

Resource Summary Report

Generated by [NIF](#) on Aug 31, 2026

pSK+KanaRpsL

RRID:Addgene_20871

Type: Plasmid

Proper Citation

RRID:Addgene_20871

Plasmid Information

URL: <http://www.addgene.org/20871>

Proper Citation: RRID:Addgene_20871

Insert Name: KanamycinResistance-RpsL

Organism: E.coli

Bacterial Resistance: Ampicillin and Kanamycin

Defining Citation: [PMID:19160076](#)

Vector Backbone Description: Backbone Size:2935; Vector Backbone:pBluescript SK+; Vector Types:Bacterial Expression; Bacterial Resistance:Ampicillin and Kanamycin

Plasmid Name: pSK+KanaRpsL

Relevant Mutation: none

Record Creation Time: 20220422T222053+0000

Record Last Update: 20260829T080159+0000

Ratings and Alerts

No rating or validation information has been found for pSK+KanaRpsL.

No alerts have been found for pSK+KanaRpsL.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Odierna GL, et al. (2024) Drosophila larval motor patterning relies on regulated alternative splicing of Dscam2. *Frontiers in molecular neuroscience*, 17, 1415207.

Onuchic L, et al. (2023) The C-terminal tail of polycystin-1 suppresses cystic disease in a mitochondrial enzyme-dependent fashion. *Nature communications*, 14(1), 1790.

Dong K, et al. (2021) Renal plasticity revealed through reversal of polycystic kidney disease in mice. *Nature genetics*, 53(12), 1649.

Maurer GW, et al. (2020) Analysis of genes within the schizophrenia-linked 22q11.2 deletion identifies interaction of night owl/LZTR1 and NF1 in GABAergic sleep control. *PLoS genetics*, 16(4), e1008727.

Ventosa M, et al. (2017) Sustained FXN expression in dorsal root ganglia from a nonreplicative genomic HSV-1 vector. *The journal of gene medicine*, 19(11), 376.

Nakayama T, et al. (2017) no privacy, a *Xenopus tropicalis* mutant, is a model of human Hermansky-Pudlak Syndrome and allows visualization of internal organogenesis during tadpole development. *Developmental biology*, 426(2), 472.

Maharana SK, et al. (2017) Identification of novel cis-regulatory elements of Eya1 in *Xenopus laevis* using BAC recombineering. *Scientific reports*, 7(1), 15033.

Ventosa M, et al. (2017) Synergistic effects of deleting multiple nonessential elements in nonreplicative HSV-1 BAC genomic vectors play a critical role in their viability. *Gene therapy*, 24(7), 433.

Schlegel P, et al. (2016) Synaptic transmission parallels neuromodulation in a central food-intake circuit. *eLife*, 5.

Fish MB, et al. (2012) Simple, fast, tissue-specific bacterial artificial chromosome transgenesis in *Xenopus*. *Genesis (New York, N.Y. : 2000)*, 50(3), 307.