$Supported_data_table$

Supported Datasets

Dataset Name	Genotypes	Environment	Phenotype (label)	Label Type	Description	Supported
Baryshnikovna_2010	6,022	1	smf	global	growth rate	✓
$Costanzo_2016_smf$	21,718	2	smf	global	growth rate	\checkmark
$Costanzo_2016_dmf$	20,705,612	1	smf	global	growth rate	\checkmark
Kuzmin_2018	91,111	1	tmf	global	growth rate	\checkmark
SGD_essential	1,101	1	viable	global	viability	\checkmark
SynthLethalDb	1,400	1	viable	global	viability	\checkmark
$scmd2_2005$	4,718	1	ellipse (e_1, e_2)	global	cell morphology	\checkmark
$scmd2_2018$	1,112	1	ellipse (e_1, e_2)	global	cell morphology	\checkmark
$scmd2_2022$	1,982	1	ellipse (e_1, e_2)	global	cell morphology	\checkmark
ODuibhir_2014	1,312	1	smf	global, node	growth rate	\checkmark
Kemmeren_2014	1,484	1	expr	global, node	sm microarray expression	\checkmark
Sameith_2015	72	1	expr	global, node	dm microarray expression	\checkmark
Zelezniak_2018	97	1	prot abund., met conc.	global, node	sm kinase deletion mutants	\checkmark
Wildenhain_2015	195	4,915	smf	global	smf drug tolerance	\checkmark
Lian_2017	18,000	1	AID	global	AID furfural tolderance	\checkmark
FitDb	6,000	1,144	smf	global	growth rate	