

**Table S1.** MSDIN family and novel cyclic peptides in *Amanita pallidorozea*

Leader Peptide	Core Peptide	Recognition Sequence	Monoisotopic Mass
MSDINATRLP	<b>IWGIGCNP</b>	CVGDEVTALITRGEA	918.3541 ( $\alpha$ -amanitin)
MSDINATRLP	<b>IWGIGCNP</b>	CVGDEVAALLTRGEA	918.3541 ( $\alpha$ -amanitin)
MSDINATRLP	<b>IWGIGCDP</b>	CVGDDVTGVLTRGEA	919.3382 ( $\beta$ -amanitin)
MSDVNATRLP	<b>AWLVDCP</b>	CVGDDINRLLTRGEK	846.3217 (phalloidin)
MSDINATRLP	<b>AWLMTCP</b>	CVGDDVNPTLTRGE	802.3506
MSDINASRLP	<b>AWLATCP</b>	CAGDDVNPTLTRGE	788.3160 (phalloidin)
MSDINATRLP	<b>DPRRLIP</b>	GGSDDVDSALTRDE	960.5869
MSDINTARLP	<b><u>EFIVFGIFP</u>★</b>	CVGDDIQTVLTRGE	1049.5586
MSDINASRLP	<b>FFPEVGFFP</b>	CVGDDTNPILTRG	1067.5117
MSDINTIRVP	<b>FPWTGPFVP</b>	CVGDDVGSVLTHGE	1028.5120
MSDINTARLP	<b><u>FVIIPPFIFP</u>★</b>	CVSDDIEMVLTRGE	1170.6841
MSDINATRLP	<b>HPFPLGLQP</b>	CAGDVDNLTTLTRGE	986.5338
MSDINATRLP	<b>HPFPLGLQP</b>	CAGDVDNLTTLTRGE	986.5338
MSDINAIRAPLP	<b>IFSLNP</b>	CVGDDVEVL	671.3643
MSDINVIRLP	<b>IFIYFP</b>	CVGDNIDNTLT	780.4210
MSDINAARLP	<b>LVYMILFP</b>	SVGDDIDVVLGRGE	976.5456
MSDINATRLP	<b>LGRPESLP</b>	CVGDDVNYILVSG	849.4709
MSDINTARLP	<b><u>MHILAPPP</u>★</b>	CVSDDIEMVLTRGE	856.4629
MSDINATRLP	<b>NWHAGPTRPP</b>	CVADDVSLTLTRGE	1113.5468
MSDINAARLP	<b>NLFVWIPP</b>	CISDDINSTLTRGE	966.5327
MSDINTARLP	<b>VFFMPPFIPP</b>	CVSDDIQMVLTRGEK	1172.6093
MSDINTTRLP	<b><u>YFFNDHPP</u>★</b>	CASDDIQMVFTTRGE	1017.4345
MSDVNATRLP	<b>MAFPEFLA</b>	CVGDDVNHTLTRGE	906.4310
MSDNASRLP	<b>YFFGFTP</b>	SVGDDVNPTLTRGE	859.3905
MSDTNATRLP	<b>SIFIVYPP</b>	CVSDDVNSTLTRGE	916.5058
MSDMNVRLP	<b>ISDPTAYP</b>	CVGDDIQAVLGRGE	844.3967
MSDMNVRLP	<b>ISDPTAYP</b>	CVGDDIQAVLGRGE	844.3967
MSDMNTACLP	<b>IFIVFPIPP</b>	CVSDDIQTVLTRGE	1023.6157
MSDLNATRLP	<b>FNLFREFYP</b>	CIGDDSGSVLTLGEG	1181.6022
MSETNAARLP	<b><u>TIHLFSAP</u>★</b>	SVGDDIEVVLGRGE	866.4650
MSNINTARLP	<b>FLVPSFPP</b>	CVSDDIQIVLTRGEK	884.4796
MADINASRLP	<b>LNILPFHLPP</b>	CVSDDATSTLTRGE	1141.6648
MYDINTTRLP	<b>HFFNLTPP</b>	CVRDDI	953.4759

<sup>1</sup>★and underlined letters indicate novel cyclic peptides detected with MS and MS/MS.

Predicted monoisotopic masses of cyclic peptides with no further modifications.

<sup>2</sup>The monoisotopic masses are for unmodified cyclic peptides based on MSDIN core peptides, except for major toxins.