

NANOG/NANOGP8 Localizes at the Centrosome and Is Spatiotemporally Associated with Centriole Maturation

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Supplementary Tables

Table S1. Validation of anti-NANOG #ab109250 antibody specificity by mass spectrometry of immunoprecipitated NTERA-2 cells.

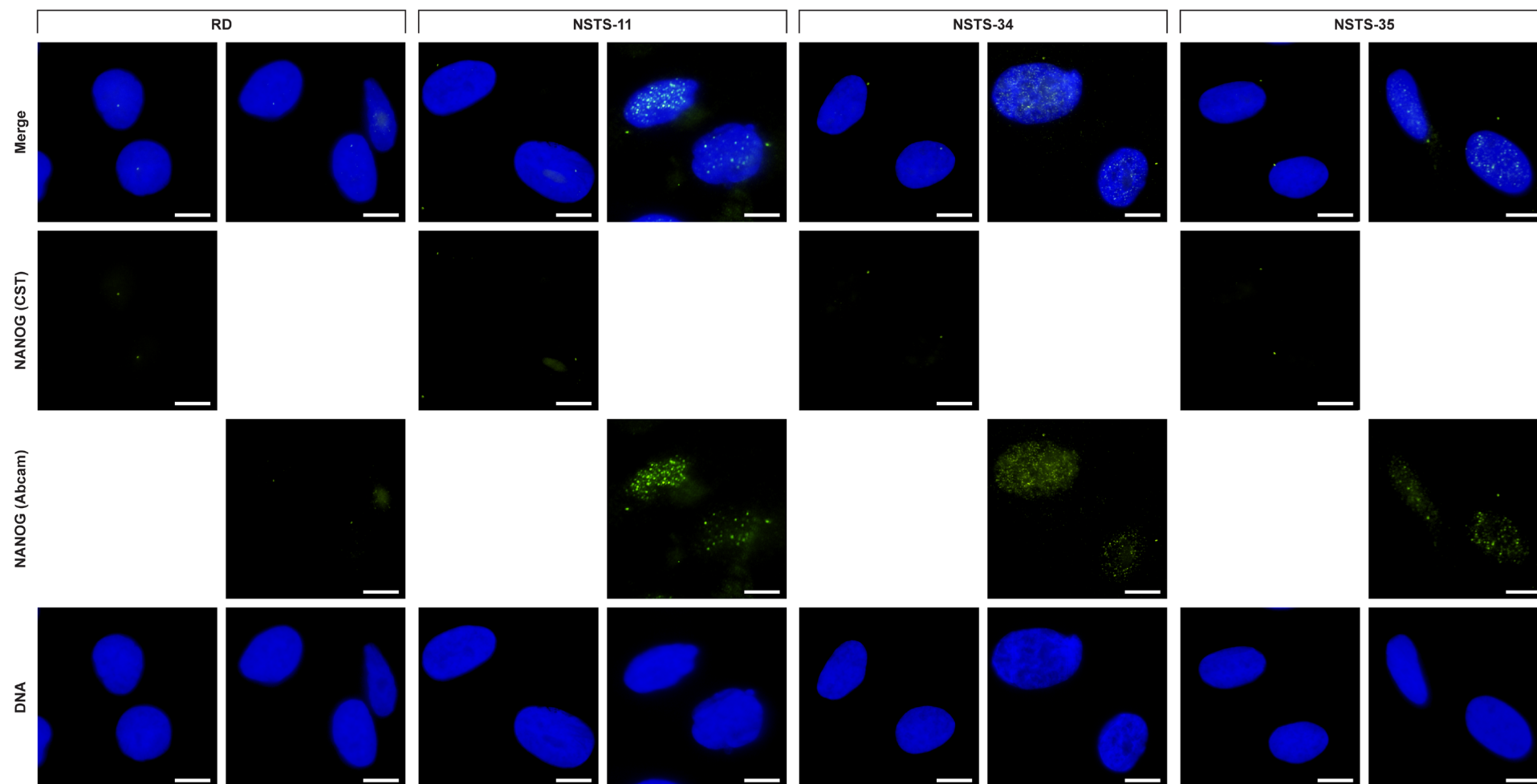


Figure S1. Unmerged images from Figure 1b. Localization of NANOG (green) was detected using two anti-NANOG antibodies (#4903, Cell Signaling Technology [CST] and #ab109250, Abcam). The nuclei were counterstained with Hoechst 33342 (blue). Scale bars, 10 μ m.

