									Patients		
dataset_id	type_of_query	omic_type	cancer_type Pancreatic	cohort_size	normalized	Log_transformed	ISample_type Mixed + in silico	:Technology Bulk	metadata	Truth	number_of_cell_types
DT1	All cell types	Transcriptome	adenocarcinoma	30	Yes	Yes	mixture	sequencing	No	simulation	5
DT2	All cell types	Mathyloma	Pancreatic adenocarcinoma	30	Yes	NA	Mixed + in silico mixture	Bulk array	No	simulation	5
	Immune cell	·	Colorectal				Mixed + in silico	Bulk			
DT3	types	Transcriptome	cancer	12	Yes	Yes		sequencing	No	simulation	6
DT4	All cell types	Methylome	Lung Adenocarcinoma	30	NA	NA	In silico mixture of cell lines and primary cells		Yes	simulation	5
DTF	All call true ca	, A a thrula wa a	Lung	100	NA	NA	In silico mixture of cell lines and		Voc		-
DT5	All cell types	Methylome	Adenocarcinoma	100	NA	NA	primary cells In vitro	Bulk array	Yes	simulation	5
DT6	All cell types	Transcriptome	Breast Cancer	32	Yes	Yes	mixture of	Bulk sequencing	No	In vitro	4
DT7	All cell types	Transcriptome	Breast Cancer	30	Yes	Yes	In silico mixture of cell lines and primary cells		No	simulation	4
2.,	cen cypes		2. cust candel	30	. 03	. 03	In silico mixture of cell lines and			2	
DT8	All cell types	Methylome	Breast Cancer	30	Yes	Yes	primary cells	Bulk array	No	simulation	4