

WATER DISINFECTION BY-PRODUCTS AND THEIR SAFETY

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Cummings Research-Disclosures

- **Role of phospholipase A₂ (PLA₂) in prostate cancer**

- Calcium independent PLA₂ (iPLA₂)
- Secretory PLA₂ (sPLA₂)
- Phospholipase A₂ receptor (PLA₂R)



- **Lipidomics**

- PLA₂ mediated changes in the lipidome in prostate tumors during transformation and treatment
- Changes in various tissues after oxidative stress
- Changes in the lipidome in different pathologies (drugs of abuse)
- Lots of ESI-MS



- **Renal Toxicology**

- Role of epigenetics in the nephrotoxicity of environmental oxidants and chemotherapeutics
- Epigenetic changes induced by nephrotoxic water disinfection by-products (DBPs)



Georgia Research Alliance



Water Disinfection Byproducts (DBPs)

- **Water disinfection**

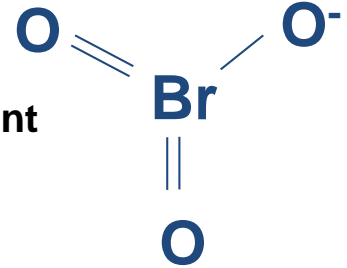
- Protects 260 million Americans from pathogens
- Oxidizes organic elements and produces byproducts

- **DBPs**

- Formed when disinfectants used in treatment plants react with naturally occurring material
 - **Chlorination = Trihalomethanes, haloacetic acid, chlorate**
 - **Chlorine Dioxide = Chlorite, chlorate**
 - **Chloramine = Chlorate**
 - **Ozonation = Bromate**

Bromate

- A source water disinfection byproduct of the ozonation process.
- Ozonation is a widely used method in Europe and Asia, and to a limited extent in the U.S.A.
- BrO_3^- is designated as a probable human carcinogen by the International Agency for Research on Cancer (IARC 1987, US EPA, 1986).
- Regulated levels established by the USEPA – **0.01 ppm** (Maximum Containment Level) – which is usually less than what is found after ozonation.

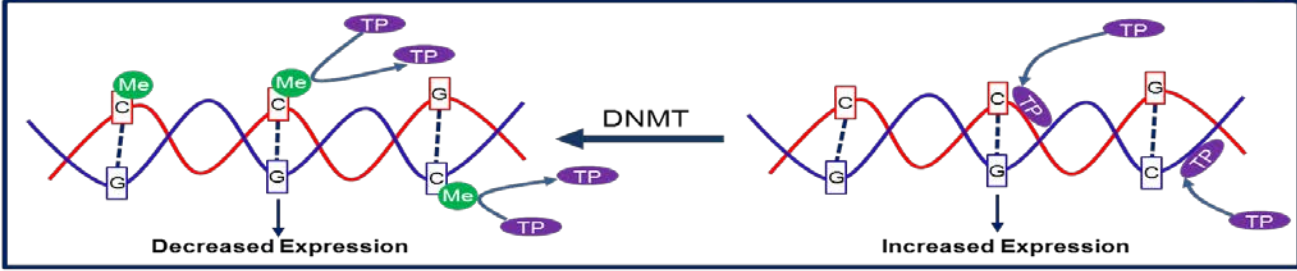
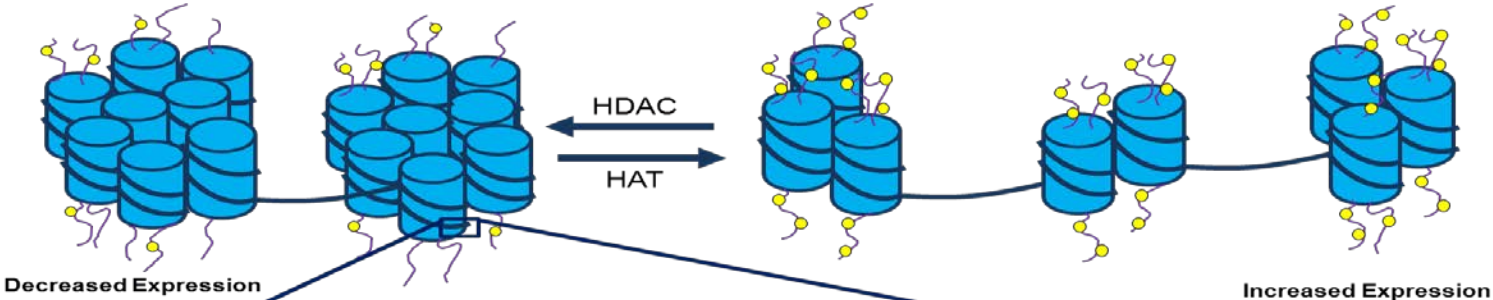


- EPA RrD = Urinary = Renal effects (urothelial hyperplasia)
- PoD = 1.1 mg/kg-day
- UF = 300

Epigenetic Alterations

- Heritable changes in gene expression that are **NOT** associated with sequence changes
- Epigenetic events include
 - DNA methylation
 - Histone modification
 - Nonfunctional RNA (microRNA)

Epigenetics and Gene Expression



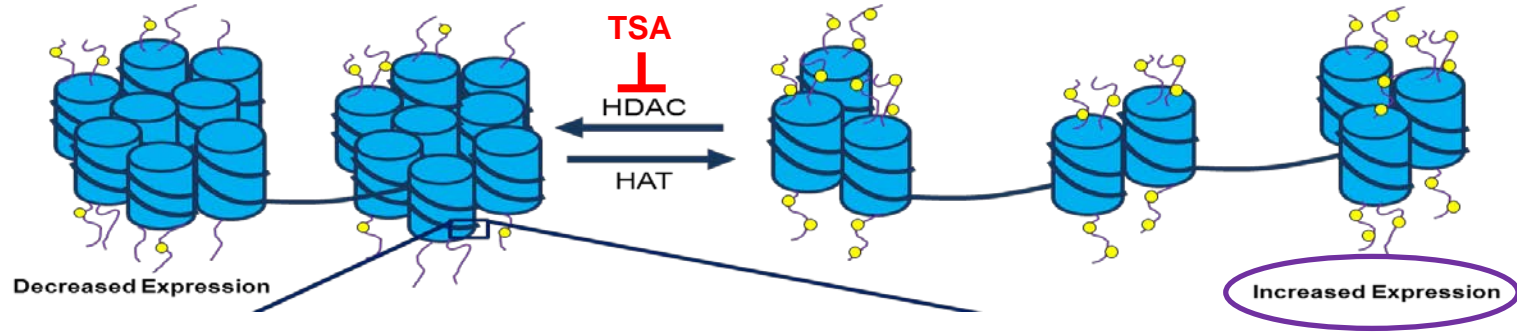
- Acetyl group
- TP Transcription Protein
- Me Methyl group

HDAC: Histone deacetylase
 HAT: Histone acetyltransferase
 DNMT: DNA methyltransferase

acetyl group: acetyl-coenzyme A
 methyl group: S-adenosyl methionine (SAM)

