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Expression, diffusion and molecular interactions determine Wnt3 distribution

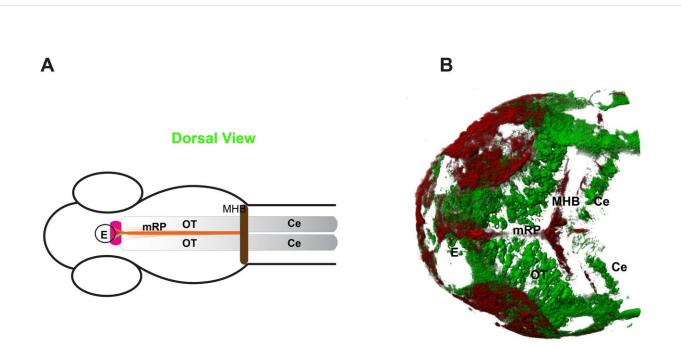


Figure shows Wnt3 source and target regions: (A) Schematic illustration of the zebrafish brain at 48 hours post fertilisation (hpf). (B) 3D projection of the source (red) and target (green) regions of the zebrafish brain at 48 hpf. (E – epithalamus; Ce – cerebellum; MHB - midbrain hindbrain boundary; mRP-midbrain roof plate; OT – optic tectum). Credit: eLife

An interdisciplinary team of physicists and biologists from NUS has characterized the mechanism by which Wnt3, a key regulator in neural patterning and brain development, spreads to long distances in the developing zebrafish brain.



The <u>development</u> of a multicellular organism requires cells to communicate with each other for coordinated growth. A common method by which cells interact during <u>embryonic development</u> is by exchanging signaling molecules. A group of cells that function as the "source" produces these signaling molecules which are secreted extracellularly and transported to distant "target" cells. At the "target" cells, these signaling molecules bind to their cognate surface receptors and activate downstream signaling to regulate vital cellular processes. Wnts are a class of signaling molecules that orchestrate cell growth, function, cell fate specification and cell death during development. In the endoplasmic reticulum, Wnts are modified with the addition of lipid moieties, a critical step for their secretion and activity. However, this modification makes Wnts hydrophobic and the transport in the aqueous extracellular spaces challenging. Therefore, the mode of transport by which Wnts accomplish long-range distribution remains an unresolved issue.

A research team led by Prof Thorsten Wohland from the Department of Biological Sciences and the Department of Chemistry, NUS has used imaging and quantitative fluorescence techniques to address how Wnt3, a member of the Wnt family, spreads in the developing zebrafish brain to achieve long-range signaling (see Figure). First, the team showed that Wnt3 tagged with enhanced green fluorescent protein (Wnt3EGFP) under the control of Wnt3 regulatory sequences spreads a considerable distance from the cells that produced it. Subsequently, the team used two characterisation techniques, fluorescence correlation spectroscopy (FCS) and fluorescence recovery after photo bleaching (FRAP) to measure the mobility of Wnt3EGFP at different length scales.

FCS is an analytical technique that measures the diffusion of fluorescent molecules by monitoring the fluctuations in fluorescence intensity emitted by the molecules in a small observation volume (femtolitre scale) whereas FRAP measures the recovery of the fluorescence



intensity in an irreversibly photo bleached region (micrometer scale). The team showed that the local mobility of Wnt3, as measured by FCS, is much larger than the long-range mobility, as measured by FRAP. They then demonstrated that this discrepancy is fully explained by Wnt3EGFP interactions with proteins present in the extracellular matrix and receptor, and co-receptor binding. This strongly reduces Wnt3's long-range mobility in the zebrafish brain. Collectively, the team's findings set the framework for future quantitative investigation of the Wnt3 interaction network during mammalian <u>brain development</u>.

Prof Wohland said, "Our capability to quantitatively measure molecular interactions and dynamics within small living organisms provides new insights into the processes that govern development at the molecular level."

More information: Veerapathiran S; Teh C; Zhu S; Kartigayen I; Korzh V; Matsudaira PT; Wohland T*, "Wnt3 distribution in the zebrafish brain is determined by expression, diffusion and multiple molecular interactions" *eLife* 2020; 9:e59489 <u>DOI: 10.7554/eLife.59489</u> Published: 2020.

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