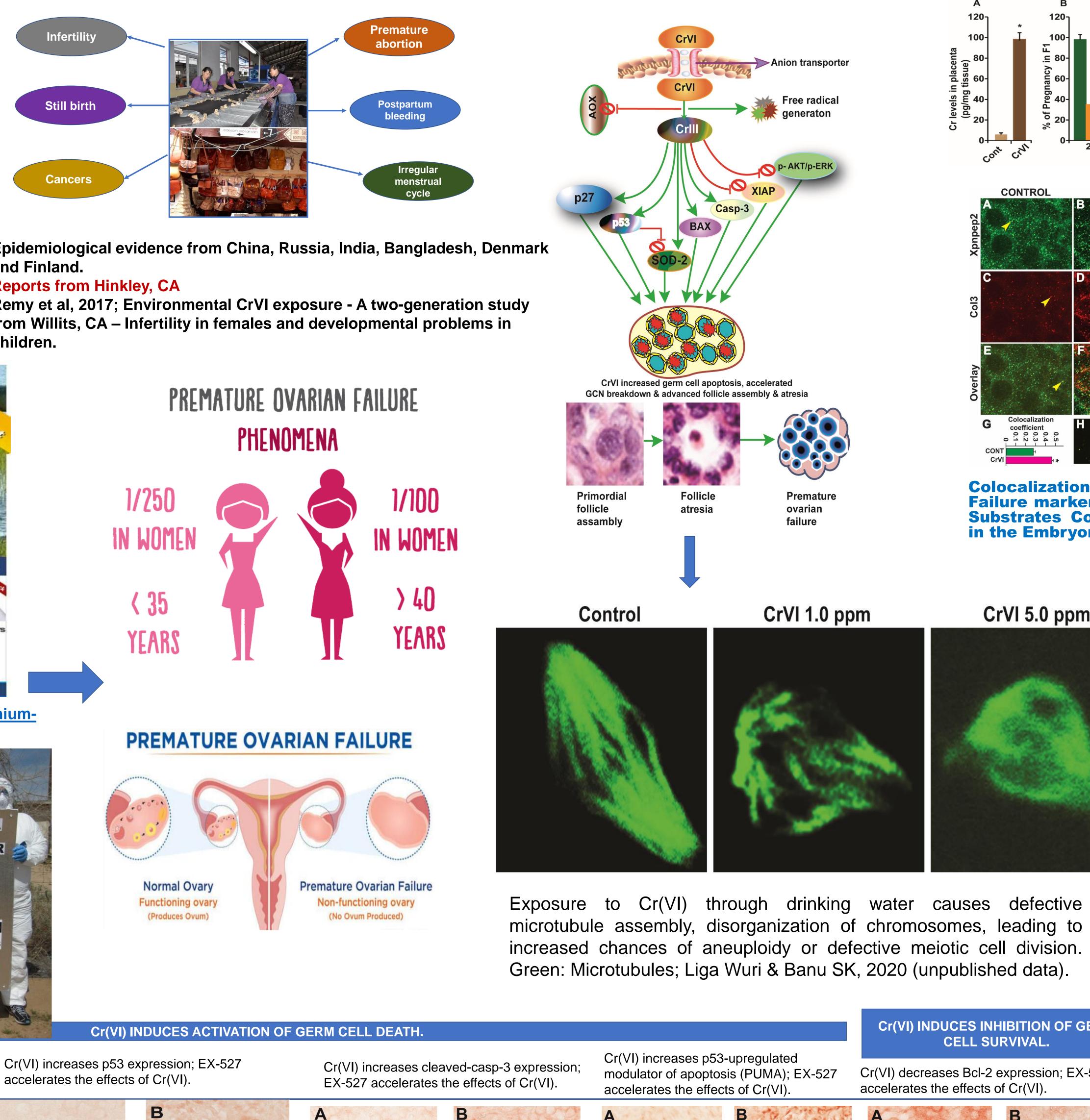
Abstract

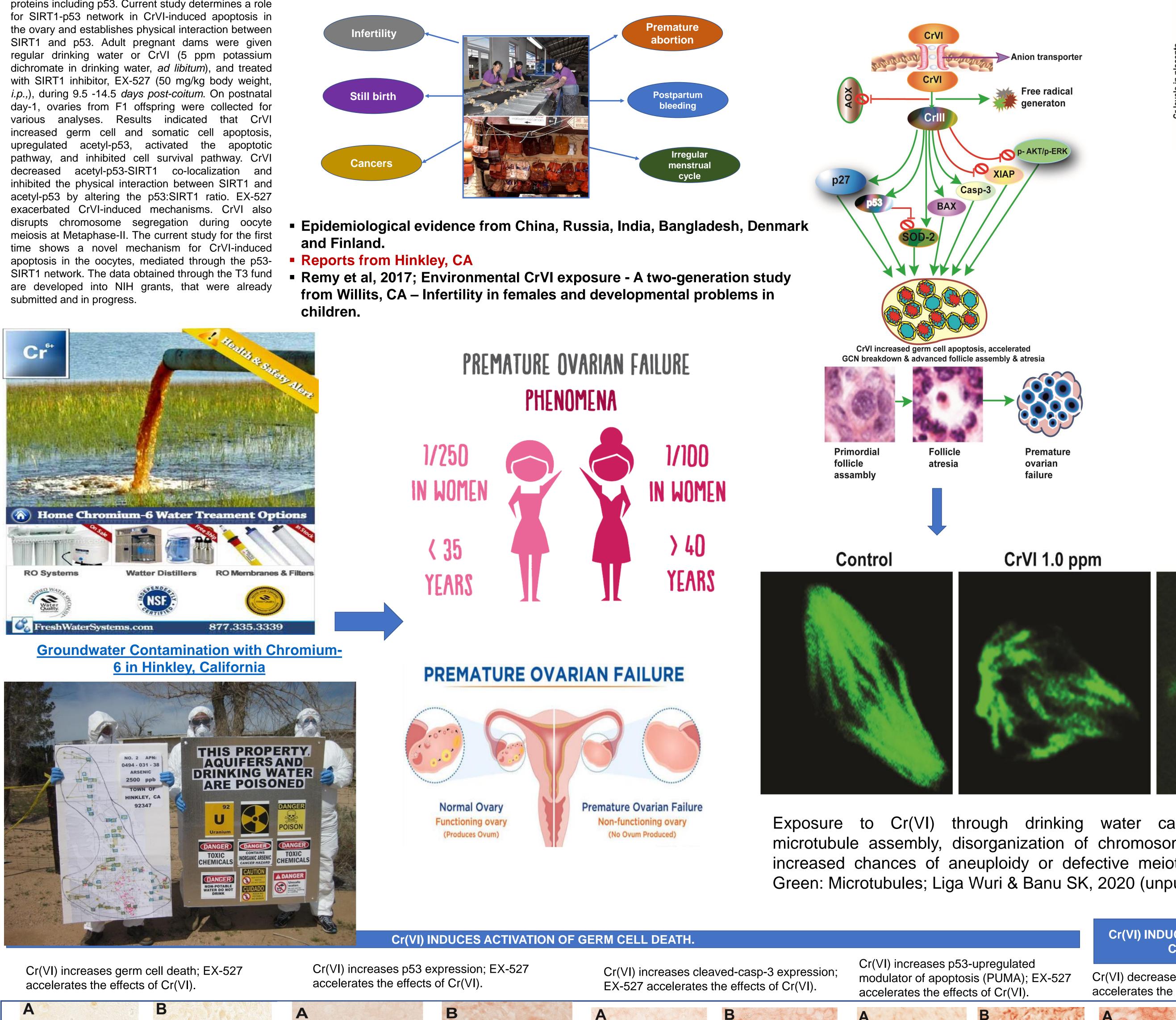
Environmental contamination with hexavalent chromium (CrVI) has been increasing in the United States as well as in the developing countries. Due to increased usage of Cr and improper disposal of Cr wastes from the industries, CrVI levels in the have been continuously increasing. Exposure to CrVI predisposes human population to various diseases, including cancer, infertility, and developmental problems in