



COBALT, ANTIMONY COMPOUNDS, AND WEAPONS-GRADE TUNGSTEN ALLOY

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Table S4.23 Evidence relevant to multiple key characteristics in human cells in vitro exposed to cobalt

End-point	Platform	Results ^a	KC associated	Tissue, cell line	Concentration, duration	Comments	Reference
<i>Soluble cobalt(II) salts</i>							
<i>Cobalt(II) chloride (CoCl₂)</i>							
Transcriptomics; protein expression	TruSeq Stranded mRNA Library Prep and NextSeq 550 sequencing; nanoLC, HF hybrid quadrupole-orbitrap MS	↑ HIF-1α activity, genes associated with cell proliferation, oxidative stress response, inflammatory response, glycolysis, and gluconeogenesis ↑/↓ Proteins associated with cell proliferation and glycolysis	KC5, KC6, KC10	Colon adenocarcinoma, differentiated Caco-2	300 μM, 24 h	Source and purity not reported, positive and negative controls included, little correlation between changes in genes and proteins (<i>n</i> = unclear).	Knyazev et al. (2021)
Transcriptomics	TruSeq Stranded mRNA Library Prep and NextSeq 550 sequencing	↑ HIF-1α mRNA (HT-29 only, no change Caco-2) ↑ Glycolysis and gluconeogenesis, MHC class I antigen presentation pathway, and ribosomal proteins	KC5, KC7, KC10	Colon adenocarcinoma, HT-29 and differentiated Caco-2	300 μM, 24 h	Source and purity not reported, positive and negative controls included (<i>n</i> = 3 replicates).	Nersisyan et al. (2021)
Transcriptomics	Affymetrix HU-gene st1.0 microarrays	↓ Cell death-related genes ↑ Carbohydrate metabolism and glycolysis, oxidoreductase activity, cell proliferation, and angiogenesis	KC5, KC10	Colon carcinoma, RKO	200 μM, 24 h	Source and purity not reported, negative controls included (<i>n</i> = 2 duplicates).	Sheffer et al. (2011)
Transcriptomics	HumanHT-12 v4.0 expression bead chip kit	↑ Cell development and death, immune response, and B-cell activation ↑/↓ Cell cycle, transcription, and kinase activity	KC7, KC10	Multiple myeloma, U266	100 μM, 24 h	Source and purity not reported, treatment caused 30% cytotoxicity, negative controls included (<i>n</i> = unclear).	Bae et al. (2012)
Transcriptomics	Global transcription response by CEA microarrays	↑ HIF-1α activity, differentiation, and proliferation ↑/↓ Signal transduction and trafficking, stress response, and immune response	KC5, KC7, KC10	Lung adenocarcinoma, A549	2 mM, 24 h	Source and purity not reported, doses chosen based upon 50% cytotoxicity by trypan blue exclusion, positive and negative controls included (<i>n</i> = 3 duplicates).	Malard et al. (2007)

Table S4.23 (continued)

