

Nonribosomal peptide synthetase gene clusters and characteristics of predicted NRPS-dependent siderophore synthetases in *Armillaria* and other species in the Physalacriaceae

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Table S1: Positions of NRPS clusters as predicted by fungiSMASH

Species	Scaffold number	Position (nucleotide)
<i>A. borealis</i>	68	133,657 – 189,842
<i>A. cepistipes</i>	261	5,409,961 – 5,466,141
<i>A. fumosa</i>	1	4,563,839 – 4,619,965
<i>A. mellea</i>	13	306,941 – 363,118
<i>A. nabsnona</i>	22	557,122 – 613,305
<i>A. novae-zelandiae</i>	11	736,489 – 792,706
<i>D. ectypa</i>	3	3,707,080 – 3,763,248
<i>D. tabescens</i>	4	273,770 – 329,930
<i>G. necrorhizus</i>	2	745,974 – 802,126
<i>C. torrendii</i>	88	4,791 – 60,480
<i>O. mucida</i>	46	99,521 – 174,178