

Summary of the curatedPCaData-package

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This is a summary of the MultiAssayExperiment-objects (MAE) offered for curated prostate cancer datasets in the *curatedPCaData*-package version 0.5.5. The following objects are currently available in the package:

```
> utils::data(package="curatedPCaData")$result[,c("Item", "Title")]
```

	Item	Title
[1,]	"mae_abida"	"Abida et al. MAE-object"
[2,]	"mae_barbieri"	"Barbieri MAE-object"
[3,]	"mae_chandran"	"Barwick MAE-object"
[4,]	"mae_friedrich"	"Friedrich et al. MAE-object"
[5,]	"mae_hieronymus"	"Hieronymus et al. MAE-object"
[6,]	"mae_icgcca"	"ICGC CA MAE-object"
[7,]	"mae_igc"	""
[8,]	"mae_kim"	""
[9,]	"mae_kunderfranco"	"Kunderfranco et al. MAE-object"
[10,]	"mae_ren"	"Ren 2017 MAE-object"
[11,]	"mae_sun"	"Sun et al. MAE-object"
[12,]	"mae_taylor"	"Taylor et al. MAE-object"
[13,]	"mae_tcgga"	"TCGA MAE-object"
[14,]	"mae_true"	"True et al. MAE-object"
[15,]	"mae_wallace"	"Wallace et al. MAE-object"
[16,]	"mae_wang"	""
[17,]	"mae_weiner"	"MultiAssayExperiment object containing gene expression (gex),"

They can be accessed via syntax *curatedPCaData::mae_name* or by first calling *library("curatedPCaData")* and then calling directly the objects *mae_name* in the workspace environment.