Webside score musabase.org

Genereret Marts 26 2025 12:47 PM

Scoren er 41/100

SEO Indhold

⚠	Titel	MusaBase **Længde : 8** Kan optimeres; Optimalt bør din titel indeholde mellem 10 og 70 karakterer (med mellemrum) Brug dette gratis redskab til at regne længden ud.
✗	Beskrivelse	**Længde : 0** Meget kritisk. Vi kan ikke finde en meta beskrivelse på dit website! Brug denne gratis meta generator til at lave beskrivelser.
✗	Nøgleord	Dårligt! Vi kan ikke finde nogle meta nøgleord på din side! Brug denne gratis online meta generator for at oprette nye nøgleord.
✗	Og Meta Egenskaber	Din side benytter ikke Og egenskaberne. Disse tags tillader sociale medier at forstå din side bedre. Brug denne gratis Og generator for at oprette tags.
●	Overskrifter	**H1** 0 · **H2** 2 · **H3** 106 · **H4** 97 · **H5** 0 · **H6** 0 • [H2] Individual Crosses: • [H2] Group of Crosses: • [H3] Thank you :-) • [H3] Breeding programs • [H3] Phenotyping • [H3] East African Highland • [H3] bananas project & Partners • [H3] New to the database? • [H3] This workflow will guide you through tissue sampling an experiment • [H3] At which level do you plan to keep track of your sampling? • [H3] Select a field trial • [H3] Plant entries in your field trial • [H3] Create tissue sample entries for this trial • [H3] Complete! You have all the entities you need to conduct your sampling.

SEO Indhold

- [H3] Complete! You have all the entities you need to conduct your sampling.
- [H3] This workflow will guide you through uploading a new trial or trials into the database
- [H3] Enter information about the experiment and upload your trial layout
- [H3] Is your trial linked with other field trials, genotyping plates, or crossing experiments in the database? If you are unsure, you can skip this. This information can be added from the trial detail page after the trial is saved.
- [H3] Fixing the missing accession(s) problem
- [H3] Trial Upload Error Messages
- [H3] Fixing the missing seedlot(s) problem
- [H3] Trial Upload Error Messages
- [H3] Submit your trial again. You should have corrected all errors by now, but if not please take a look at the errors in the red box below. You can continue to modify your file and then click Upload until it works.
- [H3] There exist these problems in your file:
- [H3] Finished! Your trial is now in the database
- [H3] Finished! Your trial is now in the database
- [H3] This workflow will guide you through designing a new trial in the database
- [H3] Enter basic information about the trial
- [H3] Design your trial layout
- [H3] Is your trial linked with other field trials, genotyping plates, or crossing experiments in the database? If you are unsure, you can skip this. This information can be added from the trial detail page after the trial is saved.
- [H3] Specify the number of rows and columns for the entire field
- [H3] If you want to change the way in which plot names will be generated by the database
- [H3] Review the generated trial layout. Make sure to click Submit at the bottom of this page if you approve of the trial!
- [H3] Complete! Your trial was saved in the database.
- [H3] Complete! Your trial was saved in the database.
- [H3] This workflow will guide you through uploading genotypes into the database
- [H3] Select the type of genotyping data being uploaded
- [H3] Select the genotyping project or create a new one. A genotyping project is a specific genotyping event. You can have many genotyping projects under the same genotyping protocol to indicate that those genotyping events used the same markers.
- [H3] Provide info about the genotyping protocol used. The genotyping protocol represents a specific instance of how genotypes were called for a set of markers in a genotyping platform. Many genotyping projects can use the same genotyping protocol.
- [H3] Provide genotype information
- [H3] Finalize and submit your genotyping data
- [H3] Complete! Your genotyping data was saved in the database.
- [H3] This workflow will guide you through adding a genotyping plate in the database

