

Singapore Developmental Biology Club

SEMINAR ANNOUNCEMENT

23 May 2012, Wednesday
Aspiration Theatrette, Level 2M, Matrix, Biopolis
5:45PM - 6:45PM



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Seminar Title: Piecing together the squint puzzle.

The earliest steps of axis formation in zebrafish are thought to be regulated by maternal factors that activate Wnt/β-catenin signaling to specify the dorsal axis. We previously showed that asymmetric localization of maternal transcripts of the conserved zebrafish TGF-β factor, Squint (Sqt/Ndr1), in 4-cell stage embryos predicts dorsal, preceding nuclear accumulation of β-catenin. Cell ablations and antisense oligonucleotides targetting Sqt lead to dorsal deficiencies, suggesting that localized maternal sqt functions in dorsal specification. However, due to discrepancies between the genetic nodal mutants and the ablations/antisense knock-down results, maternal sqt function was debated. We find that sqt RNA has activity independent of Sqt protein in dorsal specification. Surprisingly, overexpression of mutant/non-coding sqt RNA and the sqt 3' UTR, leads to ectopic nuclear b-Catenin accumulation and expands dorsal gene expression. Dorsal activity of sqt RNA requires Wnt/β-catenin but not Oep-dependent Nodal signaling, explaining the discrepancy between the nodal mutants and morpholino phenotypes. Finally, we revisited the activity of the sqt morpholinos, and show that in addition to blocking Sqt translation/splicing, they also disrupt sqt RNA localization. Depletion of maternal sqt RNA abolishes nuclear β-Catenin, providing the mechanism for the loss of dorsal in the morphants, and places maternal sqt RNA activity upstream of β-Catenin. Remarkably, loss of early dorsal gene expression can be rescued by sqt 3' UTR sequences. Our findings identify new non-coding functions for the nodal genes, and support a model wherein sqt RNA acts as a scaffold to bind and deliver/sequester maternal factors to future embryonic dorsal.