

Analyzing variant-to-function mapping

Select sample:

GSE129785_TME-All

sample_id_1 (Skin)

Traits:

BBJ_Lym_48 × CAUSALdb_BCCCEAC_F900139_11038 ×

CAUSALdb_BCCIACEAC_F900140_10855 ×

CAUSALdb_BCCM_GCST90137411_12382 ×

CAUSALdb_BCC_GD09640_11717 ×

CAUSALdb_CCSRBCB_FG03834_7097 ×

CAUSALdb_CCSRBCB_MR10820_12100 ×

CAUSALdb_CCSRBCB_PE00176_12236 ×

CAUSALdb_CRHOCTBCCN_GA475_7093 ×

CAUSALdb_LC_AT689_12050 × ... ×

Stay analysis

Reset

For example

②

①

③

Trait or disease

Basal cell carcinoma

trait_id_8687
trait_id_899
trait_id_1494
trait_id_1495
trait_id_897
trait_id_1496
trait_id_2448
trait_id_895
trait_id_894
trait_id_14561
trait_id_48
trait_id_6118
trait_id_15763

Lymphocyte

Tumor 1
Tumor 2
Tumor 3
Tumor 4
Fibroblasts

B
Th17
Memory CD8 T
Naive CD8 T
Effector CD8 T
Naive CD4 T
Th1
Endothelial
Myeloid
NK2
Plasma B
NK1
Treg
CD8 Tex
Tfh

Mean TRS
7 3.5 0

Cell type