SEO Indhold

- [H3] Select a genotyping project
- [H3] Provide info about your plate
- [H3] Provide information about the wells in your plate
- [H3] You want to upload an existing plate layout
- [H3] You want to upload a Coordinate Android Application file.
- [H3] You want to upload a Custom Android Application file.
- [H3] You want to design a completely new plate.
- [H3] Is your genotyping plate linked with field trials in the database? This information can also be added from the genotyping plate detail page once the trial is saved in the database.
- [H3] Finalize and submit your genotyping plate
- [H3] Complete! Your genotyping plate was saved in the database.
- [H3] Complete! Your genotyping plate was saved in the database.
- [H3] What is a seedlot inventory?
- [H3] Make sure you are collecting seedlot inventory in the following format
- [H3] Select your file and upload seedlot inventory
- [H3] Fixing the missing seedlot(s) problem
- [H3] Seedlot Inventory Upload Error Messages
- [H3] Submit your inventory again. You should have corrected all errors by now, but if not please take a look at the errors in the red box below. You can continue to modify your file and then click Upload until it works.
- [H3] There exist these problems in your file:
- [H3] Finished! Your seedlot inventory is in the database
- [H3] Finished! Your seedlot inventory is in the database
- [H3] The trial was saved to the database with no errors!
- [H3] What are seedlots?
- [H3] Seedlots fall into two categories
- [H3] Make sure your file matches the correct file format
- [H3] Provide basic information about the seedlots and upload your file
- [H3] Fix all errors in your file
- [H3] Seedlot Upload Error Messages
- [H3] Submit your seedlots again. You should have corrected all errors by now, but if not please take a look at the errors in the red box below. You can continue to modify your file and then click Upload until it works.
- [H3] There exist these problems in your file:
- [H3] Finished! Your seedlots are now in the database
- [H3] Finished! Your seedlots are now in the database
- [H3] Add the missing accessions to a list
- [H3] Introduction
- [H3] Select a crossing experiment for your crosses
- [H3] Enter basic information about the crosses and upload your file
- [H3] Additional options:
- [H3] Finished! Your crosses are now in the database
- [H3] Finished! Your crosses are now in the database
- [H3] What is a cross?
- [H3] Select a crossing experiment
- [H3] Enter basic information about the cross
- [H3] Enter basic information about the cross

SEO Indhold

- [H3] Optional: If you choose to record exact cross parents, you can do so.
- [H3] Optional: If you choose to record exact cross female parent, you can do so.
- [H3] If you would like to add auto-generated progeny names for this cross, you can add it here
- [H3] Optional:
- [H3] Finished! Your cross is now in the database
- [H3] Finished! Your cross is now in the database
- [H3] What are crossing experiments?
- [H3] Enter basic information about the crossing experiment
- [H3] Finished! Your crossing experiment is now in the database
- [H3] Finished! Your crossing experiment is now in the database
- [H3] Your Lists
- [H3] Elements not found:
- [H3] Optional: Add Missing Accessions to A List
- [H3] Mismatched case
- [H3] Multiple mismatched case
- [H3] List elements matching a synonym
- [H3] Multiple synonym matches
- [H3] Your Datasets
- [H3] Elements not found:
- [H3] Login
- [H3] Forgot Username
- [H3] Reset Password
- [H3] Create New User
- [H4] Old browser version detected
- [H4] This site is best viewed with:
- [H4] What are you interested in? For General Help
- [H4] Upload an experimental field trial into the database that you have saved on your computer in Excel
- [H4] Design a completely new experimental field trial in the database
- [H4] Catalog your available seed inventory into the database
- [H4] Upload phenotypic data into the database that you have saved on your computer in Excel
- [H4] Plan tissue sampling
- [H4] Upload crosses and crossing information into the database
- [H4] Print barcode labels for my experiment (for your plots or plants or tissue samples in the field, or for your 96 well plate and tissue samples)
- [H4] Analyze phenotypic performance across trials
- [H4] Prepare a 96 or 384 well plate for a genotyping experiment
- [H4] Upload VCF genotypic data
- [H4] Tissue Sampling
- [H4] Field trial is not relevant for the type of tissue sampling you selected. Go to next step.
- [H4] Plant entries not relevant for the type of tissue sampling you selected. Go to next step.
- [H4] Plant entries exist for this trial. Go to next step.
- [H4] Please create plant entries for this trial.
- [H4] Field trial tissue sample entries not relevant for the type of tissue sampling you selected. Go to next step.
- [H4] Tissue sample entries exist for this trial. Go to next step.
- [H4] Workflow for seedlot inventory

