Table S4. Parental and deletion strains used in the course of this work and respective genotypes.

|  |  |  |
| --- | --- | --- |
| **Strain** | **Genotype** | **Reference** |
| *Candida albicans*  BWP17 | *ura3∆::λimm434/ura3∆::λimm434 arg4∆::hisG*/*arg4∆::hisG* *his1∆::hisG*/*his1∆::hisG* | [[1](#_ENREF_1)] |
| CEC3785 | *ura3∆::λimm434/ura3∆::λimm434 arg4∆::hisG*/*ARG4* *his1∆::hisG*/*HIS1 ADH1/adh1::PTDH3-carTA::SAT1 RPS1/RPS1::* CIp10-P*TET*-GTW | This study |
| BWP17-GFP | *ura3∆::λimm434/ura3∆::λimm434 his1∆::hisG/HIS1 arg4∆::hisG/ARG4 RPS1/RPS1::*CIp10*-*P*TDH3*-GFP | This study |
| BWP17-mCherry | *ura3∆::λimm434/ura3∆::λimm434 his1∆::hisG/HIS1 arg4∆::hisG/ARG4 RPS1/RPS1::*CIp10*-*P*ADH1*-mCherry | This study |
| CAI4 | *ura3∆::λimm434/ura3∆::λimm434* | [[2](#_ENREF_2)] |
| SN76 | *arg4∆/arg4∆ his1∆/his1∆ ura3∆::λimm434/ura3∆::λimm434 iro1∆::λimm434/iro1∆::λimm434* | [[3](#_ENREF_3)] |
| SN148 | *arg4∆/arg4∆ leu2∆/leu2∆ his1∆/his1∆ ura3::λimm434/ura3∆::λimm434 iro1∆::λimm434/iro1∆::λimm434* | [[3](#_ENREF_3)] |
| SN152 | *his1*Δ*/his1*Δ *leu2*Δ */leu2*Δ *arg4*Δ */arg4*Δ *URA3/ura3*Δ*::*λ*imm434, IRO1/iro1*Δ*::*λ*imm434* | [[3](#_ENREF_3)] |
| CEC1121 | SN148 *GAL1/gal1::LEU2 ADH1/adh1::PADH1-cartTA::SAT1::PTET-caGFP* | This study |
| CEC1429 | CAI4 *ADH1/adh1::PADH1-cartTA::SAT1::PTET-caGFP* | This study |
| CEC2908 | *ura3∆::λimm434/ura3∆::λimm434 his1∆::hisG/HIS1 arg4∆::hisG/ARG4 ADH1/adh1::PTDH3-carTA::SAT1* | [[4](#_ENREF_4)] |
| CEC3781 | SN76 *his1∆/HIS1 ADH1/adh1::PTDH3-carTA::SAT1 Ca21chr4\_C\_albicans\_SC5314:473390 to 476401∆::PTDH3-GFP-CaARG4* | This study |
| CEC3783 | SN76 *arg4∆/ARG4 ADH1/adh1::PTDH3-carTA::SAT1 Ca21chr4\_C\_albicans\_SC5314:473390 to 476401∆::PTDH3-BFP-HIS1* | This study |
| *∆∆pga22* | BWP17 *pga22∆::ARG4/pga22∆::HIS1 RPS1/RPS1::*CIp10 | This study |
| *∆∆pga22*-GFP | BWP17 *pga22∆::ARG4/pga22∆::HIS1 RPS1/RPS1::*CIp10*-*P*TDH3*-GFP | This study |
| *∆∆pga22*-mCherry | BWP17 *pga22∆::ARG4/pga22∆::HIS1 RPS1/RPS1::*CIp10*-*P*ADH1*-mCherry | This study |
| *∆∆pga59* | BWP17 *ura3∆::λimm434* URA3 pga59Δ: : HIS1/pga59Δ: : ARG4 | [[5](#_ENREF_7)] |
| *Saccharomyces cerevisiae* | |  |
| BY4742  VIF205 | *MAT his3Δ1 leu2Δ0 lys2Δ0 ura3Δ0*  BY4742/pBC542 | [[6](#_ENREF_9)]  [7] |

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