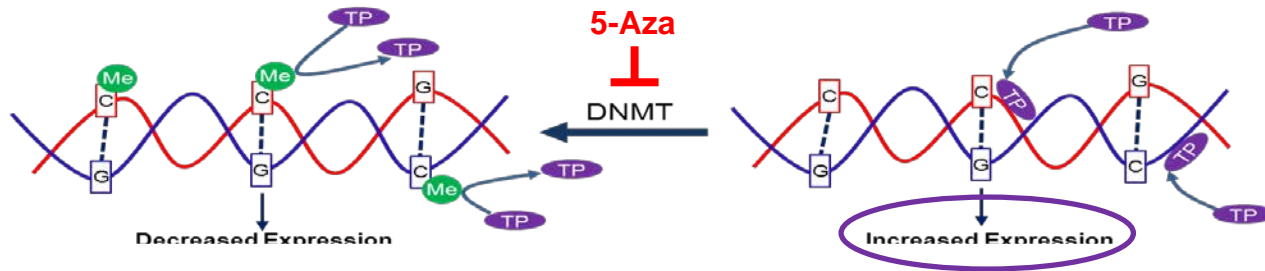
Epigenetic Inhibitors

- **Histone deacetylase (HDAC) inhibitors**
 - Trichostatin A (TSA)
 - Prevent deacetylation of histone
 - Prevents gene silencing
 - Hyperacetylation



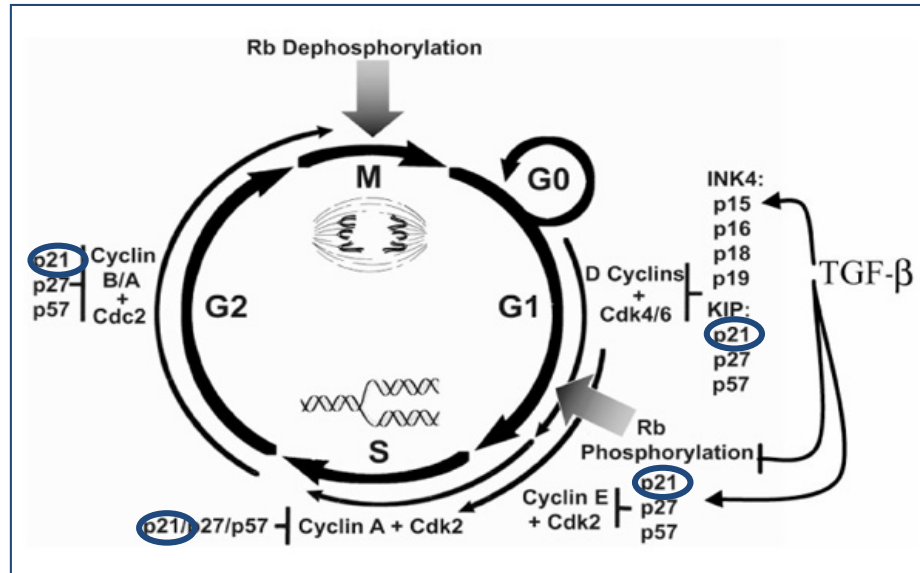
Epigenetic Inhibitors

- DNA methyltransferase (DNMT) inhibitors
 - 5-Azacytidine (5-Aza)
 - Prevents formation of 5-methylcytosine
 - Prevents gene silencing



p21

- p21 is a cyclin-dependent kinase inhibitor that regulates cell cycle progression at the G1 and S phases.
- The expression of this gene is controlled by both tumor suppressor protein p53-dependent and -independent pathways.



Bromate, p21 and Epigenetics

High doses of BrO_3^- induces DNA damage and 8-OHdG production (a measure of oxidative stress) *in vitro*.

BrO_3^- induces toxicity in human and rat kidney cells.

Inhibition of DNA methylation or histone deacetylation increased p21 expression and altered cell death.

Changes in p21 expression after exposure to low-dose BrO_3^- correlated to changes in DNA methylation within the coding region.

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Toxicological Sciences

Epigenetic Changes in p21 Expression in Renal Cells after Exposure to Bromate

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This study tested the hypothesis that bromate (KBrO_3)-induced renal cell death is mediated by epigenetic mechanisms. Global DNA methylation, as assessed by 5-methylcytosine staining, was not changed in normal rat kidney cells treated with acute cyto-

BrO_3^-	Bromate
Br^-	Bromide
Cdkn1a	Cyclin-dependent kinase inhibitor 1a
DBP	Disinfection byproduct
DNMT	DNA methyltransferase

BrO_3^- -induced increase in p21 expression is regulated by epigenetic mechanisms.

