Supplementary Table S1. (A) p-Values for Comparisons of Bone Marrow-Derived Mesenchymal Stem Cell- Versus Adipose-Derived Mesenchymal Stem Cell-Extracellular Matrix Binding in Extracellular Matrix Array (Bonferroni Adjusted α =0.00625). p-Values Comparing the Binding of (B) Bone Marrow-Derived Mesenchymal Stem Cells and (C) Adipose-Derived Mesenchymal Stem Cells on Specific Extracellular Matrix Proteins (Bonferroni Adjusted α =0.0018).

	ECM Protein	BMSC vs. ASC (p-valu ANOVA	Post-hoc (α = 0.00625)
	Collagen I		0.1165
	Collagen II		0.0243
	Collagen IV		0.0229
	Fibronectin	<0.0001	0.0077
	Laminin	10.0001	0.0970
	Tenascin		0.6992
	Vitronectin		0.2074
	BSA		0.4191

Α

B Post-hoc p-values (α = 0.0018) for BMSC binding between ECM groups [ANOVA (all groups): p < 0.0001 including BSA control, p = 0.0007 excluding control]

1	Collagen I	Collagen II	Collagen IV	Fibronectin	Laminin	Tenascin	Vitronectin	BSA
Collagen I	_	0.2860	0.5367	0.6663	0.0362	0.0406	0.0068	0.0001
Collagen II	0.2860	_	0.6137	0.4968	0.0243	0.0443	0.0004	<0.0001
Collagen IV	0.5367	0.6137	_	0.8473	0.0445	0.0537	0.0048	<0.0001
Fibronectin	0.6663	0.4968	0.8473	_	0.0448	0.0517	0.0061	<0.0001
Laminin	0.0362	0.0243	0.0445	0.0448	_	0.7375	0.0472	<0.0001
Tenascin	0.0406	0.0443	0.0537	0.0517	0.7375	10 21	0.2094	0.0001
Vitronectin	0.0068	0.0004	0.0048	0.0061	0.0472	0.2094	_	<0.0001
BSA	0.0001	<0.0001	<0.0001	<0.0001	<0.0001	0.0001	<0.0001	

C Post-hoc p-values (α = 0.0018) for ASC binding between ECM groups [ANOVA (all groups): p < 0.0001 including BSA control, p = 0.2144 excluding control]

	Collagen I	Collagen II	Collagen IV	Fibronectin	Laminin	Tenascin	Vitronectin	BSA
Collagen I		0.3372	0.1850	0.0549	0.0714	0.2333	0.1509	0.0001
Collagen II	0.3372	_	0.5860	0.1443	0.2194	0.6188	0.5071	<0.0001
Collagen IV	0.1850	0.5860	_	0.3318	0.5412	0.9466	0.9438	<0.0001
Fibronectin	0.0549	0.1443	0.3318	_	0.3745	0.5198	0.2918	<0.0001
Laminin	0.0714	0.2194	0.5412	0.3745	i i	0.7326	0.5130	<0.0001
Tenascin	0.2333	0.6188	0.9466	0.5198	0.7326		0.9855	0.0002
Vitronectin	0.1509	0.5071	0.9438	0.2918	0.5130	0.9855	_	<0.0001
BSA	0.0001	<0.0001	<0.0001	<0.0001	<0.0001	0.0002	<0.0001	_

^{*}Statistically significant values after Bonferroni-corrected post-hoc analyses are highlighted in **bold**.