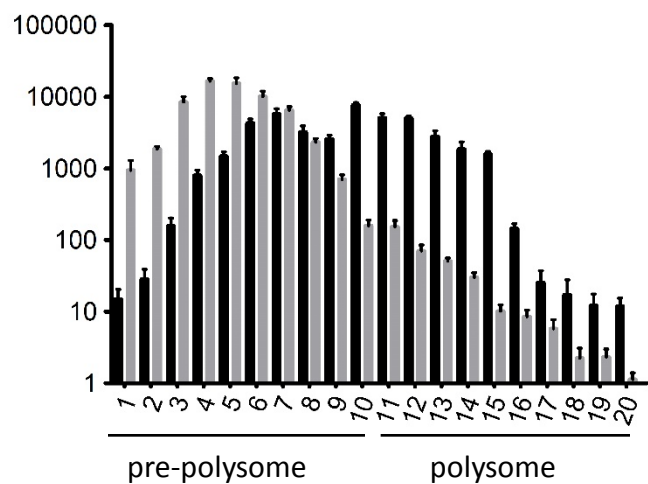
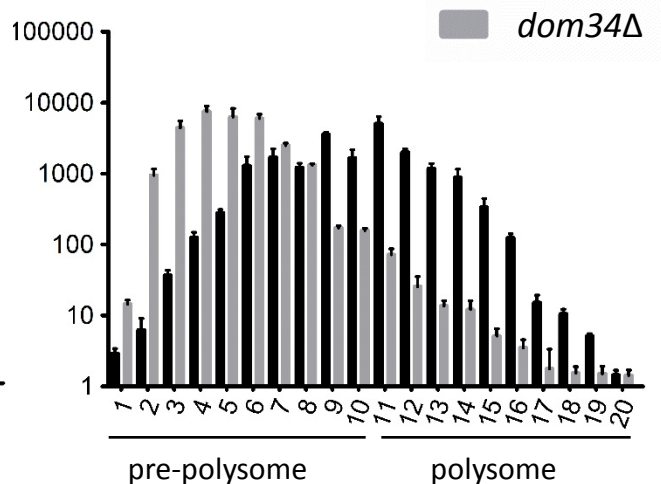


A.

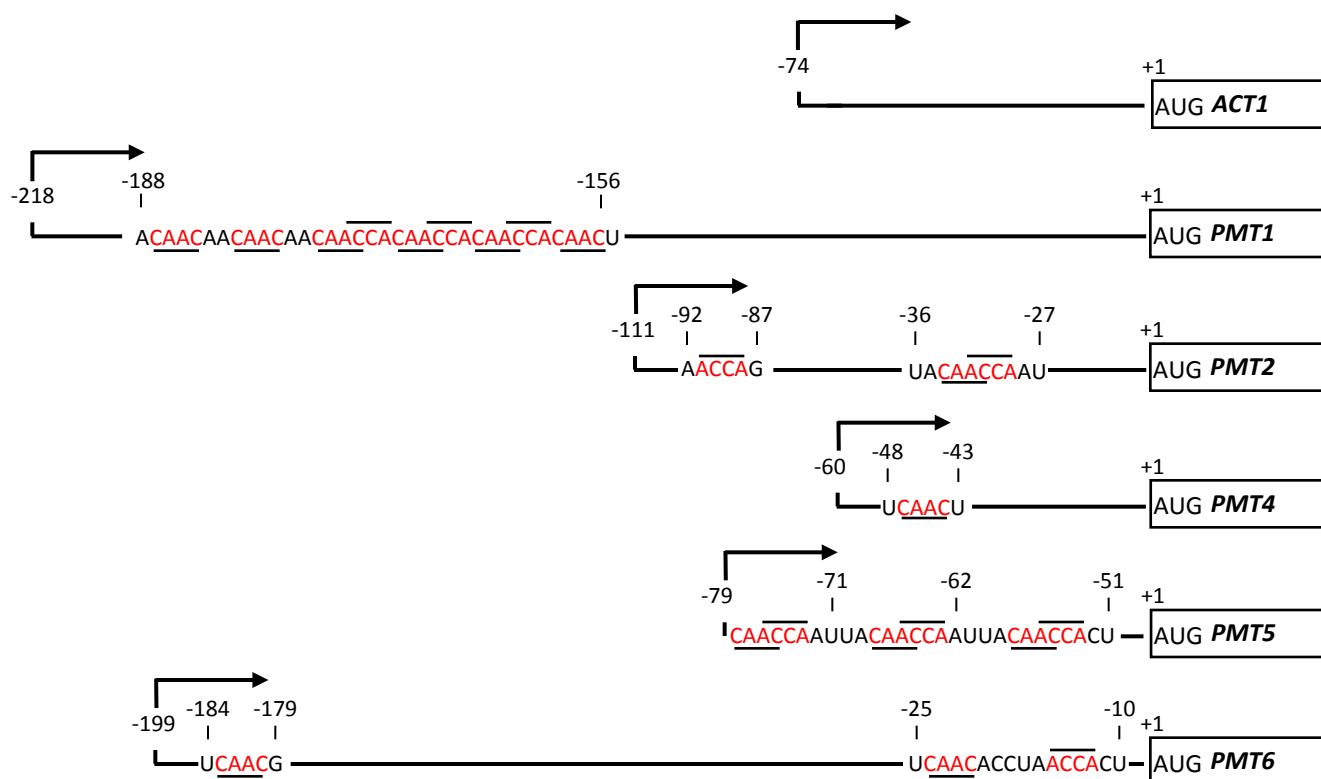
PMT4



PMT6



B.



S6 Fig. Occurrence of *PMT4* and *PMT6* transcripts in polysome-gradient fractions. **(A)** The indicated transcripts were detected by qPCR after adding a known amount of an *in vitro* generated transcript of CaCBGluc as calibrator. Each bar represents the normalized mean *PMT4* or *PMT6* transcript level of two independent experiments including the standard error of the mean. **(B)** Incidence of “CAAC” and “ACCA” motifs in 5'-UTRs of *ACT1* and *PMT*-genes. The 5'-UTR sequences of *ACT1* (reference gene) and all *C. albicans* *PMT*-genes were analyzed for occurrence of the identified “CAAC” and “ACCA” motif (Fig. 3B). Information about transcript start sites were taken from Tuch *et al.* [29] and Bruno *et al.* [30]; the transcript with the longer 5'-UTR was chosen and respective sequences were obtained from CGD (<http://www.candidagenome.org>) assembly 21. Occurrence of the “CAAC” (underlined) and “ACCA” (line on top) motifs are highlighted in red at the indicated positions. The “CAAC” motif was identified in 5'-UTRs of all *PMT*-genes, but not in the 5'-UTR sequence of the reference gene *ACT1*.