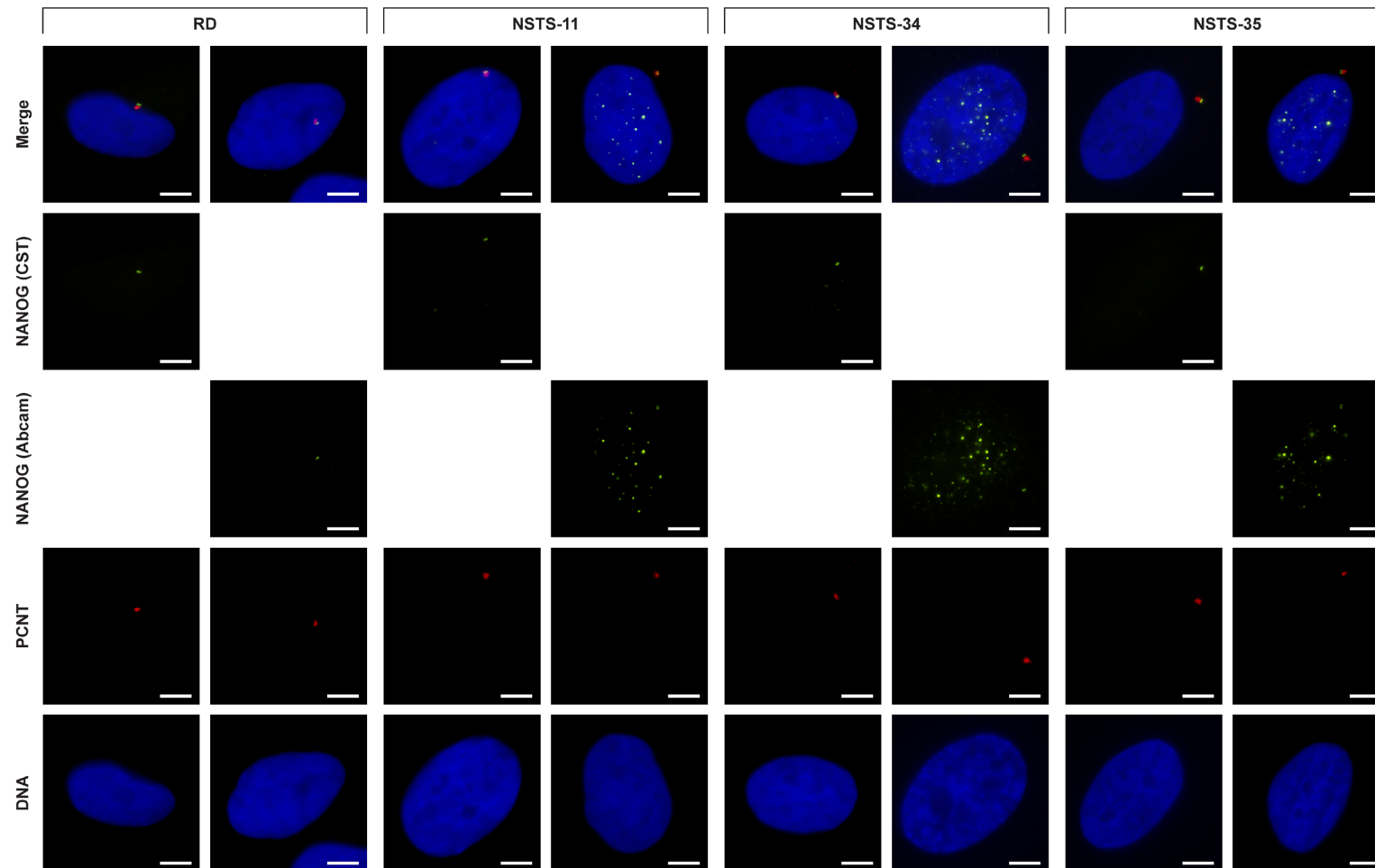


Figure S2. Unmerged images from Figure 3a. Localization of NANOG (green) was detected using two anti-NANOG antibodies (#4903, Cell Signaling Technology [CST] and #ab109250, Abcam) and anti-pericentrin antibody (PCNT; red). The nuclei were counterstained with Hoechst 33342 (blue). Scale bars, 5 μ m.