SEO Indhold

- [H4] I have new seedlots that need to be added into the database.
- [H4] I conducted an inventory (in weight(g)) and want to update the database to reflect the current state of the inventory.
- [H4] Workflow for uploading phenotypes
- [H4] Workflow for trial barcoding
- [H4] Workflow for comparing one or many trials
- [H4] Upload Existing Trial(s)
- [H4] Upload Template Information
- [H4] Upload Template Information
- [H4] Upload Trial Metadata
- [H4] Upload Trial Metadata Template Information
- [H4] Design New Trial
- [H4] Which accessions will be in the field?
- [H4] Which crosses will be in the field?
- [H4] Which family names will be in the field?
- [H4] Number of Plants:
- [H4] Number of Columns (required):
- [H4] Number of columns between two check columns (Optional):
- [H4] Which seedlots will you grow in the field? This is optional and can be decided later. If you do not know exactly which seedlot packets you will be planting at this time, you can add this information on the Trial Detail Page after the trial has been saved in the database.
- [H4] Add Field Management Factor to Design
- [H4] Add Field Management Factor to Design
- [H4] Partially Replicated Design Usage Help
- [H4] Background:
- [H4] Design Parameters:
- [H4] NOTE:
- [H4] Upload Genotypes
- [H4] Upload VCF Template Information
- [H4] Upload Intertek Template Information
- [H4] Upload Tassel HDF5 Template Information
- [H4] Upload SSR Marker Info Template Information
- [H4] Upload SSR Marker Info Error
- [H4] Success
- [H4] Upload SSR Protocol (Marker Info)
- [H4] Upload SSR Data Template Information
- [H4] Upload KASP data Template Information
- [H4] Add Genotyping Plate
- [H4] Upload Template Information
- [H4] Upload Template Information
- [H4] Upload Template Information
- [H4] Upload Seedlot Inventory
- [H4] Upload Template Information
- [H4] Upload Seedlots
- [H4] Upload Template Information For Named Accessions
- [H4] Upload Template Information For Harvested Seedlots
- [H4] Create New Seedlot
- [H4] OR
- [H4] Add Accessions
- [H4] Upload Accessions Template Information
- [H4] Accessions to be Added

SEO Indhold

		• [H4] Fuzzy Matches • [H4] Found Accessions • [H4] Accessions Saved • [H4] Upload Crosses • [H4] Upload Crosses File Error • [H4] Template Information • [H4] Add New Cross • [H4] Template Information • [H4] Success • [H4] Add New Crossing Experiment • [H4] Please Note: Website Data Usage Policy • [H4] MusaBase adheres to the Toronto agreement on prepublication data release • [H4] Featured Publication • [H4] Public Lists • [H4] List Contents • [H4] List Validation Report: Failed • [H4] Fuzzy Search Results • [H4] Synonym Search Results • [H4] Available Seedlots • [H4] Public Datasets • [H4] Dataset Contents • [H4] Dataset Validation Failed • [H4] Your Calendar • [H4] Add New Event • [H4] Event Info • [H4] Edit Event • [H4] Working • [H4] Progress
	Billeder	Vi fandt 49 billeder på denne side. 41 alt tags mangler eller er tomme. Tilføj alternativ tekst til dine billeder for at gøre siden mere brugervenlig, og for at optimere din SEO i forhold til søgemaskinerne.
	Text/HTML balance	Balance : **36%** Optimalt! Denne sides text til HTML fordeling er mellem 25 og 70 procent.
	Flash	Perfekt, ingen Flash objekter er blevet fundet på siden.
	iFrame	Perfekt, der er ikke nogen iFrames på din side!

SEO Links

	URL Omskrivning	Godt. Dine links ser venlige ud!

