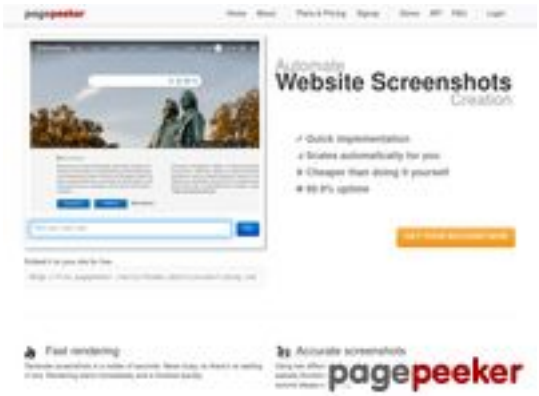

Analisi sito web musabase.org

Generato il Marzo 26 2025 12:47 PM

Il punteggio e 41/100

SEO Content

⚠	Title	MusaBase **Lunghezza : 8** Idealmente, il tuo title dovrebbe contenere tra 10 e 70 caratteri (spazi inclusi). Usa questo strumento free per calcolare la lunghezza del testo.
✗	Description	**Lunghezza : 0** Molto male. Non abbiamo trovato meta description nella tua pagina. Usa questo generatore online gratuito di meta tags per creare la descrizione.
✗	Keywords	Molto male. Non abbiamo trovato meta keywords nella tua pagina. Usa questo generatore gratuito online di meta tags per creare keywords.
✗	Og Meta Properties	Questa pagina non sfrutta i vantaggi Og Properties. Questi tags consentono ai social crawler di strutturare meglio la tua pagina. Use questo generatore gratuito di og properties per crearli.
●	Headings	**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 Content

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 Content

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 Content

- [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 Content

- [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

SEO Content

		• [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
✗	Images	Abbiamo trovato 49 immagini in questa pagina web. 41 attributi alt sono vuoti o mancanti. Aggiungi testo alternativo in modo tale che i motori di ricerca possano comprendere meglio il contenuto delle tue immagini.
✓	Text/HTML Ratio	Ratio : **36%** Ideale! Il rapporto testo/codice HTML di questa pagina e tra 25 e 70 percento.
✓	Flash	Perfetto, non e stato rilevato contenuto Flash in questa pagina.
✓	Iframe	Grande, non sono stati rilevati Iframes in questa pagina.

SEO Links

	URL Rewrite	Buono. I tuoi links appaiono friendly!

SEO Links

		
	Underscores in the URLs	Abbiamo rilevato underscores nei tuoi URLs. Dovresti utilizzare trattini per ottimizzare le pagine per il tuo SEO.
	In-page links	Abbiamo trovato un totale di 121 links inclusi 14 link(s) a files
	Statistics	External Links : noFollow 0% External Links : Passing Juice 28.1% Internal Links 71.9%

In-page links

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

In-page links

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

In-page links

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

In-page links

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

In-page links

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

SEO Keywords

Keywords Cloud	file field database trial name genotyping number information plot upload

Consistenza Keywords

Keyword	Contenuto	Title	Keywords	Description	Headings
trial	159	✖	✖	✖	✔
file	139	✖	✖	✖	✔
database	107	✖	✖	✖	✔
name	95	✖	✖	✖	✔
field	86	✖	✖	✖	✔

Usabilita

	Url	Dominio : musabase.org Lunghezza : 12
	Favicon	Grande, il tuo sito usa una favicon.
	Stampabilita	Non abbiamo riscontrato codice CSS Print-Friendly.
	Lingua	Non hai specificato la lingua. Usa questo generatore gratuito di meta tags generator per dichiarare la lingua che intendi utilizzare per il tuo sito web.
	Dublin Core	Questa pagina non sfrutta i vantaggi di Dublin Core.

Documento

	Doctype	XHTML 1.0 Transitional
	Encoding	Perfetto. Hai dichiarato che il tuo charset e UTF-8.
	Validita W3C	Errori : 38 Avvisi : 52
	Email Privacy	Attenzione! E stato trovato almeno un indirizzo mail in plain text. Usa antispam protector gratuito per nascondere gli indirizzi mail agli spammers.

Documento

Deprecated HTML	**Deprecated tags** / **Occorrenze** — 110 <u> — 2 Tags HTML deprecati sono tags HTML che non vengono piu utilizzati. Ti raccomandiamo di rimuoverli o sostituire questi tags HTML perche ora sono obsoleti.
Suggerimenti per velocizzare	✖ Attenzione! Cerca di evitare di utilizzare nested tables in HTML. ✖ Molto male, il tuo sito web utilizza stili CSS inline. ✖ Molto male, il tuo sito web ha troppi file CSS files (piu di 4). ✖ Molto male, il tuo sito web ha troppi file JS (piu di 6). ✔ Perfetto, il vostro sito si avvale di gzip.

Mobile

Mobile Optimization	✖ Apple Icon ✔ Meta Viewport Tag ✔ Flash content

Ottimizzazione

XML Sitemap	Non trovato Il tuo sito web non ha una sitemap XML - questo può essere problematico. A elenca sitemap URL che sono disponibili per la scansione e possono includere informazioni aggiuntive come gli ultimi aggiornamenti del tuo sito, frequenza delle variazioni e l'importanza degli URL. In questo modo i motori di ricerca di eseguire la scansione del sito in modo più intelligente.
Robots.txt	http://musabase.org/robots.txt Grande, il vostro sito ha un file robots.txt.
Analytics	Grande, il vostro sito ha uno strumento di analisi dei dati. Google Analytics

Ottimizzazione