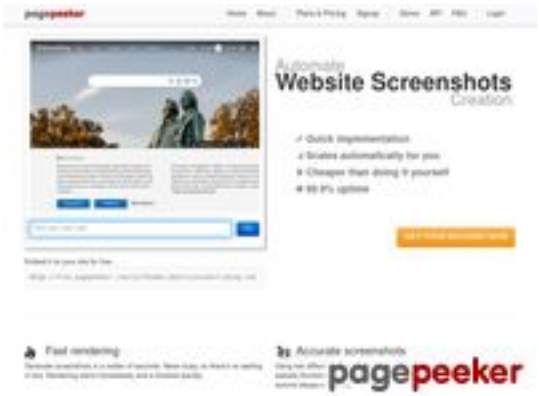

Webbplats analys musabase.org

Genereras på Mars 26 2025 12:47 PM

Ställningen är 41/100

SEO Innehåll

⚠	Titel	MusaBase **Längd : 8** Idealisk, din titel bör innehålla mellan 10 och 70 tecken (mellanslag räknas som tecken). Använd denna gratis verktyg för att räkna ut textlängden.
✗	Beskrivning	**Längd : 0** Mycket dåligt. Vi har inte lyckats hitta någon metabeskrivning på din sida. Använd denna online meta-taggar generator, gratis för att skapa beskrivningar.
✗	Nyckelord	Mycket dåligt. Vi har inte lyckats hitta några meta-taggar på din sida. Använd denna meta-tag generator, gratis för att skapa nyckelord.
✗	Og Meta Egenskaper	Den här sidan drar inte nytta utav Og. Deras taggar möjliggör sociala sökrobotar att bättre strukturera strukturera din sida. Använd denna og generatorn gratis för att skapa dom.
●	Rubriker	**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

SEO Innehåll

your sampling.
- [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

SEO Innehåll

plate in the database
- [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

SEO Innehåll

- [H3] Enter basic information about the cross
- [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.

SEO Innehåll

- [H4] Workflow for seedlot inventory
- [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

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		• [H4] Accessions to be Added • [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
	Bilder	Vi hittade 49 bilder på denna webbsida. 41 alt attribut är tomma eller saknas. Lägg till alternativ text så att sökmotorer enklare kan förstå innehållet i dina bilder.
	Text/HTML Ratio	Ratio : **36%** Idealisk! Den här sidans text till HTML-kod förhållande är mellan 25 och 70 procent.
	Flash	Perfekt, inga Flash-innehåll har upptäckts på denna sida.
	Iframe	Bra, vi upptäckte inga Iframes på den här sidan.

SEO Länkar

	URL Rewrite	Bra. Dina adressfält ser bra ut!

SEO Länkar

	Understreck i URLen	Vi har upptäckt understreck i din webbadress. Du bör hellre använda bindestreck för att optimera din SEO.
	In-page länkar	Vi hittade totalt 121 länkar inklusive 14 länk(ar) till filer
	Statistics	Externa Länkar : noFollow 0% Externa Länkar : Passing Juice 28.1% Interna Länkar 71.9%

In-page länkar

Anchor	Typ	Juice
order	Interna	Passing Juice
MusaBase	Interna	Passing Juice
Wizard	Interna	Passing Juice
Accessions and Plots	Interna	Passing Juice
Organisms	Interna	Passing Juice
Progenies and Crosses	Interna	Passing Juice
Field Trials	Interna	Passing Juice
Genotyping Plates	Interna	Passing Juice
Genotyping Data Projects	Interna	Passing Juice
Genotyping Protocols	Interna	Passing Juice
Accessions Using Genotypes	Interna	Passing Juice
Traits	Interna	Passing Juice
Markers	Interna	Passing Juice
Images	Interna	Passing Juice
People	Interna	Passing Juice
FAQ	Interna	Passing Juice
FTP Data	Interna	Passing Juice
User Roles	Interna	Passing Juice

In-page länkar

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

In-page länkar

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

In-page länkar

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

In-page länkar

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

SEO Nyckelord

Nyckelord Moln	information database plot trial file genotyping upload name number field

Nyckelord Konsistens

Nyckelord	Innehåll	Titel	Nyckelord	Beskrivning	Rubriker

Nyckelord Konsistens

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

Användbarhet

	Url	Domän : musabase.org Längd : 12
	Favikon	Bra, din webbplats har en favicon.
	Utskriftbart	Vi kunde inte hitta CSS för utskrifter.
	Språk	Du har inte angett språk. Använd denna meta-tag generator helt gratis för att ange språk på din webbplats.
	Dublin Core	Denna sida drar inte nytta utav Dublin Core.

Dokument

	Doctype	XHTML 1.0 Transitional
	Encoding	Perfekt. Din deklarerade teckenuppsättning är UTF-8.
	W3C Validity	Errors : 38 Varningar : 52
	E-post Sekretess	Varning! Minst en e-postadress har påträffats i klartext. Använd gratis antispam skydd för att dölja e-post från spammare.
	Föråldrad HTML	**Föråldrade taggar** / **Förekomster** / 110

Dokument

		<u> 2 Föråldrade HTML-taggar är HTML-taggar som inte längre används. Vi rekommenderar att du tar bort eller ersätter dessa eftersom dom nu är föråldrade.
	Hastighets Tips	✖ Varning! Försök att undvika nästlade tabeller i HTML. ✖ Synd, din webbplats använder sig utav inline stilar. ✖ Synd, din webbplats har för många CSS-filer (fler än 4 stycken). ✖ Synd, din webbplats har för många JS filer (fler än 6 stycken). ✔ Perfekt, din webbplats utnyttjar gzip.

Mobil

	Mobiloptimering	✖ Apple Ikon ✔ Meta Viewport Tagg ✔ Flash innehåll

Optimering

	XML Sitemap	Saknas Din webbplats saknar en XML sitemap - detta kan orsaka ett problem. Eftersom en sitemap listar webbadresser som är tillgängliga för genomsökning och kan innehålla ytterligare information såsom din webbplats senaste uppdatering, antal förändringar och betydelsen av webbadresserna. Detta gör det möjligt för sökmotorer att på ett mer intelligent sätt genomsöka webbplatsen.
	Robots.txt	http://musabase.org/robots.txt Bra, din webbplats har en robots.txt fil.
	Analytics	Bra, din webbplats har ett analysverktyg. Google Analytics