

Supplemental Information

Route of Arsenic Exposure Differentially Impacts the Expression of Genes Involved in Gut-Mucosa-Associated Immune Responses and Gastrointestinal Permeability

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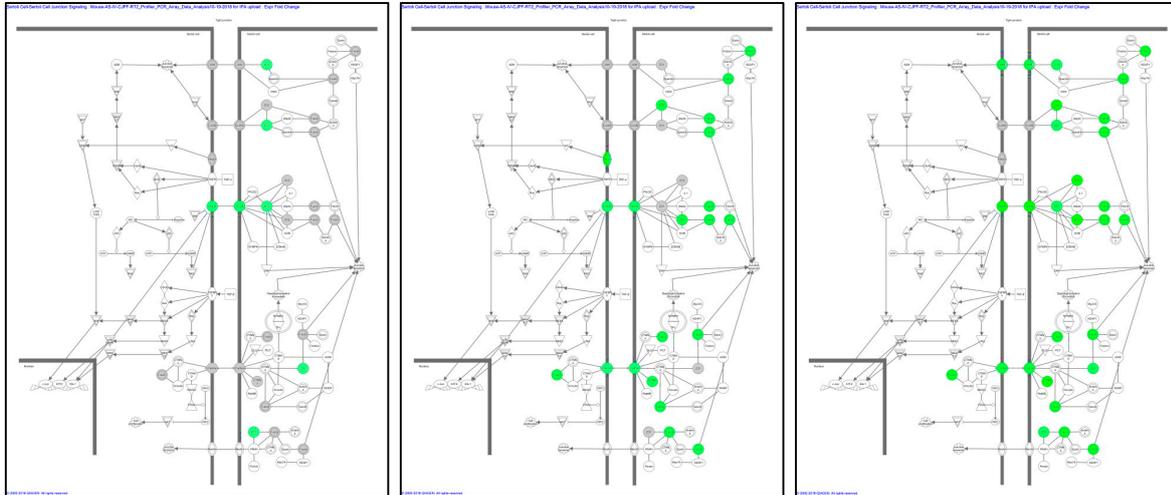
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1 h

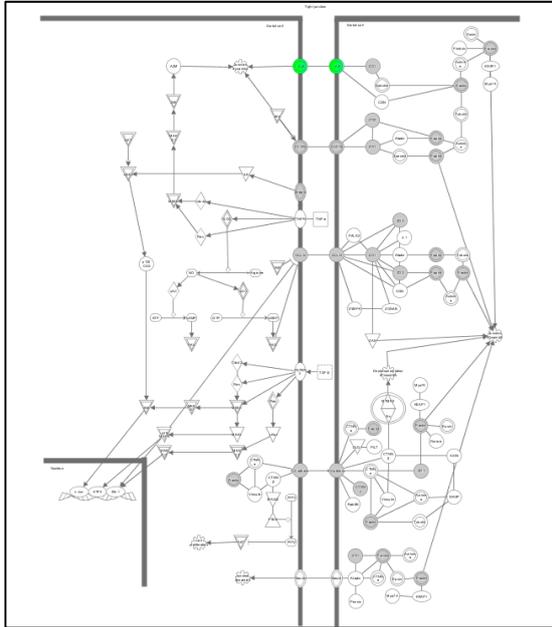
4 h

48 h

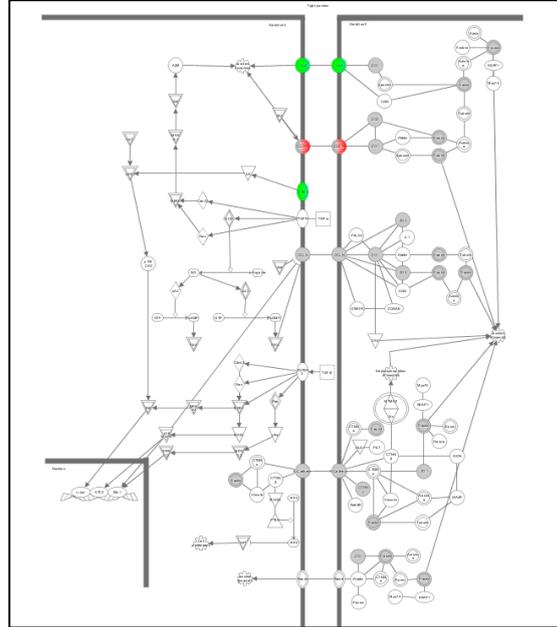


Supplemental Figure S1. Ingenuity Pathway Analysis of differential expressed genes in the cell-cell junction pathway during sodium arsenite IV exposure. This schematic diagram shows the interaction and expression of genes involved in the cell-cell junction pathway in the intestinal mucosa during sodium arsenite exposure. Sodium arsenite IV exposure initiate a cascade of reaction for the perturbation of mRNA expression of cell junction related genes as early as 1 h post exposure, with a maximum perturbation at 48 h post exposure. A fold change 1.5 with a p-value ≤ 0.05 were considered significant. Green circles represent down regulation, red circles represent upregulation, and gray circles represent no perturbation.

