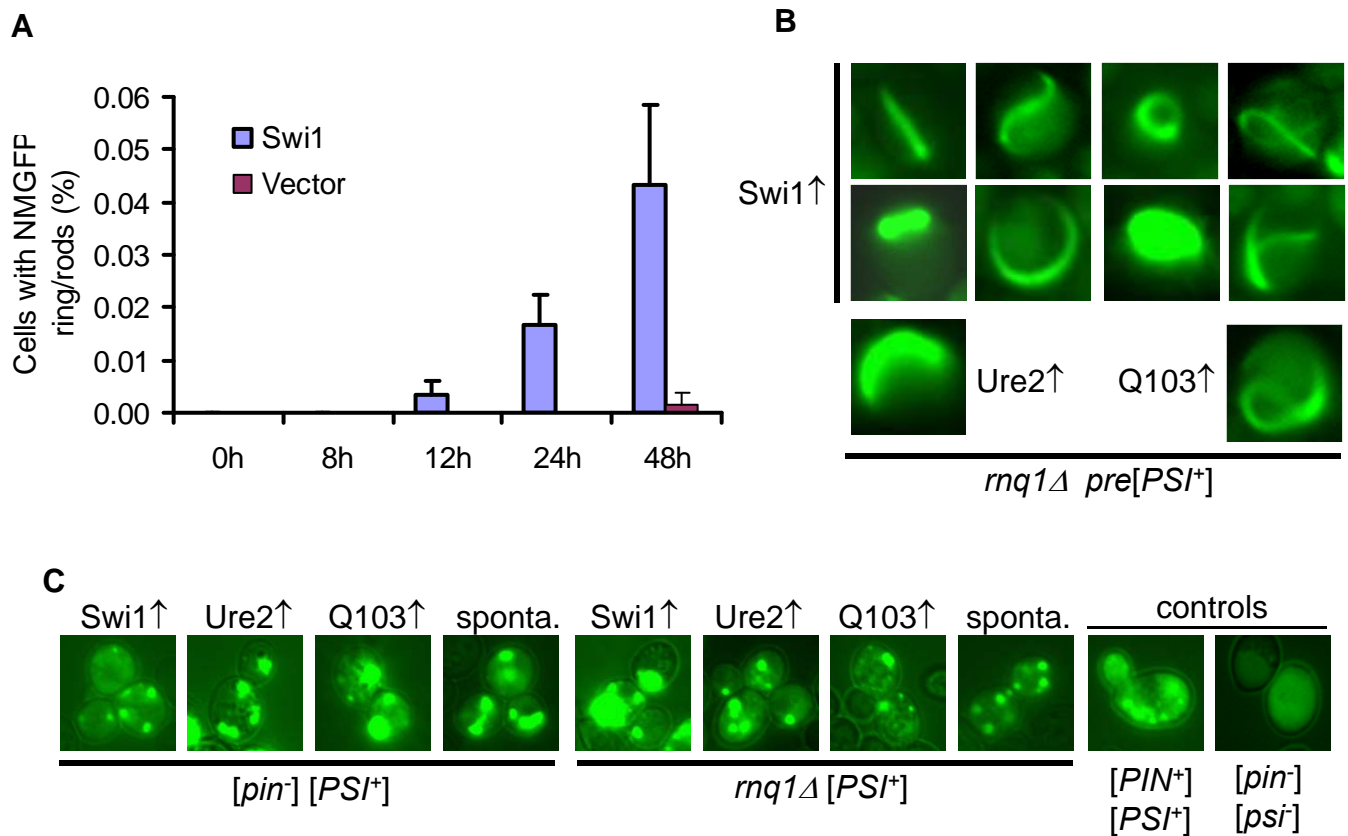
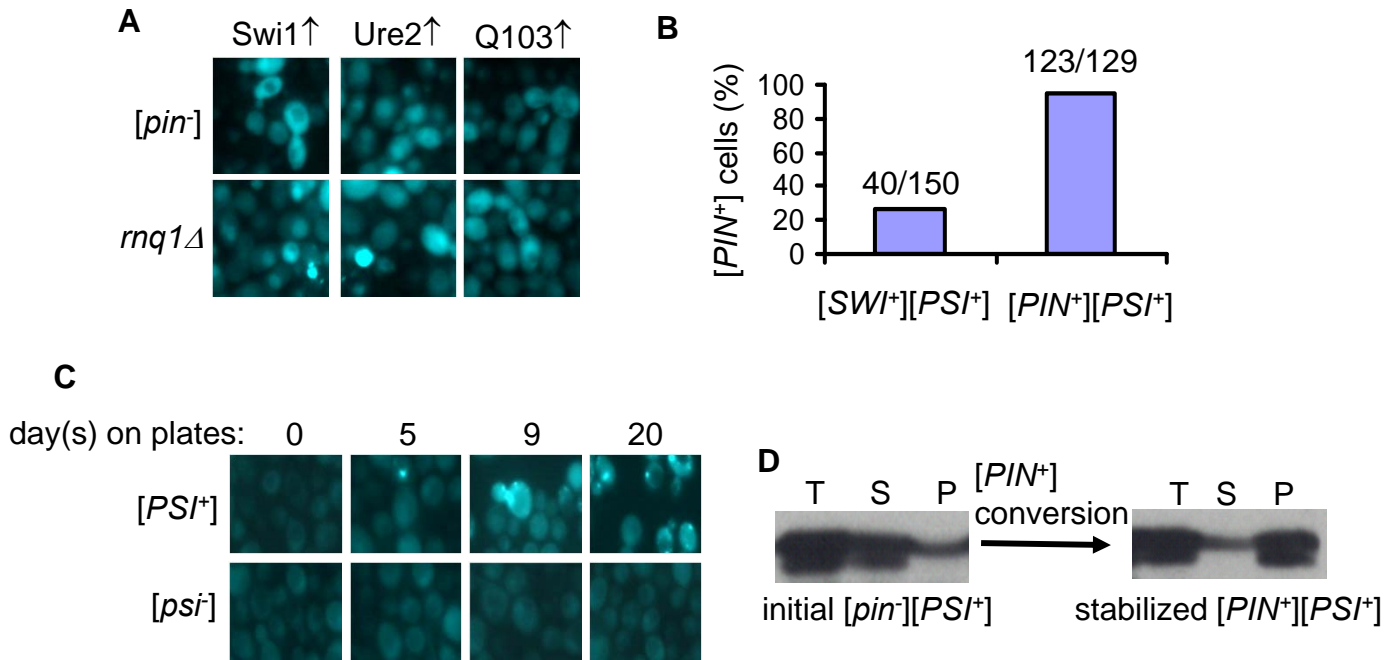


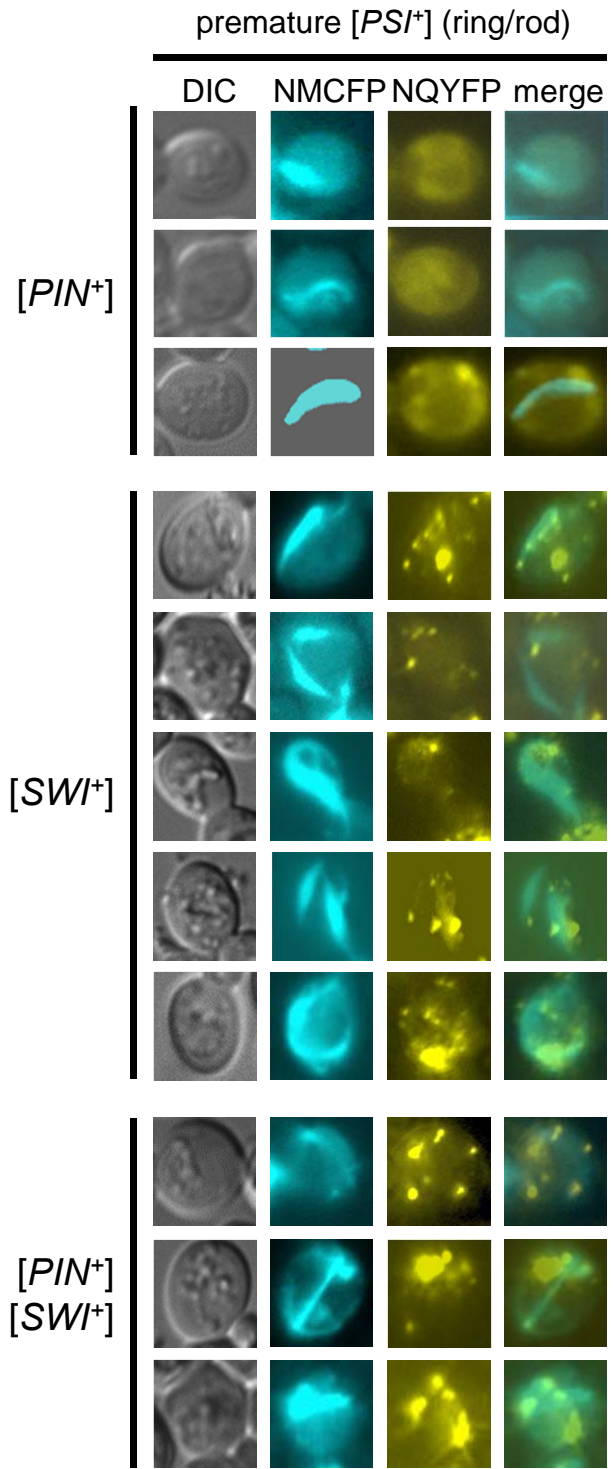
**Figure S1** **A.** Yeast cells with indicated strain background were transformed with plasmid *p413CUP1-NMGFP* (NMGFP↑) or *pRS413CUP1-NM* (NM↑), and also one of the following plasmids: *p426GPD-SWI1* (Swi1↑), *p416GPD-URE2NPDGFP* (Ure2↑), *p426GPD-Q103GFP* (Q103↑) or *p426GPD-GFP* (GFP↑). [*PSI*<sup>+</sup>] induction was carried out as described in Materials and Methods. After 48 hours of expression upon addition of 100 μM CuSO<sub>4</sub>, cultures were spotted onto the indicated plates. Note: The appearance of Ade<sup>+</sup> colonies by Swi1, Ure2, or Q103 overproduction in [*pin*<sup>-</sup>] cells was delayed ~2 to 3 days when compared to that obtained from [*PIN*<sup>+</sup>]. **B.** Randomly selected [*PSI*<sup>+</sup>] isolates shown in panel A were replica-plated onto GdnHCl-containing plates for up to three times (3x) and then patched back to YPD plates to see the curability and to determine [*PSI*<sup>+</sup>] variants. Spontaneously formed [*PSI*<sup>+</sup>] isolates were also included. Shown are representative results. Note: overproduction of NM and NMGFP gave similar [*PSI*<sup>+</sup>] induction efficiency.



**Figure S2** **A.