

Additional Data 6: Kruskal-Wallis analysis within the American ancestral group. For each population the synthesis of the *TTR* expression scores, in terms of median, minimum and maximum, for each tissue involved in TTR amyloidosis is reported. Detailed information about population definitions is available at <http://www.1000genomes.org/about>.

TISSUE	CLM			MXL			PEL			PUR			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>	
Colon-Transverse	-0.27	0.00	0.49	-0.09	0.00	0.27	-0.09	0.09	0.36	-0.09	0.00	0.36	3.1971E-07*
Colon-Sigmoid	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.26	0.52	0.00	0.00	0.52	0.0002*
Esophagus-Muscularis	-0.13	0.00	0.15	-0.34	0.00	0.15	-0.34	0.00	0.00	-0.34	0.00	0.00	0.1224
Esophagus-Mucosa	0.00	0.00	0.41	0.00	0.00	0.41	0.00	0.00	0.41	0.00	0.00	0.41	0.0531
Heart - Atrial Appendage	-0.20	0.00	0.37	-0.20	0.00	0.19	-0.40	0.00	0.19	-0.20	0.00	0.73	0.0054
Heart - Left Ventricle	-0.20	0.00	0.35	-0.20	0.00	0.20	-0.41	0.00	0.20	-0.30	0.00	0.70	0.0511
Liver	-0.20	0.00	0.00	-0.20	0.00	0.00	-0.35	0.00	0.27	-0.20	0.00	0.00	0.0427
Muscle - Skeletal	-0.09	-0.04	0.17	-0.09	-0.04	0.16	-0.09	-0.04	0.32	-0.09	-0.04	0.17	0.0009*
Nerve-Tibial	-0.16	-0.08	0.00	-0.16	-0.08	0.00	-0.16	-0.08	0.00	-0.16	-0.08	0.00	0.2607
Stomach	-0.16	0.00	0.00	-0.16	0.00	0.00	-0.16	0.00	0.00	-0.31	0.00	0.00	0.0204
Small Intestine - Terminal Ileum	-0.21	0.00	0.31	-0.21	0.00	0.31	-0.21	0.00	0.00	-0.42	0.00	0.31	0.1332
Adipose - Subcutaneous	-0.50	0.00	0.00	-0.50	0.00	0.00	0.00	0.00	0.00	-0.50	0.00	0.00	0.4438
Cells - Transformed fibroblasts	-0.06	0.03	0.21	-0.06	0.00	0.21	0.00	0.03	0.17	-0.06	0.00	0.17	0.0969
Skin - Sun Exposed (Lower leg)	-0.08	0.00	0.23	-0.08	0.00	0.24	-0.04	0.00	0.24	-0.08	0.00	0.33	0.0006*
N samples	94			64			85			104			

* $p < 0.0035$ according to Bonferroni adjustment