

**Table 12. Cells lacking CDC4 exhibit increases in primary constriction gaps**

Treatment	Number*	Cohesed <sup>†</sup>	Primary constriction gap (PCG)				
			PCG <sub>I</sub> <sup>†</sup>	PCG <sub>II</sub> <sup>†</sup>	PCG <sub>III</sub> <sup>†</sup>	Total <sup>‡‡</sup>	FI <sub>Total</sub> <sup>§</sup>
HCT116 control	215	93.95	5.58	0.47	0.00	6.05	NA
CDC4 <sup>+/-</sup>	253	91.70	7.90	0.40	0.00	8.30	1.37×
CDC4 <sup>-/-</sup>	298	84.56	13.76	1.01	0.67	15.44	2.55×

\*Number of mitotic spreads included in the analyses.

<sup>†</sup>Values represent percentages.

<sup>‡‡</sup>Total represents the sum of PCG<sub>I</sub> + PCG<sub>II</sub> + PCG<sub>III</sub>.

<sup>§</sup>FI<sub>Total</sub> represents the relative fold increase of the total PCG for the indicated condition over that of the HCT116 parental control cells.