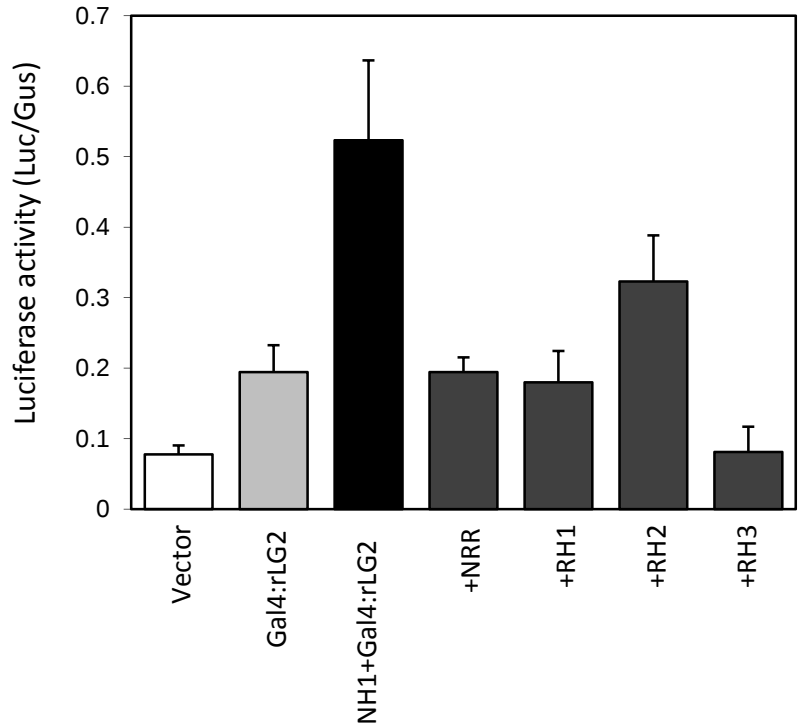
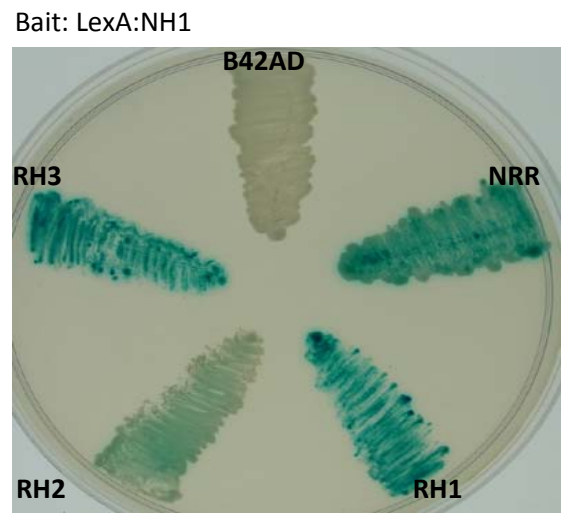


A



C



B

NRR	37	VEEFYAILRRMRDAT	RRLG-----ARPPPPRAPAWRPSFSWEDF
RH1	45	VERFYALLANIRALR	GMYSRYNGEEGAAGDGDGASG-RKRARRAEPPWRPAFRMEDF
RH2	45	MERFYALVANVRALR	AMFKAAALPSCREDDVSGGGGGEQRQKRPAAPWRPAFEMAVF
RH3	81	VDRFYALLDEVREMR	ELWRRNGDCVATKRTSVDGG----QKKQDRQQLWRPTFVMEDF
wNRR	33	VEEFYAILRRMRDAS	RRLVSGGVAAAT-----ARATPAPRAPAWCPSFSWEDF
Consensus		VERFYALL---	RWRP-F-w/MEDF

NPR1 interact.

NH1 interact.