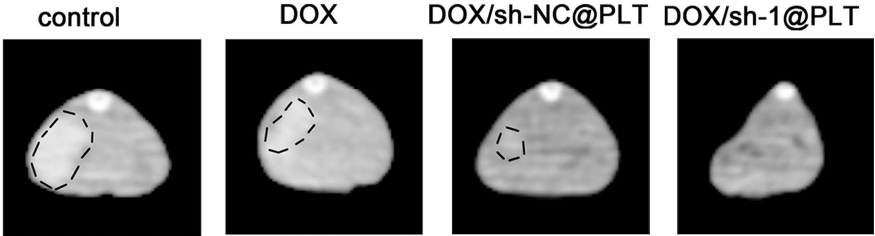
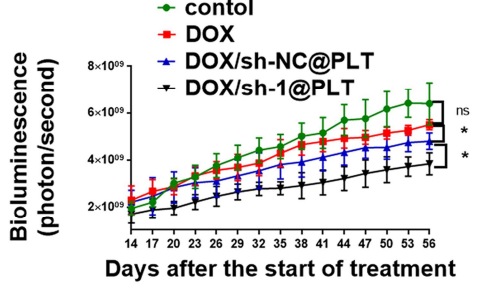
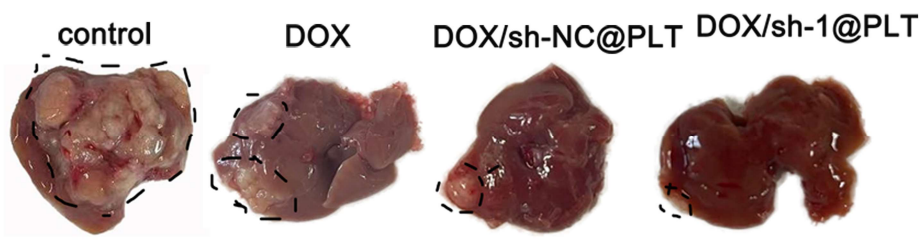
B

pathwayName	↕	log10(pval)
GO_Activation_Of_Gtpase_Activity	↑	-3.277011
GO_Adherens_Junction_Assembly		-3.07185
GO_Alpha_Beta_T_Cell_Differentiation		-2.71895
GO_Anatomical_Structure_Formation_Involved_In_Morphogenesis		-3.46456
GO_Camp_Mediated_Signaling		-2.78799
GO_Cell_Cycle		-4.45041
GO_Cell_Cycle_Checkpoint		-4.07137
GO_Cell_Cycle_Process		-3.26651
GO_Cell_Death		-3.12568

A**B**

A**B****C****D**



A**B****C****D**