We are IntechOpen, the world's leading publisher of Open Access books Built by scientists, for scientists

6,900

186,000

200M

Downloads

154
Countries delivered to

Our authors are among the

TOP 1%

most cited scientists

12.2%

Contributors from top 500 universities



WEB OF SCIENCE

Selection of our books indexed in the Book Citation Index in Web of Science™ Core Collection (BKCI)

Interested in publishing with us? Contact book.department@intechopen.com

Numbers displayed above are based on latest data collected.

For more information visit www.intechopen.com



Immunocytochemistry in Early Mammalian Embryos

Hesam Dehghani

Embryonic and Stem Cell Biology and Biotechnology Research Group, Research Institute of Biotechnology, and Department of Basic Science, Faculty of Veterinary Medicine, Ferdowsi University of Mashhad, Mashhad,

Iran

1. Introduction

The preimplantation period of mammalian development hosts very important cellular and molecular events. This period starts with the fertilization of oocyte by sperm, a process that reprograms the highly differentiated nuclei of these germ cells, and leads to the generation of a totipotent one-cell embryo. Then, the embryo performs cleavage divisions with short cell cycles to quickly increase its cell number. During this period, the genome of the preimplantation embryo manifests profound changes in nuclear and chromatin organization, histone modifications, and transcriptional activity. These genome alterations are also coupled to cell signaling pathways and their regulatory effects. The final product of the preimplantation development is a multi-cellular blastocyst containing three types of cells, epiblasts, hypoblasts, and trophoblast cells [1].

To study and understand the biology of preimplantation embryos, different techniques have been used. The paucity of cells and the difficulties associated with the preparation and production of preimplantation embryos have been the main limiting factors for the application of a wide range of experimental techniques. Thus, what is known about early embryos today is mainly the results of the use of a few experimental techniques and their adapted modifications. These include DNA and RNA amplification techniques, transcript labeling, in situ hybridization of DNA and RNA, gene manipulation studies, and light, electron, and immunofluorescence microscopy techniques.

The application of each technique has revealed a specific aspect of preimplantation developmental biology. Table 1 summarizes and compares the contributions of different experimental techniques applied on preimplantation mammalian embryos. In the rest of this chapter, I will focus on the immunocytochemical staining of embryos and its different applications in preimplantation development.

2. Contribution of immunocytochemistry to understanding the biology of preimplantation mammalian embryos

Application of immunocytochemistry (ICC) on preimplantation embryos has provided invaluable information on different aspects of preimplantation development. I will briefly

Technique	Knowledge contribution	Example references
Conventional and quantitative RT-PCR	Evaluation of the transcription of individual genes	[2-8]
Gene expression profiling (microarray)	Large-scale evaluation of the expression of genes	[9-14]
Electron microscopy techniques	Studying the ultrastructural organization of the embryonic cells	[15-19]
Labeling of nascent transcripts	Quantification of transcriptional activity	[20-24]
In situ hybridization of DNA and RNA	Intracellular localization of chromosomes and transcripts	[25-28]
Gene knockout and knock down techniques	Studying the function of individual genes	[29-38]
Immunocytochemistry	Intracellular localization of proteins Quantitative evaluation of the expression of proteins Identification of protein modifications Evaluation of the activity of certain signaling pathways	[20, 31, 39-45]

Table 1. The major Experimental techniques applied to study the preimplantation embryos.

review the applications of ICC for localization of proteins, for studying the modifications of chromatin and alteration of chromatin organization, and for analyzing cell signaling pathways in preimplantation embryos.

2.1 Cellular and intra-cellular localization of proteins

During preimplantation development, it is very important to identify whether a given protein is expressed, where in the cell it is localized, in which blastomeres it is expressed, and when its expression is eliminated. All of this information relate to the function of protein during preimplantation development. Immunocytochemistry has been an indispensable technique to reveal this information. Application of an alternative Western blotting will not provide any information on the intracellular localization of the protein or the types of expressing cells.

Looking at more than two decades of research on Oct4 clearly shows that what we know on the role of this transcription factor in pluripotency, has all started from this immunocytochemical observation that this protein is differentially expressed in the mouse preimplantation embryonic cells [46]. While it had been previously revealed that it has a strong transcriptional activator effect in the inner cell mass of the preimplantation embryo [47] and it is transcribed in these cells [48], it was its protein localization (using specific antibodies and ICC procedure) that convincingly illustrated its relationship to stemness and pluripotency. A number of later functional studies also used ICC to reveal the function of Oct4 during preimplantation development and pluripotency [49, 50]. The same route of discovery has been traveled for other stemness genes [51].

Using immunocytochemistry and confocal microscopy we have been able to reveal the subcellular distribution and to analyze the relative amount of ten isozymes of PKC (alpha, betaI, betaII, gamma, delta, epsilon, eta, theta, zeta, iota/lambda) and a PKC-anchoring protein, receptor for activated C-kinase 1 (RACK1), between the two-cell and blastocyst stages of mouse preimplantation development [39]. In a functional study, we used the same principle to analyze the relative amount of each PKC isozyme within each blastomere and relate this to the transcriptional activity of the 4-cell mouse embryo [20]. Thus for a given protein in the preimplantation embryo, ICC technique can be applied to study its differential expression between embryonic blastomeres, to identify its intracellular localization within individual blastomeres, and also to semi-quantitate its expression. Recently, using fluorescently-labeled specific antigen binding fragments (Fabs), it has been shown that it is possible to monitor the distribution and global level of endogenous histone modifications in living blastomeres without disturbing cell growth and embryo development [52].

2.2 Identification of histone modifications and the study of nuclear organization

The last two decades has witnessed a considerable number of research efforts using ICC to identify a variety of post-translational modifications on histones and to analyze the expression of chromatin-remodeling factors in preimplantation embryos (Table 2). Immunocytochemical detection and localization of nuclear subdomains (Figure 1), histone modifications, enzymes responsible for these modifications, different histone variants, distinct chromatin remodeling factors, and the status of transcription in preimplantation stages of development (Figure 2), has provided ample evidence and knowledge on the biology of chromatin during preimplantation development (Table 2).

In a very close subject, ICC procedure has also been applied to investigate the organization of chromatin, the architecture of nucleus, and the formation of sub-nuclear compartments by ultra-structural studies in preimplantation embryos. In fact, the correlative fluorescence and electron microscopy technique has allowed the ultra-structural identification of nuclear entities which are identified and tagged by immunocytochemistry [15, 53, 54] (Figure 3). As it has been shown in the figure, immunocytochemical detection of a chromocenter domain immuno-stained with CREST antibody is indispensable for finding and imaging it under the electron microscope. The same principle has been used to identify a sub-nuclear compartment immunocytochemically, and to study its ultra-structure, e.g. localizing fibrillarin by ICC to identify nucleolus in the nucleus of preimplantation embryos for ultra-structural analysis [19, 55-57].

