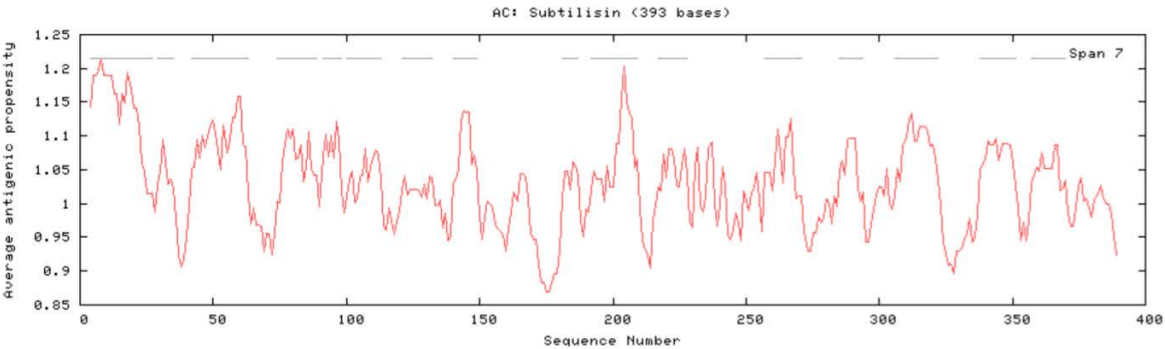


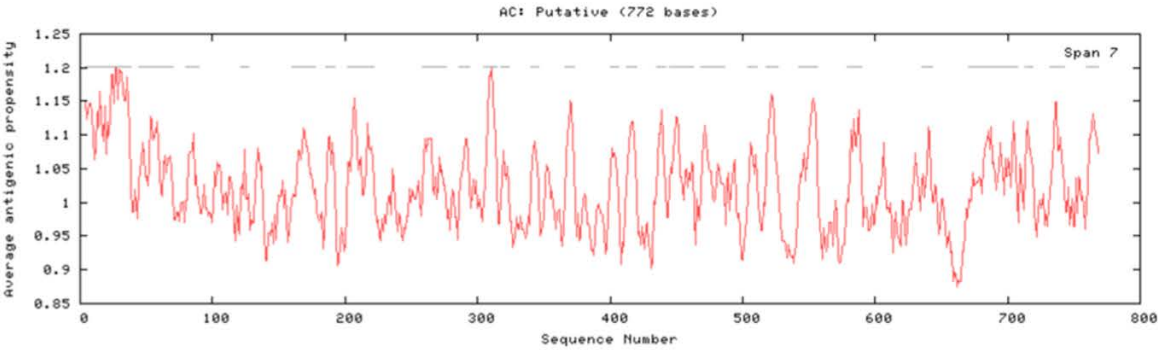
Figure S1

NCBI Accession no. XP_018384475.1
(subtilisin-like serine protease-like protein PR1A (*Alternaria alternata*))



n	Start Position	Sequence	End Position
1	4	SLLLALLPVALALPAPVIVPRAGS	27
2	29	IPGRYIV	35
3	42	LQALIDAALKVLKKDPVHVYKF	63
4	74	ADDIVELLQNLPGVDY	89
5	91	EQDAVVKA	98
6	100	LGEIHLEKKAYVTQ	113
7	121	GRISHQAKGISS	132
8	140	GAGTCAYVID	149
9	181	HGTHCAG	187
10	192	KTYGVAKKTKLYAVKVLD	209
11	217	SGVIAGINYVAN	228
12	257	AANAVTAGVFMATAA	271
13	285	SEPTVYTVGA	294
14	306	NYGSIVDILAPGVSILS	322
15	338	ATPHVAGLGAYILG	351
16	357	TPAALSSRLTALSL	370

NCBI Accession no. KZN92610.1
Putative dipeptidyl-peptidase (*Penicillium chrysogenum*)



n	Start Position	Sequence	End Position
1	4	SSLCVWSPSIYISVTVRSAVRLLFCAFLTVCVAGY	38
2	44	RVFQLTTALGSIIVSATALTPEQLIAA	70
3	80	PSGEVALFSTS	90
4	121	NVSELTW	127
5	160	ANSAYKAASLPAPLSGLKT	178
6	185	DINFLVYG	192
7	202	NKELVAAPLSSARIHDSIYVR	222
8	258	SLKNLVAGVKNLESPYPPF	276
9	286	SNGKLVAFK	294
10	305	HTASYIYVVP	315
11	318	SKKAVAI	324
12	340	SSSPVFS	346
13	366	RRTVYVYT	373
14	400	GKSLVLN	406
15	412	RSRLFLPA	420
16	435	GSVAAYYG	442
17	445	NGEYLVSSSAIHTSRTVYT	463
18	469	GVTGVIFSANEVDPALKN	486
19	504	IHSWIIYP	511
20	518	KSYPLLFYIH	527
21	543	NYKVFADQGYYVIA	556
22	579	SYPYDDLKVCWE	590
23	635	DGTFVADAK	643
24	671	DPSAPEHIKQFATPQLIHSDDLRYRLPISEGLALFNI	707
25	713	VPSRLLS	719
26	732	NSLVWHQQVLGW	743
27	760	DTTIPVVDY	768