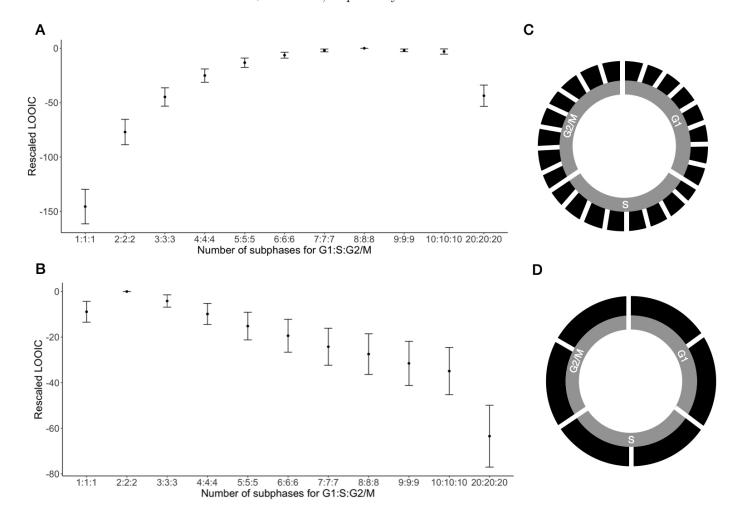
Fig. S4 Using LOOIC to determine number of stages m. LOOIC comparison among different m for (A) -DOX and (B) +DOX cells. The values were rescaled: The best model was set to zero. Having more negative values means the model has worse prediction. (A) shows m=8 is the best model for -DOX and (B) shows m=2 is the best model for +DOX in terms of model predictions. (C) and (D) are the cartoon-like figures for visualizing the most probable number of subphases of each phase for -DOX and +DOX cells, respectively.



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