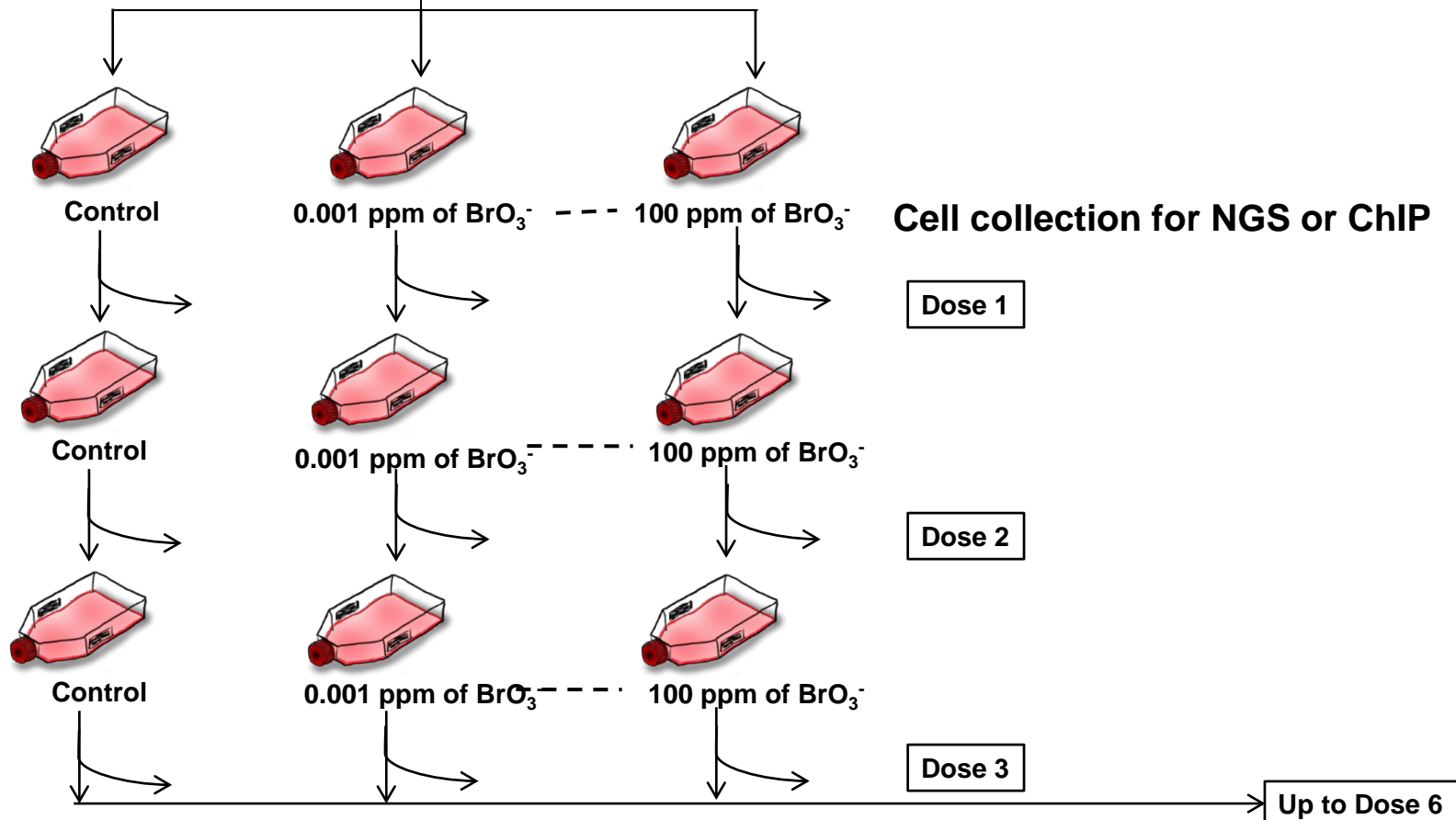
Questions

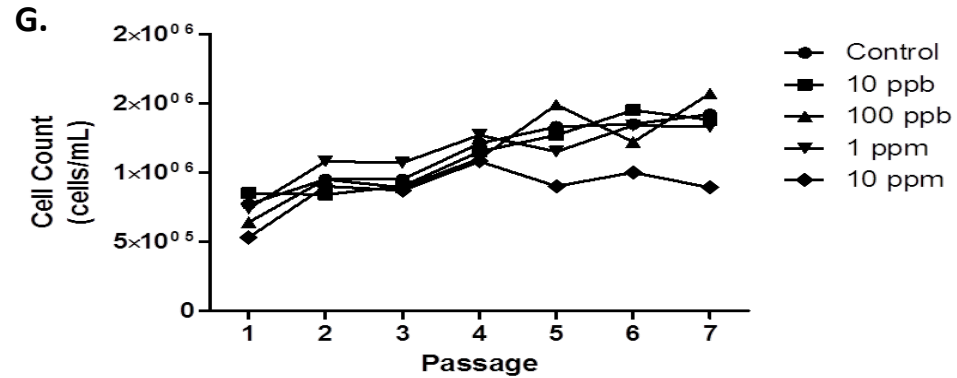
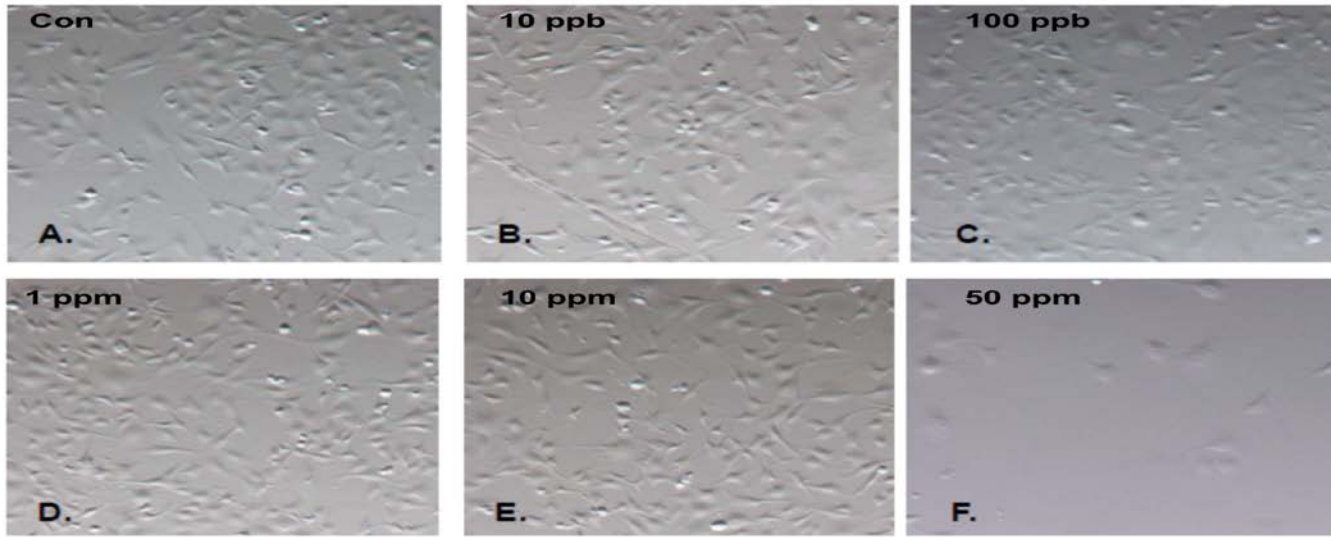
- 1. Should epigenetic be used to assess the safety of DBPs?**
- 2. What are the difference between rats and humans with regards to epigenetics?**
- 3. What does this mean for risk assessment?**