children. Findings from our laboratory reported that prenatal exposure to CrVI caused premature ovarian failure through p53mediated mechanisms. Sirtuin 1 (SIRT1) is an NAD+dependent histone deacetylase class III. SIRT1 deacetylates several histones and non-histone proteins including p53. Current study determines a role acetvl-p53. activated the apoptotic

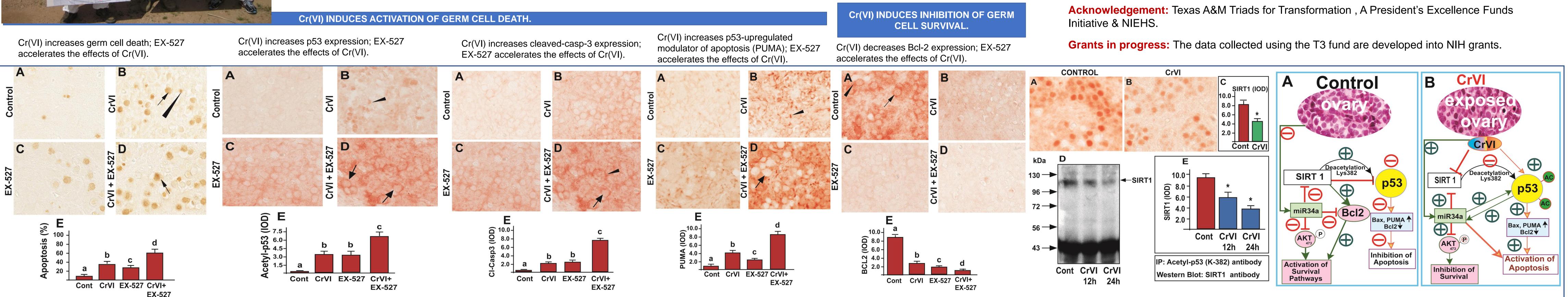
Cr(VI) Toxicity Targets p53-SIRT1-mediated germ cell death that Leads to Premature Ovarian Failure.

