## Figure S3

A. The expression of 10 genes in both subconfluent and confluent toxin-treated cells was measured by qRT-PCR. Fold changes shown are relative to untreated samples.

B. Densitometry was performed on immunoblots of lysates from toxin-treated, confluent cells. The intensity of each band was normalized to the intensity of GAPDH, the loading control. The values above each row indicate the amount of protein relative to the amount in untreated cells. The relative intensity for p57 was not calculated because the protein was not detectable in untreated cells. The blots show the presence of p57 after toxin treatment.

C. A Rac1 antibody (BD#610650) that recognizes non-glucosylated Rac1 (protein that has not been glucosylated by either toxin) shows the activity of TcdA and TcdB in HCT-8 cells.

glucosylated by either toxin) shows the activity of IcdA and IcdB in HCI-8 cells.																							
A A. 6 hr			D.	6 hr	A, 24 hr		ъ.	24 hr		В													
Confluence			High		High	Low			ng/ml		1	1.2	1.0	0.9	1.0	0.8	0.7	1	<0.1	0.6	<0.1	<0.1	
CCND1	1.1	1.6	-2.9	-4.2	-1.5	-2.1	-1.4	-1.4	10	Cyclin A2 GAPDH	_	_	_	_	_	_		-	-				
	1.0	1.8	-1.1	-1.9	1.0	-1.3	-1.4	-1.8	1		1	0.3	0.7	1.1	0.3	0.3	0.5	1	0.1	0.7	0.2	0.4	
	1.2	1.5	1.0	1.1	1.1	-1.3	1.0	-1.9	0.1	Cyclin D1		0.3	0.7		0.5	0.5	0.5	•	0.1	0.7	0.5	0.4	
CCNE2	-1.1	3.9	-1.9	1.4	-1.9	-3.7	-6.8	-3.1	10	GAPDH	-	-	-	-	-	-	-	-	-	-	Mariant	against the	
	1.0	4.0	-1.1	2.4	1.0	-1.3	-2.1	-3.6	1	Cyclin E2	1	0.6	0.7	1.0	0.5	0.5	0.5	1	<0.1	0.3	<0.1	<0.1	
	1.0	3.8	-1.1	2.8	1.1	-1.2	-1.1	-2.2	0.1					=					_				
CCNA2	1.0	4.3	-1.1	4.3	-1.4	-3.3	-11.7	-8.9	10		1	1.3	1.0	1.0	2.2	1.7	1.8	1	2.2	1.3	2.1	2.5	
	1.0	4.0	-1.1	4.3	1.0	-1.1	-1.5	-13.4	1	p27	****	-	-	-	-	-	-	-		-			
	1.1	3.6	1.0	4.2	-1.1	1.2	1.0	-1.4	0.1	GAPDH	-	-	-	-	-	-	-	-	-	-	-	**********	
CDKN1C	-0.5	5.1	3.2	51.6	2.3	3.2	3.8	1.9	10	p57	1	52	2.4	2.3	125	97	63	1	2.9	<0.1	0.9	3.3	
	1.0	3.8	-1.1	18.1	1.2	1.3	2.5	4.1	1	GAPDH	*****	-	-	-	-	-	-	-	-	-	-	-	
	-1.4	3.5	-1.1	4.7	1.3	1.0	1.0	2.4	0.1	Toxin	-	Α	Α	Α	В	В	В	-	Α	Α	В	В	
CDKN1B CDC25A	1.1	4.5	1.8	4.9	1.1	1.8	2.0	2.8	10	ng/ml Time	-	1000	100	10 6 hr		100	10	-	1000	100   24 h		10	
	1.0	4.1	1.3	4.6	1.0	1.3	1.2	2.6	1	Time			7					1		24 N	r		
	1.1	4.3	1.1	3.1	-1.1	-1.1	-1.2	1.4	0.1	C		untrea	ite <sup>O</sup> in	nin					<b>1</b> 01				
	1.2	1.7	-1.7	-1.2	-2.0	-3.6	-6.1	-4.9	10	non-		NULL 1	50,30	1,14	, 5 W	V/1, P,	ν, <sup>8</sup> ν,	, 15,	Σ <sub>V.</sub> qι	uratio	n of		
	1.2	1.7	1.2	1.3	1.0	-1.1	-2.2	-7.9	1	glucosyla	ted		-	-	-	-	Toxin A exposure						
	1.3	1.7	1.2	2.4	1.0	1.1 -2.3	-1.1	-1.5	0.1	Rac1		- Allerin		4 - 4	210			1					
DUSP6	-1.2	2.9	-6.7	-4.6	1.1		-2.3	-1.2	10			in.	n mir		-< · v	×. ×		m.	m.				
	1.0	4.0	-1.4	-1.5 3.3	1.0	1.1 -1.1	-1.4	-1.76	1	non-		15 min nin 2m 4m 6m 8m 2m 2m duration of Toxin B exposure											
JUN	1.1	4.1 6.1	-1.1 4.3		<b>-1.1</b>	14.3	1.0	-1.4 37.1	0.1	glucosyla	ted	-								oxin E	s exp	osure	
	-1.1	4.2	1.8	11.0 9.6	2.0	2.9	11.8	34.4	10 1	Rac1					549	197							
	-1.1	3.5	-1.2	3.7	1.2	-1.1	2.4	7.9	0.1														
	2.1	12.3	8.1	34.2	6.3	10.2	15.9	19.1	10														
RHOB	1.2	5.4	3.3	29.3	1.7	3.1	8.1	16.5	1														
	1.4	J. <del>T</del>	٥.5	27.5	1.7	ا . ا	0.1	10.5	1														

6.0

-1.34 1.9 -1.0 -1.1 -1.1

1.0 | -1.1

-3.5

1.0 4.1 1.5

-1.6 2.7

1.1 | 2.0 | -1.34 | -4.0 | 1.0 | -1.3

-1.3

CTGF

2.1

-5.4

-22.7 -38.1

6.0 0.1

-20.1

-30.9

10