

Table S2 | Frequency of the T allele of rs3741845 in different populations

Population	Frequency	Source
AFDS	0.11	This paper
JPT [‡]	0.24	International HapMap Consortium [1]
Japanese	0.25	Kim <i>et al.</i> , 2005 [2]
CEU [†]	0.35	International HapMap Consortium [1]
Hungarian	0.56	Kim <i>et al.</i> , 2005 [2]
African American	0.65	Entrez SNP [3]
Pygmy	0.70	Kim <i>et al.</i> , 2005 [2]
YRI [§]	0.77	International HapMap Consortium [1]
Amerindian	0.80	Kim <i>et al.</i> , 2005 [2]
Cameroonian	0.86	Kim <i>et al.</i> , 2005 [2]

[†], Utah residents of European Ancestry

[‡], Japanese in Tokyo, Japan

[§], Yoruba in Ibadan, Nigeria

1. The International HapMap Consortium (2005) A haplotype map of the human genome. *Nature* 437: 1299-1320.
2. Kim U, Wooding S, Ricci D, Jorde LB, Drayna D (2005) Worldwide haplotype diversity and coding sequence variation at human bitter taste receptor loci. *Hum Mutat* 26: 199-204.
3. Sherry ST, Ward MH, Kholodov M, Baker J, Phan L, et al. (2001) dbSNP: the NCBI database of genetic variation. *Nucleic Acids Res* 29: 308-311.