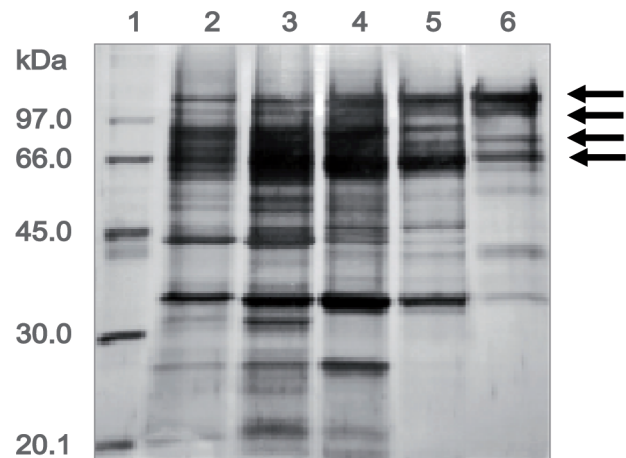
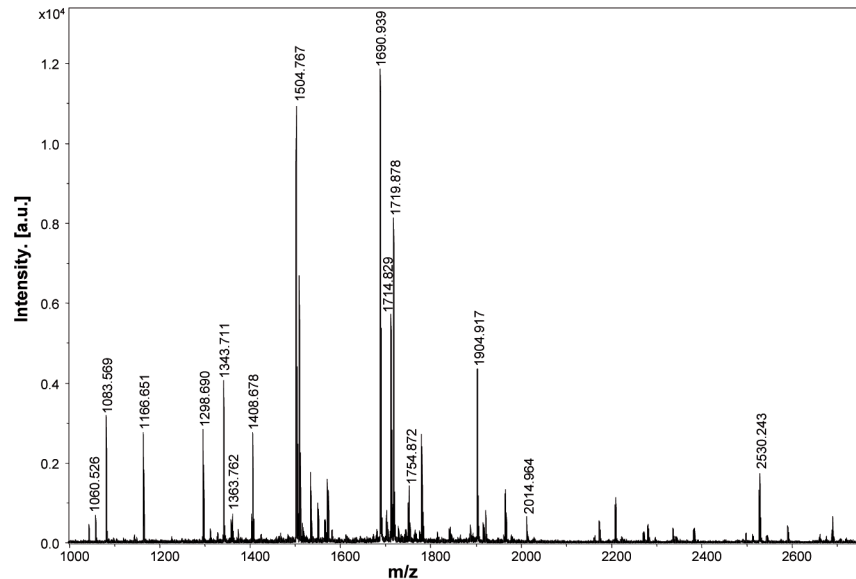


A**B****C**

gi|67541975 Mass: 100148 Score: 209 Expect: 4e-15 Queries matched: 15

hypothetical protein AN7151.2 [*Aspergillus nidulans* FGSC A4]

Observed	Mr (expt)	Mr (calc)	Delta	Start	End	Miss	Peptide
1060.5257	1059.5184	1059.5308	-0.0124	808	- 818	0	K.VGGGLSSAQER.F
1083.5694	1082.5622	1082.5720	-0.0098	15	- 25	0	R.SALGIGEPSPR.I
1166.6512	1165.6439	1165.6455	-0.0016	298	- 309	0	K.SVLAAPDAPAVR.R
1298.6896	1297.6823	1297.6601	0.0222	440	- 449	0	K.LHENIWSMR.G
1343.7111	1342.7039	1342.6782	0.0257	466	- 477	0	R.LGWTGDIHAFAR.T
1363.7621	1362.7548	1362.7394	0.0154	695	- 705	1	R.LVDIIKENEYK.V
1408.6784	1407.6712	1407.6517	0.0194	137	- 148	0	K.DFDVDEEILSAR.L
1504.7665	1503.7592	1503.7357	0.0235	478	- 490	0	R.TANFIYDTSGFLR.G
1690.9389	1689.9317	1689.9049	0.0267	843	- 859	0	K.VPPNSEAVVSLPGAPTR.G
1714.8285	1713.8213	1713.7998	0.0215	819	- 833	0	R.FLSPYGSAESSWEVR.D
1719.8783	1718.8710	1718.8951	-0.0240	802	- 818	1	R.FAVDVKVGGLSSAQER.F
1754.8725	1753.8652	1753.8345	0.0307	560	- 575	0	K.AMLSEQYAGATAWLDK.G
1904.9169	1903.9096	1903.8846	0.0250	450	- 465	1	R.GNFLSIPTDCPQORDER.L
2014.9637	2013.9564	2013.9431	0.0133	819	- 836	1	R.FLSPYGSAESSWEVRDGK.F
2530.2426	2529.2354	2529.2645	-0.0291	348	- 370	0	R.GSTISFVHTEVMENGEVATRPLR.N