Description	Dhp1-FTP Purification <u>Abundance</u> ** (% of bait)	
	Abundance**	Abundance*
5'-3' exoribonuclease	(% of bait)	
5'-3' exoribonuclease		(% of bait)
0 0 00000000000000000000000000000000000	100	34.75
Dhp1p-interacting protein	50.47	100
CwfJ family protein, splicing factor (predicted)	36.55	6.25
Franslation initiation RNA helicase (Moc2)	12.62	4.1
Franscription elongation factor	8.64	5.63
Histone H2B	5.77	0.70
RNA polymerase II complex subunit	5.65	6.95
RNA polymerase II large subunit	5.43	11.17
nistone H2A alpha	4.57	0.062
DNA-directed RNA polymerase complex II subunit	4.51	9.76
DNA-directed RNA polymerase I, II and III subunit	4.14	3.95
nRNA export factor	3.78	0.55
DNA-directed RNA polymerase II complex subunit	2.46	2.11
RNA binding protein Mlo3	1.23	0.05
TH family RNA binding protein	0.50	0.014
Exosome 3'-5' exoribonuclease subunit	0.042	0
Silencing Protein – CIrC subunit	0.017	0
	<ul> <li>wfJ family protein, splicing factor (predicted)</li> <li>ranslation initiation RNA helicase (Moc2)</li> <li>ranscription elongation factor</li> <li>listone H2B</li> <li>RA polymerase II complex subunit</li> <li>RA polymerase II large subunit</li> <li>istone H2A alpha</li> <li>NA-directed RNA polymerase complex II subunit</li> <li>NA-directed RNA polymerase I, II and III subunit</li> <li>nRNA export factor</li> <li>NA-directed RNA polymerase II complex subunit</li> <li>RNA binding protein Mlo3</li> <li>TH family RNA binding protein</li> <li>ixosome 3'-5' exoribonuclease subunit</li> </ul>	bhp1p-interacting protein50.47cwfJ family protein, splicing factor (predicted)36.55translation initiation RNA helicase (Moc2)12.62transcription elongation factor8.64listone H2B5.77tNA polymerase II complex subunit5.65tNA polymerase II large subunit5.43istone H2A alpha4.57tNA-directed RNA polymerase I, II and III subunit4.14nRNA export factor3.78tNA-directed RNA polymerase II complex subunit2.46tNA-directed RNA polymerase I, II and III subunit1.23that have a subunit protein Mlo31.23that family RNA binding protein0.50translation grotein State0.042

S3 Table: Dhp1-FTP- and Din1-FTP-interacting proteins identified by LC-MS/MS\*

\*All proteins listed were not detected in the purification of the untagged control strain \*\*Abundance values for each identified protein are normalized to the bait values