Improvement of the reverse tetracycline transactivator by single amino acid substitutions that reduce leaky target gene expression to undetectable levels

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The following blots were cropped to produce the inset of Figure 2a.



Supplemental figure S1. Glycine to valine substitution at residue 72 does not affect steady state protein levels. **a**, Western blot comparing the rtTA protein abundances of the rtTA-M2 and rtTA-M2(G72V) variants when expressed from theTDH3 promoter. **b**, Loading control comparing protein abundances of Clb2 in strains expressing the rtTA-M2 variant or rtTA-M2(G72V) variant.

The following blots were cropped to produce Figure 3B.



Supplemental Figure S2. Substitution of glycine for non-polar side chains at residue 72 does not affect steady state protein levels. **a**, Western blot comparing the rtTA protein abundances of all four rtTA variants when cells were induced by 1000nM of β -estradiol or no induction. **b**, Loading control comparing protein abundances of Cdk1 when cells expressing the four rtTA variants were induced by 1000nM of β -estradiol or no induction.



Supplemental Figure S3. yeGFP expression from the P_{TET4} promoter and GEV cassette alone is indistinguishable from autofluorescence of wildtype BY4742 yeast cells when cells were cultured in 1000nM ß-estradiol.



Supplemental Figure S4. **a,b**, The basal activity of rtTA variants as a function of their ß-estradiol-dependent expression level (mean +/- s.e.m of three technical replicates). In the inset of **b**, histograms of single cell fluorescence of the G72P variant and a SE-G72P variant cultured in 1000nM ß-estradiol.



Supplemental Figure S5. Surface plots of the data presented in Figure 4 (c-f). a-d, Surface plots displaying reporter expression (Arbitrary units) as a function of β -estradiol-dependent transactivator expression and doxycycline induction.