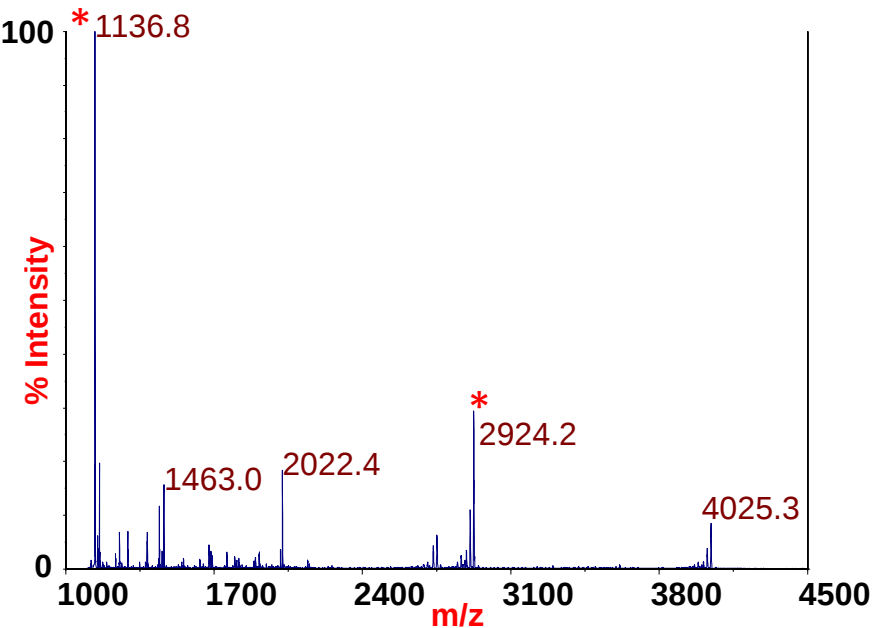


A



B

1	MPKINSFNNDPVNDNTILYIKPGGCQQFYKSFNIMKNIWIIIPERNVIGTIPQNFLPPTS	60
61	LKNGDSSYYDPNYLQNDQEKDRFLKIVTKVFNRINDNLSGRILLEELSKANPYLGNDNTR	120
121	DDDFIINDGSAVPIQFSNGSQSILLPTVIIMGAEPDLFETNSSNVSLINNYSPSNHGFGS	180
181	IAIVTFSPEYSFRFNDNSMNEFIQDPALTLIHELIIHSLHGLYGAKGITTKYTITQQQNPL	240
241	ITNIRGINIEEFLTFTGGNNLNIITSSQLNDIYTNLLDDYKKIASKLSKVQVSNPQLNPYK	300
301	DVFQEKYGLDKNASGIYSVNINKFNDIFKKLYSFEFDLATKFKQVKCRETYIGQYKYFKL	360
361	SNLLNDSIYNISEGYNINTLNVNFRGQNPNLNPRIITPITDRGLVKKIIRFCKNIVSVKG	420
421	IRKSICIEVNNGDLFFVASEKSYNNDSINIPKEIDDTVTLLNNNYENDLDQVILNFNESA	480
481	PGLSDKKLNIISIQDDVYIPKYDSNGTSDIEQYDVSELNVFFYLDAQKVPEGENNVNLTSS	540
541	IDTALLEQSKIYTTFFSSEFINNVNKPVQAALFVGWIQQVLVDFTTTEATQKSTVDKIADIS	600
601	IVVPYIGLALNIGNESQKGNFKDALELLGAGILLEFVPELLIPTILVFTIKSFLGSSDNK	660
661	NKVIKAINNALKERDEKWKEVYSFIVSNWITKINTQFNKRKEQMYQALQNQVNALKTIIE	720
721	SKYNSYTLEEKNELTNKYDIEQIENELNQKVSIAMNNIDRFLTESSISYLMKLINEVKIN	780
781	KLREYDENVKTYLLDYITKHGSILGESQQELNSMIIDTLNNSIPFKLSSYTDDKILISYF	840
841	NKFFKTIKSSSVLSMRYKNDKYIDTSGYDSNININGDVFIYPTNKNQFGIYNSKLSEVNI	900
901	SQNDYIIYDNKYKNFSISFWVRIPNYNNKIVNVNNEYTIINCMRDNNSGWKISLNHNEII	960
961	WTLQDNAGINQKLVFYKGNANGISDYINKWIFVTITNDRLGYSKLYINGHLIDQKSILNL	1020
1021	GNIHVSDNIFKIVNCSYTRYIGMRYFNIFDKELDETEIQTLYNNEPNANVLKDFWGNYL	1080
1081	LYNKEYYLLNMLKPSKITISHNRDLTFSIYNNRNIVNGLYRLYSGIKVKIQKINDSDTRDN	1140
1141	IVRDNDQVYVNYINGNVYYSLYADTNATNKEKTIKSSTSGNRFNQVVMNSVRNCTMNF	1200
1201	KNNNGHDIGLLGFKSNALVASTWYYTNMRDHTNSNGCFWSFIPEENGWQEH	1260