Will address the above questions using p21 and bromate as an model

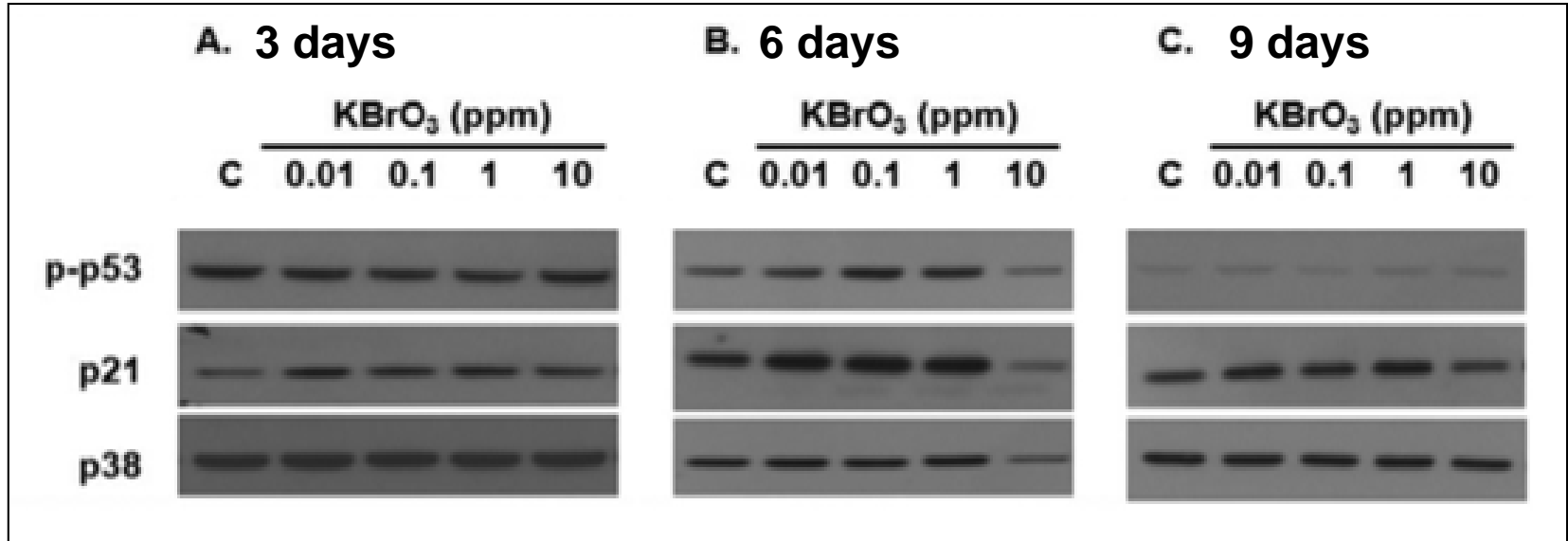
Kidney cells

Sub-Chronic Dosing Regime



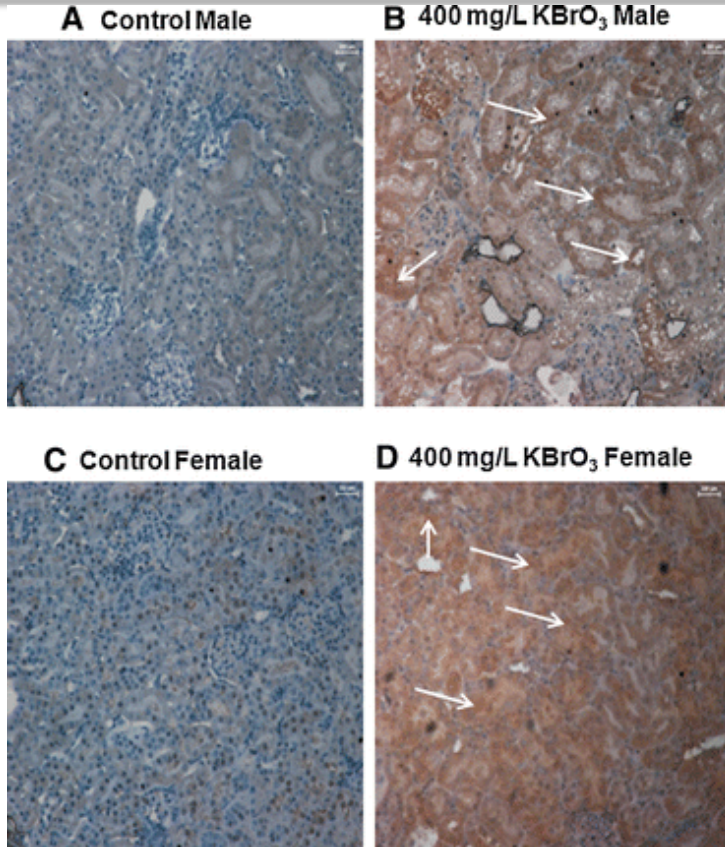


Effect of Bromate on p21 Expression in Renal Cells

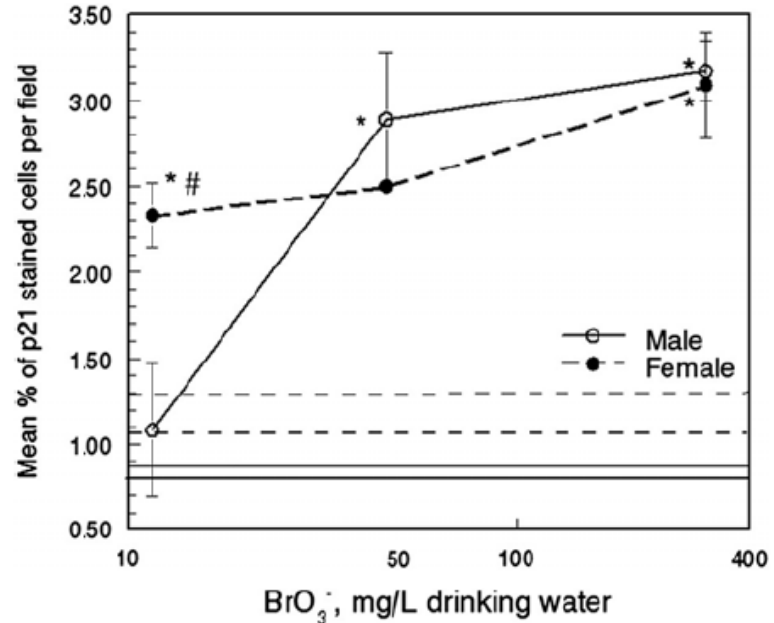


Normal Rat Kidney Cells
Bromate MCL = 0.01 ppm
Scholpa et. al, *Toxicol Sci*, 2014

Effect of Bromate on p21 Expression in Renal Cells



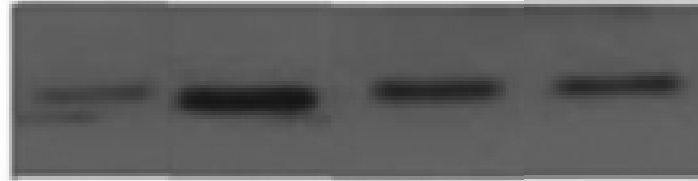
28 day exposure



Effect of Epigenetic Inhibitors on p21 Expression in Renal Cells

NRK Cells
72 hour Exposure

p21



KBrO ₃ (100 ppm)	-	+	-	-
5-Aza (20 μM)	-	-	+	-
TSA (100 nM)	-	-	-	+

5-Aza = DNA Methyltransferase

TSA = Histone deacetylase inhibitor

Summary Part 1

- Bromate increases the expression of p21 in renal cells after both acute and sub-chronic exposures
- Increases in p21 expression *in vitro* occurred at doses that are as low as the MCL (0.01 ppm)
- Bromate-induced p21 expression was altered by epigenetic inhibitors
- What is mechanism mediating bromate-induced epigenetic changes in p21?
- What are the differences between rats and humans?
- What does this mean for risk assessment?

How Does Bromate Alter p21 Expression?

- **Identify the specific changes in DNA induced by bromate**
- **Epigenetic changes**
 - DNA Methylation?
 - Histone acetylation?
- **Problem**
 - Assessment of DNA methylation is laborious and time consuming
 - Need a high through put more accurate way to assess DNA methylation of targeted genes

Targeted-Bisulfite Next Generation Sequencing (TB-NGS)

Log phase cells

Dose with BrO_3^- for 72 hrs

Extract DNA and convert by bisulfite treatment

Gene-specific PCR for p21 coding and promoter region (fusion primers with adapters for Illumina)

Elute from gel and normalize

Limited cycle PCR with tagged primers (Tag A-H i5 and Tag 1-12 i7)

Pool and sequence on Illumina MiSeq (next-generation)

