Mobile DNA

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Correction to: Upregulation of selected HERVW loci in multiple sclerosis

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   Following the publication of the original article [1] the authors noticed the published Table 1 has a distorted headings and columns not in proper order.
   The correct Table 1 is shown below. The original article [1] has been updated.

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Reference
1.   Macías-Redondo S, Strunk M, Cebollada-Solanas A, et al. Upregulation of selected HERVW loci in multiple sclerosis. Mob DNA. 2021;12:18. https://doi.org/10.1186/s13100-021-00243-1.

Table 1 Clinical features of MS patients included in this study

Case	MS subtype	Age	Sex	Status	RRMS designation
RSM-49	CIS	25	F	NA	NO
RSM-58	CIS	44	F	A	YES
RSM-70	CIS	47	F	A	YES
RSM-73	CIS	39	M	NA	YES
RMS-103	CIS	53	F	NA	YES
RMS-114	CIS	44	M	NA	NO

No MS Controls	Age	Sex
RMS-3	36	F
RMS-15	31	M
RMS-23	46	F
RMS-30	46	M
RSM-42	39	M
RSM-46	44	M
RSM-62	29	F
RSM-72	46	M
RSM-80	48	M
RSM-98	47	M
RSM-99*	17	M
RSM-102*	56	F
RSM-104	44	M
RMS-106*	59	M
RMS-109	45	M
RMS-112	47	F
RMS-113	28	M
RMS-118	30	F

Clinical data of patients whose PBMCs were analyzed for HERVW expression. Median ages for both patients and controles groups were 44 years (mean and SEM are 42,0 ± 4,25 and 40,4 ± 1,94 for patient and control groups, respectively)

A/NA status refers to active and non-active patients respectively. Posterior progression towards MS diagnosis (RRMS) is indicated for all CIS cases. Samples analyzed by NGS are marked in blue. median ages in these groups are 44 year for patients and 47 years for controls. * indicates samples only analyzed by NGS