

SUPPLEMENTAL METHODS

Recombinant DNA. Plasmids to express recombinant proteins were generated by PCR with the following primer pairs and template DNAs, and gel-eluted PCR product and indicated plasmid were digested with the indicated restriction enzymes. The primer sequences are provided in Supplementary Table 1. All plasmid constructions were verified by restriction mapping and DNA sequencing. pGex2TN-term: primers KB1109, KB1110 and pcDNAFLRHA (C.G Lee, University of Dentistry and Medicine of New Jersey), EcoRI digestion and ligation to pGEX2T (GE). pGex2TDEIH: primers KB1733, KB 1734 and template pcDNAFLRHA, EcoRI and BamHI digestion and ligation to pGex2T. pT7.7HisC-term: primers KB1661, KB1662 and template pcDNAFLRHA, NdeI and HindIII digestion and ligation to pT7.7 (USB). pFLAG-N-term: primers KB1726, KB1727 and template pcDNAFLRHA, EcoRV and NotI digestion and ligation to pcDNA3.1. PCR-based site-directed mutagenesis (Stratagene) introduced indicated mutations by the indicated primers and template DNA: pGEX2TK54A/55A: KB1674, KB1675 and pGex2TN-term; pGEX2TN-termK236E: KB1440, KB1441 and pGex2TN-term; pGEX2T N-termK54A/55A/236E: KB1440, KB1441 and pGEX2TN-termK54A/55A. To generate pMH110, junD sequences were amplified by PCE with primers KB1377, KB1378 and template genomic DNA from HEK293 cells; gel-elution, EcoRI digestion and ligation to pCRII-TOPO (Invitrogen).

In vitro transcription. T7 RNA polymerase was used to synthesize EMSA RNA probes (Promega). Template DNA was generated by PCR with the following primer/template DNA combinations. SNV PCE: primers KB588, KB 664/pYW100 (15); mutant SNV PCE: KB1352, KB588/pAC'Aall (1); junD PCE: KB 1702, KB 1378/pMH110.

LEGENDS SUPPLEMENTAL FIGURE

Fig S1. **Evaluation of equilibrium binding conditions.** Representative electrophoretic mobility shift assays to establish equilibrium conditions upon varied incubation time and temperature. (A) Comparison of incubation time on binding of N-term to ³²P-SNV PCE; 30 minutes (lanes 1-7) versus 60 minutes (lanes 8-14) (B) Comparison of incubation temperature on binding of N-term to ³²P-SNV PCE; ice (lanes 1-7) versus room temperature (lanes 8-14).

Fig S2. **RHA C-term and DEIH exhibit weak and undetectable interaction with PCE RNA, respectively.** Representative results of 3 or more electrophoretic mobility shift assays with indicated recombinant protein and ³²P-labeled RNA. (A) DEIH and ³²P-junD PCE. (B) DEIH and ³²P-SNV PCE (C) C-term and ³²P-junD PCE (D) C-term and ³²P-SNV PCE (E) C-term and ³²P-mutant SNV PCE.

Fig S3. **DEIH domain lacks detectable binding to PCE RNA.** Representative FA assays with DEIH and indicated RNA. (A) PCE^{AB}. (B) PCE^{AC}. (C) mutAC.

	Primer name	Primer Sequence
IN VITRO TRANSCRIPTION	KB588	GTACTACGGATTCAGTCCGG
	KB 664	AAATCTAGACGATAGAATTCTAATACGACTCACTATAGGGGGTGCCTCGCCGTCCTA CACATTGTTGTT
	KB 1352	AGACGATAGAATTCTAATACGACTCACTATAGGGGGTGCCTGCAACACAAACA ATGTGAC
	KB 1702	AAGGTAATACGACTCACTATAGGGAGGAGCCGCCGCCAGTGG
RECOMBINANT DNA	KB 1109	TTTTGCGAATCCCATGGGTGACGTTAAAAATTTTCTG
	KB 1110	AAAACGGAATTCGGGGGCAAATCTCAAGATTTAGC
	KB 1674	GGGAAATTCCACCAATGCAGCAGATGCACAAAGC
	KB 1675	GCTTTGTGCATCTGCTGCATTGGTGGAATTTCCC
	KB 1440	GGATCAAATAAGGAATTGGCAGCACAGTCC
	KB 1441	GGACTGTGCTGCCAATCCTTATTTGATCC
	KB 1733	AAAGGATCCGGGGCTACTGGATGTGGGC
	KB 1734	AAAGAATTCTCATCACCGGCCAGCTCGCCCTTTCC
	KB 1661	CTTAGGATCTAAGCTTGAATTCCTTATTAATAGCCGCCACCTCCTCT
	KB 1662	CGTATCCAGCCATATGCATCATCATCATCATCCTCCCAAGATGGCCCGA
	KB 1377	CCGAGGCTATAAGAGGGCGC
KB 1378	CCTCCGCTCCCCCGCCGCG	
RNA IP PCR	KB 1303	GTCGCCCCATCGACATGGACACG
	KB 1304	GCCGCTGTTGACGTGGCTGAGG
	KB 750	CTTTGGTATCGTGGAAGGACTC
	KB 752	GTTGCTGTAGCAAATTCGTTG
REAL TIME pcr	KB1252	TCACCCACACTGTGCCCATCTACGA
	KB1253	CAGCGGAACCGCTCATTGCCAATGG
	KB1614	GTAAGAAAAAGGCACAGCAAGCAG
	KB1615	CATTTGCCCTGGAGGTTCTG