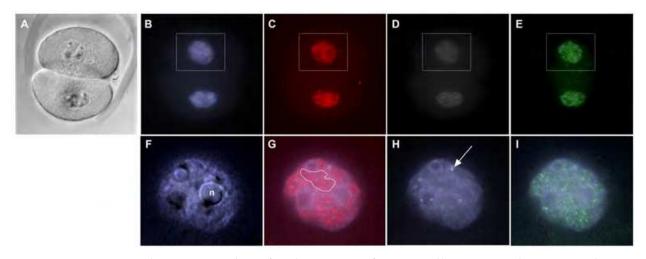


Fig. 1. Immunocytochemistry and confocal imaging of a two-cell mouse embryo to evaluate the function of nucleus. Top row contains confocal images from an optical slice of a two-cell mouse embryo which has been immunolabeled and stained with different antibodies and imaged in different channels. The bottom row contains the merged images of the top nucleus in different channels. A two-cell embryo (A; DIC image) contains two nuclei that are not very chromatin-condensed by DAPI staining (B). Nucleoli (n) in the magnified nucleus in F (the merged image of A and B) show very thin rim of fairly condensed chromatin. Immunolabeling of RNA polymerase II (phosphorylated at serine 5 of its CTD) shows a hyperactive transcription (C). A highly transcribed region of nucleus has been marked in G (the merged image of B and C). Immunolabeling with CREST antibody reveals centromeres (D), which are mainly located at the edge of nucleoli in H (the merged image of B and D). White arrow in H, shows a CREST-labeled spot. Immunolabeling with the antibody against acetylated lysine of H3 histone reveals regions of "open" chromatin (E) which are distributed throughout the nucleus (I, the merged image of B and E).

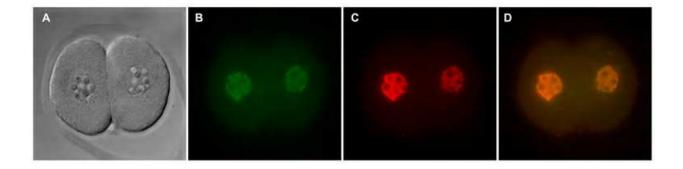


Fig. 2. Immunocytochemical localization of hyperactive transcription domains in a two-cell stage mouse embryo. A) DIC image; B) Immunolabeling with the antibody against the acetylated lysine of histone H3; C) Immunolabeling with the antibody against RNA polymerase II (phosphorylated at serine 5 of its CTD); D) A merged image of B and C. The yellow color in D represents nuclear domains which contain acetylated H3K9 and RNA pol II, indicating that transcription is occurring in chromatin domains with a relaxed state, where a large number of acetylated histone H3K9 moieties are present.

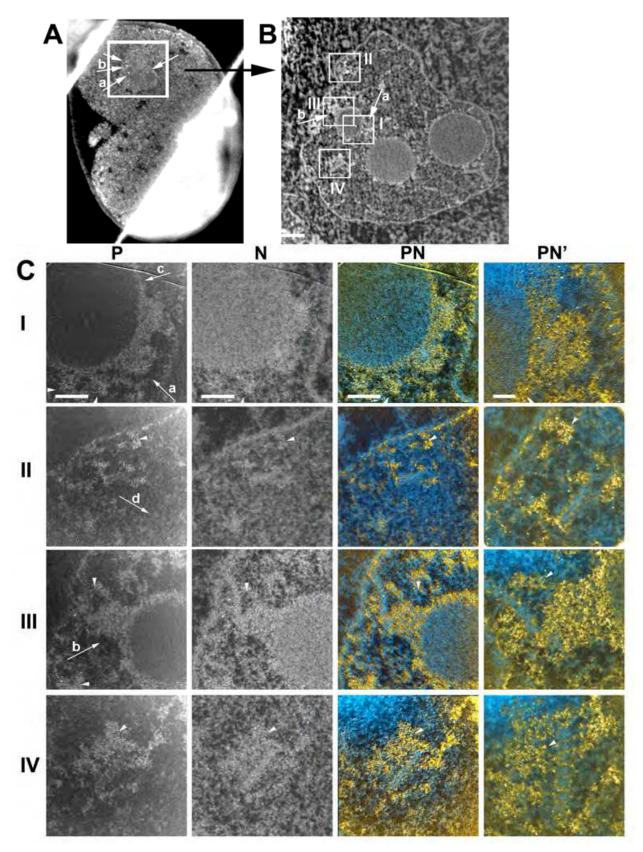


Fig. 3. Chromatin organization in the two-cell stage preimplantation mouse embryo. **A)** Fluorescence image of a physical section spanning through the nucleus (shown by the white

box) of a two-cell stage embryo. White arrows point to the centromeres which are immunostained with CREST antibody. B) The rectangular region in panel A has been imaged by low magnification electron spectroscopic imaging (ESI; 155 keV phosphorusenriched)[15, 53, 54, 58, 59]. Three different-sized nucleoli with very homogenous mass are noticeable in this nucleus. Arrows 'a' and 'b' point to the centromeres designated similarly in panel A. The scale bar is 2 nm. C) Different regions of the nucleus in panel B have been imaged with higher magnification ESI. Columns P, N, PN, and PN' denote images of phosphorus map, nitrogen map, overlay of phosphorus and nitrogen maps, and higher magnification overlays of phosphorus and nitrogen maps, respectively. The segmentation of signals in PN and PN' permits the visualization of chromatin fibers as yellow, while nonchromosomal proteins due to their relatively low N:P ratio content are in blue color. White arrowheads point to the representative gold-tagged histone H3 (methylated at lysine 9) molecules which are accumulated at different areas of the nucleus. Scale bars for columns P, N, and PN are 500 nm, and scale bar for column PN' is 200 nm. (C-I) Nucleoli comprise a homogeneous structure with scarce amounts of ribonucleoprotein (weak signal in P map), but large amounts of protein (strong signal in N map). A more condensed patch of chromatin at the edge of nucleolus (arrow a) is highly positive for K9-methylated H3, while a very thin layer of chromatin at the edge of nucleolus (arrow c) does not show accumulation of this signal. The area shown by 'arrow a' which is designated similarly in panels A and B corresponds to a chromocenter. (C-II) A very thin layer of condensed chromatin (as 30nm fibers) at the nuclear envelope which in some parts is positive for K9methylated H3 blends in with the open lattice of 10nm chromatin fibers (shown by arrow d). The open lattice is filled with large amounts of non-chromosomal proteins shown as blue in PN image. The relation of chromatin and non-chromosomal proteins is better visualised in the higher resolution/magnification image of PN'. (C-III) Patches of condensed chromatin at the edge of nucleolus and in the vicinity of nucleolus (white arrowheads) are positive for K9-methylated H3, but only the area at the edge of nucleolus (shown by arrow b) corresponds to the chromocenter 'b' in panels A and B. (C-IV) Non-centromeric condensed chromatin (as 30 nm fibers and positive for K9-methylated H3) is surrounded by dispersed network of 10nm fibers.

