

## germinatebase

id int  
general\_identifier int  
number varchar(255)  
name varchar(255)  
bank\_number varchar(255)  
breeders\_code char(50)  
subtaxa\_id int  
taxonomy\_id int  
institution\_id int  
breeder\_id int  
plant\_passport varchar(255)  
donor\_code int  
donor\_number varchar(255)  
acqdate varchar(255)  
collnumb varchar(255)  
colldate date  
collcode int  
duplsite varchar(255)  
biologicalstatus\_id int  
collsrc\_id int  
location\_id int  
created\_on datetime  
updated\_on timestamp

## markers

id int  
markertype\_id int  
marker\_name varchar(45)  
created\_on datetime  
updated\_on timestamp

## genotypes

id int  
marker\_id int  
dataset\_id int  
germinatebase\_id int  
sample\_name varchar(45)  
allele1 varchar(3)  
allele2 varchar(3)  
y double  
x double

## datasets

id int  
experiment\_id int  
location\_id int  
description text  
date date  
source\_file varchar(255)  
contact varchar(255)  
version char(10)  
created\_by int  
dataset\_state\_id int  
is\_external tinyint  
hyperlink varchar(255)  
created\_on datetime  
updated\_on timestamp

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