

1 **Table S1. Yeast deletion strains screened for effects on Acm1 stabilization.**

Non-essential E3s		Non-essential E2s	Essential E3s
<i>ASI3</i>	<i>PEX12</i>	<b><i>UBC2</i></b>	<i>RSP5</i>
<i>ASR1</i>	<b><i>PIB1</i></b>	<b><i>UBC4</i></b>	<i>PRP19</i>
<b><i>BRE1</i></b>	<i>PSH1</i>	<b><i>UBC5</i></b>	<i>SCF*</i>
<b><i>CUL3</i></b>	<i>RAD5</i>	<i>UBC6</i>	<i>APC*</i>
<b><i>CUL8</i></b>	<i>RAD16</i>	<b><i>UBC7</i></b>	
<b><i>DMA1</i></b>	<i>RAD18</i>	<b><i>UBC8</i></b>	
<b><i>DMA2</i></b>	<i>RCO1</i>	<b><i>UBC10</i></b>	
<i>ETP1</i>	<b><i>RKR1</i></b>	<b><i>UBC11</i></b>	
<i>FAP1</i>	<i>RTC1</i>	<b><i>UBC12</i></b>	
<i>FAR1</i>	<i>SAN1</i>	<b><i>UBC13</i></b>	
<i>HEL1</i>	<b><i>SLX5</i></b>		
<i>HEL2</i>	<b><i>SLX8</i></b>		
<b><i>HRD1</i></b>	<i>SNT2</i>		
<b><i>HUL4</i></b>	<b><i>SSM4</i></b>		
<b><i>HUL5</i></b>	<b><i>TOM1</i></b>		
<i>IRC20</i>	<i>TUL1</i>		
<i>ITT1</i>	<b><i>UBR1</i></b>		
<i>MAG2</i>	<b><i>UBR2</i></b>		
<b><i>MOT2(NOT4)</i></b>	<b><i>UFD2</i></b>		
<i>NFI1</i>	<b><i>UFD4</i></b>		
<i>PEP3</i>	<i>ULS1</i>		
<i>PEP5</i>	<i>VPS8</i>		
<i>PEX2</i>	<i>YBR062C</i>		
<i>PEX10</i>			

2 For all non-essential genes, deletion strains were obtained from the Open Biosystems gene  
 3 deletion library and crossed to YKA404 to generate combined deletions with  
 4 *acm1Δ::KanMX4*.

5 For essential genes RSP5 and PRP19, tetracycline-repressible strains were obtained from the  
 6 Open Biosystems Tet-promoter collection.

7 \*Effects of SCF on Acm1 stability were tested using conditional *cdc34* and *cdc53* mutant  
 8 strains (not shown and [32]). No effect was found. Lack of dependence on APC can  
 9 be found in Figure 1 and [31].

10 Strains in bold were tested in both Figure S1 and Figure S3 screens. Non-essential E3 and  
 11 E2 strains not in bold were only tested in the Figure S3 screen.

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