2.3 Evaluation of the activity of certain signaling pathways

Immunocytochemsitry has also been used to discover the presence of many components of signaling pathways including Wnt, hedgehog, receptor tyrosine kinase, and PKC in preimplantation embryos. These studies based on imaging and localization of specific proteins has clearly established a framework for future functional studies. In Table 3 some of these studies have been summarized.

3. Immunocytochemistry of oocytes and preimplantation mammalian embryos

3.1 Harvesting oocytes and preimplantation embryos

Depending to the species, oocytes can be acquired and preimplantation embryos can be produced in different ways. In mouse, it is very easy to harvest from oviduct and uterus, the oocytes and embryos grown in vivo to certain stages of preimplantation development. It is also possible to harvest embryos at early cleavage stages and grow them in culture medium

Findings	Implication	Example references
Lack of the constitutive heterochromatin markers histone H4 trimethyl Lys20 (H4K20me3) and chromobox homolog 5 (HP1α); the presence of heterochromatin markers, H3K9me3, 5-methyl cytosine (5MC), HP1β, H3K27me3, H4K20me1 and H4K20me2	Heterochromatin is in an immature state in mouse preimplantation embryos	[60]
Presence of the acetylated forms of H3K9 and H3K27	H3K27 acetylation is important for normal embryonic development	[52]
Relatively higher expression in oocytes and early cleavage stage embryos of methionine adenosyltransferase 1A protein up to the 8-cell stage compared with the morulae and blastocyst stages	nutrient-sensitive epigenetic regulation and perturbation may be performed through specific enzymes at the earliest stages of preimplantation development	[61]
Embryos at 2-, 4-, and 8-cell stages lack macroH2A except in residual polar bodies. MacroH2A protein expression reappears in embryos after the 8-cell stage and persists in morulae and blastocysts, where nuclear macroH2A is present in both the trophectodermal and inner cell mass cells.	Normal embryos execute three to four mitotic divisions in the absence of macroH2A prior to the onset of embryonic macroH2A expression. Embryos made by somatic nuclear transfer utilize the same chromatin remodeling mechanisms.	[62, 63]
HDAC1 is expressed in preimplantation embryos, where its expression inversely correlates with changes in the acetylation state of histone H4K5 during preimplantation development	HDAC1 is involved in the formation of a chromatin-mediated transcriptionally repressive state that initiates in the late two-cell embryo	[31]
ICC of late zygotes shows that constitutive heterochromatin is only maternally labeled by H3K9me3 and HP1 β	In early embryos, Suv39h-mediated H3K9me3 constitutes the dominant maternal transgenerational signal for pericentric heterochromatin formation	[34]
After fertilization, level of H3K79me2 and H3K79me3 modifications rapidly decrease, and the hypomethylated state is maintained at the interphase (before the blastocyst stage), except for a transient increase in H3K79me2 at mitosis (M phase). H3K79me3 is not detected throughout preimplantation, even at M phase	Elimination of H3K79 methylation after fertilization is involved in genomic reprogramming	[64]
p150CAF-1 is expressed in preimplantation embryos and loss of p150CAF-1 function leads to early developmental arrest and alteration of heterochromatin organization	Chromatin assembly machinery is involved in controlling the spatial organization and epigenetic marking of the genome in early embryos	[42]

Table 2. Immunocytochemical identification and analysis of some histone modifications and chromatin remodeling factors in preimplantation embryos.

Findings	Pathway*	Example references
Expression of protein kinase C isoforms in each stage of preimplantation development	Activation of PKC through G-protein coupled receptors	[39, 65]
Expression of Hh receptor PTCH1 and co-receptor SMO	Signaling events mediated by the Hedgehog family	[66]
Expression of β-catenin	Wnt signaling network	[67-69]
Presence of Aurora C in cleavage- stage embryos	Signaling by Aurora kinases	[70]
Expression of proteins in MAPK pathway	p38 MAPK signaling pathway	[71]
IRS-1 is expressed in all cell lineages of the peri-implantation mouse embryo and mediates some effects of insulin and IGFs at this stage.	Insulin pathway	[72]
Expression and localization of beta 1, beta 5 and alpha 6 integrins and ZO-1 and E-cadherin proteins	E-cadherin signaling pathway & integrin family cell surface interactions	[73, 74]
Strong expression of c-MYC signal in the nucleus of growing and fully grown oocytes as well as in preimplantation embryos before the morulae stage	C-MYC pathway	[75]
The p 85 and p110 subunits of PI3K and Akt are expressed from the 1-cell through the blastocyst stage of murine preimplantation embryo development	The PI3K/Akt pathway	[76]

*Name of pathways have been adapted from NCI-Nature Pathway Interaction Database [77]

Table 3. Components of signaling pathways immunocytochemically identified in preimplantation embryos.

(in vitro culture; IVC). In addition, the early embryos could be produced by in vitro fertilization (IVF) of oocytes, and subsequently cultured in vitro. The most practical method to acquire bovine embryos is through IVF followed by IVC.

Superovulation of female mouse: In mouse, whether we need oocytes, in vivo grown embryos, or in vitro fertilized and cultured embryos, the female mice required to be

superovulated. In response to a hormonal regimen, 3 weeks-old female mice produce the highest number of oocytes (metaphase II stage) and embryos. This is believed to be related to the lack of reproductive cycles and an inactive state of hypothalamic-hypophysial-gonadal axis at this age. The acquired number of harvested oocytes and embryos after superovulation is also largely affected by the strain and maintenance (nutritional and light-dark cycle) conditions.

To induce superovulation of female mice, the following steps need to be taken.

- 1. Mice should be kept in a 12 hour light-dark cycle in a properly ventilated room with a temperature of 22-26°C.
- 2. Administration of hormones is performed by intra-peritoneal injection of female mice at 3-weeks age. If the mice are bred in the same facility, then the first injection time would be two days after weaning from mother. However, if mice will be transferred to the facility from another location, then the first injection time would be after a two-day acclimatization period.
- 3. Human chorionic gonadotropin (hCG) and pregnant mare serum gonadotropin (PMSG), which are in the lyophilized powder form, should be dissolved in sterile saline solution (0.9% NaCl) under a laminar hood. The final concentration is 5 IU per 0.1 ml. Once all the powder in each vial has been dissolved, 0.5 ml of each solution should be drawn into individual insulin syringes and immediately placed in -80°C freezer.
- 4. Each 3 weeks-old female mouse is injected intra-peritoneally at 14pm on day -3 with 0.1 ml of PMSG (5 IU). The syringe containing the hormone should be removed from freezer and brought to the ambient temperature 15 minutes before injection.
- 5. On day -1 at 12pm (46 hours after PMSG injection), each injected mouse will be injected again with 0.1 ml of hCG (5 IU) intra-peritoneally. If harvesting of embryos is intended, each female mouse after injection should be placed in the cage of individual males (Note 1) for overnight mating. However if oocyte recovery is anticipated, the females are returned to their own cage after second injection.

