

CORRECTION

Correction: RNA Sequence Analysis of Human Huntington Disease Brain Reveals an Extensive Increase in Inflammatory and Developmental Gene Expression

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Table 1 and Table 2 appear incorrectly in the published article. Please see the correct Tables [1](#) and [2](#) and their captions here.



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Table 1. HD sample statistics.

Sample ID	PMI	Age of Death	RIN	mRNA-Seq reads	Age of Onset	Duration	CAG	Vonsattel Grade	H-V Striatal Score	H-V Cortical Score
H_0001	37.25	55	7.1	74,635,390	44	11	45	3	2.661	0.922
H_0002	5.75	69	7.5	71,015,288	63	6	41	3	2.644	1.081
H_0003	20.5	71	7.0	77,385,918	52	19	43	3	2.428	1.707
H_0005	19.15	48	6.9	82,366,794	25	23	48	4	3.820	1.939
H_0006	NA	40	6.2	77,123,676	34	6	51	4	3.522	1.431
H_0007	8	72	8.5	63,294,390	55	17	41	3	2.593	0.849
H_0008	21.3	43	7.4	71,056,116	28	15	49	3	2.701	1.701
H_0009	3.73	68	7.8	66,169,262	45	23	42	3	2.668	1.701
H_0010	6.16	59	8.3	65,341,820	35	24	46	3	2.621	1.200
H_0012	12.75	68	6.0	83,110,358	52	16	42	3	2.661	1.077
H_0013	25.1	57	6.1	71,320,688	40	17	49	3	2.911	1.491
H_0539	14.5	54	6.5	124,222,130	42	12	45	3	2.132	0.401
H_0657	24.3	61	8.1	136,764,622	36	25	45	4	3.290	1.604
H_0658	11	48	7.8	85,591,704	42	6	44	3	2.410	0.978
H_0681	19.06	69	7.0	78,493,314	50	19	42	3	2.484	1.088
H_0695	16.15	55	7.9	86,412,654	36	19	45	4	3.581	2.062
H_0700	15.66	50	8.0	78,329,378	33	17	47	3	2.741	1.202
H_0726	14.75	50	9.2	86,025,890	27	23	48	4	3.598	1.201
H_0740	13.58	75	6.4	101,997,434	60	15	42	3	2.621	2.361
H_0750	16.16	53	6.0	122,909,122	38	15	48	4	3.260	1.010

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Table 2. Control sample statistics.

Sample ID	PMI	Age of Death	RIN	mRNA-Seq reads
C_0012	19	66	7.1	118,327,116
C_0013	15	69	7.8	89,478,160
C_0014	21	79	8.0	65,377,604
C_0015	10	61	8.2	123,746,070
C_0016	20	58	8.4	67,758,208
C_0017	21	70	8.2	72,238,818
C_0018	17	66	8.5	64,688,322
C_0019	19	73	7.5	108,683,922
C_0020	24	60	7.9	83,696,384
C_0021	26	76	7.3	79,487,172
C_0022	17	61	7.8	73,133,936
C_0023	18	62	6.6	94,493,436
C_0024	26	69	8.7	62,989,822
C_0025	25	61	8.1	55,810,684
C_0026	11	88	7.1	72,581,752
C_0029	13	93	6.4	59,386,108
C_0031	24	53	7.3	73,283,170
C_0032	24	57	8.3	70,994,352
C_0033	15	43	7.5	69,505,712
C_0035	21	46	7.6	62,300,754
C_0036	17	40	7.5	63,961,372
C_0037	28	44	8.3	60,288,132

(Continued)

Table 2. (Continued)

Sample ID	PMI	Age of Death	RIN	mRNA-Seq reads
C_0038	20	57	7.7	61,019,098
C_0039	15	80	7.3	74,892,650
C_0050	2	74	8.5	85,310,070
C_0053	2	69	8.4	167,044,880
C_0060	2	76	7.5	103,952,680
C_0061	3	78	7.6	95,393,100
C_0062	2	87	8.7	83,773,400
C_0065	2	86	8.7	115,714,502
C_0069	24	54	8.3	128,459,102
C_0070	19	68	6.3	145,087,692
C_0071	21	106	7.6	86,840,836
C_0075	23	52	7.4	99,946,984
C_0076	30	46	8.2	85,890,116
C_0077	21	36	8.5	80,103,722
C_0081	26	55	7.6	82,917,984
C_0082	18	57	7.8	123,118,398
C_0083	32	66	8.4	80,696,360
C_0087	19	64	8.7	77,198,978
C_0002	2	73	7.7	120,108,434
C_0003	2	91	7.9	38,420,004
C_0004	2	82	8.6	75,850,406
C_0005	2	97	9.1	150,661,916
C_0006	5	86	8.6	63,607,838
C_0008	2	91	8.7	66,131,458
C_0009	3	81	6.0	69,284,092
C_0010	2	79	8.4	60,542,776
C_0011	2	63	6.5	93,702,684

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Reference

1. Labadorf A, Hoss AG, Lagomarsino V, Latourelle JC, Hadzi TC, Bregu J, et al. (2015) RNA Sequence Analysis of Human Huntington Disease Brain Reveals an Extensive Increase in Inflammatory and Developmental Gene Expression. PLoS ONE 10(12): e0143563. doi:[10.1371/journal.pone.0143563](https://doi.org/10.1371/journal.pone.0143563) PMID: [26636579](https://pubmed.ncbi.nlm.nih.gov/26636579/)