## chr4: HMG box

AAINQILGRRWHELSREEQSKYYEKARQERQLHMELYPGWSARDNYGYVS

pan

MPHTHSRHGSSGDDLCSTDEVKLFKDEGDREDEKLSSENLLVEEKSSLID Beta-catenin binding site (By similarity) MPHTHSRHGSSGDDLCSTDEVKIFKDEGDREDEKISSENLLVEEKSSLID 0

0

50

50

100

100

150

150

200

200

250

250

300

300

350

0

LTESEEKGHKI SRPDHSPVFNKLDTHAPSFNMGYLVSPYSYANGSPSGLP

<u>LTESEEKGHKI SRPDHSPVFNKLDTHAPSFNMGYLVSPYSYANGTPSGLP</u> VTMANKI GLPPFFCHNADPLSTPPPAHCGI PPYQLDPKMGLTRPALYPFA VTMANKI GLPSFFCHNADPLSTPPPAHCGI PPYQLDPKMGLTRPALYPFA GGQYPYPMLSSDMSQVASWHTPSVYSASSFRTPYPSSLPINTTLASDFPF

GGQYPYPMLSSDMSQVASWHTPSVYSASSFRTPYPSSLPINTTLARNLEA RFSPSLLPSVHATSHHVI NAHSAI VGVSSKQECGVQDPTTNNRYPRNLEA

sec

sec sim

mel

sim

mel

mel

KHSSNAQSNESKEATNDKKKPHIKKPLNAFMLYMKEMRAKVVAECTLKES

KHTSNAQSNESKETTNDKKKPHIKKPLNAFMLYMKEMRAKVVAECTLKES

HMG box

AINQILGRRWHELSREEQSKYYEKARQERQLHMELYPGWSARDNYGYVS

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sec sim

## pan chr4: HMG box

+. HIVIG DOX

sec sim mel	KKKKRKKDRSTTDSGGNNMKKCRARFGLDQQSQWCKPCRRKKKCI RYMEAYMEA KKKKRKKDRSTTDSGGNNMKKCRARFGLDQQSQWCKPCRRKKKCI RYMEA  NLS	350 4 400
sec	L NGNGPAEDGSCFDEHGSQL SDDDEDDYDDDKL GGSCGSADETNKI EDED	400
sim	L NGNGPAEDGSCFDEHGSQL SDDDEDDYDDDKL GGSCGSADETNKI EDED	54
mel	L NGNGPAEDGSCFDEHGSQL SDDDEDDYDDDKL GGSCGSADETNKI EDED	450
sec	SESLNQSMPSPGCLSGLSSLQSPSTTMSLASPLNI NANSATNVI FPASSN	450
sim	SESLNQSMPSPGCLSGLSSLQSPSTTMSLASPLNMNANSATNVI FPASSN	104
mel	SESLNQSMPSPGCLSGLSSLQSPSTTMSLASPLNMNANSATNVI FPASSN	500
sec	AVLI VGADQSTVQQRPTLVSTSGSSSGSTSSI STTPNTSSTVSPVTCMTG	500
sim	AVLI VGADQSTVQQRPTLVSTSGSSSGSTSSI STTPNTSSTVSPVTCMTG	154
mel	ALLI VGADQPTAQQRPTLVSTSGSSSGSTSSI STTPNTSSTVSPVTCMTG	550
sec	PCPGSSQERAMMLGNRFSHLGMGLSPPVVSTSTSKSEPFFKPHPTVSNNP	550
sim	PCPGSSQERAMMLGNRFSHLGMGLSPPVVSTSTSKSEPFFKPHPTVSNNP	204
mel	PCLGSSQERAMMLGNRFSHLGMGLSPPVVSTSTSKSEPFFKPHPTVCNNP	600
sec	I FSLPSI GNCSLNI SSMPNTSRNPI GANPRDI NNPLSI NQLTKRRDYKNV	600
sim	VFSLPSI GNCSLNI SSMPNTSRNPI GANPRDI NNPLSI NQLTKRRDYKNV	254
mel	I FALPSI GNCSLNI SSMPNTSRNPI GANPRDI NNPLSI NQLTKRREYKNV	650
sec	DLI EASESKTI VAHAASSI I QHVAVNGYHANHSLLNSNLSQLHHQLNSRT	650
sim	DLI EASESKTI VAHAASSI I QHVAVNGYHANHSLLNSNLSQLHHQLNSRT	304
mel	ELI EASESKTI VAHAATSI I QHVAVNGYHANHSLLNSNLGHLHHQLNNRT	700

## pan

chr4: HMG box

sec sim mel	EKANRSEKAMLSVSNHSVNSNECHKESDSQAI VSSNPPNAGSSDNGVI SV EKANRSEQAMLSVSNHSVNSNECHKESDSQAI VSSNPPNAGSSDNGVI FV ENPNRSEQTMLSVSNHSVNSSECHKESDSQAI VSSNPPNAGSSDNGVI SV 75	4
sec sim mel	S 701 S 355 S 751	