

Supplementary Table S11. Cellular component gene ontology terms for KC-altered genes responsive to TGFβ1 treatment (FDR ≤ 0.05).

	Homo sapiens (REF)	Client Text Box Input (▼ Hierarchy NEW! ?)					
GO cellular component complete	#	#	expected	Fold Enrichment	+/-	raw P value	FDR
basement membrane	101	9	.97	9.31	+	1.05E-06	3.05E-04
↳ collagen-containing extracellular matrix	434	24	4.15	5.78	+	1.23E-11	8.37E-09
↳ extracellular matrix	575	31	5.50	5.63	+	1.75E-14	3.58E-11
↳ external encapsulating structure	576	31	5.51	5.62	+	1.83E-14	1.87E-11
↳ cell periphery	6460	102	61.81	1.65	+	4.36E-09	1.78E-06
anchored component of membrane	171	9	1.64	5.50	+	5.69E-05	1.45E-02
extracellular space	3430	65	32.82	1.98	+	2.52E-08	8.57E-06
↳ extracellular region	4396	82	42.06	1.95	+	2.38E-10	1.21E-07