

Supplementary Materials: An Inflammatory Nucleus Pulpous Tissue Culture Model to Test Molecular Regenerative Therapies: Validation with Epigallocatechin 3-Gallate

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Table S1. Average cycle threshold values ($C_q \pm SD$) for all expressed genes in inflammation group. GAPDH, glyceraldehyde-3-phosphate dehydrogenase; MMP, matrix metalloproteinase; IL, interleukin; EGCG, epigallocatechin 3-gallate; TIMP1, tissue inhibitor of matrix metalloproteinases; PTGS2, prostaglandin-endoperoxide synthase 2; ADAMTS, a disintegrin and metalloproteinase with thrombospondin motifs; iNOS, inducible nitric oxide synthase.

Condition	Gene	Mean $C_q \pm SD$	Condition	Gene	Mean $C_q \pm SD$
Control	<i>S18</i>	27.49 ± 5.12	IL-1 β	<i>S18</i>	27.61 ± 5.19
	<i>GAPDH</i>	27.74 ± 0.95		<i>GAPDH</i>	27.83 ± 1.17
	<i>Collagen 1</i>	43.46 ± 3.43		<i>Collagen 1</i>	43.38 ± 3.43
	<i>Collagen 2</i>	34.13 ± 3.29		<i>Collagen 2</i>	38.01 ± 3.55
	<i>Aggrecan</i>	36.14 ± 3.53		<i>Aggrecan</i>	35.59 ± 4.35
	<i>MMP1</i>	41.37 ± 8.87		<i>MMP1</i>	32.51 ± 12.11
	<i>MMP3</i>	43.61 ± 2.91		<i>MMP3</i>	32.47 ± 8.21
	<i>MMP13</i>	37.50 ± 10.18		<i>MMP13</i>	33.92 ± 5.44
	<i>TIMP1</i>	35.71 ± 9.91		<i>TIMP1</i>	31.41 ± 2.94
	<i>IL-6</i>	45 ± 0		<i>IL-6</i>	36.35 ± 4.49
	<i>IL-8</i>	45 ± 0		<i>IL-8</i>	38.76 ± 4.12
	<i>PTGS2</i>	43.13 ± 3.23		<i>PTGS2</i>	37.21 ± 2.21
	<i>ADAMTS4</i>	45 ± 0		<i>ADAMTS4</i>	43.57 ± 2.85
	<i>iNOS</i>	45 ± 0		<i>iNOS</i>	35.31 ± 1.54
EGCG	<i>S18</i>	27.69 ± 4.49	IL-1 β + EGCG	<i>S18</i>	27.61 ± 5.25
	<i>GAPDH</i>	27.91 ± 0.45		<i>GAPDH</i>	27.84 ± 1.14
	<i>Collagen 1</i>	45 ± 0		<i>Collagen 1</i>	45 ± 0
	<i>Collagen 2</i>	36.62 ± 5.01		<i>Collagen 2</i>	37.86 ± 4.09
	<i>Aggrecan</i>	36.31 ± 4.43		<i>Aggrecan</i>	37.01 ± 3.71
	<i>MMP1</i>	39.28 ± 11.15		<i>MMP1</i>	37.95 ± 8.34
	<i>MMP3</i>	40.79 ± 5.05		<i>MMP3</i>	35.82 ± 7.13
	<i>MMP13</i>	37.84 ± 8.86		<i>MMP13</i>	35.23 ± 8.58
	<i>TIMP1</i>	36.31 ± 7.25		<i>TIMP1</i>	32.83 ± 5.05
	<i>IL-6</i>	41.62 ± 4.65		<i>IL-6</i>	38.06 ± 3.95
	<i>IL-8</i>	44.41 ± 1.45		<i>IL-8</i>	42.25 ± 4.35
	<i>PTGS2</i>	41.67 ± 3.83		<i>PTGS2</i>	38.94 ± 2.61
	<i>ADAMTS4</i>	42.76 ± 4.46		<i>ADAMTS4</i>	43.33 ± 3.32
	<i>iNOS</i>	42.77 ± 4.44		<i>iNOS</i>	38.31 ± 4.57