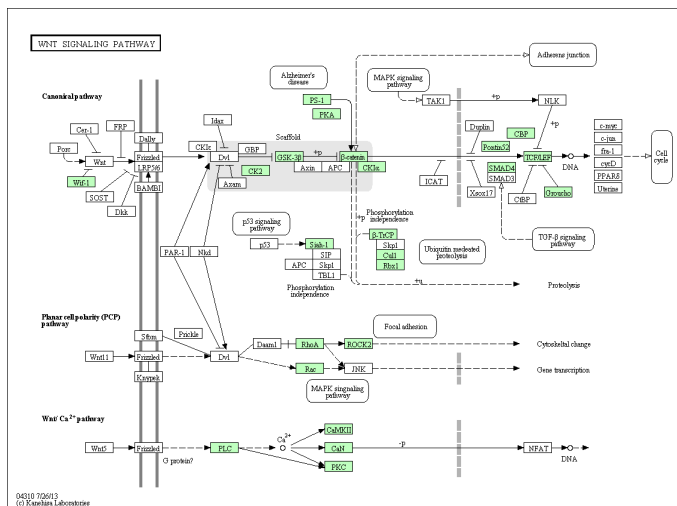
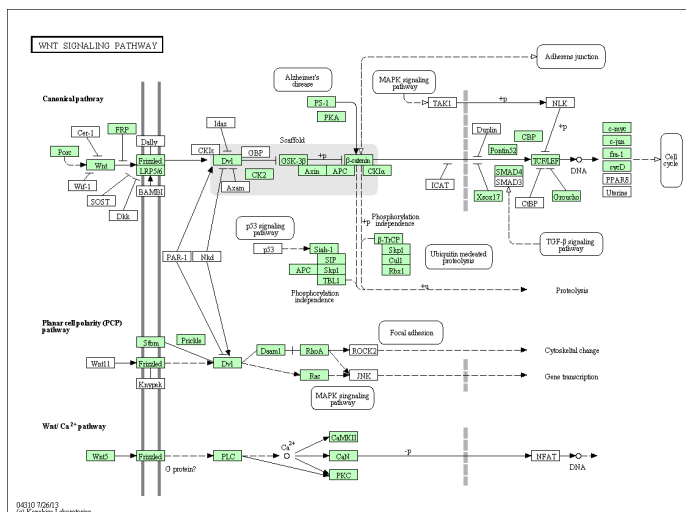


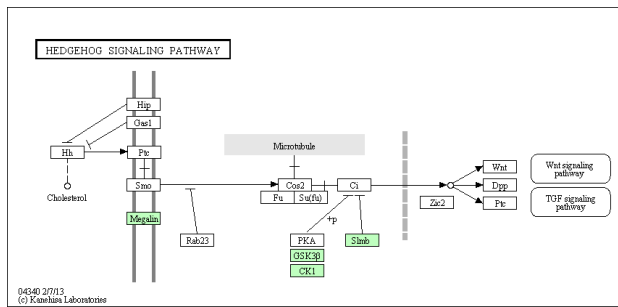
Wnt pathway components identified in the genome and transcriptome of *Kudoa iwatai* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



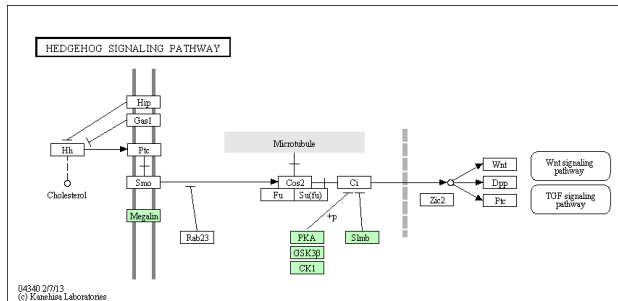
Wnt pathway components identified in the genome and transcriptome of *Myxobolus cerebralis* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



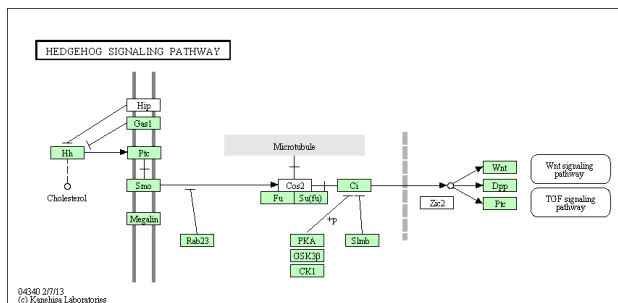
Wnt pathway components identified in the genome and transcriptome of *Polypodium hydriiforme* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



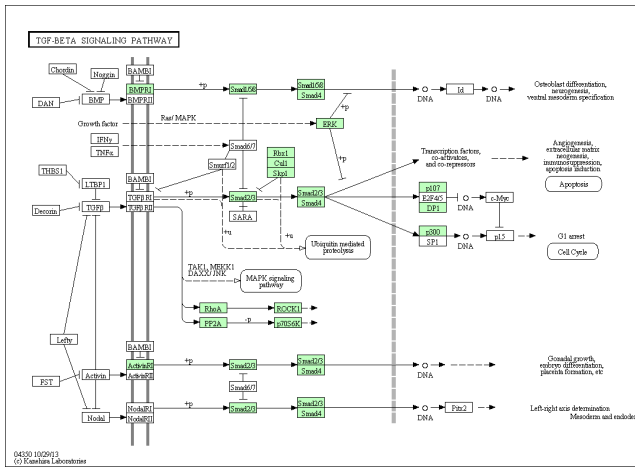
Hedgehog signaling pathway components identified in the genome and transcriptome of *Kudoa iwatai* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



