slbo 2L:bZip

sec	ML NMESPQMYADAVQTL AQL DL KKEPPL QQATI GQI TLTAMSTAQQQQQQ	50
sim	ML NMESPQMYADAVQTL AQL DL KKEPPL QQATI GQI TLTAMSTAQQQQQQ	50
mel	ML NMESPQMYADAVQTL AQL DL KKEPPL QQATI GQI TLTAMSTAQQQQQQ	50
sec	QQQQQQQQQQQQQQQQQQQTTDANNNTSQDAALLVKQHAMHQMQQV	100
sim	QQQQQQQQQQQQQQQQQQQTTDANNNTSQDAALLVKQHAMHQMQQV	100
mel	QQQQQQQQQQQQQQQQPQFQFQQTTDANNNTSQDAALLVKQHAMHQMQQV	99
sec sim mel	AALGNNNNLLQKQMLQQYSTQTDLDELTTQEITLDLQHLIDDQFRDTETLAALGNNNNLLQKQMLQQYSTQTDLDELTTQEITLDLQHLIDDQFRDTETLAALGSNNNLLQKQMLQQYSTQTDLDELTTQEITLDLQHLIDDQFRDTETL	150 150 149
sec sim mel	GI FSDMVTSPGGLSATLPPTGMVSAAAKVLQQQTLRNQHGYGGRGGGG-AGI FSDMVTSPGGLSATLPPSGMVSAAAKVLQQQTLRNQHGYGGRGGGG-AGI FSDMVTSPGGLSATLPPSGMVSAAAKVLQQQTLRNQHGYGGRGGGGA	199 199 199
sec	GGAL AYMPQPVHATYNNSSDENSSVGSDSSTI KEEPI DPEYRRHL QEAAS	249
sim	GGAL AYMPQPVHATYNNSSDENSSVGSDSSTI KEEPI DPEYRRHL QEAAS	249
mel	GGAL AYMPQPVHATYNNSSDENSSVGSDSSTI KEEPI DPEYRRHL QEAAS	249
sec sim mel	QQAAFMGNGAGLYNGYGSGANGLSGGGNPLNGGNTTPSSNGSNGSTGSSN QQAAFMGNGAS	299 260 299
sec	GSQFTNLTTANVLAHHNLPHLAAAAGAHNLLKQHSKLHAQQQHQQHQQQQ	349
sim	LKQHSKLHAQQQHQQHQQQQ	280
mel	GSQFTNLTTANVLAHHNLPHLAAAAGAHNLLKQHSKLHAQQQHQQHQQQQ	349

slbo

2L:bZip

sec	QHRKHSNKHVDKGTDEYRRRERNNI A			399
sim	QHRKHSNKHVDKGTDEYRRRRERNNI A		_	330
mel	QHRKHSNKHVDKGTDEYRRRERNNI A	VRKSREKAK <u>VRSRI</u>	VEERVKSLL	399
	basic (DNA	binding)	leucine-	
sec	KEKDALI RQLGEMTNELQLHKQI YMQL			449
sim	KEKDALI RQLGEMTNELQLHKQI YMQL			380
mel	KEKDALI RQLGEMTNELQLHKQI YMQL	MNHANPEVSRVCR	SFLNTNEHSL	449
	zipper			