Exploring microglial functions under physiological and pathological conditions by genome-wide expression profiling
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Oosterhof N, **Holtman IR**, van der Linde H, Boddeke HWGM, Eggen BJL, van Ham TJ. Genome-wide RNA-sequencing of zebrafish microglia reveals a conserved vertebrate microglial genetic signature and in situ proliferation of activated microglia in vivo. (in prep)


Janssen L, Dubbelaar ML, **Holtman IR**, Eggen BJL, Boddeke HWGM, De Deyn P, Van Dam D Aging, microglia and cytoskeletal regulation are key factors in the pathological evolution of the APP23 mouse model for Alzheimer’s disease. (submitted)