

Additional Data 2: Kruskal-Wallis analysis among ancestral groups. For each ancestral group the synthesis of the *TTR* expression scores, in terms of median, minimum and maximum, for each tissue involved in TTR amyloidosis is reported.

	AFRICA			EUROPE			EAST_ASIA			SOUTH_ASIA			AMERICA			Kruskal-Wallis p-value
	<i>min</i>	<i>median</i>	<i>max</i>	<i>min</i>	<i>median</i>	<i>max</i>	<i>min</i>	<i>median</i>	<i>max</i>	<i>min</i>	<i>median</i>	<i>max</i>	<i>min</i>	<i>median</i>	<i>max</i>	
Colon-Transverse	-0.22	-0.11	0.30	-0.28	-0.11	0.15	-0.22	-0.06	0.00	-0.22	-0.06	0.04	-0.22	-0.06	0.15	5.448E-11*
Colon-Sigmoid	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.00	0.52	1.8339E-17*
Esophagus-Muscularis	0.00	0.00	0.23	-0.32	0.00	0.23	0.00	0.00	0.00	0.00	0.00	0.46	-0.32	0.00	0.23	0.4912
Esophagus-Mucosa	0.00	0.10	0.29	0.00	0.00	0.38	0.00	0.10	0.38	0.00	0.00	0.38	0.00	0.00	0.38	1.3638E-58*
Heart - Atrial Appendage	-0.60	0.00	0.88	0.00	0.00	0.05	-0.45	0.00	0.02	-0.60	0.00	0.05	-0.60	0.00	0.88	4.5521E-28*
Heart - Left Ventricle	-0.86	0.00	0.95	0.00	0.00	0.10	-0.43	0.00	0.06	-0.68	0.00	0.13	-0.60	0.00	0.86	6.5427E-35*
Liver	-0.83	0.00	0.11	-0.09	0.00	0.23	-0.58	0.00	0.00	-0.83	0.00	0.00	-0.83	0.00	0.11	7.0173E-66*
Muscle - Skeletal	-0.09	-0.09	0.32	-0.09	-0.04	0.17	-0.09	-0.04	0.32	-0.09	0.00	0.33	-0.09	-0.04	0.32	5.0174E-105*
Nerve-Tibial	-0.16	-0.08	0.00	-0.16	-0.08	0.32	-0.16	-0.08	0.00	-0.16	-0.08	0.32	-0.16	-0.08	0.00	2.8912E-09*
Stomach	-0.31	0.00	0.00	-0.24	0.00	0.00	-0.08	0.00	0.00	-0.16	0.00	0.00	-0.31	0.00	0.00	1.4815E-100*
Small Intestine - Terminal Ileum	-0.42	-0.21	0.31	-0.21	0.00	0.63	0.00	0.00	0.31	-0.21	0.00	0.63	-0.42	0.00	0.31	2.0985E-203*
Adipose - Subcutaneous	-0.94	0.00	0.00	-0.50	0.00	0.00	-0.94	0.00	0.00	-0.50	0.00	0.00	-0.50	0.00	0.00	6.2184E-11*
Cells - Transformed fibroblasts	-0.16	0.00	0.20	-0.10	0.00	0.29	-0.20	0.00	0.20	-0.10	0.00	0.20	-0.07	0.04	0.24	1.8783E-25*
Skin - Sun Exposed (Lower leg)	-0.09	0.00	0.57	-0.09	0.00	0.21	-0.07	0.00	0.21	-0.07	0.00	0.10	-0.07	0.00	0.51	1.0020E-86*
N samples	661			503			504			489			347			

* $p < 0.0035$, according to Bonferroni adjustment