**Camoco style sheet**

one to ten; 11+

1000

10 kb

box plot

*cis* gene

*cis* interaction

*cis*-regulatory element

co-expression

copy-number variants

dataset

dominant-negative (a)

ent-kaurene synthase

Eq. 1

Fig. 1A

joint-linkage (a)

Markov Cluster (MCL) algorithm

multigenic

null distribution

ortholog

protein-protein

*p*-value

RNA-Seq

-specific (a; as pa, no hyphen)

Supp. Figs. 1–3

*tgd1*

*t*-test

USA

vs.

wild-type (n, a)

z-score

ABC

BLUP

DELLA domain

FCR

FDR

FGS

FPKM

GA

GBS

GFF

GRAS domain

HPO, HPO+

ICP-MS

LD

MCL

MCR

NAM

PCC

PRC2

RBRs

RILs

RMIP

SNP

SRA