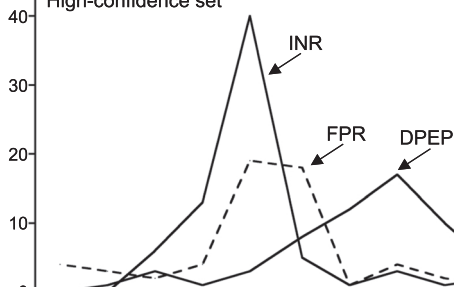


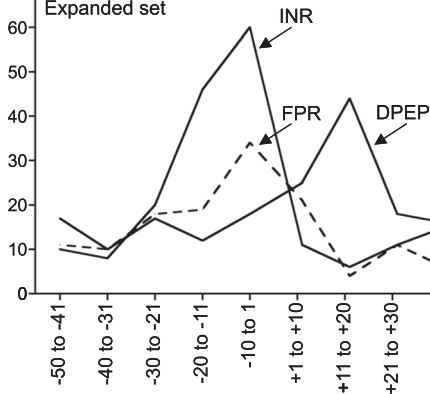
A

High-confidence set

Number of occurrences



Expanded set



Bases from predicted transcription start site

B

	-64	INR	FPR	DPEP
PINF	ACAGG	CCACTTT	GGAATTTGC	GTCCA-----CAAGCCC
PSOJ	TCGGC	TCACTTT	GGAATTTCGC	GTCTAGAAGCC-CAAGCAC
PRAM	ACTGC	TCACTTT	GGAATTTCGC	TCTTCAGTAGCCCAAGCAC

	-31	start codon
PINF	AGCCACCTTTTACCCCCCTACTTTCAACA	ATG
PSOJ	CGCC---TTCACCCCCTCTTCTTCCAACA	ATG
PRAM	C-----TTTTACCCCCCAGTCAACTAACC	ATG