Clean with Speedbeads and normalize

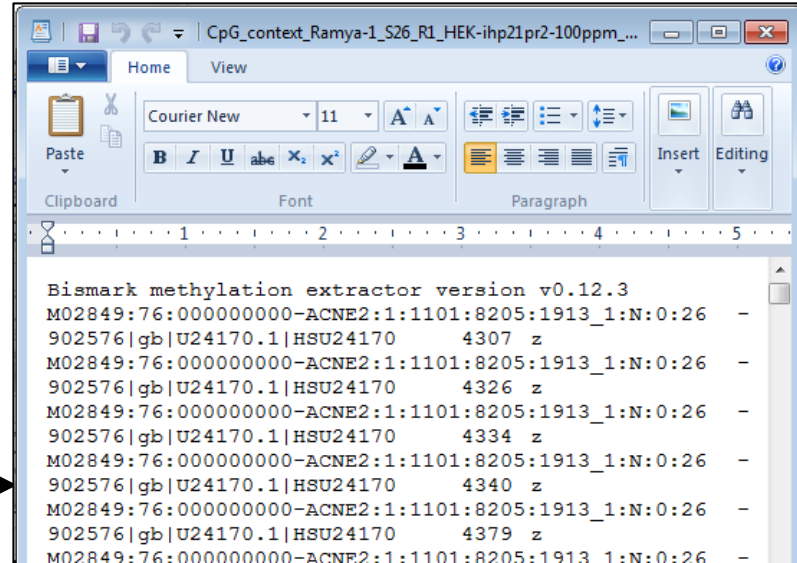
DNA Methylation Analysis Using Bismark Bisulfite Mapper

Prepare reference genome folder

Convert reference genome to bisulfite converted version.

Align the sequence reads from MiSeq to reference.

Extract percent methylation output.



The screenshot shows a text editor window with the following text:

```
Bismark methylation extractor version v0.12.3
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
902576|gb|U24170.1|HSU24170 4307 z
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
902576|gb|U24170.1|HSU24170 4326 z
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
902576|gb|U24170.1|HSU24170 4334 z
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
902576|gb|U24170.1|HSU24170 4340 z
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
902576|gb|U24170.1|HSU24170 4379 z
M02849:76:000000000-ACNE2:1:1101:8205:1913_1:N:0:26 -
```



B)

Tagged i5

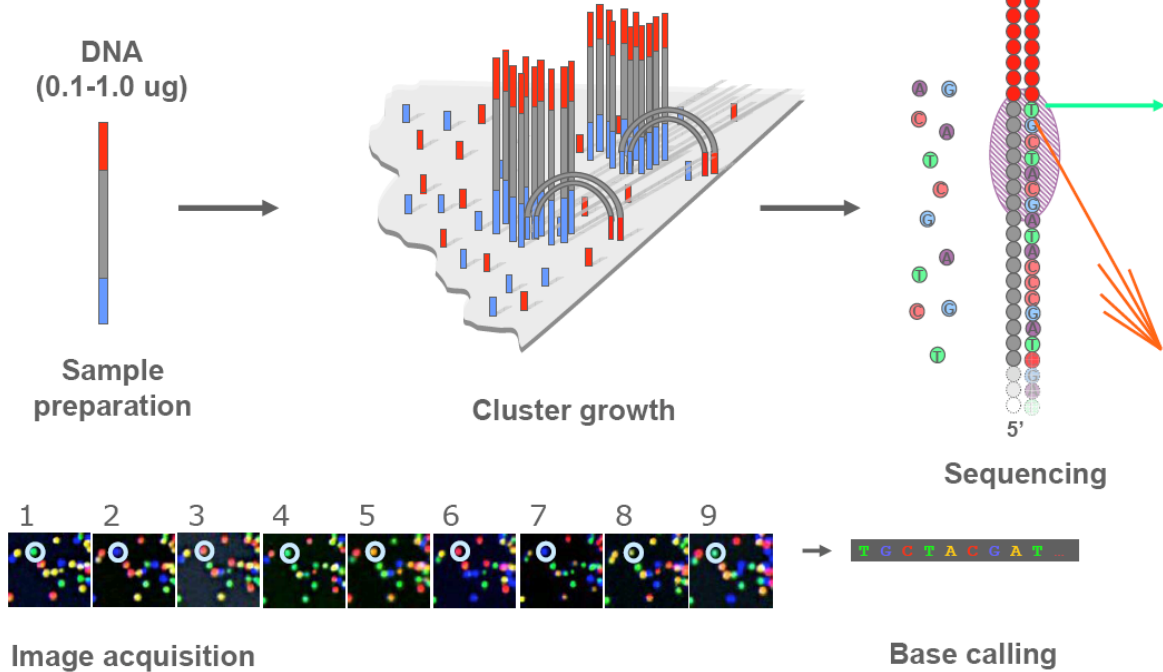
Tagged i7

	A	B	C	D
1	A-1	B-1	C-1	D-1
2	A-2	B-2	C-2	D-2
3	A-3	B-3	C-3	D-3
4	A-4	B-4	C-4	D-4

Sequencing using Illumina MiSeq

Illumina Sequencing Technology

Robust Reversible Terminator Chemistry Foundation



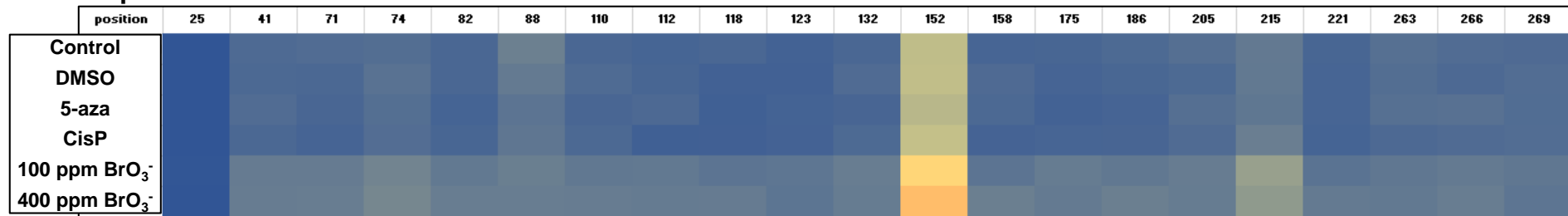
Heat-Map of Site-Specific Percent DNA Methylation Changes

