

### New insights on winter-associated regulators of the circadian clock in poplar.

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Winter dormancy is an adaptive mechanism that allows trees from temperate and cold regions to survive the harsh conditions of this season. This process is strongly influenced by daylength and temperature, which are signals that are integrated by the circadian clock. Low temperatures cause the disruption of the circadian clock in trees, which consists in a transcriptional activation followed by an arrhythmic expression. In this work we uncover winter-associated regulators of the circadian clock in poplar. We generated transgenic poplars that express a transcriptional fusion between the promoter of a circadian clock gene and the luciferase gene. Setting up a circadian luminiscence assay by which we register the luciferase activity from leaf discs with a luminometer, we have characterized the expression of this promoter under different photoperiods. Then, we carried out a Yeast One Hybrid (Y1H) screening with a library enriched in winter-associated factors using this promoter as bait. Candidate regulators are tested *in vivo* by transient assays in poplar, overexpressing and silencing the candidate genes. Our results indicate the selected promoter region contains the circadian elements as well as the luciferase activity shows the expected expression under both long and short days. In the Y1H screening, we found several candidates that are classified either as transcription factors or chromatin remodelers. We discuss the putative role of these proteins as regulators of the poplar circadian clock.

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