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Corrigendum: Motivational, proteostatic and transcriptional deficits precede synapse loss, gliosis and neurodegeneration in the B6.*Htt*$^{Q111/+}$ model of Huntington's disease

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This Article contains a typographical error in the Methods section, under the subheading "Library construction, RNA Sequencing and RNASeq analysis", where:

"The fastq files were aligned using the default parameters of SNAPR[50] (https://github.com/PriceLab/snapr) against the GRCh38 genome assembly, along with the transcriptome assembly gtf file from Ensembl, GRCh38.75".

should read:

"The fastq files were aligned using the default parameters of SNAPR[50] (https://github.com/PriceLab/snapr) against the GRCm38 genome assembly, along with the transcriptome assembly gtf file from Ensembl, GRCm38".