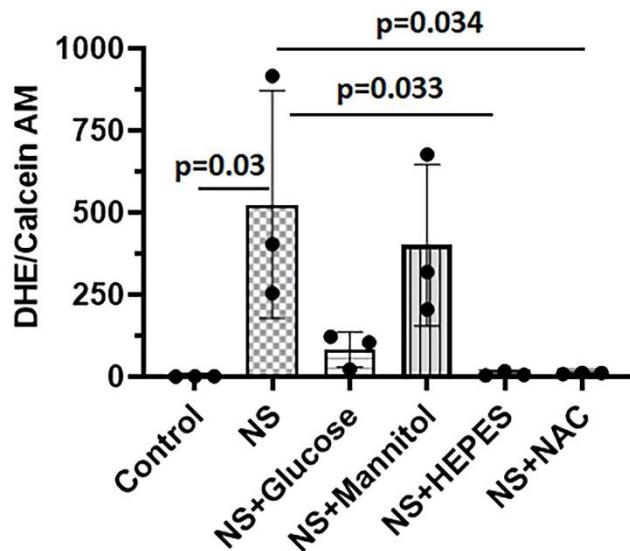


Supplementary Figure S1. Effect of normal saline (NS) and buffers (25 mM HEPES or 0.5 mM sodium bicarbonate) on chondrocyte death using a flow cytometer. (a-c) Representative plots for annexin V (FL1-H; x-axis)/propidium iodide (PI) (FL3-H; y-axis) staining after irrigant incubation for 30 minutes (a), 1 hour (b), and 3 hours (c). Q1: annexin V negative/PI positive, Q2: annexin V positive / PI positive, Q3: annexin V positive / PI negative, Q4: annexin V negative / PI negative. (d-f) Representative FSC vs SSC plots for cell morphology after irrigant incubation for 30 minutes (d), 1 hour (e), and 3 hours (f). FSC-H (forward scatter height; x-axis) and SSC-H (side scatter height; y-axis) represent cell size and granularity, respectively.



Supplementary Figure S2. Effect of 0.9% normal saline (NS) and various supplements on oxidative stress in monolayer primary bovine chondrocytes. Glucose (Sigma-Aldrich): 31.52 mM, Mannitol (Research Products International): 31.52 mM, HEPES: 25 mM, NAC (Sigma-Aldrich): 2.5 mM with 26 mM sodium bicarbonate (Research Products International) ($n = 3$).

Supplementary Table S1. List of metabolites showing \geq 2-fold changes between control and 0.9% normal saline (NS). Chondrocytes were treated with each irrigation solution for 30 minutes (n = 6). Positive values of fold change (up): NS or NS + HEPES > control, negative values of fold change (down): NS or NS + HEPES < control.

	NS versus Control		NS + HEPES versus Control	
Metabolites (up)	Fold change	p-value	Fold change	p-value
Glucose 6-phosphate	70.3	5.9E-09	23.4	1.5E-04
Ribose 5-phosphate	34.3	5.8E-09	6.8	1.2E-02
Fructose 6-phosphate	7.5	5.9E-09	2.8	7.6E-04
O-octanoyl-R-carnitine	5.0	1.1E-07	7.0	6.2E-09
D-Ribulose 5-phosphate	4.2	5.8E-09	2.0	1.6E-05
O-propanoyl-carnitine	4.2	1.9E-08	3.9	6.1E-08
Inosine	3.2	5.8E-09	0.6	4.8E-03
Mannose	3.0	6.3E-08	1.9	6.8E-04
Guanosine	2.9	5.9E-09	1.7	1.2E-04
Adenosine monophosphate (AMP)	2.3	5.8E-09	0.8	9.1E-03
Nicotinamide adenine dinucleotide phosphate (NADP+)	2.2	7.7E-08	1.7	8.1E-05
Deoxyguanosine diphosphate (dGDP)	2.2	5.8E-09	0.6	3.3E-05
O-hexanoyl-R-carnitine	2.1	2.7E-08	2.1	2.3E-08
Metabolites (down)	Fold change	p-value	Fold change	p-value
Reduced nicotinamide adenine dinucleotide phosphate (NADPH)	-31.1	6.6E-09	-0.8	7.2E-04
Alpha-Ketoglutarate (KG)	-22.0	5.8E-09	-15.5	5.8E-09
Alpha-Ketoisovalerate (KIV)	-12.4	6.1E-09	-4.8	8.3E-09
Cytidine triphosphate (CTP)	-11.8	5.8E-09	-1.4	5.7E-06
Alpha-Keto-beta-Methylvalerate (KMV)	-10.2	5.8E-09	-10.2	5.8E-09
Nicotinamide adenine dinucleotide + hydrogen (NADH)	-9.6	1.2E-07	-0.8	7.0E-03
O-oleylcarnitine	-9.1	1.2E-05	-1.0	9.8E-01
Alpha-Ketoisocaproate (KIC)	-9.0	5.8E-09	-6.8	5.8E-09
Lactate	-7.2	5.8E-09	-3.3	5.8E-09
Xanthosine	-5.7	0.020	-0.6	4.4E-02
Pyruvic acid	-5.5	5.8E-09	-2.6	6.4E-09
Cytidine diphosphate (CDP)	-4.8	5.8E-09	-2.2	5.8E-09
Oxidized glutathione (GSSG)	-4.8	7.9E-08	-1.7	1.4E-04
O-Phosphoethanolamine	-4.4	7.1E-09	-3.6	9.1E-09

Dihydroxyacetone phosphate (DHAP)	-4.1	5.8E-09	-2.2	5.8E-09
Beta-Hydroxybutyrate (3-Hydroxybutyrate)	-3.8	5.8E-09	-2.3	5.9E-09
Inosine monophosphate (IMP)	-3.8	1.4E-07	-1.8	8.2E-05
Malate	-3.7	1.7E-08	-7.3	6.9E-09
Citrulline	-3.7	5.8E-09	-3.6	5.8E-09
Deoxythymidine triphosphate (dTTP)	-3.6	7.4E-07	-2.4	9.4E-06
Succinic acid	-3.5	8.1E-09	-4.2	6.7E-09
Guanosine triphosphate (GTP)	-3.2	6.0E-09	-0.9	4.5E-01
Hypoxanthine	-3.1	4.8E-07	-1.0	0.838
Fumarate	-3.0	8.3E-08	-3.7	2.7E-08
Glutaryl carnitine	-3.0	5.9E-09	-1.8	2.8E-08
Pyruvate	-2.9	5.8E-09	-2.0	5.9E-09
Adenosine diphosphate (ADP) ribose	-2.9	6.1E-09	-1.6	6.3E-07
Carnitine	-2.9	5.8E-09	-1.4	1.7E-08
Lysine	-2.8	5.8E-09	-2.3	5.8E-09
Malic acid	-2.8	1.8E-06	-4.0	2.3E-07
Fumaric acid	-2.8	2.9E-06	-4.1	3.4E-07
Uridine triphosphate (UTP)	-2.5	5.8E-09	-1.0	0.506
Deoxyadenosine triphosphate (dATP)	-2.4	5.8E-09	-1.6	2.2E-08
Citric acid	-2.4	5.7E-04	-2.5	4.8E-04
Deoxyguanosine triphosphate (dGTP)	-2.1	6.2E-09	-0.9	3.6E-04