

This Zenodo deposit contains a publicly available description of the Dataset:

"Bulk RNAseq of microglia in vitro cultures".

Dataset Description:

This dataset includes bulk RNAseq (2 x 150bps reads; Illumina TruSeq Stranded mRNA library prep kit with poly-A selection) of differentiated human pluripotent stem cells (hPSCs) into microglia. Briefly, hES (H9) and hiPSCs (KOLF2.1) underwent mesodermal induction via embryoid body formation, followed by hemogenic endothelium induction and myeloid differentiation to generate primitive monocyte-like precursors. These precursors were then matured into microglia by supplementing with key neuron-derived factors that promote microglial identity, thereby recapitulating human microglia in monoculture. A detailed protocol can be found at doi: [dx.doi.org/10.17504/protocols.io.14egnr8ezl5d/v1](https://doi.org/10.17504/protocols.io.14egnr8ezl5d/v1).

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Dataset Name: jakobsson-inviro-bulk-rnaseq-microglia, v1.0

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ASAP Project: Activation of transposable elements as a trigger of neuroinflammation in Parkinson's disease

Project Description: A range of genetic clinical and pathological studies have established that inflammation is a central component of neurodegenerative disorders including Parkinson's disease (PD). However what still remains unclear is the underlying cause for this inflammation. Here we propose a novel

hypothesis involving the aberrant activation of transposable elements (TEs) as the trigger for neuroinflammation in PD. TEs are viral-like mobile genetic elements that comprise nearly 50% of the human genome. We propose that the combination of age and the underlying pathogenic processes in PD result in their aberrant activation and with this the expression of viral-like sequences in the brain which in turn activates an immune response (as would be the case for any viral infection in the brain) and then drives further neurodegeneration. This theory by invoking an entirely new pathogenic mechanism will open up a number of new avenues of PD-research with clear clinical relevance. This includes novel diagnostic biomarkers that could be developed based on the presence of TE-derived transcripts or peptides as well as new drug targets that could be exploited for example by targeting viral-like mechanisms or viral-induced inflammation. In summary this project has the potential to identify a novel disease-contributing mechanism in PD with a high clinical potential for development of much needed diagnostics and therapies for this condition.

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This dataset is made available to researchers via the ASAP CRN Cloud: cloud.parkinsonsroadmap.org. Instructions for how to request access can be found in the [User Manual](#).

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This Zenodo deposit was created by the ASAP CRN Cloud staff on behalf of the dataset authors. It provides a citable reference for a CRN Cloud Dataset