** Sup35NMGFP and Swi1 were co-overproduced in a non-prion strain containing the plasmids *p413CUP1-NMGFP* and *p426GPD-SWI1* (blue) in the presence of 100  $\mu$ M CuSO<sub>4</sub>. The frequency of Sup35NMGFP ring/rod-like aggregates was analyzed at the indicated times. Cells containing *p413CUP1-NMGFP* and *p426GPD* were treated under identical conditions as the vector control (red). **B.** Representative images of ring/rod Sup35NMGFP aggregates in pre-mature [*PSI*<sup>+</sup>] cells generated by co-overproduction of Sup35NMGFP and one of the indicated proteins. This experiment was carried out in a *rnq1Δ* strain with conditions similar to what described for Figure S1A. Images were taken after 24h induction. **C.** Sup35NMGFP aggregates of mature [*PSI*<sup>+</sup>] isolates obtained in experiments shown in Figure S1A-1B. Ure2: Ure2<sub>1-65</sub>-GFP; Q103: polyQ103-GFP; and sponta.: spontaneously formed [*PSI*<sup>+</sup>] isolates.



**Figure S3** **A.** Majority [*PSI*<sup>+</sup>] isolates obtained by co-production of Sup35NMGFP and Swi1 (Swi1 ↑), Ure2<sub>1-65</sub>-GFP (Ure2 ↑) or polyQ103-GFP (Q103↑) remained [*pin*-]. As shown, Rnq1CFP remained diffused in stabilized [*PSI*<sup>+</sup>] cells obtained under the indicated co-overproduction conditions. **B.