

Additional Data 4: Kruskal-Wallis analysis within the European 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	CEU			FIN			GBR			IBS			TSI			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	-0.14	0.00	0.13	-0.05	0.00	0.07	-0.16	-0.02	0.13	-0.05	-0.02	0.07	-0.16	-0.02	0.56	0.0024*
Colon-Sigmoid	0.00	0.00	0.52	0.00	0.00	0.26	0.00	0.00	0.52	0.00	0.00	0.52	0.00	0.00	0.52	0.2923
Esophagus-Muscularis	-0.34	0.00	0.15	0.00	0.00	0.15	-0.34	0.00	0.15	-0.34	0.00	0.15	-0.34	0.00	0.00	0.5622
Esophagus-Mucosa	0.00	0.00	0.38	0.00	0.00	0.38	0.00	0.00	0.29	0.00	0.00	0.38	0.00	0.00	0.38	0.3718
Heart - Atrial Appendage	0.00	0.00	0.02	0.00	0.00	0.02	0.00	0.00	0.04	0.00	0.00	0.04	-0.04	0.00	0.04	0.0055
Heart - Left Ventricle	0.00	0.00	0.07	0.00	0.00	0.07	0.00	0.00	0.07	0.00	0.00	0.07	-0.06	0.00	0.07	0.5261
Liver	0.00	0.00	0.11	0.00	0.00	0.11	0.00	0.00	0.23	-0.09	0.00	0.11	-0.09	0.00	0.11	0.2178
Muscle - Skeletal	-0.13	-0.07	0.00	-0.13	-0.07	0.00	-0.13	-0.07	0.00	-0.13	-0.07	0.00	-0.13	-0.07	0.00	6.7949E-06*
Nerve-Tibial	-0.16	-0.08	0.32	-0.16	-0.08	0.32	-0.16	-0.08	0.24	-0.16	-0.08	0.32	-0.16	-0.08	0.00	0.0359
Stomach	-0.08	0.00	0.00	-0.08	0.00	0.00	-0.16	0.00	0.00	-0.24	0.00	0.00	-0.16	0.00	0.00	0.0039
Small Intestine - Terminal Ileum	0.00	0.00	0.63	0.00	0.00	0.63	0.00	0.00	0.63	-0.21	0.00	0.63	0.00	0.00	0.31	0.9489
Adipose - Subcutaneous	0.00	0.00	0.00	0.00	0.00	0.00	-0.47	0.00	0.00	-0.50	0.00	0.00	-0.50	0.00	0.00	0.7008
Cells - Transformed fibroblasts	-0.07	0.00	0.20	-0.07	0.00	0.14	-0.10	0.00	0.29	-0.07	0.00	0.24	-0.07	0.00	0.20	0.0251
Skin - Sun Exposed (Lower leg)	-0.08	0.00	0.28	-0.06	0.00	0.14	-0.08	0.00	0.28	-0.06	0.00	0.22	-0.06	0.00	0.28	0.4988
N samples	99			99			91			107			107			

* $p < 0.0035$, according to Bonferroni adjustment