A. Human p21 promoter region

CpG Site-Specific % Methylation

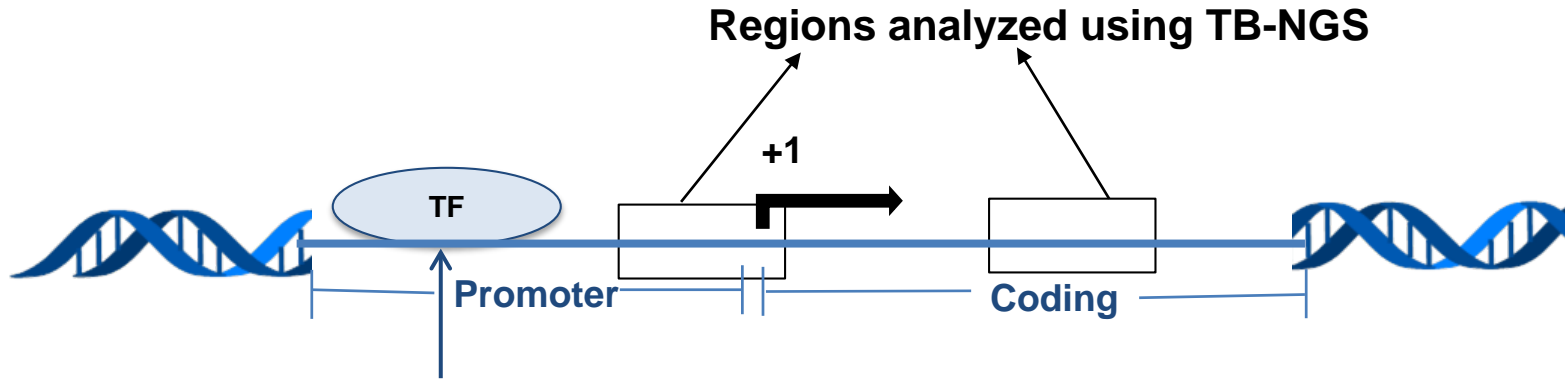


B. Rat p21

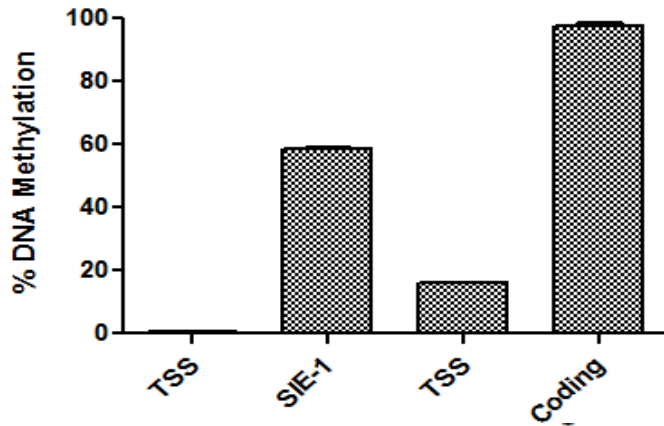


0%  100%

Percent Methylation

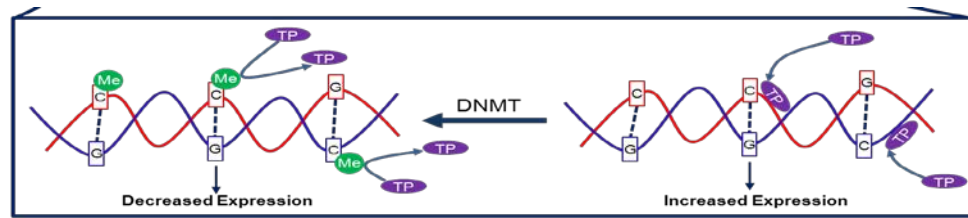


Transcription Factor
Binding Site

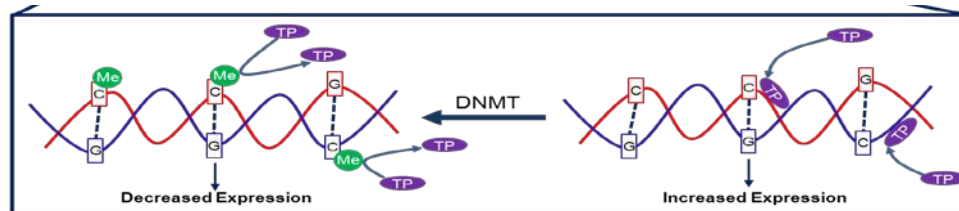
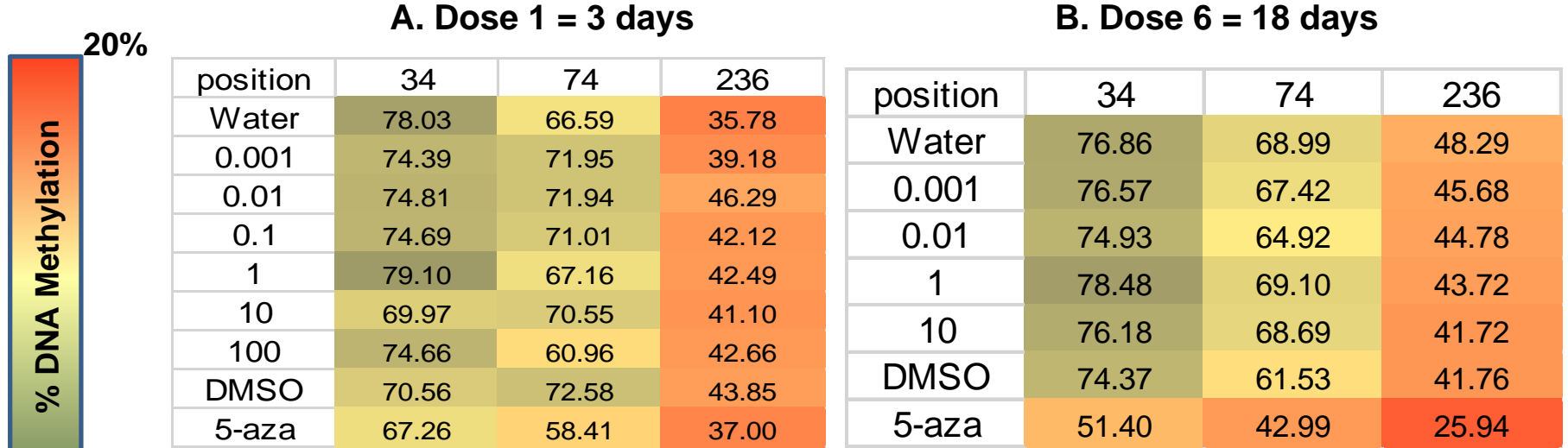