** Heritable Rnq1CFP aggregates (indicative of [*PIN*<sup>+</sup>]) were analyzed after eliminating the plasmid *pRS413CUP1-NMGFP* and *p426GPD-SWI1*. Numbers shown are [*PIN*<sup>+</sup>] isolates versus total examined isolates. **C.** A [*PSI*<sup>+</sup>] isolate acquired by co-overproduction of Sup35NMYFP and Swi1 was compared with an isogenic [*pin*-][*psi*<sup>-</sup>] strain for their ability in promoting spontaneous [*PIN*<sup>+</sup>] conversion after incubation at 4°C for the indicated days. The method used in this study is similar to that described in an earlier paper (Derkatch *et al.*, 2001, cell 106: 171-182). Shown are Rnq1CFP signals. **D.** The acquired Rnq1CFP aggregate-containing [*PSI*<sup>+</sup>] cells (20 days in panel C) were further stabilized by passages and compared with the initial [*PSI*<sup>+</sup>] strain (0 day cells) using a centrifugation assay. T, total lysate; S, supernatant; P, pellet. An anti-Rnq1 polyclonal antibody was used in the Western Blotting. Note: all results shown in this figure were obtained using cells carry an integrated copy of *TEF1-RNQ1CFP*.



**Figure S4** Representative images of Swi1-NQYFP and Sup35NM-CFP fluorescence patterns in pre-mature [*PS*<sup>+</sup>] cells. Sup35NM-YFP was overproduced under the *CUP1* promoter. In strains containing the indicated prion(s), the *SWI1*-NQYFP was expressed under the *TEF1* promoter.