

Additional Data 5: Kruskal-Wallis analysis within the Eastern Asian 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	CDX			CHB			JPT			KHV			CHS			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	-0.03	0.00	0.10	-0.03	0.00	0.05	-0.03	0.00	0.05	-0.03	0.00	0.10	-0.03	0.00	0.10	0.6429
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	0.0388
Esophagus-Muscularis	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	-0.13	0.00	0.00	0.00	0.00	0.00	0.3938
Esophagus-Mucosa	0.00	0.13	0.41	0.00	0.13	0.34	0.00	0.00	0.34	0.00	0.13	0.41	0.00	0.13	0.34	0.0042
Heart - Atrial Appendage	-0.09	0.00	0.00	-0.09	0.00	0.04	-0.09	0.00	0.00	-0.09	0.00	0.04	-0.09	0.00	0.04	0.0555
Heart - Left Ventricle	-0.15	0.00	0.10	-0.10	0.00	0.05	-0.10	0.00	0.05	-0.10	0.00	0.10	-0.15	0.00	0.05	0.4324
Liver	-0.17	0.00	0.00	-0.17	0.00	0.00	-0.17	0.00	0.00	-0.35	0.00	0.00	-0.17	0.00	0.00	0.0308
Muscle - Skeletal	-0.09	-0.04	0.16	-0.09	-0.04	0.17	-0.09	-0.04	0.16	-0.09	-0.04	0.32	-0.09	-0.04	0.16	0.1752
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	0.00007*
Stomach	0.00	0.00	0.00	-0.08	0.00	0.00	0.00	0.00	0.00	-0.08	0.00	0.00	-0.08	0.00	0.00	0.7475
Small Intestine - Terminal Ileum	0.00	0.00	0.00	0.00	0.00	0.31	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.4206
Adipose - Subcutaneous	-0.47	0.00	0.00	-0.94	0.00	0.00	-0.47	0.00	0.00	-0.47	0.00	0.00	-0.47	0.00	0.00	0.5963
Cells - Transformed fibroblasts	-0.10	0.00	0.14	-0.20	0.00	0.10	-0.10	0.00	0.14	-0.10	0.00	0.20	-0.10	0.00	0.14	0.0173
Skin - Sun Exposed (Lower leg)	-0.07	0.00	0.21	-0.07	0.00	0.10	-0.07	0.00	0.10	-0.07	0.00	0.21	-0.07	0.00	0.10	0.9463
N samples	93			103			104			99			105			

* $p < 0.0035$ according to Bonferroni adjustment