End-point	Platform	Results ^a	KC associated	Tissue, cell line	Concentration, duration	Comments	Reference
Transcriptomics	Whole human genome 44 K oligonucleotide array	↑ HIF-1α activity, glycolysis and gluconeogenesis, and focal adhesion	KC5, KC10	Keratinocytes, HaCaT	3 µg/mL [\sim 23 µM], 3 h or 3 days	Purity not reported, negative controls included ($n = 5$ independent replicates).	Busch et al. (2010)
Transcriptomics	Affymetrix U95A version 1 gene chip	↑ HIF-1α activity ↑/↓ Carbohydrate metabolism and glycolysis, oxidoreductase activity, cell proliferation, and angiogenesis	KC5, KC10	Liver carcinoma, Hep3B	100 µM, 24 h	Purity not reported, positive and negative controls included ($n = 2$ replicates).	Vengellur et al. (2005)
Extracellular proteins in conditioned media	Shotgun proteomics, nanoLC-MS/MS using LTQ-Orbitrap XL hybrid mass spectrometer	24/66 secreted proteins were affected, 23/24 downregulated; effects on connective tissue regulation, histone acetylation, lymphocyte proliferation, and protein processing	KC4, KC10	Bronchial cells, BEAS-2B	20 µg/mL [\sim 150 µM], 24 h	Doses chosen based upon < 5% cytotoxicity by trypan blue exclusion, positive and negative controls included ($n = 3$ duplicates).	Malard et al. (2012)
Intracellular and extracellular metabolites	MCF derivatization and GC-MS analysis	↓ Intracellular fatty acids, glutathione, amino acids, and TCA intermediates ↓ Extracellular secretion of methionine, citramalic acid, gamma-linolenic acid, and conjugated linoleic acid	KC5, KC10	Placenta trophoblast cells, HTR8/Svneo	300 µM, 24 h	Source and purity not reported, dose induced 50% cytotoxicity, negative and various control groups for oxidative stress and/or hypoxia included ($n = 6$ replicate wells).	Chen et al. (2020)
Intracellular protein expression	2-DGE, silver staining, MALDI-TOF/TOF MS and/or MS/MS	↑ HIF-1α and N-Myc downstream regulated 1 ↓ Proteins associated with cellular metabolism, cell proliferation, and differentiation	KC5, KC10	Acute promonocytic leukaemia cells, U937	50 µM, 24 h	Purity not reported, positive and negative controls included ($n =$ unclear).	Han et al. (2006)
<i>Cobalt(II) sulfate (CoSO₄)</i>							
Hypoxia-responsive gene induction	Evaluation of 36 hypoxia-regulated promoters and long-range transcriptional regulatory elements	↑ HIF-1α activity (both), VEGF secretion (ME-180), glycolysis and gluconeogenesis, nutrient supply, and metals transport (HCT 116)	KC5, KC7, KC10	Human colon carcinoma, HCT 116; cervical carcinoma, HRE-bla ME-180	100 µM, 20 h 31 µM, 20 h	No significant cytotoxicity observed over dose range evaluated, negative and positive controls included ($n = 3$ replicate wells).	Xia et al. (2009)

Table S4.23 (continued)

End-point	Platform	Results ^a	KC associated	Tissue, cell line	Concentration, duration	Comments	Reference
<i>Insoluble cobalt (II or II,III) compounds</i>							
<i>Cobalt(II,III) oxide NPs (Co₃O₄)</i>							
Transcriptomics	Whole human genome 4 × 44 K oligonucleotide array (G4112F)	> 100× DETs in BEAS-2B versus A549 cells ↑ Cellular metabolism and energetics ↓ Immune signalling and antigen presentation	KC8, KC7, KC10	Bronchial cells, BEAS-2B; lung adenocarcinoma, A549	6.09 µg/mL [\sim 80 µM], 3, 6, 10, 24 h 60.9 µg/mL [\sim 800 µM], 3, 6, 10, 24 h	NP size: d ₅₀ , 7 nm. Doses chosen based upon < 30% cytotoxicity by neutral red uptake, positive and negative controls included (<i>n</i> = 3 independent replicates).	Verstraelen et al. (2014)

2-DGE, two-dimensional gel electrophoresis; CEA, carcinoembryonic antigen; d₅₀, mean particle diameter; DETs, differentially expressed transcripts; GC-MS, gas chromatography-mass spectrometry; HIF-1 α , hypoxia-inducible factor-1 alpha; KC, key characteristic of carcinogens; KC4, induces epigenic alterations; KC5, induces oxidative stress; KC6, induces chronic inflammation; KC7, is immunosuppressive; KC8, modulates receptor-mediated effects; KC10, alters cell proliferation, cell death, or nutrient supply; MALDI-TOF, matrix-assisted laser desorption/ionization, time-of-flight mass analyser; MCF, methylchloroformate; MHC, major histocompatibility complex; mRNA, messenger RNA; MS, mass spectrometry; MS/MS, tandem mass spectrometry; nanoLC, nanoscale liquid chromatography; TOF, time-of-flight.

^a ↓, decreased; ↑, increased.

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