Harvesting oocytes: In the morning of day 0, oocytes can be recovered from oviducts of injected females. Oviducts are flushed with M2 medium (Sigma-Aldrich, St Louis, MO, USA) as previously described [78]. It should be noted that the oocytes at this stage are surrounded by layers of granulosa cells. Thus, to perform ICC and properly localize and image specific proteins in oocytes, the granulosa cells need to be digested away. Otherwise, it will not be possible to properly image oocyte itself, especially when an epi-fluorescence microscope is used for imaging.

Harvesting embryos: To harvest embryos, the injected female is placed in male's cage for overnight mating. Presence of a copulation (vaginal) plug the next morning (on day 0), would be an indication for mating. Embryos at different stages of preimplantation development can be harvested at different time points. Table 4 represents approximate time points for the recovery of embryos at different stages of mouse development.

3.2 Immunocytochemistry

Oocytes or embryos do not attach to the slides or coverslips. Thus, the ICC procedure on harvested oocytes or embryos is somewhat different from the ICC procedure performed on cells grown on coverslips or cells attached to slides. During the procedure, oocytes and

Stage	Day	Time
One-cell stage	0 (The day after hCG injection and mating*)	10-12 am
Two-cell stage (most likely at G2 phase of cell cycle)[79] Four-cell stage (G1 or S phase of the cell cycle)[79]		9 am (45 hours after hcG injection) (33 hours post coitum*) 4 pm (52 hours after hcG injection) (40 hours post coitum)
Eight-sixteen cell stage	2	9 am (69 hours after hcG injection) (57 hours post coitum)
Morulae stage	2	4 pm (76 hours after hcG injection) (64 hours post coitum)
Early blastocyst	3	9 am (93 hours after hcG injection) (81 hours post coitum)

^{*} When male and females are placed in a cage for mating in an evening, the 12:00 midnight is arbitrarily chosen as the time of mating.

Table 4. Approximate time points for the recovery of embryos at different stages of mouse preimplantation development.

embryos should be manually transferred between different media containing fixative, permeabilizing agent, or antibodies. Use of depression slides as container and a stereomicroscope would facilitate the procedure. Pipettors (e.g. 20 µl) or mouth-controlled pipet devices [78] are used for the transfer, while embryos are watched under the stereomicroscope.

The following procedure is a prototype to perform ICC (using fluorescent secondary antibodies) on oocytes and embryos. For simplicity, only embryos (not oocytes) are referred to in the procedure.

1. **Washing:** Wash embryos in 200 µl of PBS twice. This will involve the quick transfer of the harvested embryos into the depression slides containing PBS. Under the stereomicroscope, the embryos could be counted and screened for fragmented or abnormal morphology.

- 2. **Fixation:** Transfer embryos into 200 μl of 4% paraformaldehyde in PBS and incubate at room temperature for 20 minutes. After fixation, the embryos are washed in PBS three times (of 5 minutes each) at room temperature. At this step, the embryos can be stored in PBS at 4°C overnight.
- 3. **Permeabilization** (Note 2): Transfer fixed embryos into 200 µl of 0.5% Triton X 100 in PBS and incubate for 5 minutes at room temperature. Wash the permeabilized embryos in PBS three times (of 5 minutes each) at room temperature.
- 4. **Incubation in primary antibody:** Transfer embryos into 200 μl of primary antibody (diluted in PBS). Incubate in a humid chamber for 2 hours at room temperature or overnight at 4°C. Wash the embryos in PBS three times (of 5 minutes each) at room temperature (Note 3).
- 5. **Incubation in secondary antibody:** Transfer embryos into 200 μl of secondary antibody (diluted in PBS). Incubate in a humid chamber for 1 hour at room temperature or overnight at 4°C. Wash the embryos in PBS three times (of 5 minutes each) at room temperature. If the antibody is conjugated to a fluorescent tag, then the incubation and washing steps should be performed at dark (Note 4).
- 6. **Mounting:** During the mounting procedure, the embryos should be placed in a small volume (20 μl) of mounting medium in the circle on the slide (Figure 4) (Note 5). First, place the mounting or anti-fade medium in the circle. Then, transfer the embryos into the middle of medium. Eyelash probe could be used to move embryos into the middle of circle. Let the embryos sink to the bottom of the medium. Place a coverslip very carefully on the circle on the slide, trying not to move embryos toward the edges of the circle. Seal around the edges of coverslip with nail polish. The mounted embryos can be examined right away or stored at 4°C.

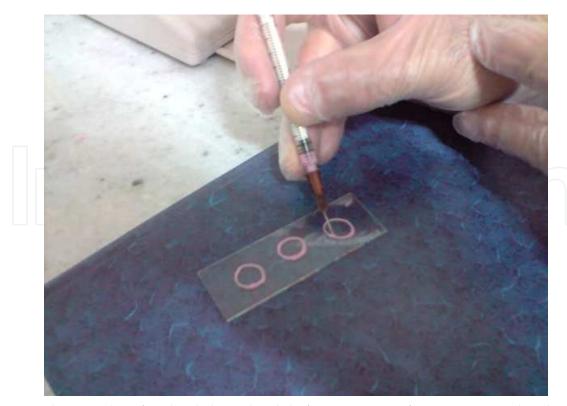


Fig. 4. Making circles of nail polish on the slide for mounting of immunostained embryos.

7. **Microscopic examination:** Depending to the type of secondary antibody used and the available equipment, the embryos can be imaged using light, epi-fluorescence, or confocal laser scanning fluorescence microscopy.

3.3 Notes

Note 1. Male mice reach sexual maturity at the age of 8 weeks. It is important that after weaning the individual male pups to be kept in separate cages. It is believed that keeping several male pups together in one cage, except in the dominant male, may suppress their hormonal maturity. It is also important to place one injected female in the male's cage. Male mouse should not be placed in female's cage. Only one female and not more should be placed in the male's cage. The day after mating males and females should be separated again.

Note 2. Permeabilization is only necessary when an intracellular antigen or protein is to be detected. For immunocytochemical detection of proteins or antigens which are localized on the cellular membrane, a permeabilization step is not performed.