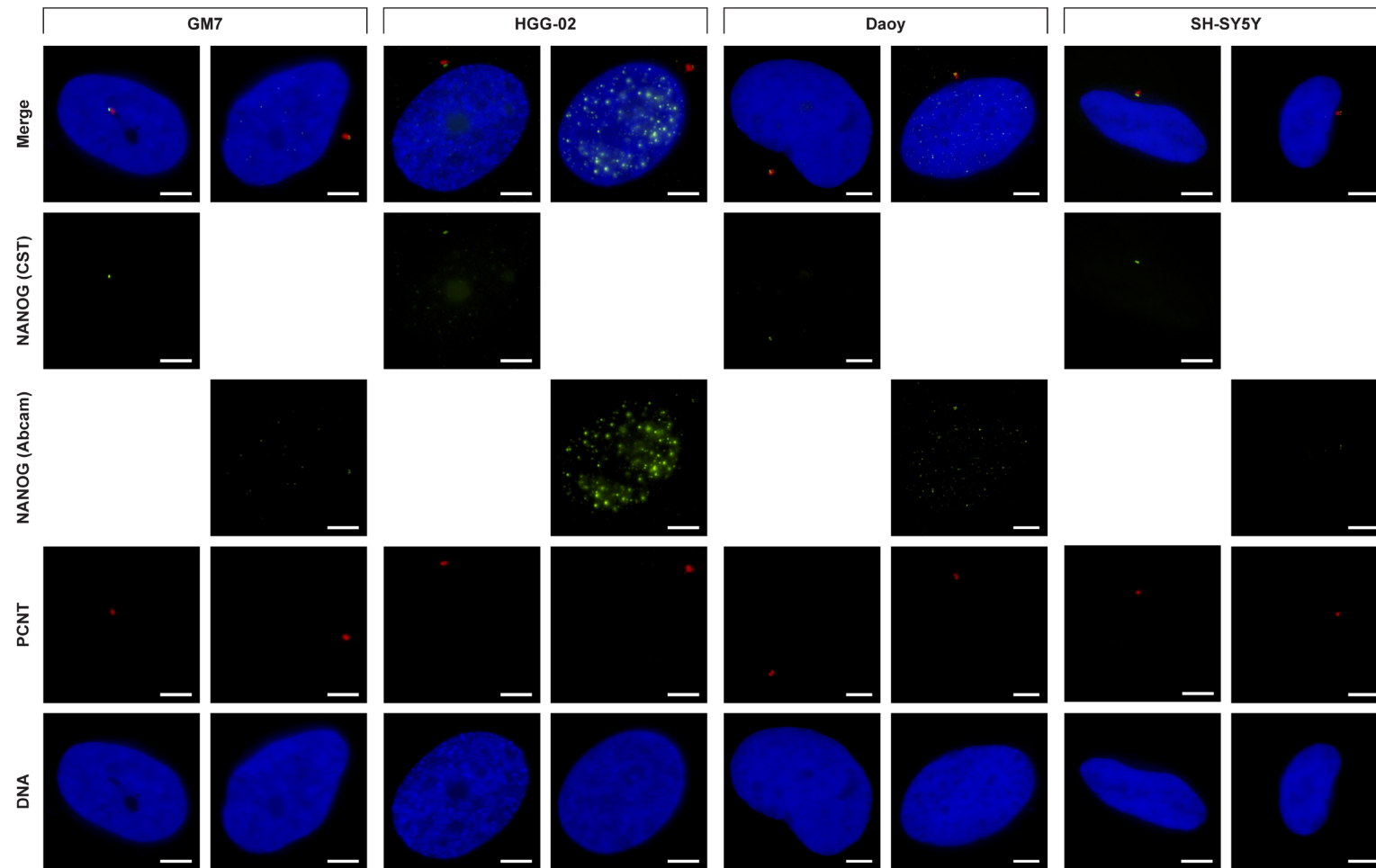


Figure S3. Unmerged images from Figure 3b. Localization of NANOG (green) was detected using two anti-NANOG antibodies (#4903, Cell Signaling Technology [CST] and #ab109250, Abcam) and anti-pericentrin antibody (PCNT; red). The nuclei were counterstained with Hoechst 33342 (blue). Scale bars, 5 μ m.