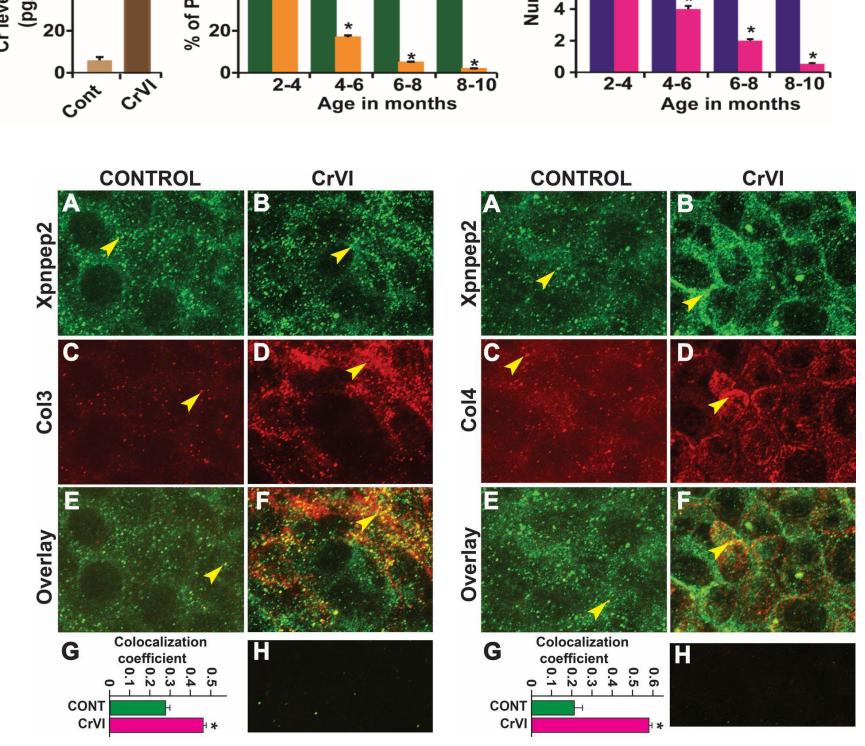
Sakhila K. Banu



- and Finland.

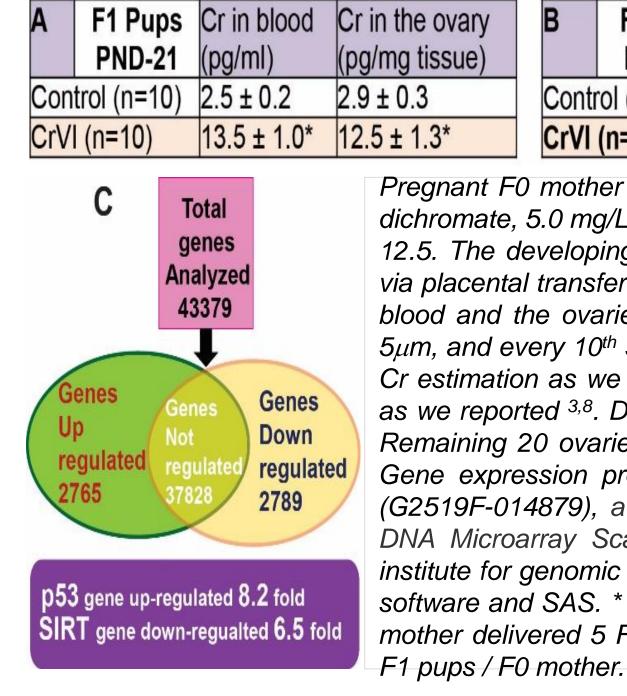






Colocalization of Xpnpep2, a Premature Ovarian Failure marker candidate in the human, and it's **Substrates Collagen-3 and Collagen-4 proteins** in the Embryonic (day 17.5) ovaries

