Apr 11th, 3:00 PM - 4:00 PM

The Role of Chromatin Modification in Germ Cell Specification and Development

Jenessa Olson
jenessarolson@hotmail.com

Let us know how access to this document benefits you.
Follow this and additional works at: https://scholarworks.umt.edu/umcur

https://scholarworks.umt.edu/umcur/2014/poster_2/7

This Poster is brought to you for free and open access by ScholarWorks at University of Montana. It has been accepted for inclusion in University of Montana Conference on Undergraduate Research (UMCUR) by an authorized administrator of ScholarWorks at University of Montana. For more information, please contact scholarworks@mso.umt.edu.
The Role of Chromatin Modification in Germ Cell Specification and Development

Jenessa Olson, Ekaterina Voronina, Ph.D
Division of Biological Sciences, Center for Biomolecular Structure and Dynamics, University of Montana

Chromatin Modifications and Fertility

Chromatin modifications are modifications of proteins called histones that regulate the compaction of DNA so it can fit in the nucleus of the cell. These modifications include methylation or acetylation of histones, and can result in activation or repression of transcription of the DNA, which is essential for the cell to synthesize proteins and pass on genetic information. Some chromatin modifications have been linked to fertility and specification of reproductive cells called germ cells.

Objectives

- Study chromatin modification H3 lysine-9 trimethylation (H3K9me3) that methylates histones causing transcriptional repression and has not been previously identified in germ cells
- Identify the enzyme that regulates this modification by testing various mutant strains
- Determine whether the loss of H3K9me3 is linked with loss of fertility

Hypothesis

- The loss or decline of H3K9me3 in germ cells will affect reproductive cells specification and fertility

Methods

- *C. elegans*, a eukaryotic nematode, was used as the model organism
- *C. elegans* shares similar reproductive regulation mechanisms as humans do, so results from this study can contribute to a better understanding of the mechanism in the development of germ cells in humans
- Used mutant strains with different methyltransferase mutations, some linked to sterility
- Examined presence of H3K9me3 in germ cells compared to somatic cells by using indirect immunofluorescence
- Compared the loss of H3K9me3 in germ cells with mutants known to be sterile to determine correlation

Immunofluorescence

- Specific antibodies used to stain and identify the germ cells and the H3K9me3 methylation
  1. Primary antibody binds to antigen
  2. Secondary antibody with fluorescence binds to primary antibody, "highlighting" it

Results

- Continue to test other mutants known to be sterile
- Investigate why Set-25 (supposed to be disrupted in H3K9me3) had a phenotype like the wild-type
- Analyze what stage in embryonic development H3K9me3 is lost and the regulator of it

Correlation of Loss of H3K9 Trimethylation with Sterility

<table>
<thead>
<tr>
<th>Mutant</th>
<th>Methylation disrupted (according to literature)</th>
<th>Excess of H3K9me3 in Germ Cells compared to Somatic Cells</th>
<th>Sterile?</th>
</tr>
</thead>
<tbody>
<tr>
<td>Met-1</td>
<td>H3K36m e3</td>
<td>+</td>
<td>No</td>
</tr>
<tr>
<td>Met-2</td>
<td>H3K9 me2</td>
<td>-</td>
<td>Yes, overtime</td>
</tr>
<tr>
<td>Met-1/Met-2</td>
<td>H3K36m e3 &amp; H3K9 me2</td>
<td>-</td>
<td>Yes</td>
</tr>
<tr>
<td>Mett-10</td>
<td>unknown</td>
<td>-</td>
<td>Partial</td>
</tr>
<tr>
<td>Set-2</td>
<td>H3K4 me3</td>
<td>-</td>
<td>Yes, overtime</td>
</tr>
<tr>
<td>Set-25</td>
<td>H3K9 me3</td>
<td>+</td>
<td>No</td>
</tr>
</tbody>
</table>

Future Research

- Mutants tested so far reveal a strong correlation between loss of excess H3K9me3 in germ cells and increasing sterility
- In Set-2 and Met-2 mutants, H3K9me3 decreases in germ cells over time, as the worms become more sterile
- Set 2 and Met 1/ Met 2 mutants appear to not lose H3K9me3 until the later 100 cell stage when 2 germ cells are present