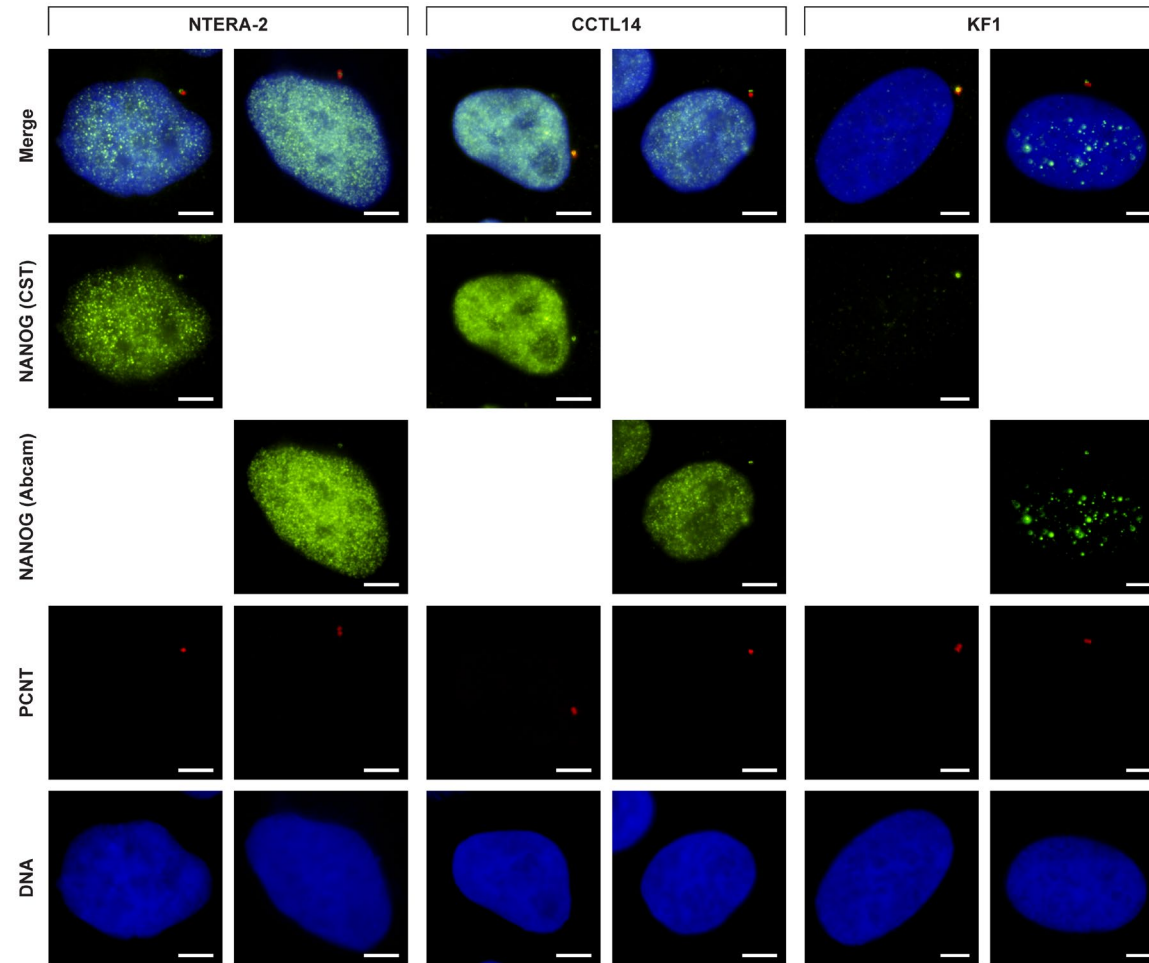


Figure S4. Unmerged images from Figure 3c and Figure 4. Localization of NANOG (green) was detected using two anti-NANOG antibodies (#4903, Cell Signaling Technology [CST] and #ab109250, Abcam) and anti-pericentrin antibody (PCNT; red). The nuclei were counterstained with Hoechst 33342 (blue). Scale bars, 5 μ m.

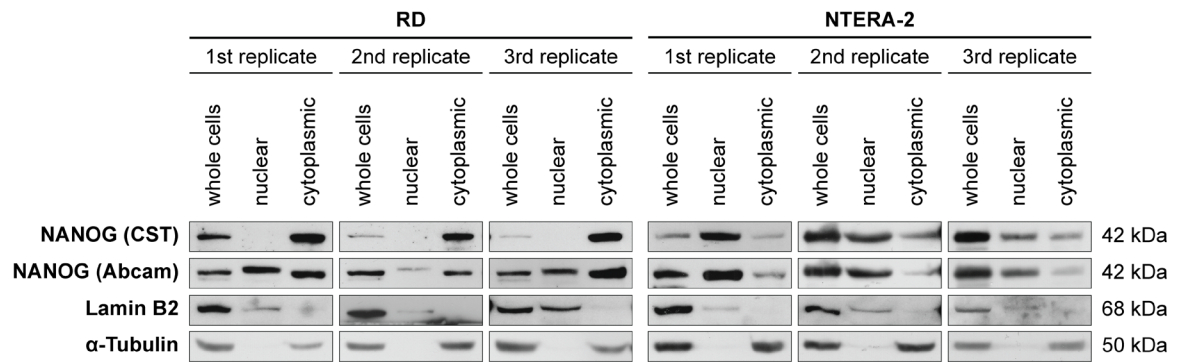


Figure S5. Western blot images of NANOG immunodetection in three biological replicates of the nuclear and cytoplasmic fractions of RD and NTERA-2 cells. Nuclear/cytoplasmic fractionation followed by immunoblotting was performed using two independent anti-NANOG commercial antibodies (#4903, Cell Signaling Technology [CST] and #ab109250, Abcam). Lamin B2 and α -tubulin served as controls of the nuclear fraction and cytoplasmic fraction purity, respectively.

