

Additional Data 7: Kruscal-Wallis analysis within the African 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	ASW			ACB			ESN			GWD			LWK			MSL			YRI			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>	<i>min</i>	<i>median</i>	<i>max</i>	<i>min</i>	<i>median</i>	<i>max</i>	
Colon-Transverse	-0.25	-0.06	0.07	-0.25	-0.07	0.10	-0.25	-0.08	0.09	-0.23	-0.05	0.10	-0.25	-0.06	0.09	-0.25	-0.07	0.10	-0.23	-0.08	0.05	0.0096
Colon-Sigmoid	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.26	0.52	0.00	0.26	0.52	0.00	0.00	0.52	0.00	0.00	0.52	0.0822
Esophagus-Muscularis	0.00	0.00	0.00	-0.13	0.00	0.15	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.4331
Esophagus-Mucosa	0.00	0.14	0.27	0.00	0.14	0.31	0.00	0.14	0.27	0.00	0.14	0.31	0.00	0.14	0.27	0.00	0.14	0.27	0.00	0.14	0.27	0.0423
Heart - Atrial Appendage	-0.44	0.00	0.35	-0.59	0.00	0.35	-0.59	0.00	0.55	-0.59	0.00	0.55	-0.59	0.00	0.55	-0.59	0.00	0.55	-0.59	0.00	0.35	0.2389
Heart - Left Ventricle	-0.20	0.07	1.00	-0.20	0.07	0.73	-0.40	0.07	0.78	-0.20	0.07	0.80	-0.20	0.08	0.80	-0.34	0.07	0.73	-0.20	0.08	0.80	0.5295
Liver	-0.43	0.00	0.00	-0.56	0.00	0.20	-0.56	0.00	0.00	-0.56	0.00	0.00	-0.56	0.00	0.00	-0.56	0.00	0.00	-0.56	0.00	0.00	0.0356
Muscle - Skeletal	-0.09	-0.09	0.16	-0.09	-0.09	0.32	-0.09	-0.09	0.32	-0.09	-0.09	0.32	-0.09	-0.09	0.32	-0.09	-0.09	0.12	-0.09	-0.09	0.32	0.0337
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	-0.16	-0.08	0.00	-0.16	-0.08	0.00	-0.16	-0.08	0.00	0.4619
Stomach	-0.24	0.00	0.00	-0.31	-0.08	0.00	-0.31	-0.08	0.00	-0.31	0.00	0.00	-0.31	0.00	0.00	-0.31	-0.16	0.00	-0.31	-0.08	0.00	0.0111
Small Intestine - Terminal Ileum	-0.42	0.00	0.31	-0.42	-0.11	0.31	-0.42	-0.21	0.00	-0.42	0.00	0.00	-0.42	-0.21	0.00	-0.42	-0.21	0.00	-0.42	-0.21	0.00	0.0432
Adipose - Subcutaneous	-0.94	0.00	0.00	-0.50	0.00	0.00	-0.47	0.00	0.00	-0.47	0.00	0.00	-0.94	0.00	0.00	-0.47	0.00	0.00	-0.47	0.00	0.00	0.0852
Cells - Transformed fibroblasts	-0.17	0.00	0.14	-0.12	0.00	0.14	-0.12	0.00	0.14	-0.12	0.00	0.14	-0.24	0.00	0.14	-0.12	0.00	0.07	-0.12	0.00	0.12	0.2337
Skin - Sun Exposed (Lower leg)	-0.11	0.00	0.35	-0.04	0.00	0.23	0.00	0.00	0.45	-0.04	0.00	0.45	-0.04	0.00	0.45	-0.04	0.00	0.39	0.00	0.00	0.28	0.1661
N samples	61			96			99			113			99			85			108			

** $p < 0.0035$, according to Bonferroni adjustment*