SEO Links

	Underscores i links	Dårligt! Vi har fundet underscores i dine links, du bør benytte bindestreg istedet for underscores for at optimere din SEO.
	On-page links	Vi fandt et total af 121 links inkluderende 14 link(s) til filer
	Statistics	Eksterne Links : noFollow 0% Eksterne Links : Sender Juice 28.1% Interne Links 71.9%

On-page links

Anker	Type	Juice
order	Intern	Sender Juice
MusaBase	Intern	Sender Juice
Wizard	Intern	Sender Juice
Accessions and Plots	Intern	Sender Juice
Organisms	Intern	Sender Juice
Progenies and Crosses	Intern	Sender Juice
Field Trials	Intern	Sender Juice
Genotyping Plates	Intern	Sender Juice
Genotyping Data Projects	Intern	Sender Juice
Genotyping Protocols	Intern	Sender Juice
Accessions Using Genotypes	Intern	Sender Juice
Traits	Intern	Sender Juice
Markers	Intern	Sender Juice
Images	Intern	Sender Juice
People	Intern	Sender Juice
FAQ	Intern	Sender Juice
FTP Data	Intern	Sender Juice
User Roles	Intern	Sender Juice

On-page links

Breeding Programs	Intern	Sender Juice
Locations	Intern	Sender Juice
Accessions	Intern	Sender Juice
Seed Lots	Intern	Sender Juice
Crosses	Intern	Sender Juice
Field Trials	Intern	Sender Juice
Genotyping Plates	Intern	Sender Juice
Tissue Samples	Intern	Sender Juice
Field Book App	Intern	Sender Juice
Phenotyping	Intern	Sender Juice
Barcodes	Intern	Sender Juice
Label Designer	Intern	Sender Juice
NIRS	Intern	Sender Juice
Markerset	Intern	Sender Juice
Download	Intern	Sender Juice
Upload	Intern	Sender Juice
ODK Data Collection	Intern	Sender Juice
Identifier Generation	Intern	Sender Juice
Stored Analyses	Intern	Sender Juice
Compare Trials	Intern	Sender Juice
Graphical Filtering	Intern	Sender Juice
Selection Index	Intern	Sender Juice
Genomic Selection	Intern	Sender Juice
Accession Usage	Intern	Sender Juice
Mixed Models	Intern	Sender Juice
Heritability	Intern	Sender Juice
Stability AMMI	Intern	Sender Juice
GWAS	Intern	Sender Juice

On-page links

BoxPlotter	Intern	Sender Juice
Image Analysis	Intern	Sender Juice
BLAST	Intern	Sender Juice
Ontology Browser	Intern	Sender Juice
Compose a New Trait	Intern	Sender Juice
Musa acuminata genome	Intern	Sender Juice
Nematode screening	Intern	Sender Juice
Weevil screening	Intern	Sender Juice
Sigatoka resistance screening	Intern	Sender Juice
Xvm resistance screening	Intern	Sender Juice
FOC-R1 resistances screening (Glasshouse)	Intern	Sender Juice
Phenotyping for FOC-R1	Intern	Sender Juice
Phenotyping for Sigatoka	Intern	Sender Juice
Phenotyping for BXW	Intern	Sender Juice
Sigatoka and Fusarium Collection	Intern	Sender Juice
About	Intern	Sender Juice
Contact	Intern	Sender Juice
Cite Musabase	Intern	Sender Juice
Manual	Ekstern	Sender Juice
Video tutorials	Ekstern	Sender Juice
Database statistics	Intern	Sender Juice
Forum	Intern	Sender Juice
Twitter	Ekstern	Sender Juice
Facebook	Ekstern	Sender Juice
Design and create breeding trials	Intern	Sender Juice
Upload accessions	Intern	Sender Juice
Make crosses	Intern	Sender Juice
Manage trials tutorials @ SGN	Ekstern	Sender Juice

On-page links

Use search & list tutorials @ SGN	Ekstern	Sender Juice
MGIS	Ekstern	Sender Juice
Search accessions	Ekstern	Sender Juice
Search germplasm collection	Ekstern	Sender Juice
Taxonomy browser	Ekstern	Sender Juice
The banana (Musa acuminata) genome and the evolution of monocotyledonous plants	Ekstern	Sender Juice
Browse the banana genome	Ekstern	Sender Juice
BLAST search	Ekstern	Sender Juice
Download reference genome	Ekstern	Sender Juice
NARO Uganda	Ekstern	Sender Juice
Mueller lab @BTI	Ekstern	Sender Juice
For General Help	Ekstern	Sender Juice
Single Trial Design	Intern	Sender Juice
Multiple Trial Designs	Intern	Sender Juice
Inventory	Ekstern	Sender Juice
Using Lists	Intern	Sender Juice
Uploading a File	Intern	Sender Juice
data usage policy	Intern	Sender Juice
Zoom Link	Ekstern	Sender Juice
NARO: New Banana Breed Are Disease Resistant	Ekstern	Sender Juice
See all news...	Intern	Sender Juice
Genomic Prediction in a Multiploid Crop: Genotype by Environment Interaction and Allele Dosage Effects on Predictive Ability in Banana	Ekstern	Sender Juice
See all publications...	Intern	Sender Juice
BreedBase Workshop at PAG 32	Ekstern	Sender Juice
PAG 32	Ekstern	Sender Juice
See all events...	Intern	Sender Juice
BTRACT	Ekstern	Sender Juice

