

TIG-3.B	-----NAPNFDVMVFQP-NTVTAGTSDCVGCCVIPFYVNFTEIGWNDWILSPPGF	49
INHA_HUMAN	-----STPLMSWPWSPSALRLLQRPPEEPAAHANCHRVALNISFQELGWERWIVYPPSF	54
GDF9_HUMAN	-GQETVSSSELKK-PLGPASFNLSEYFRQFLPQNECELHDFRLSFSQLKWDNWIVAPHRY	58
BMP15_HUMAN	-QADGISAEVT-----ASSSKHSGPENNQCSLHPFQISFRQLGWDHWI IAPPFY	48
INHBA_HUMAN	-----GLECDGKVNICCKKQFFVFSFKDIGWNDWI IAPSGY	35
TIG-2	-----SESAYFEKPNENERCQRKGLYVDFDILGWKQWVIAPEGF	39
BMP2_HUMAN	-----QAKHKQRKRLKSSCKRHPLYVDFSDVGWNDWIVAPPGY	38
DBL-1	SAQTGNSEKKNR-----KKGRKHHNTEAESNLCRRTDFYVDFDDLNWQDWIMAPKGY	52
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TIG-3.B	YANVCSDTV-----ST-----ESDEVYQFMKAAISDLPEPKCA--PNYYGSVDMIV	94
INHA_HUMAN	IFHYCHGGCGLHIPNLSLP----VPGAPPTPAQPYSLPGAQPCCAALPGTMRPLHVRT	110
GDF9_HUMAN	NPRYCKGDCPRAVGHRYGSP----VHTMVQNI IYEKLDSSVPRPSCV--PAKYSPLSVLT	112
BMP15_HUMAN	TPNYCKGTCLRVLRLDGLNSP----NHAI IQNLINQLVDQSVPRPSCV--PYKYVPISVLM	102
INHBA_HUMAN	HANYCEGECPSHIAGTSGSSLSFHSTVINHYRMRGHSPFANLKS CCV--PTKLRPMSMLY	93
TIG-2	SAFYCSGDCSAPFSKEMNAT----SHAI VQSTLHRVRPNSTTPAKCA--PSSLGSYKILF	93
BMP2_HUMAN	HAFYCHGECPFPLADHLNST----NHAI VQTLVNSVN-SKIPKACC V--PTELSAISMLY	91
DBL-1	DAYQCQGSCPNMPAQLNAT----NHAI IQSLLHSLRPDEVPPPCCV--PTETSPLSILY	106
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TIG-3.B	ALS-PRDIRKTRVHGLRALSCSCT	117
INHA_HUMAN	TSDGGYSFKYETVPNLLTQHCACI	134
GDF9_HUMAN	IEP-DGSIAYKEYEDMIATKCTCR	135
BMP15_HUMAN	IEA-NGSILYKEYEGMIAESCTCR	125
INHBA_HUMAN	YDD-GQNI IKKDIQNMIVEECGCS	116
TIG-2	VDQ-NKQVQIKRYRDMVVDECGCH	116
BMP2_HUMAN	LDE-NEKVVLKQYQDMVVEGCGCR	114
DBL-1	MDV-DKVIVIREYADMRVESCGR	129
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