Human p21

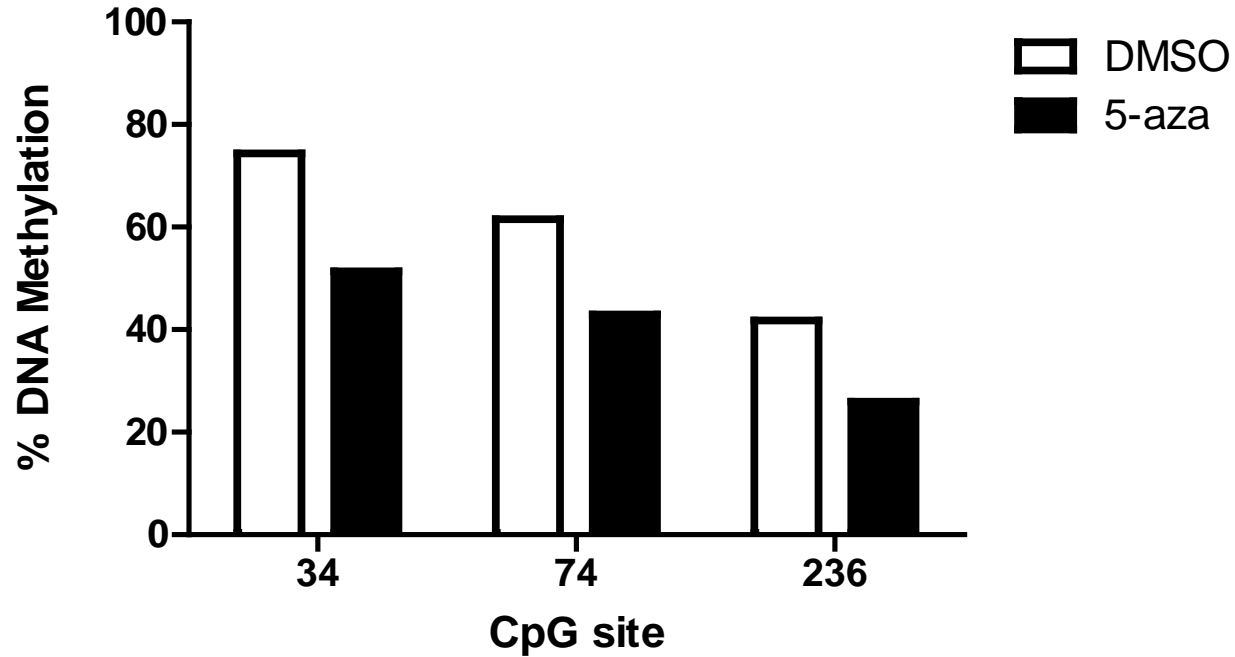
Rat p21



Bromate-Induced Change in DNA Methylation in Human p21



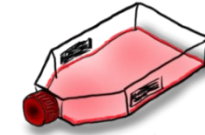
5-Aza-Induced Change in DNA Methylation in Human p21



Bromate does not appear to induce change in DNA methylation in p21

Bromate-Induced Change in p21 Histone Acetylation (ChIP Assay)

