

Supplementary Table 4 – Descriptive statistics of admixture proportions of NTR participants

	AFFY6						AXIOM						ILLGSA						HARMONIZED					
	Mean	SD	Min	Max	Range		Mean	SD	Min	Max	Range		Mean	SD	Min	Max	Range		Mean	SD	Min	Max	Range	
Q1	0.1869	0.0587	0.0000	0.7468	0.7468		0.1908	0.0663	0.0039	0.7405	0.7366		0.1948	0.0596	0.0000	0.7194	0.7194		0.1878	0.0658	0.0000	0.7608	0.7608	
Q2	0.0112	0.0527	0.0000	0.8731	0.8731		0.0114	0.0538	0.0000	0.8164	0.8164		0.0106	0.0505	0.0000	0.8937	0.8937		0.0113	0.0511	0.0000	0.8807	0.8807	
Q3	0.0725	0.0220	0.0000	0.4332	0.4332		0.0726	0.0212	0.0000	0.3204	0.3204		0.0756	0.0206	0.0000	0.6184	0.6184		0.0702	0.0312	0.0000	0.6213	0.6213	
Q4	0.6949	0.0919	0.0000	0.9314	0.9314		0.6868	0.1041	0.0000	0.8588	0.8588		0.6833	0.0954	0.0000	0.9218	0.9218		0.6936	0.1024	0.0000	0.9575	0.9575	
Q5	0.0054	0.0100	0.0000	0.4484	0.4484		0.0054	0.0138	0.0000	0.5075	0.5074		0.0055	0.0117	0.0000	0.5525	0.5525		0.0060	0.0126	0.0000	0.5576	0.5575	
Q6	0.0036	0.0154	0.0000	0.4681	0.4681		0.0049	0.0216	0.0000	0.4586	0.4585		0.0041	0.0186	0.0000	0.5581	0.5581		0.0043	0.0185	0.0000	0.5584	0.5584	
Q7	0.0171	0.0419	0.0000	0.8840	0.8840		0.0189	0.0530	0.0000	0.8491	0.8491		0.0174	0.0400	0.0000	0.9001	0.9001		0.0173	0.0442	0.0000	0.9330	0.9330	
Q8	0.0039	0.0167	0.0000	0.5172	0.5171		0.0049	0.0229	0.0000	0.5952	0.5952		0.0043	0.0197	0.0000	0.6412	0.6412		0.0044	0.0199	0.0000	0.6672	0.6672	
Q9	0.0045	0.0144	0.0000	0.5465	0.5465		0.0042	0.0123	0.0000	0.3438	0.3438		0.0043	0.0205	0.0000	0.7515	0.7515		0.0050	0.0183	0.0000	0.7733	0.7733	

Q1-Q9 represent each of the nine ancestry populations as determined by ADMIXTURE, SD=standard deviation, Min=minimum, Max=maximum