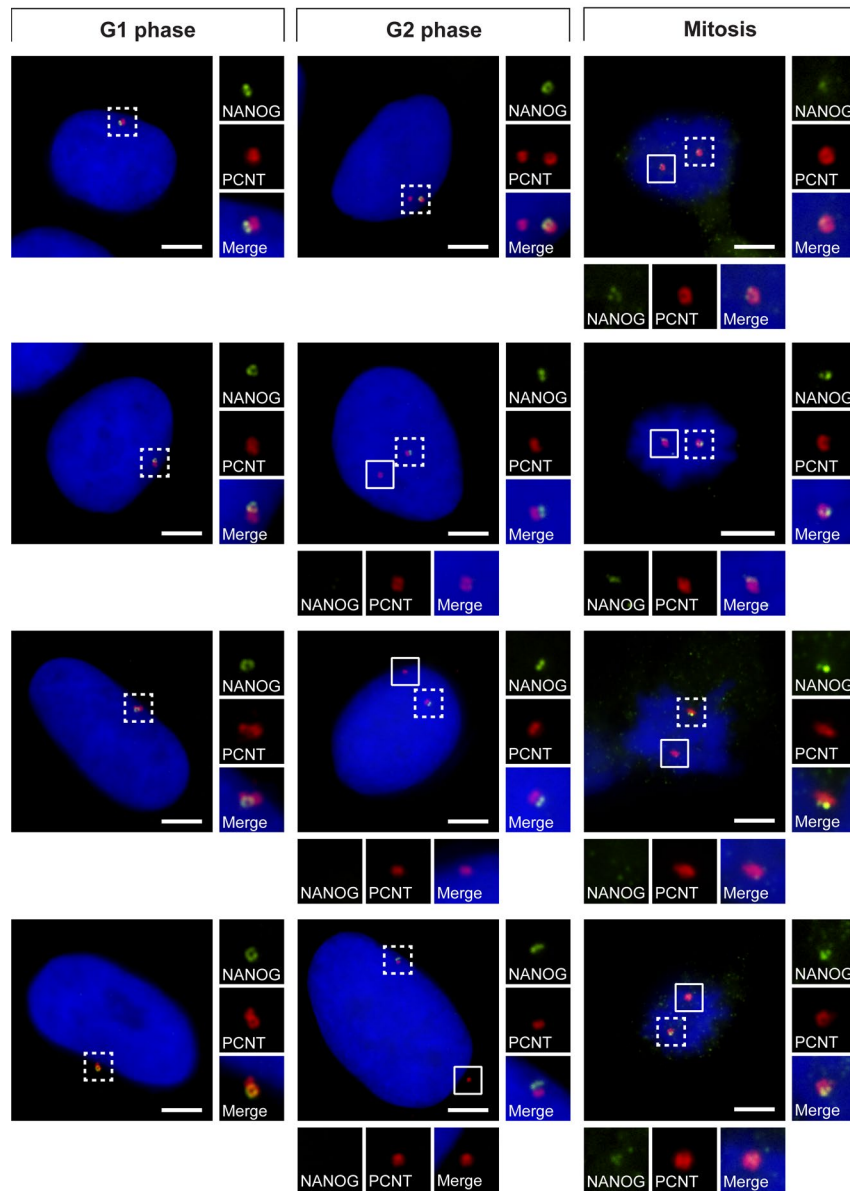


Figure S6. NANOG localization during the cell cycle. NANOG (green; #4903, Cell Signaling Technology [CST]) colocalized with one centrosome (pericentrin, PCNT; red) only during G1 and G2 phases but with both centrosomes (pericentrin, PCNT; red) during mitosis. The nuclei were counterstained with Hoechst 33342 (blue). For each image, regions of interest are indicated by the dashed and solid boxes, and the close-ups are provided on the right and below, respectively. Scale bars, 5 μm .

Table S1. Validation of anti-NANOG #ab109250 antibody specificity by mass spectrometry of immunoprecipitated NTERA-2 cells.

Accession	Gene	Gene symbol	Peptides Count	Spectra Count
P35580	Myosin-10	<i>MYH10</i>	45	144
Q13813	Spectrin alpha chain, non-erythrocytic 1	<i>SPTAN1</i>	31	70
Q01082	Spectrin beta chain, non-erythrocytic 1	<i>SPTBN1</i>	23	51
P35579	Myosin-9	<i>MYH9</i>	14	42
Q16643	Drebrin	<i>DBN1</i>	13	38
P11142	Heat shock cognate 71 kDa protein	<i>HSPA8</i>	11	41
Q8TEW0	Partitioning defective 3 homolog	<i>PARD3</i>	10	23
P05787	Keratin, type II cytoskeletal 8	<i>KRT8</i>	9	30
P10809	60 kDa heat shock protein, mitochondrial	<i>HSPD1</i>	8	20