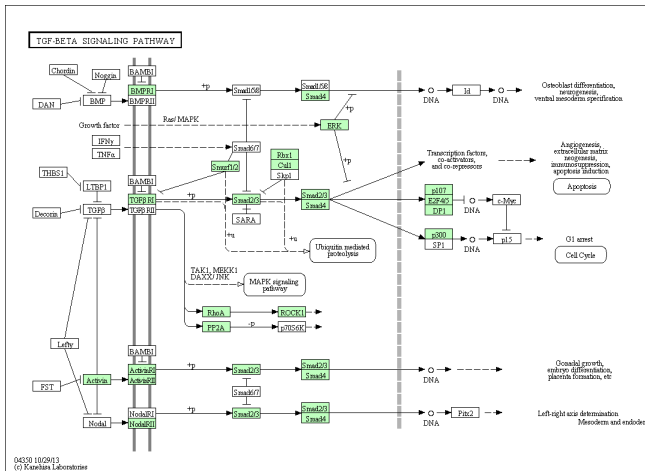
Hedgehog signaling pathway components identified in the genome and transcriptome of *Myxobolus cerebralis* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



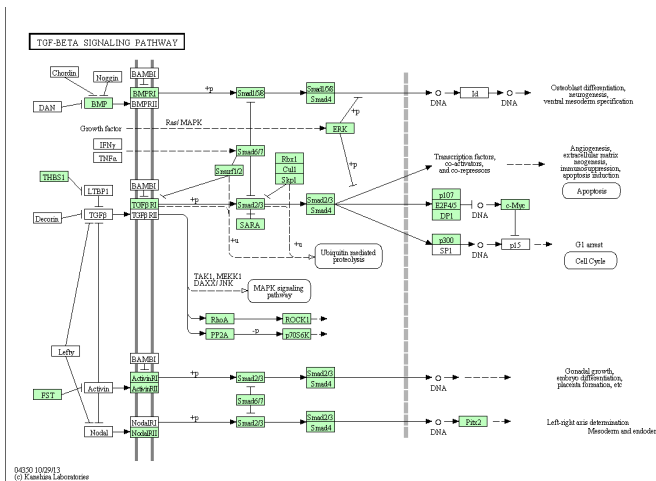
Hedgehog signaling pathway components identified in the genome and transcriptome of *Polypodium hydriforme* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



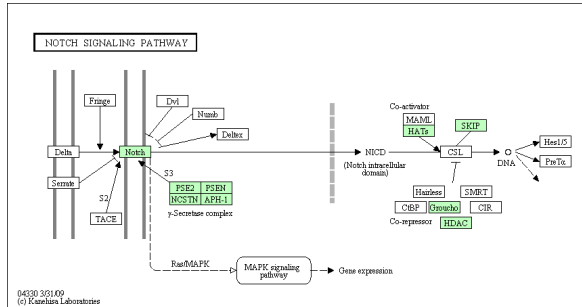
TGF- β signaling pathway components identified in the genome and transcriptome of *Kudoa iwatai* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



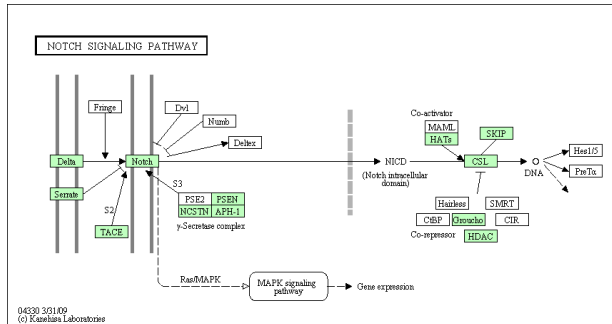
TGF- β signaling pathway components identified in the genome and transcriptome of *Myxobolus cerebralis* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



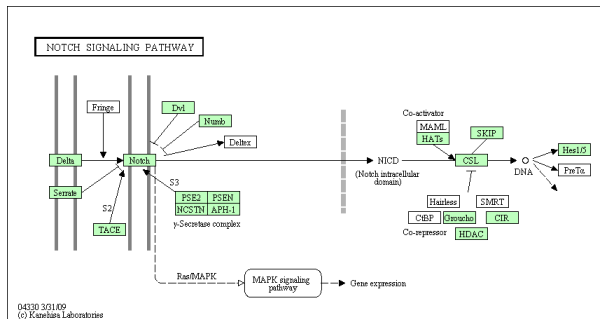
TGF- β signaling pathway components identified in the genome and transcriptome of *Polypodium hydriforme* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



Notch signaling pathway components identified in the genome and transcriptome of *Kudoa iwatai* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



Notch signaling pathway components identified in the genome and transcriptome of *Myxobolus cerebralis* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.



Notch signaling pathway components identified in the genome and transcriptome of *Myxobolus cerebralis* through assignment of KEGG orthology terms. Components identified in this species are highlighted in green.