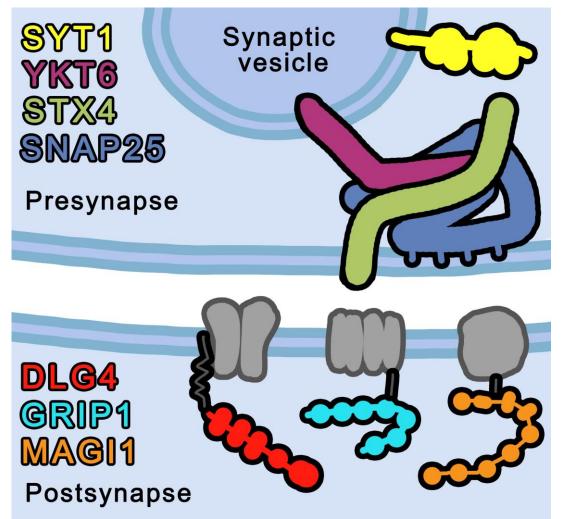
## MOL231: Revealing pre- and postsynaptic gene expressions in *Oikopleura dioica*

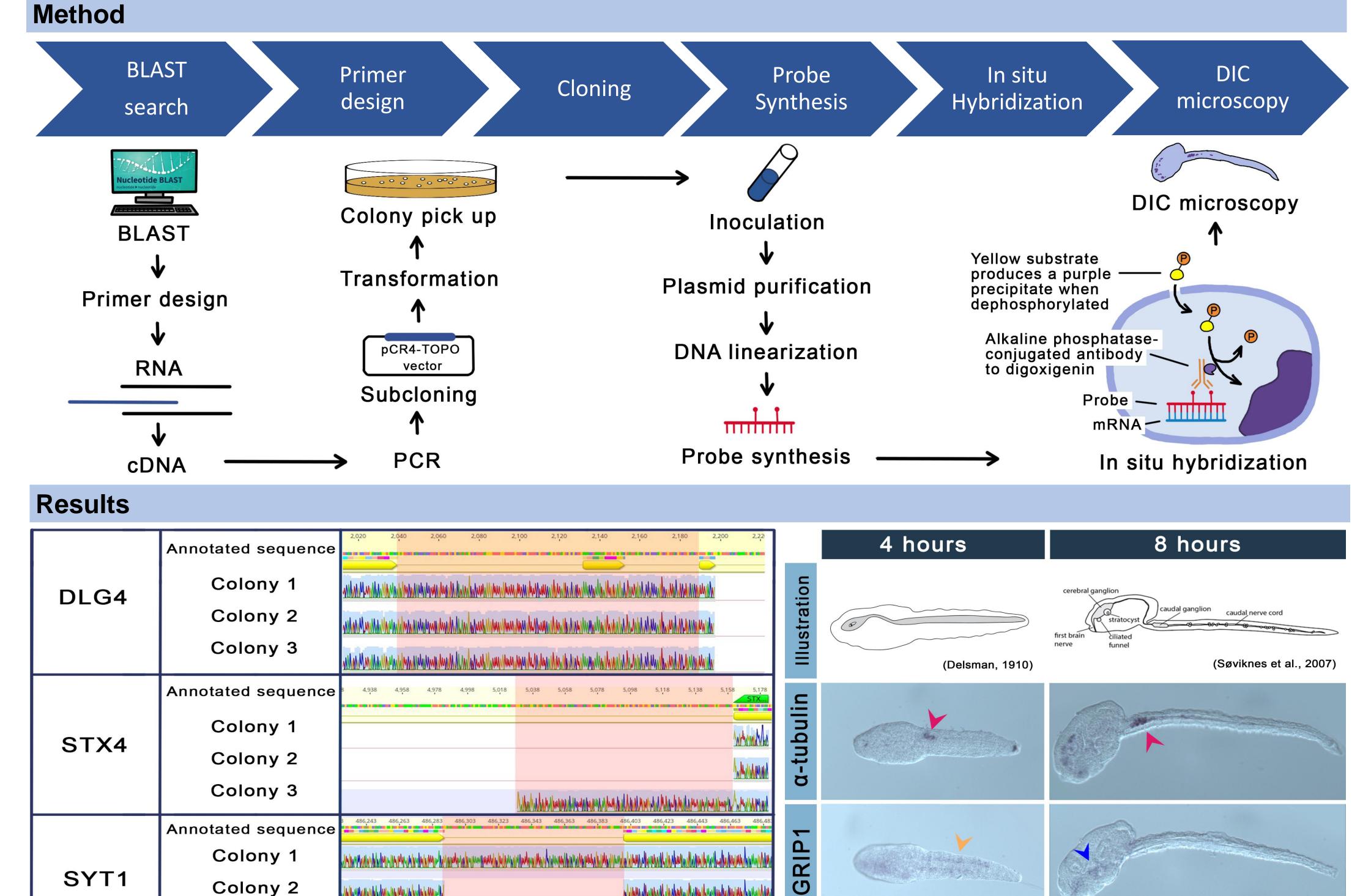
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## **Abstract & Aim**

Molecular cross-talk between nerve synapses is necessary to maintain important bodily functions. Studies regarding pre- and postsynaptic genes are therefore important for further understanding of the nervous system. In this study we took advantage of the simplified nervous system of *Oikopleura dioica*, a planktonic tunicate closely related to vertebrates, to study the synapsis. The **aim** of this study was to find marker genes for pre- and postsynaptic cells in *O. dioica*. A set of 18 genes were chosen; 11 of them are well-known presynaptic genes in other species, and 7 are postsynaptic genes. A total of 17 genes were successfully cloned and used for probe synthesis to do in situ hybridization in *O. dioica*.



**Fig. 1**: Scheme of synapse showing the selected genes for this study.



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**Figure 2:** Gene sequences compared to their annotated sequence from Oikobase (yellow boxes), with red areas showing discrepancies. DLG4 demonstrates wrong annotation in Oikobase; STX4, alternative splicing; SYT1, intron retention.

**Figure 3**: In situ hybridization *O.diocia.* The left panel shows a ventral view, while right panels shows a left lateral view. Top panels illustrate morphology and nervous system. Pink arrows indicate caudal ganglion; orange arrows, stained muscle cells; blue arrows, digestive tract expression.

## **Discussion & Conclusion**

- The presence of different splice variants and uncomplete processed transcripts show the high transcription levels of the synaptic genes in *O. dioica* in the selected developmental stages.
- Preliminary expression analysis revealed the presence of putative postsynaptic cells in the muscles, mouth, and ciliary
  rings confirming in situ hybridization as a good approximation to map cell types.



## **References:**

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