Note 3. Permeabilization and incubation of embryos in primary antibody causes them to sink toward and occasionally adhere the bottom of depression slides. This makes the transfer of embryos between different containers very difficult. Eyelash probe (commercially supplied or homemade by gluing an eyelash to a needle) would be an indispensible device for these situations. With this device under a stereomicroscope, it would be very easy to detach the embryos from the bottom of depression slides and guide them toward the transfer pipette.

Note 4. Different secondary antibodies may be used. If the secondary antibody is conjugated to biotin, alkaline phosphatase, or horseradish peroxidase, different substrates are used to reveal antigen-primary antibody-secondary antibody complexes and different procedures are followed before the mounting step.

Note 5. Placing coverslip directly onto a slide with embryos in between will cause the physical rupture and burst of embryos. Thus, it is very important to produce a space between slide and coverslip. For this purpose, small circles (with a diameter of 5mm) are made on the slide by nail polish. We use an insulin syringe (attached to its needle) filled with nail polish to make the circles with defined edges. When the circle of nail polish is dried, the space in the middle will be used for mounting of embryos.

4. Conclusion

The mammalian preimplantation development contains a highly regulated series of cellular and molecular events that are necessary for normal cell growth, cell division and differentiation. Our understanding of the mechanisms involved in these events has significantly increased in recent years, while much remains to be learned about the mechanisms involved in controlling growth and proliferation, transcriptional control and cell fate decisions. Immunocytochemistry has had and remains to have a significant role for the discovery of these events. In this chapter, its contribution to our current understanding of the different aspects of preimplantation development has succinctly reviewed. In addition, the ICC procedure has been elaborated.

5. Acknowledgments

Work in the author's laboratory is supported by grants from Ferdowsi University of Mashhad. I would like to thank Professor David Bazett-Jones and members of his laboratory, especially Ren Li, Reagan Ching, and Kashif Ahmed for making it possible to begin the ultra-structural analysis of preimplantation embryos, which with no doubt will be a source of important clues on the regulation of transcription and differentiation. I apologize to those colleagues whose publications due to space limitations could not be cited.

6. References

- [1] Rossant, J. Lineage development and polar asymmetries in the peri-implantation mouse blastocyst. *Semin Cell Dev Biol*, 15, 5 (Oct 2004), 573-581.
- [2] Falco, G., Stanghellini, I. and Ko, M. S. Use of Chuk as an internal standard suitable for quantitative RT-PCR in mouse preimplantation embryos. *Reprod Biomed Online*, 13, 3 (Sep 2006), 394-403.
- [3] Fiddler, M., Abdel-Rahman, B., Rappolee, D. A. and Pergament, E. Expression of SRY transcripts in preimplantation human embryos. *Am.J.Med.Genet.*, 55, 1 1995), 80-84.
- [4] Fiorenza, M. T. and Mangia, F. Quantitative RT-PCR amplification of RNA in single mouse oocytes and preimplantation embryos. *Biotechniques*, 24, 4 (Apr 1998), 618-623.
- [5] Nowak-Imialek, M., Wrenzycki, C., Herrmann, D., Lucas-Hahn, A., Lagutina, I., Lemme, E., Lazzari, G., Galli, C. and Niemann, H. Messenger RNA expression patterns of histone-associated genes in bovine preimplantation embryos derived from different origins. *Mol.Reprod.Dev.*, 75, 5 2008), 731-743.
- [6] May, A., Kirchner, R., Muller, H., Hartmann, P., El Hajj, N., Tresch, A., Zechner, U., Mann, W. and Haaf, T. Multiplex RT-PCR Expression Analysis of Developmentally Important Genes in Individual Mouse Preimplantation Embryos and Blastomeres. *Biol Reprod.* 2009 Jan; 80(1): 194-202.
- [7] Cui, X. S., Shen, X. H. and Kim, N. H. High mobility group box 1 (HMGB1) is implicated in preimplantation embryo development in the mouse. *Mol Reprod Dev*, 75, 8 (Aug 2008), 1290-1299.
- [8] Ebrahimian, M., Mojtahedzadeh, M., Bazett-Jones, D. and Dehghani, H. Transcript isoforms of promyelocytic leukemia in mouse male and female gametes. *Cells Tissues Organs*, 192, 6 2010), 374-381.
- [9] Beyhan, Z., Ross, P. J., Iager, A. E., Kocabas, A. M., Cunniff, K., Rosa, G. J. and Cibelli, J. B. Transcriptional reprogramming of somatic cell nuclei during preimplantation development of cloned bovine embryos. *Dev. Biol.*, 305, 2 2007), 637-649.
- [10] Cheon, Y. P., Li, Q., Xu, X., DeMayo, F. J., Bagchi, I. C. and Bagchi, M. K. A Genomic Approach to Identify Novel Progesterone Receptor Regulated Pathways in the Uterus during Implantation. *Mol.Endocrinol.*, 16, 12 2002), 2853-2871.
- [11] Dobson, A. T., Raja, R., Abeyta, M. J., Taylor, T., Shen, S., Haqq, C. and Pera, R. A. The unique transcriptome through day 3 of human preimplantation development. *Hum Mol Genet*, 13, 14 (Jul 15 2004), 1461-1470.
- [12] Hamatani, T., Daikoku, T., Wang, H., Matsumoto, H., Carter, M. G., Ko, M. S. and Dey, S. K. Global gene expression analysis identifies molecular pathways distinguishing

