Table S1. Primers used in this study.

Primers	Nucleotide sequence (5' to 3')	Employment
spt1-1820-UF1	GCTCGTAGCCAAGTAGGG	*spt1* deletion strain construction
spt1-78-UR1	CTCCTTCAATATCAGTTAAGGTCGAAAACGCAGCGAAACAGA	*spt1* deletion strain construction
spt1-325-DF1	AAATTCCGTCACCAGCCCTGGGTTGCCCAATCCTCGCAATCTT	*spt1* deletion strain construction
spt1-1557-DR1	GGTCGCCATTATCCGTTG	*spt1* deletion strain construction
hph-F	CGACCTTAACTGATATTGAAGGAG	*hph* markers amplifcation
hph-R	CAACCCAGGGCTGGTGAC	*hph* markers amplifcation
spt1-1458-UF2	ATTCTGTCGCAGTATGGG	strain construction
spt1-1287-DR2	GAGGAGTCTGGGTCTTGTT	strain construction
spt1-1820-UF1	GCTCGTAGCCAAGTAGGG	*spt1* complement strain construction
spt1-1498-R1	TTTGATGATTTCAGTAACGTTAAGTTTAATGGTGATGGTGATG ATGCTCGGAGAGCTCAGAGATGAA	*spt1* complement strain construction
spt1-325-DF2	TTATGTGTAACAAGAAAGACAGTATAATACAAACAAAGATGC AAGAGCGGCCCAATCCTCGCAATCTT	*spt1* complement strain construction
spt1-1557-DR1	GGTCGCCATTATCCGTTG	*spt1* complement strain construction
T*trpC*-F1	ACTTAACGTTACTGAAATCAT	*spt1* complement strain construction
T*trpC*-R1	ACAATTCCACACAACATACGAGCCGGAAGCATAAAGTGTAAA GCCTGGGGGAGTGGAGATGTGGAGTGGG	*spt1* complement strain construction
ptrA-F	CCCCAGGCTTTACACTTTAT	*ptrA* markers amplifcation
ptrA-R	CCGCTCTTGCATCTTTGTT	*ptrA* markers amplifcation
spt1-1820-UF1	GCTCGTAGCCAAGTAGGG	Spt1-eGFP fusion strain construction
spt1-1498-R2	TCGACCAGGATGGGCACCACCCCGGTGAACAGCTCCTCGCCC TTGCTCACCTCGGAGAGCTCAGAGATGAA	Spt1-eGFP fusion strain construction
spt1-325-DF2	TTATGTGTAACAAGAAAGACAGTATAATACAAACAAAGATGC AAGAGCGGCCCAATCCTCGCAATCTT	Spt1-eGFP fusion strain construction
spt1-1557-DR1	GGTCGCCATTATCCGTTG	Spt1-eGFP fusion strain construction
T*trpC*-F2	TCGTGACCGCCGCCGGGATCACTCTCGGCATGGACGAGCTGT ACAAGTAAACTTAACGTTACTGAAATCATCAAA	Spt1-eGFP fusion strain construction
T*trpC*-R1	ACAATTCCACACAACATACGAGCCGGAAGCATAAAGTGTAAA GCCTGGGGGAGTGGAGATGTGGAGTGGG	Spt1-eGFP fusion strain construction
egfp-F	GTGAGCAAGGGCGAG	Spt1-eGFP fusion strain construction
egfp-R	TTACTTGTACAGCTCGTCCATGCC	Spt1-eGFP fusion strain construction
spt1-1820-UF1	GCTCGTAGCCAAGTAGGG	*spt1* overexpression strain SOE construction
spt1-1498-R1	TTTGATGATTTCAGTAACGTTAAGTTTAATGGTGATGGTGATG ATGCTCGGAGAGCTCAGAGATGAA	*spt1* overexpression strain SOE construction
T*trpC*-F1	ACTTAACGTTACTGAAATCAT	*spt1* overexpression strain SOE construction
T*trpC*-R2	AGACGAGACACGACGGCGGTGCATTCTGGGTAAACGACT	*spt1* overexpression strain SOE construction
pyrG-F	GATGCCAACGGTATCCTCA	*pyrG* markers amplifcation
pyrG-R	TGCGGTCAATCTCCTCCT	*pyrG* markers amplifcation
P*cdna1*-F	TGGTACATGGATCTCGAACTGA	*spt1* overexpression strain SOD construction
P*cdna1*-R	GAGAGAAGTTGTTGGATTGATC	*spt1* overexpression strain SOD construction
spt1-1-F	GATCAATCCAACAACTTCTCTCATGCGGTCCGTTGTCGCCC	*spt1* overexpression strain SOD construction
spt1-1498-R1	TTTGATGATTTCAGTAACGTTAAGTTTAATGGTGATGGTGATG ATGCTCGGAGAGCTCAGAGATGAA	*spt1* overexpression strain SOD construction
T*trpC*-F1	ACTTAACGTTACTGAAATCAT	*spt1* overexpression strain SOD construction
T*trpC*-R2	AGACGAGACACGACGGCGGTGCATTCTGGGTAAACGACT	*spt1* overexpression strain SOD construction