

Huntington disease mutation in Venezuela: age of onset, haplotype analyses and geographic aggregation

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In Table 2, a column heading was inadvertently omitted.
Corrected table is as follows:

Table 2 Complete haplotypes in carriers according to geographic origin

Haplotypes in mutation carrier chromosomes SNPs;VNTR;CCG; Δ 2642	Geographical origin	Frequency (%)	Number of chromosomes
1;G;C;7;(A)	Zulia, Táchira, Lara, Miranda, Apure, Guárico, Anzoátegui, Falcón, Trujillo, Mérida, Sucre, Carabobo, Nueva Esparta, Vargas, Colombia, Perú, Spain, Portugal	95.6	43
1;G;C;7;(B)	Perú (Piura)	2.2	1
4;G;C;7;(A)	France	2.2	1
Haplotypes in normal patient chromosomes SNPs;VNTR;CCG; Δ 2642	Geographical origin	Frequency (%)	Number of chromosomes
1;G;C;7;(A)	Zulia, Táchira, Lara, Miranda, Apure, Guárico, Anzoátegui, Falcón, Trujillo, Mérida, Sucre, Carabobo, Nueva Esparta, Vargas, Colombia, Spain, Portugal, France	62.2	28
1;G;C;7;(B)	Perú (Piura)	2.2	1
1;G;T;7;(A)	Lara, Falcón	4.4	2
1;A;C;7;(A)	Sucre	2.2	1
2;A;C;7;(A)	Trujillo	2.2	1
1;G;C;10;(A)	Apure, Falcón, Vargas	6.7	3
2;G;C;10;(A)	Miranda	2.2	1
4;G;C;7;(A)	Lara, Táchira, Miranda, Carabobo, Trujillo, Spain	17.8	8

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