- blastocyst dormancy and activation. *Proc.Natl.Acad.Sci.U.S.A.*, 101, 28 2004), 10326-10331.
- [13] Hamatani, T., Ko, M., Yamada, M., Kuji, N., Mizusawa, Y., Shoji, M., Hada, T., Asada, H., Maruyama, T. and Yoshimura, Y. Global gene expression profiling of preimplantation embryos. *Hum Cell*, 19, 3 (Aug 2006), 98-117.
- [14] Zeng, F. and Schultz, R. M. RNA transcript profiling during zygotic gene activation in the preimplantation mouse embryo. *Dev. Biol.*, 283, 1 2005), 40-57.
- [15] Ahmed, K., Dehghani, H., Rugg-Gunn, P., Fussner, E., Rossant, J. and Bazett-Jones, D. P. Global chromatin architecture reflects pluripotency and lineage commitment in the early mouse embryo. *PLoS One*, 5, 5 2010), e10531.
- [16] Cremer, T. and Zakhartchenko, V. Nuclear architecture in developmental biology and cell specialisation. *Reprod Fertil Dev*, 23, 1 2011), 94-106.
- [17] Svarcova, O., Strejcek, F., Petrovicova, I., Avery, B., Pedersen, H. G., Lucas-Hahn, A., Niemann, H., Laurincik, J. and Maddox-Hyttel, P. The role of RNA polymerase I transcription and embryonic genome activation in nucleolar development in bovine preimplantation embryos. *Mol Reprod Dev*, 75, 7 (Jul 2008), 1095-1103.
- [18] Kikuchi, K., Ekwall, H., Tienthai, P., Kawai, Y., Noguchi, J., Kaneko, H. and Rodriguez-Martinez, H. Morphological features of lipid droplet transition during porcine oocyte fertilisation and early embryonic development to blastocyst in vivo and in vitro. *Zygote.*, 10, 4 2002), 355-366.
- [19] Laurincik, J., Thomsen, P. D., Hay-Schmidt, A., Avery, B., Greve, T., Ochs, R. L. and Hyttel, P. Nucleolar proteins and nuclear ultrastructure in preimplantation bovine embryos produced in vitro. *Biol.Reprod.*, 62, 4 2000), 1024-1032.
- [20] Dehghani, H., Reith, C. and Hahnel, A. C. Subcellular localization of protein kinase C delta and epsilon affects transcriptional and post-transcriptional processes in four-cell mouse embryos. *Reproduction*, 130, 4 (Oct 2005), 453-465.
- [21] Aoki, F., Worrad, D. M. and Schultz, R. M. Regulation of transcriptional activity during the first and second cell cycles in the preimplantation mouse embryo. *Dev. Biol.*, 1811997), 296-307.
- [22] Aoki, F., Hara, K. T. and Schultz, R. M. Acquisition of transcriptional competence in the 1-cell mouse embryo: requirement for recruitment of maternal mRNAs. *Mol Reprod Dev*, 64, 3 (Mar 2003), 270-274.
- [23] Harrouk, W., Khatabaksh, S., Robaire, B. and Hales, B. F. Paternal exposure to cyclophosphamide dysregulates the gene activation program in rat preimplantation embryos. *Mol.Reprod.Dev.*, 57, 3 2000), 214-223.
- [24] Kanzler, B., Haas-Assenbaum, A., Haas, I., Morawiec, L., Huber, E. and Boehm, T. Morpholino oligonucleotide-triggered knockdown reveals a role for maternal Ecadherin during early mouse development. *Mech Dev*, 120, 12 (Dec 2003), 1423-1432.
- [25] Baart, E. B., van den Berg, I., Martini, E., Eussen, H. J., Fauser, B. C. and Van Opstal, D. FISH analysis of 15 chromosomes in human day 4 and 5 preimplantation embryos: the added value of extended aneuploidy detection. *Prenat Diagn*, 27, 1 (Jan 2007), 55-63.
- [26] Rubio, C., Rodrigo, L., Perez-Cano, I., Mercader, A., Mateu, E., Buendia, P., Remohi, J., Simon, C. and Pellicer, A. FISH screening of aneuploidies in preimplantation embryos to improve IVF outcome. *Reprod.Biomed.Online.*, 11, 4 2005), 497-506.

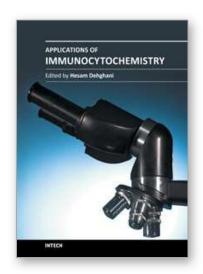
- [27] Baart, E. B., Van Opstal, D., Los, F. J., Fauser, B. C. and Martini, E. Fluorescence in situ hybridization analysis of two blastomeres from day 3 frozen-thawed embryos followed by analysis of the remaining embryo on day 5. *Hum Reprod*, 19, 3 (Mar 2004), 685-693.
- [28] Verlinsky, Y., Cieslak, J., Evsikov, S., Galat, V. and Kuliev, A. Nuclear transfer for full karyotyping and preimplantation diagnosis for translocations.

 **Reprod.Biomed.Online., 5, 3 2002), 300-305.
- [29] Fedoriw, A. M., Stein, P., Svoboda, P., Schultz, R. M. and Bartolomei, M. S. Transgenic RNAi reveals essential function for CTCF in H19 gene imprinting. *Science*, 303, 5655 (Jan 9 2004), 238-240.
- [30] Lykke-Andersen, K., Gilchrist, M. J., Grabarek, J. B., Das, P., Miska, E. and Zernicka-Goetz, M. Maternal Argonaute 2 Is Essential for Early Mouse Development at the Maternal-Zygotic Transition. *Mol Biol Cell* (Aug 13 2008).
- [31] Ma, P. and Schultz, R. M. Histone deacetylase 1 (HDAC1) regulates histone acetylation, development, and gene expression in preimplantation mouse embryos. *Dev Biol*, 319, 1 (Jul 1 2008), 110-120.
- [32] Nganvongpanit, K., Muller, H., Rings, F., Gilles, M., Jennen, D., Holker, M., Tholen, E., Schellander, K. and Tesfaye, D. Targeted suppression of E-cadherin gene expression in bovine preimplantation embryo by RNA interference technology using double-stranded RNA. *Mol Reprod Dev*, 73, 2 (Feb 2006), 153-163.
- [33] Dehghani, H., Narisawa, S., Millan, J. L. and Hahnel, A. C. Effects of disruption of the embryonic alkaline phosphatase gene on preimplantation development of the mouse. *Dev Dyn*, 217, 4 (Apr 2000), 440-448.
- [34] Puschendorf, M., Terranova, R., Boutsma, E., Mao, X., Isono, K., Brykczynska, U., Kolb, C., Otte, A. P., Koseki, H., Orkin, S. H., van Lohuizen, M. and Peters, A. H. PRC1 and Suv39h specify parental asymmetry at constitutive heterochromatin in early mouse embryos. *Nat Genet*, 40, 4 (Apr 2008), 411-420.
- [35] Zou, G. M., Thompson, M. A. and Yoder, M. C. RNAi knockdown of transcription factor Pu.1 in the differentiation of mouse embryonic stem cells. *Methods Mol Biol*, 4072007), 127-136.
- [36] Yagi, R., Kohn, M. J., Karavanova, I., Kaneko, K. J., Vullhorst, D., DePamphilis, M. L. and Buonanno, A. Transcription factor TEAD4 specifies the trophectoderm lineage at the beginning of mammalian development. *Development*, 134, 21 (Nov 2007), 3827-3836.
- [37] Xie, Y., Wang, Y., Sun, T., Wang, F., Trostinskaia, A., Puscheck, E. and Rappolee, D. A. Six post-implantation lethal knockouts of genes for lipophilic MAPK pathway proteins are expressed in preimplantation mouse embryos and trophoblast stem cells. *Mol Reprod Dev*, 71, 1 (May 2005), 1-11.
- [38] Truchet, S., Chebrout, M., Djediat, C., Wietzerbin, J. and Debey, P. Presence of permanently activated signal transducers and activators of transcription in nuclear interchromatin granules of unstimulated mouse oocytes and preimplantation embryos. *Biol.Reprod.*, 71, 4 2004), 1330-1339.
- [39] Dehghani, H. and Hahnel, A. C. Expression profile of protein kinase C isozymes in preimplantation mouse development. *Reproduction.*, 130, 4 2005), 441-451.