Cell collection and DNA cross-linking



Cell lysis and DNA shearing



Protein/DNA Immunoprecipitation



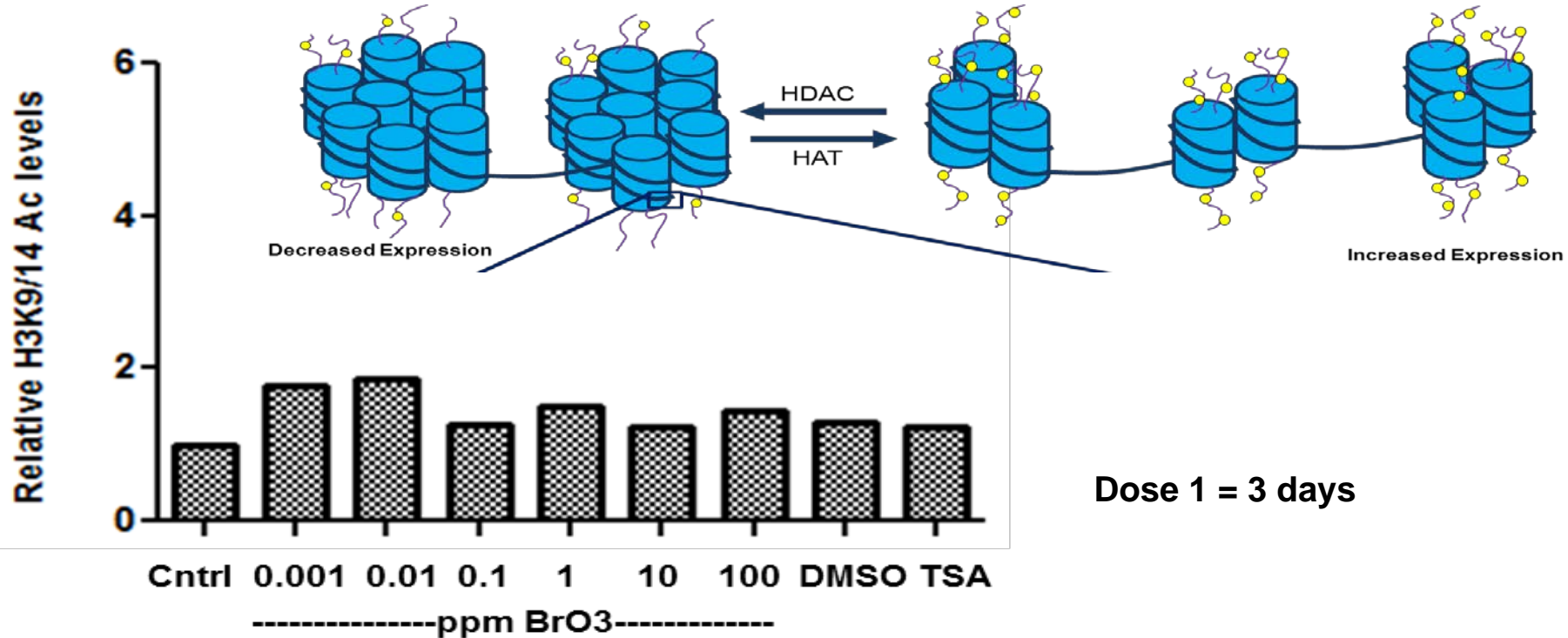
Reverse cross-link



DNA purification and PCR

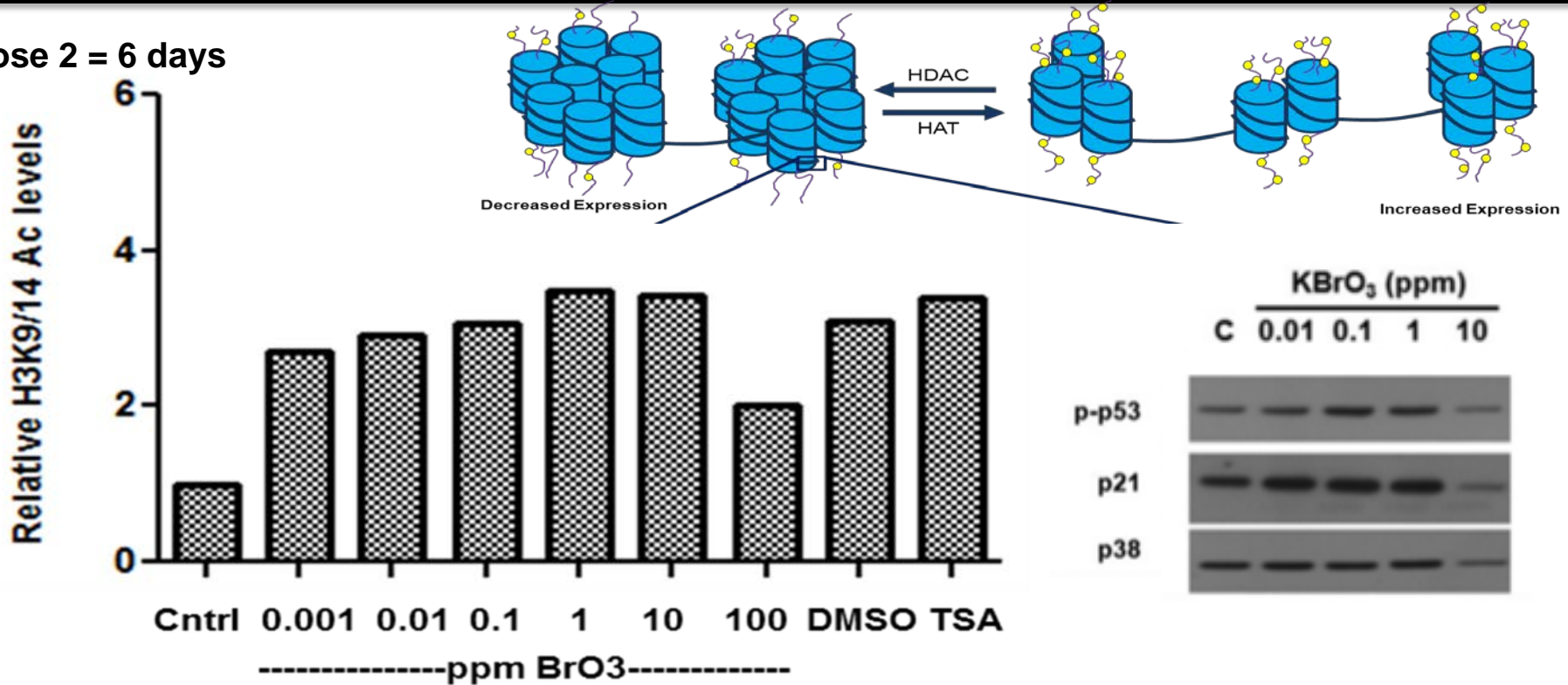


Acetylation of H3K9/14 in the p21 Promoter Region in Rat Kidney Cells

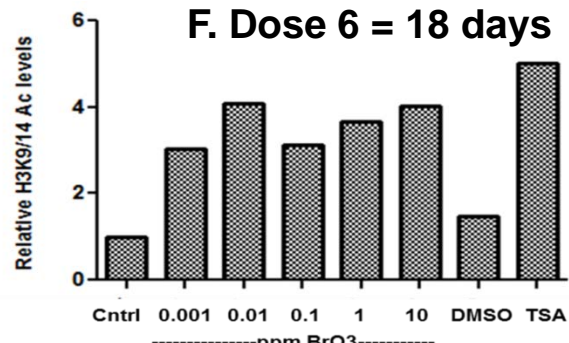
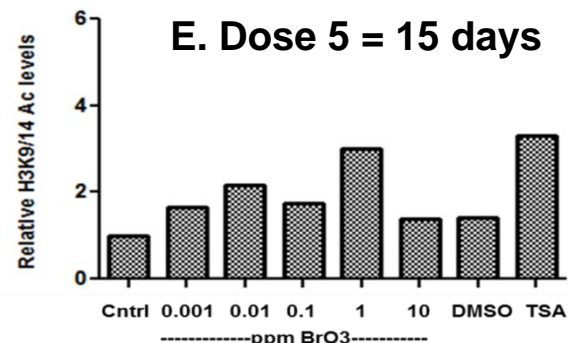
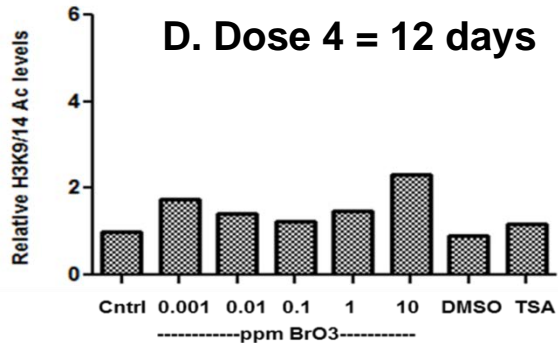
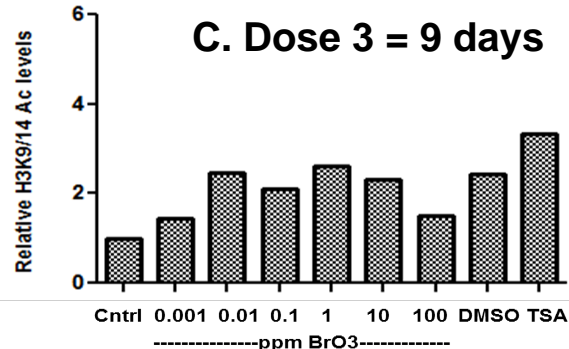
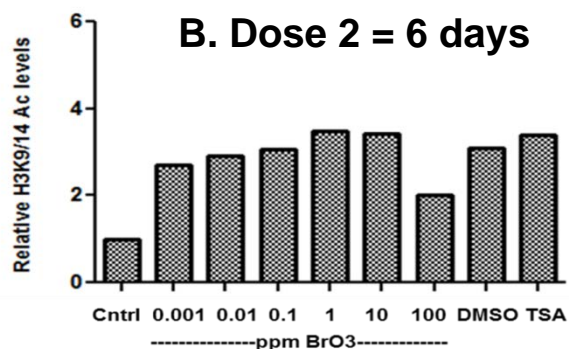
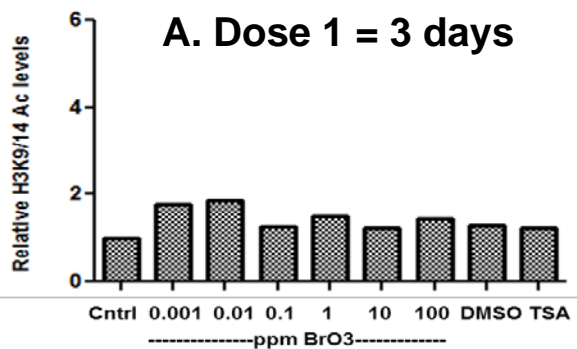


Acetyl Histone of H3K9/14 in the p21 Promoter Region in Rat Kidney Cells

Dose 2 = 6 days



Acetyl Histone of H3K9/14 in the p21 Promoter Region in Rat Kidney Cells



Summary Part 2

- **Bromate increases the acetylation of histone in p21 in renal cells after both acute and chronic exposures**
- **Increases in p21 histone acetylation in vitro occurred at doses that are as low as the MCL (0.01 ppm)**
- **Bromate increase in p21 expression appear to be regulated by histone acetylation not DNA methylation**

Questions

- **What is mechanism mediating bromate-induced epigenetic changes in p21?**
 - **Histone acetylation**
- **What are the difference between rats and humans?**
 - **Significance difference in basal DNA methylation in p21 between rats and humans**
 - **Acetylation data is still being tabulated**

Questions

- **What does this mean for risk assessment?**
 - **Differences exist in the epigenetic regulation of p21 between rats and humans**
 - **Increases in p21 expression in the kidney is protective against ischemia reperfusion and chemotherapeutic toxicity**
 - **Epigenetic changes in p21 may be protective and not part of the mechanism of toxicity**

Key Takeaways-Epigenetics and DBPs

- Epigenetic changes in rats do not always translate to humans
 - **At this moment, you can't use epigenetic changes in rats to assess the risk of bromate in humans**
- An epigenetic change isn't always an adverse change
 - EPA Bromate PoD = 1.1 mg/kg-day Urothelial hyperplasia
 - Epigenetic change = 0.01 mg/L-3 day = p21 expression
 - **My big worry is if someone uses this epigenetic marker *in vivo* as a PoD**

Cummings Research-Acknowledgements-People

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