Table III. Late gene NF-κB prediction and human-mouse promoter mapping.

Late Promoter	Chr.	Human Location (bp)	NFkB	MM	Sequence	Chr.	Mouse Location (bp)	NFkB	MM	Sequence
Cyb5	18	70110091-70111159	129 (-)	0.871	gtggtaTTTCCtctgt	18	85362396-85363465	497 (-)	0.869	GGAAAccacg
			568 (+)	0.988	caataGGAAActccg			982 (-)	0.972	GGAATgtccc
			963 (-)	0.861	gtggggTTCCCggcgc					
			981 (-)	0.972	GGAATgtccc					
ECE1	1	21362214-21363213	100 (-)	0.856	cagggaTTCCCgcctt	4	136319312-136363170	NF	NA	NA
			859 (-)	0.829	ggggtcTCCCCgggag					
ICAM1	19	10241779-10242835	448 (-)	0.851	GAAATgcccg	9	20941067-20942141	174 (-)	0.999	GGAAAccccc
			518 (+)	0.86	ggaggATTCC			431 (-)	0.808	caggacTTTCTcacag
			832 (-)	0.986	GGAAAttccg			828 (-)	0.977	GGAAAttcct
			937 (+)	0.869	cgaggTTTCC			933 (+)	0.869	cgaggTTTCC
IL27RA	19	14002560-14003684	138 (+)	0.968	tggccTTTCC	NF	NA	NA	NA	NA
			947 (+)	0.992	GGGACtttcc					
KLRC2	12	10479853-10480897	NF	NA	NA	6	130353839-130354838	NF	NA	NA
NAF-1	5	150424850-150441744	150 (+)	0.885	gtctaGGAAAtcccag	11	54580586-54603649	268 (+)	0.955	tgggtCTTCC
			218 (-)	0.881	ctggatTTGCCcaccc			588 (+)	0.879	ggagaTTTCC
			452 (+)	0.86	gggtcTTTTC			615 (-)	0.981	GGAATccccc
			497 (-)	0.814	ctggagTGCCCtggac			913 (+)	0.983	ggggaATTCC
			570 (-)	0.919	ttggcagGTCCCct					
			765 (+)	0.844	tagagGGGAAggccgc					
			914 (-)	0.998	ggggacTTTCCcaggg					
			953 (+)	0.94	cggcaCTTCC					
NFkB2	10	104143320-104145706	60 (+)	0.828	cgtggTGAAAcccca	19	45645210-45646595	80 (+)	0.973	cgggaATTCC
			616 (+)	0.991	GGGGAtcccc			109 (+)	1	ggggcTTTCC
			675 (-)	0.957	GGAAGctccc			715 (+)	0.931	gagggGGAAActcctc
			898 (-)	0.977	tgggaaTTCCCccctc			793 (+)	0.95	GGAAGaaccc
			928 (+)	1	ggggcTTTCC			842 (-)	0.972	GGAATgtccc
								890 (+)	0.863	ctggcTGAAAatccca
								989 (-)	0.923	GGAACcgcct
								1088 (+)	0.992	GGGACtttcc

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NK4	16	3054660-3055813	142 (+)	0.939	tggtaCTTCC	NF	NA	NA	NA	NA
			645 (+)	0.947	caGGGAGtttcccc					
			677 (+)	0.977	tggacTTTCC					
PTGES	9	129594846-129595898	375 (-)	0.948	GGAAGggcc	2	30863242-30864320	199 (-)	0.936	GGAAGggcca
			459 (-)	0.868	GAAAAgtccc					
			799 (-)	0.867	gaggatTTGCCtggaa					
			985 (+)	0.866	tgtgcTTTCC					
TAP1	6	32929572-32930601	583 (-)	0.962	ggtaaGTCCC	NF	NA	NA	NA	NA
TRAF1	9	120765569-120769727	1328 (+)	1	ggggaTTTCC	2	34918072-34922576	280 (+)	0.844	tacacGGGAAgacctt
			1385 (+)	0.965	gggtaATTCC			1177 (+)	0.913	accctGGGAGattccca
			1743 (+)	0.852	cagagGGGAAgacctc			1181 (-)	0.945	tggagaTTCCCa
TRAF3	14	102312569-102406291	255 (-)	0.86	GGAAAgtgcc	12	105963380-105982148	NF	NA	NA
			938 (+)	1	ggggaTTTCC					
			959 (-)	0.924	GGAAGtgcct					
			1050 (+)	0.996	cGGGACtttcca					
TRIM16	17	18544451-18545450	NF	NA	NA	11	62432053-62433052	659 (-)	0.923	GGAAGagcct

Table III. Late gene NF-κB prediction and human-mouse promoter mapping. Same as in Table II for the Late gene promoters.