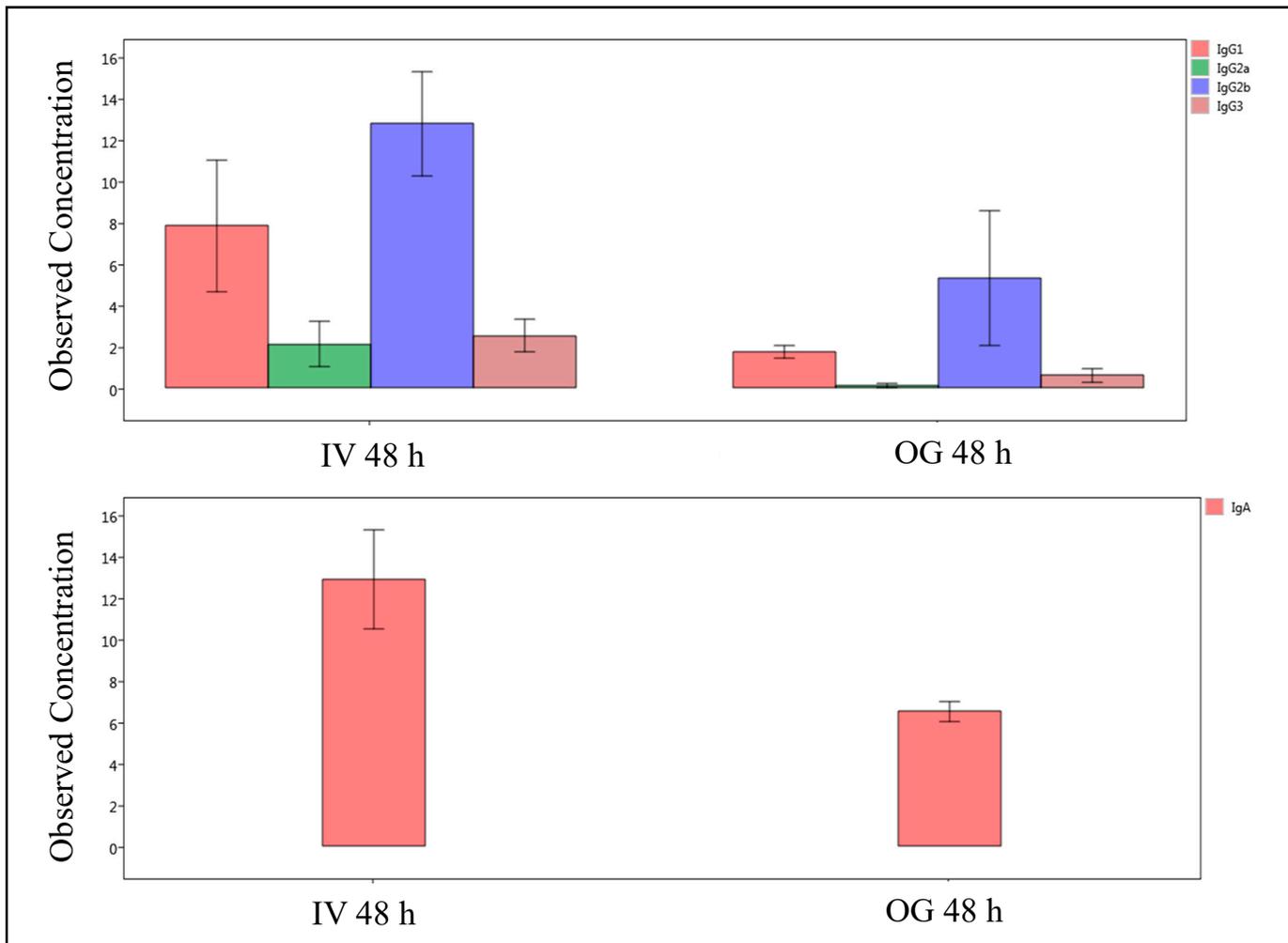
OG 24 h



OG 48 h



Supplemental Figure S2. Ingenuity pathway analysis of genes in the cell-cell junction pathway during sodium arsenite oral gavage. This schematic diagram shows the interaction and expression of genes involved in the cell-cell junction pathway in the intestinal mucosa during sodium arsenite oral exposure. A fold change 1.5 with a p-value ≤ 0.05 were considered significant. Sodium arsenite oral exposure had very limited perturbation of mRNA expression of cell junction related genes. Green circles represent down regulation, red circles represent upregulation, and gray circles represent no perturbation.



Supplemental Figure S3: Comparison of IgG Isotypes and IgA in Ileal Tissue. Plot of IgG isotypes (Top panel; a) or IgA (lower panel; b) in ileal tissue of mice after 48 h exposure to sodium arsenite intravenously (IV 48 h) versus orally (OG 48 h). Significance is represented as a star, where significance indicates $p \leq 0.05$. P-values were determined using the Mann-Whitney method of statistical analysis.