

Additional Data 8: Kruscal-Wallis analysis within the Southern 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	BEB			GIH			ITU			PJI			STU			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	-0.06	0.00	0.06	-0.06	0.00	0.08	-0.06	0.00	0.08	-0.06	0.00	0.08	-0.06	0.00	0.06	0.0597
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	0.1705
Esophagus-Muscularis	0.00	0.00	0.15	-0.13	0.00	0.15	0.00	0.00	0.31	0.00	0.00	0.00	0.00	0.00	0.00	0.5843
Esophagus-Mucosa	0.00	0.00	0.34	0.00	0.00	0.34	0.00	0.00	0.34	0.00	0.00	0.26	0.00	0.00	0.34	0.1124
Heart - Atrial Appendage	-0.17	0.00	0.08	-0.17	0.00	0.04	-0.17	0.00	0.04	-0.09	0.00	0.08	-0.17	0.00	0.08	0.4693
Heart - Left Ventricle	-0.34	0.00	0.20	-0.34	0.00	0.00	-0.34	0.00	0.00	-0.17	0.00	0.00	-0.34	0.00	0.20	0.6019
Liver	-0.38	0.00	0.00	-0.38	0.00	0.00	-0.38	0.00	0.00	-0.20	0.00	0.00	-0.35	0.00	0.00	0.2695
Muscle - Skeletal	-0.09	0.00	0.32	-0.09	0.00	0.33	-0.09	0.00	0.32	-0.09	0.00	0.33	-0.09	0.00	0.33	0.0778
Nerve-Tibial	-0.16	-0.08	0.24	-0.16	-0.08	0.00	-0.16	-0.08	0.32	-0.16	-0.08	0.00	-0.16	-0.08	0.00	0.1365
Stomach	-0.16	0.00	0.00	-0.08	0.00	0.00	-0.08	0.00	0.00	-0.16	0.00	0.00	-0.16	0.00	0.00	0.1479
Small Intestine - Terminal Ileum	-0.21	0.00	0.63	-0.21	0.00	0.31	-0.21	0.00	0.63	-0.21	0.00	0.31	-0.21	0.00	0.31	0.4384
Adipose - Subcutaneous	-0.50	0.00	0.00	-0.50	0.00	0.00	-0.50	0.00	0.00	-0.50	0.00	0.00	-0.47	0.00	0.00	0.7584
Cells - Transformed fibroblasts	-0.08	0.00	0.14	-0.12	0.00	0.14	0.00	0.00	0.14	-0.08	0.00	0.17	-0.12	0.00	0.17	0.1595
Skin - Sun Exposed (Lower leg)	-0.03	0.00	0.10	-0.07	0.00	0.10	-0.03	0.00	0.10	-0.07	0.00	0.10	-0.03	0.00	0.10	0.0151
N samples	86			103			102			96			102			

* $p < 0.0035$, according to Bonferroni adjustment