

Table S3. Chou–Fasman secondary structure prediction for calculating ig-score of *D. circinale*'s akinete-specific protein.

No.	Secondary structure prediction	
1	SS	0044444444444411111144441111222222221222220444444
	CP	72899999999986454326652134478899998863567899999999
51	SS	2444444111222244422222224442222211112204444112224
	CP	998604599899999999999999997326898621010389999999851
101	SS	222222111122011222221121244211111111211111140222
	CP	66641025316776407855899999998468998888899989999988
151	SS	22222204114442222011444022244411111444221422221114
	CP	75379999999999998578887599999999999999997601666666
201	SS	4422222222222211222442442442222222224442222212222
	CP	76788999999999999999998899999998587888899999998557899
251	SS	2242444444244422211112222111122224444111141111222
	CP	9999999999999999864277089999999999999999642111268999
301	SS	111112222114222222222211111222222211441112222224
	CP	75187047899998668889987899313126887665103420147899
351	SS	00444411111112444444222222222222221111111414444222
	CP	99999998647999899999999999999999999982199188987789999
401	SS	22422111444411112222221141122444444444444411444421
	CP	9999999999999999999873858899999999999965799988999999
451	SS	11114140144111122221111214444444444422211122224411
	CP	999999998615997511104552589999999975305788786888864
501	SS	122444111112221111444422222244422244411112222222
	CP	17714678888735777778998888322676466641453246789999
551	SS	4222222211122222244222244442411222222224222111444
	CP	9999999778997899999999999999999998236613445433368999

SS indicates secondary structure prediction (0: lack of secondary structure, 1: turn, 2: helix, 4: sheet), whereas CP indicates the confidence of the prediction [low (0) to high (9)].