CrVI 5.0 ppm

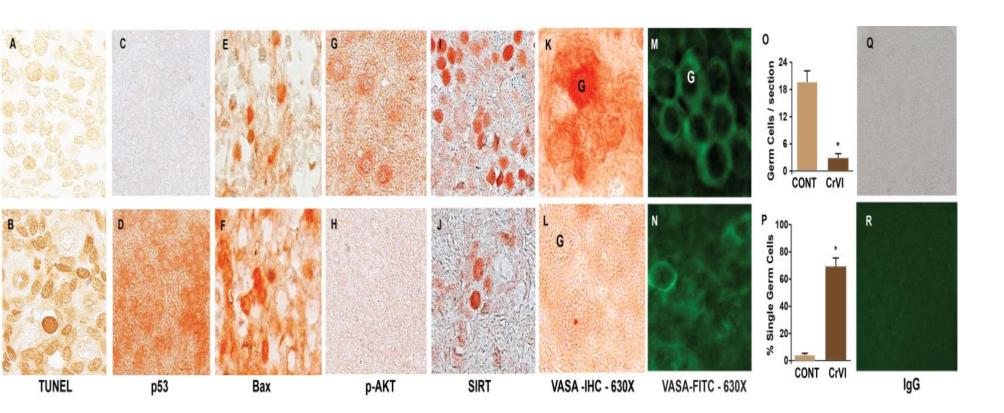


Conclusion: Exposure to hexavalent Cr(VI) through drinking water causes premature ovarian failure by causing germ cell death, disrupting cell survival, activating p53, and inhibiting Sirtuin-1. Cr(VI) disrupts chromosome segregation during oocyte meiotic division. Cr(VI) alters extracellular matrix, and disrupts premature ovarian failure markers. All these molecular changes lead to premature ovarian failure, premature abortions, and infertility in females.





T3: TEXAS A&M TRIADS FOR TRANSFORMATION A President's Excellence Fund Initiative



Pregnant F0 mother rats were given regular drinking water (n=5) or exposed to CrVI (potassium dichromate, 25 mg/L) (n=5) through drinking water during gestational days (GD) 8.5 – 12.5. The developing F1 fetuses were exposed to the respective treatment in utero via placental transfer. On PND-1 the female F1 pups (n=20/group) were euthanized and ovaries collected. Randomly, 10 ovaries were fixed in 4% PFA and processed for TUNEL assay and IHC, 10 ovaries were fixed as whole mount to analyze germ cell nest breakdown, and 20 ovaries were used for RNA extraction for miRNA and/or gene expression analyses. (A-B) TUNEL assay was performed using in situ TUNEL assay kit as we reported (3). (B-J) Expression of proapoptotic and apoptotic proteins was performed using IHC and quantified using Image Pro Plus as we reported (3). Note: histograms were not shown due to page limit. Expression of VASA-positive germ cells by IHC (K-L) and by whole mounts and confocal microscopy (M-N). Germ cell nest break down was measured by z-stack imaging as described by our collaborator (39). Number of VASA-positive germ cells/section was counted in 32 random fields from 8 ovaries (4 fields/ovary). Percentage of single oocytes was counted in each section and expressed as mean ± SEM (O-P). Data were analyzed by ANOVA. * P<0.05, Control vs. CrVI. Internal control IgG (Q-R).

В	F1 Pups	No. of primordial	No. of primary	No. of secondary	No. of atretic
	PND-21	follicles	follicles	follicles	follicles
Cont	rol (n=10)	501 ± 25	61 ± 3.8	31 ± 3.8	34 ± 3.0
CrVI	(n=10)	241 ± 11*	40 ± 2.5*	6.1 ± 0.5*	99 ± 7.0*

Pregnant F0 mother rats were given drinking water (n=5) or exposed to (potassium) dichromate, 5.0 mg/L) (n=5) through drinking water during gestational days (GD) 8.5 – 12.5. The developing F1 fetuses were exposed to the respective treatment in utero via placental transfer. On PND-21 the female F1 pups (n=10/group) were euthanized, blood and the ovaries collected. Ten ovaries were fixed in 4% PFA, serially cut at $5\mu m$, and every 10th section was examined using H&E. (A) Ten ovaries were used for Cr estimation as we described ^{3,8}. (B) The healthy and atretic follicles were counted as we reported ^{3,8}. Data were analyzed by ANOVA. * P<0.05, Control Vs. CrVI. (C) Remaining 20 ovaries were used for RNA extraction and processed for microarray. Gene expression profiling was performed using Rat Gene Expression Microarray (G2519F-014879), a glass slide formatted with four high-definition 44K arrays and DNA Microarray Scanner, (Agilent Technologies). The protocol provided by "The institute for genomic research" was followed. Data were analyzed using 'R' Statistical software and SAS. * P<0.05, Control Vs. CrVI. Note: Our data indicate that each F0 mother delivered 5 F1 female pups; therefore, the estimated sex ratio is five female