- [40] Yamazaki, T., Kobayakawa, S., Yamagata, K., Abe, K. and Baba, T. Molecular dynamics of heterochromatin protein 1beta, HP1beta, during mouse preimplantation development. *J.Reprod.Dev.*, 53, 5 2007), 1035-1041.
- [41] Ohnuma-Ishikawa, K., Morio, T., Yamada, T., Sugawara, Y., Ono, M., Nagasawa, M., Yasuda, A., Morimoto, C., Ohnuma, K., Dang, N. H., Hosoi, H., Verdin, E. and Mizutani, S. Knockdown of XAB2 enhances all-trans retinoic acid-induced cellular differentiation in all-trans retinoic acid-sensitive and -resistant cancer cells. *Cancer Res*, 67, 3 (Feb 1 2007), 1019-1029.
- [42] Houlard, M., Berlivet, S., Probst, A. V., Quivy, J. P., Hery, P., Almouzni, G. and Gerard, M. CAF-1 is essential for heterochromatin organization in pluripotent embryonic cells. *PLoS.Genet.*, 2, 11 2006), e181.
- [43] Dodge, J. E., Kang, Y. K., Beppu, H., Lei, H. and Li, E. Histone H3-K9 methyltransferase ESET is essential for early development. *Mol.Cell Biol.*, 24, 6 2004), 2478-2486.
- [44] Burns, K. H., Viveiros, M. M., Ren, Y., Wang, P., Demayo, F. J., Frail, D. E., Eppig, J. J. and Matzuk, M. M. Roles of NPM2 in chromatin and nucleolar organization in oocytes and embryos. *Science*, 300, 5619 2003), 633-636.
- [45] Sarmento, O. F., Digilio, L. C., Wang, Y., Perlin, J., Herr, J. C., Allis, C. D. and Coonrod, S. A. Dynamic alterations of specific histone modifications during early murine development. *J Cell Sci*, 117, Pt 19 (Sep 1 2004), 4449-4459.
- [46] Palmieri, S. L., Peter, W., Hess, H. and Scholer, H. R. Oct-4 transcription factor is differentially expressed in the mouse embryo during establishment of the first two extraembryonic cell lineages involved in implantation. *Dev Biol*, 166, 1 (Nov 1994), 259-267.
- [47] Scholer, H. R., Balling, R., Hatzopoulos, A. K., Suzuki, N. and Gruss, P. Octamer binding proteins confer transcriptional activity in early mouse embryogenesis. *Embo J*, 8, 9 (Sep 1989), 2551-2557.
- [48] Scholer, H. R. Octamania: the POU factors in murine development. *Trends Genet.*, 7, 10 1991), 323-329.
- [49] Nichols, J., Zevnik, B., Anastassiadis, K., Niwa, H., Klewe-Nebenius, D., Chambers, I., Scholer, H. and Smith, A. Formation of pluripotent stem cells in the mammalian embryo depends on the POU transcription factor Oct4. *Cell*, 95, 3 (Oct 30 1998), 379-391.
- [50] Haraguchi, S., Saga, Y., Naito, K., Inoue, H. and Seto, A. Specific gene silencing in the pre-implantation stage mouse embryo by an siRNA expression vector system. *Mol Reprod Dev*, 68, 1 (May 2004), 17-24.
- [51] Rossant, J. The impact of developmental biology on pluripotent stem cell research: successes and challenges. *Dev Cell*, 21, 1 (Jul 19 2011), 20-23.
- [52] Hayashi-Takanaka, Y., Yamagata, K., Wakayama, T., Stasevich, T. J., Kainuma, T., Tsurimoto, T., Tachibana, M., Shinkai, Y., Kurumizaka, H., Nozaki, N. and Kimura, H. Tracking epigenetic histone modifications in single cells using Fab-based live endogenous modification labeling. *Nucleic Acids Res*, 39, 15 (Aug 1 2011), 6475-6488.
- [53] Bazett-Jones, D. P., Li, R., Fussner, E., Nisman, R. and Dehghani, H. Elucidating chromatin and nuclear domain architecture with electron spectroscopic imaging. *Chromosome Res*, 16, 3 2008), 397-412.

- [54] Fussner, E., Ahmed, K., Dehghani, H., Strauss, M. and Bazett-Jones, D. P. Changes in Chromatin Fiber Density as a Marker for Pluripotency. *Cold Spring Harb Symp Quant Biol* (Dec 7 2010).
- [55] Biggiogera, M., Burki, K., Kaufmann, S. H., Shaper, J. H., Gas, N., Amalric, F. and Fakan, S. Nucleolar distribution of proteins B23 and nucleolin in mouse preimplantation embryos as visualized by immunoelectron microscopy. *Development.*, 110, 4 1990), 1263-1270.
- [56] Baran, V., Mercier, Y., Renard, J. P. and Flechon, J. E. Nucleolar substructures of rabbit cleaving embryos: an immunocytochemical study. *Mol.Reprod.Dev.*, 48, 1 1997), 34-44.
- [57] Hyttel, P., Laurincik, J., Rosenkranz, C., Rath, D., Niemann, H., Ochs, R. L. and Schellander, K. Nucleolar proteins and ultrastructure in preimplantation porcine embryos developed in vivo. *Biol Reprod*, 63, 6 (Dec 2000), 1848-1856.
- [58] Dehghani, H., Dellaire, G. and Bazett-Jones, D. P. Organization of chromatin in the interphase mammalian cell. *Micron*, 36, 2 2005), 95-108.
- [59] Dellaire, G., Nisman, R. and Bazett-Jones, D. P. Correlative light and electron spectroscopic imaging of chromatin in situ. *Methods Enzymol*, 3752004), 456-478.
- [60] Wongtawan, T., Taylor, J. E., Lawson, K. A., Wilmut, I. and Pennings, S. Histone H4K20me3 and HP1alpha are late heterochromatin markers in development, but present in undifferentiated embryonic stem cells. *J Cell Sci*, 124, Pt 11 (Jun 1 2011), 1878-1890.
- [61] Ikeda, S., Namekawa, T., Sugimoto, M. and Kume, S. Expression of methylation pathway enzymes in bovine oocytes and preimplantation embryos. *Journal of experimental zoology. Part A, Ecological genetics and physiology*, 313, 3 (Mar 1 2010), 129-136.
- [62] Chang, C. C., Ma, Y., Jacobs, S., Tian, X. C., Yang, X. and Rasmussen, T. P. A maternal store of macroH2A is removed from pronuclei prior to onset of somatic macroH2A expression in preimplantation embryos. *Dev Biol*, 278, 2 (Feb 15 2005), 367-380.
- [63] Chang, C. C., Gao, S., Sung, L. Y., Corry, G. N., Ma, Y., Nagy, Z. P., Tian, X. C. and Rasmussen, T. P. Rapid elimination of the histone variant MacroH2A from somatic cell heterochromatin after nuclear transfer. *Cellular reprogramming*, 12, 1 (Feb 2010), 43-53.
- [64] Ooga, M., Inoue, A., Kageyama, S., Akiyama, T., Nagata, M. and Aoki, F. Changes in H3K79 methylation during preimplantation development in mice. *Biol Reprod*, 78, 3 (Mar 2008), 413-424.
- [65] Pauken, C. M. and Capco, D. G. The expression and stage-specific localization of protein kinase C isotypes during mouse preimplantation development. *Dev.Biol.*, 223, 2 2000), 411-421.
- [66] Nguyen, N. T., Lo, N. W., Chuang, S. P., Jian, Y. L. and Ju, J. C. Sonic hedgehog supplementation of oocyte and embryo culture media enhances development of IVF porcine embryos. *Reproduction*, 142, 1 (Jul 2011), 87-97.
- [67] Xie, H., Tranguch, S., Jia, X., Zhang, H., Das, S. K., Dey, S. K., Kuo, C. J. and Wang, H. Inactivation of nuclear Wnt-beta-catenin signaling limits blastocyst competency for implantation. *Development*, 135, 4 (Feb 2008), 717-727.
- [68] Kemler, R., Hierholzer, A., Kanzler, B., Kuppig, S., Hansen, K., Taketo, M. M., de Vries, W. N., Knowles, B. B. and Solter, D. Stabilization of beta-catenin in the mouse