P14866	Heterogeneous nuclear ribonucleoprotein L	<i>HNRNPL</i>	8	43
P25705	ATP synthase subunit alpha, mitochondrial	<i>ATP5F1A</i>	8	33
Q9UQ35	Serine/arginine repetitive matrix protein 2	<i>SRRM2</i>	8	12
O15164	Transcription intermediary factor 1-alpha	<i>TRIM24</i>	7	30
P05783	Keratin, type I cytoskeletal 18	<i>KRT18</i>	7	26
O00264	Membrane-associated progesterone receptor component 1	<i>PGRMC1</i>	5	13
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	<i>RPN1</i>	5	11
P11021	Endoplasmic reticulum chaperone BiP	<i>HSPA5</i>	5	7
P60660	Myosin light polypeptide 6	<i>MYL6</i>	5	18
P78527	DNA-dependent protein kinase catalytic subunit	<i>PRKDC</i>	5	8
Q00839	Heterogeneous nuclear ribonucleoprotein U	<i>HNRNPU</i>	5	8
Q07157	Tight junction protein ZO-1	<i>TJP1</i>	5	9
Q92734	Protein TFG	<i>TFG</i>	5	22
P01870	Ig gamma chain C region		4	23
P04406	Glyceraldehyde-3-phosphate dehydrogenase	<i>GAPDH</i>	4	16
P08670	Vimentin	<i>VIM</i>	4	8
P09493	Tropomyosin alpha-1 chain	<i>TPM1</i>	4	28
P35221	Catenin alpha-1	<i>CTNNA1</i>	4	5
P43243	Matrin-3	<i>MATR3</i>	4	5
P61978	Heterogeneous nuclear ribonucleoprotein K	<i>HNRNPK</i>	4	8
P62805	Histone H4	<i>HIST1H4A</i>	4	32
Q9H444	Charged multivesicular body protein 4b	<i>CHMP4B</i>	4	14