On-page links

Breeding Better Bananas Project	Ekstern	Sender Juice
Southgreen Banana Genome Hub	Ekstern	Sender Juice
MGIS	Ekstern	Sender Juice
Musapedia	Ekstern	Sender Juice
Farm Radio	Ekstern	Sender Juice
IITA banana program	Ekstern	Sender Juice
Ugandan banana research portal	Ekstern	Sender Juice
Product profiles	Intern	Sender Juice
Int'l Society for Tropical Root Crops	Ekstern	Sender Juice
PDF	Intern	Sender Juice
Documentation	Ekstern	Sender Juice
Videos	Intern	Sender Juice
'+response[i].title+'	Intern	Sender Juice
'+d.seedlot[0]+'	Intern	Sender Juice
'+d.contents[0]+'	Intern	Sender Juice
" + event.title + "	Intern	Sender Juice
" + event.property + "	Intern	Sender Juice
" + event.event_url + "	Intern	Sender Juice
Export	Ekstern	Sender Juice
directory search	Intern	Sender Juice

SEO Nøgleord

Nøgleords cloud

database plot number file upload information field genotyping name trial

Nøgleords balance

Nøgleord	Indhold	Titel	Nøgleord	Beskrivelse	Overskrifter

Nøgleords balance

trial	159	✖	✖	✖	✔
file	139	✖	✖	✖	✔
database	107	✖	✖	✖	✔
name	95	✖	✖	✖	✔
field	86	✖	✖	✖	✔

Brugervenlighed

	Link	Domæne : musabase.org Længde : 12
	FavIkon	Godt, din side har et FavIcon!
	Printervenlighed	Vi kunne ikke finde en printer venlig CSS skabelon.
	Sprog	Du har ikke tildelt din side et sprog! Brug denne gratis meta tag generator til at tildele din side dit valgte sprog.
	Dublin Core	Denne side benytter IKKE Dublin Core principperne.

Dokument

	Dokumenttype	XHTML 1.0 Transitional
	Kryptering	Perfekt. Dit Charset er tildelt UTF-8.
	W3C Validering	Fejl : 38 Advarsler : 52
	Email Privatliv	Advarsel! Mindst én email adresse er blevet fundet i rå tekst. Brug gratis antispam beskytter for at hemme din email fra spammere.
	Udgået HTML	**Udgåede tags** / **Forekomster** / 110

Dokument

		<u> 2 Fejl! Vi har fundet udgåede HTML tags i din kildekode. Udgåede tags bliver ikke længere understøttet af alle browsere.
	Hastigheds Tips	✖ Advarsel! Prøv at undgå at benytte nestede tabeller i HTML. ✖ Advarsel! Din webside benytter inline CSS kode! ✖ Dårligt, din webside har for mange CSS filer (mere end 4). ✖ Dårligt, din webside har for mange JavaScript filer (mere end 6). ✔ Perfekt, din hjemmeside udnytter gzip.

Mobil

	Mobil Optimering	✖ Apple Ikon ✔ Meta Viewport Tag ✔ Flash indhold

Optimering

	XML Sitemap	Mangler Dit websted ikke har en XML sitemap - det kan være problematisk. Et sitemap lister URL'er, der er tilgængelige for gennemgang og kan indeholde yderligere oplysninger som dit websted seneste opdateringer, hyppigheden af ændringer, og betydningen af de webadresser. Dette gør det muligt søgemaskiner til at gennemgå webstedet mere intelligent.
	Robots.txt	http://musabase.org/robots.txt Stor, din hjemmeside har en robots.txt-fil.
	Analytics	Stor, din hjemmeside har et analyseværktøj. Google Analytics