

Plasmid	Description
pV1393	<i>C. albicans</i> Solo CRISPR vector flanked by Neut5+FRT targeting regions; FLP expression driven by <i>SAP2</i> promoter
pV1524	<i>C. albicans</i> Solo CRISPR vector flanked by Neut5+FRT targeting regions; FLP expression driven by <i>MAL2</i> promoter
pV1539	<i>C. albicans</i> Gene Drive vector - derived from pV1524 Kpn1/Xma1 backbone replaced with Kpn1/Xma1 fragment from pV1093
pV1543	<i>C. albicans</i> Gene Drive vector - derived from pV1539 with second Sac2 site placed between BsmBI sites
pV1326	<i>S. cerevisiae</i> and <i>C. glabrata</i> Solo CRISPR vector, marked with <i>URA3</i> and <i>Nat<sup>R</sup></i>
pV1338	pV1326 + sgScADE2
pV1382	<i>S. cerevisiae</i> and <i>C. glabrata</i> Solo CRISPR vector, marked with <i>URA3</i> and <i>Nat<sup>R</sup></i>
pV1386	pV1382 + sgScADE2
pV1464	<i>N. castellii</i> Solo CRISPR vector, marked with <i>URA3</i> and <i>Nat<sup>R</sup></i>
pVG1	pV1382 + sgScADE2 + ScADE2 stop codon repair template
pV1534	<i>C. albicans</i> Gene drive vector for <i>ADE2</i> , with BsmBI <i>in vitro</i> linearization sites
pV1535	<i>C. albicans</i> Gene drive vector for <i>ADE2</i> with sgADE2 <i>in vivo</i> linearization site
pV1418	pV1393 + sgCPH1
pV1420	pV1393 + sgEFG1
pV1528	pV1524 + sgCaLEU2
pV1531	pV1524 + sgCaMET15
pV1487	pV1382 + sgCgMet15
pV1518	pV1382 + sgCgLeu2
pV1435	pV1382 + sgCgADE2
pV1329	pV1326 +sgCgADE2
pV1465	pV1464 + sgNcADE2
pGB9	pV1382 + sgScLEU2
pV1563	<i>C. albicans</i> Inducible Gene Drive vector - <i>MAL2</i> driven <i>CaCas9</i> , <i>SAP2-FLP</i> (with intron) - second Sac2 site for cloning
pV1571	<i>C. albicans</i> maltose-inducible Gene Drive vector for <i>ADE2</i> , pV1563 + guide/targeting/linearization regions as in pV1534
pV1465	pV1464 with sgNcADE2
pV1612	pV1465 with stop repair template
pV1613	pV1465 with delta repair template