Accession	Gene	Gene symbol	Peptides Count	Spectra Count
Q9NYL9	Tropomodulin-3	<i>TMOD3</i>	4	5
P05023	Sodium/potassium-transporting ATPase subunit alpha-1	<i>ATP1A1</i>	3	5
P23396	40S ribosomal protein S3	<i>RPS3</i>	3	9
P35613	Basigin	<i>BSG</i>	3	12
P46779	60S ribosomal protein L28	<i>RPL28</i>	3	5
P49411	Elongation factor Tu, mitochondrial	<i>TUFM</i>	3	7
P52272	Heterogeneous nuclear ribonucleoprotein M	<i>HNRNPM</i>	3	7
P56747	Claudin-6	<i>CLDN6</i>	3	10
Q07020	60S ribosomal protein L18	<i>RPL18</i>	3	9
Q07955	Serine/arginine-rich splicing factor 1	<i>SRSF1</i>	3	5
Q15393	Splicing factor 3B subunit 3	<i>SF3B3</i>	3	4
Q16629	Serine/arginine-rich splicing factor 7	<i>SRSF7</i>	3	13
Q2KHR3	Glutamine and serine-rich protein 1	<i>QSER1</i>	3	6
Q5PRF9	Protein Smaug homolog 2	<i>SAMD4B</i>	3	4
Q6NSW7	Homeobox protein NANOGP8	NANOGP8	3	6
	GKQPTSAENSVAK (NANOGP8)			
	QPTSAENSVAK (NANOGP8)			
	TVFSSTQLCVLNDR (NANOG, NANOGP1 and NANOGP8)			
O75569	Interferon-inducible double-stranded RNA-dependent protein kinase activator A	<i>PRKRA</i>	2	3
P04792	Heat shock protein beta-1	<i>HSPB1</i>	2	8
P06753	Tropomyosin alpha-3 chain	<i>TPM3</i>	2	11
P09496	Clathrin light chain A	<i>CLTA</i>	2	4
P17302	Gap junction alpha-1 protein	<i>GJA1</i>	2	8
P35232	Prohibitin	<i>PHB</i>	2	3
P36578	60S ribosomal protein L4	<i>RPL4</i>	2	5
P46776	60S ribosomal protein L27a	<i>RPL27A</i>	2	7
P46782	40S ribosomal protein S5	<i>RPS5</i>	2	5
P63244	Receptor of activated protein C kinase 1	<i>RACK1</i>	2	3
P78310	Coxsackievirus and adenovirus receptor	<i>CXADR</i>	2	8
Q00796	Sorbitol dehydrogenase	<i>SORD</i>	2	8
Q14257	Reticulocalbin-2	<i>RCN2</i>	2	5
Q15392	Delta(24)-sterol reductase	<i>DHCR24</i>	2	12
Q9BWH2	FUN14 domain-containing protein 2	<i>FUNDC2</i>	2	6