- zygote leads to premature epithelial-mesenchymal transition in the epiblast. *Development*, 131, 23 (Dec 2004), 5817-5824.
- [69] Wang, Q. T., Piotrowska, K., Ciemerych, M. A., Milenkovic, L., Scott, M. P., Davis, R. W. and Zernicka-Goetz, M. A genome-wide study of gene activity reveals developmental signaling pathways in the preimplantation mouse embryo. *Dev Cell*, 6, 1 (Jan 2004), 133-144.
- [70] Avo Santos, M., van de Werken, C., de Vries, M., Jahr, H., Vromans, M. J., Laven, J. S., Fauser, B. C., Kops, G. J., Lens, S. M. and Baart, E. B. A role for Aurora C in the chromosomal passenger complex during human preimplantation embryo development. *Hum Reprod*, 26, 7 (Jul 2011), 1868-1881.
- [71] Wang, Y., Wang, F., Sun, T., Trostinskaia, A., Wygle, D., Puscheck, E. and Rappolee, D. A. Entire mitogen activated protein kinase (MAPK) pathway is present in preimplantation mouse embryos. *Dev.Dyn.*, 231, 1 2004), 72-87.
- [72] Puscheck, E. E., Pergament, E., Patel, Y., Dreschler, J. and Rappolee, D. A. Insulin receptor substrate-1 is expressed at high levels in all cells of the peri-implantation mouse embryo. *Mol Reprod Dev*, 49, 4 (Apr 1998), 386-393.
- [73] Bloor, D. J., Metcalfe, A. D., Rutherford, A., Brison, D. R. and Kimber, S. J. Expression of cell adhesion molecules during human preimplantation embryo development. *Mol.Hum.Reprod.*, 8, 3 2002), 237-245.
- [74] Dubey, A. K., Cruz, J. R., Hartog, B. and Gindoff, P. R. Expression of the alphav integrin adhesion molecule during development of preimplantation human embryos. *Fertil.Steril.*, 76, 1 2001), 153-156.
- [75] Suzuki, T., Abe, K., Inoue, A. and Aoki, F. Expression of c-MYC in nuclear speckles during mouse oocyte growth and preimplantation development. *J Reprod Dev*, 55, 5 (Oct 2009), 491-495.
- [76] Riley, J. K., Carayannopoulos, M. O., Wyman, A. H., Chi, M., Ratajczak, C. K. and Moley, K. H. The PI3K/Akt pathway is present and functional in the preimplantation mouse embryo. *Dev Biol*, 284, 2 (Aug 15 2005), 377-386.
- [77] Schaefer, C. F., Anthony, K., Krupa, S., Buchoff, J., Day, M., Hannay, T. and Buetow, K. H. PID: the Pathway Interaction Database. *Nucleic Acids Res*, 37, Database issue (Jan 2009), D674-679.
- [78] Nagy A, Gertsentein M, Vintersten K and R, B. *Manipulating the Mouse Embryo: A Laboratory Manual*. Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY, 2003.
- [79] Pratt, H. P. M. and Monk, M. Isolation, culture and manipulation of preimplantation mouse embryos. IRL Press, City, 1987.



Applications of Immunocytochemistry

Edited by Dr. Hesam Dehghani

ISBN 978-953-51-0229-8 Hard cover, 320 pages Publisher InTech Published online 09, March, 2012 Published in print edition March, 2012

Immunocytochemistry is classically defined as a procedure to detect antigens in cellular contexts using antibodies. However, over the years many aspects of this procedure have evolved within a plethora of experimental setups. There are different ways to prepare a given specimen, different kinds of antibodies to apply, different techniques for imaging, and different methods of analyzing the data. In this book, various ways of performing each individual step of immunocytochemistry in different cellular contexts are exemplified and discussed. Applications of Immunocytochemistry offers technical and background information on different steps of immunocytochemistry and presents the application of this technique and its adaptations in cell lines, neural tissue, pancreatic tissue, sputum cells, sperm cells, preimplantation embryo, arabidopsis, fish gonads, and Leishmania.

How to reference

In order to correctly reference this scholarly work, feel free to copy and paste the following:

Hesam Dehghani (2012). Immunocytochemistry in Early Mammalian Embryos, Applications of Immunocytochemistry, Dr. Hesam Dehghani (Ed.), ISBN: 978-953-51-0229-8, InTech, Available from: http://www.intechopen.com/books/applications-of-immunocytochemistry/immunocytochemistry-in-early-mammalian-embryos



InTech Europe

University Campus STeP Ri Slavka Krautzeka 83/A 51000 Rijeka, Croatia Phone: +385 (51) 770 447

Fax: +385 (51) 686 166 www.intechopen.com

InTech China

Unit 405, Office Block, Hotel Equatorial Shanghai No.65, Yan An Road (West), Shanghai, 200040, China 中国上海市延安西路65号上海国际贵都大饭店办公楼405单元

Phone: +86-21-62489820 Fax: +86-21-62489821 © 2012 The Author(s). Licensee IntechOpen. This is an open access article distributed under the terms of the <u>Creative Commons Attribution 3.0</u> <u>License</u>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.



