

Evaluation du site musabase.org

Généré le 26 Mars 2025 12:47

Le score est de 41/100



Optimisation du contenu

	Titre	MusaBase Longueur : 8 Idéalement, votre titre devrait contenir entre 10 et 70 caractères (espaces compris). Utilisez cet outil gratuit pour calculer la longueur du texte.												
	Description	Longueur : 0 Très mauvais. Nous n'avons pas trouvé de balise META description sur votre page. Utilisez ce générateur gratuit de balises META en ligne pour créer une description.												
	Mots-clefs	Très mauvais. Nous n'avons pas trouvé de balise META keywords sur votre page. Utilisez ce générateur gratuit de balises META en ligne pour créer des mots-clés.												
	Propriétés Open Graph	Cette page ne profite pas des balises META Open Graph. Cette balise permet de représenter de manière riche n'importe quelle page dans le graph social (environnement social). Utilisez ce générateur gratuit de balises META Open Graph pour les créer.												
	Niveaux de titre	<table><tr><th>H1</th><th>H2</th><th>H3</th><th>H4</th><th>H5</th><th>H6</th></tr><tr><td>0</td><td>2</td><td>106</td><td>97</td><td>0</td><td>0</td></tr></table> • [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	H1	H2	H3	H4	H5	H6	0	2	106	97	0	0
H1	H2	H3	H4	H5	H6									
0	2	106	97	0	0									

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- [H3] Create tissue sample entries for this trial
- [H3] Complete! You have all the entities you need to conduct 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

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database.

- [H3] This workflow will guide you through adding a genotyping 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?

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- [H3] Select a crossing experiment
- [H3] Enter basic information about the cross
- [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

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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
- [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

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		• [H4] Add Accessions • [H4] Upload Accessions Template Information • [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	Nous avons trouvé 49 image(s) sur cette page Web. 41 attribut(s) alt sont vides ou manquants. Ajouter un texte alternatif permet aux moteurs de recherche de mieux comprendre le contenu de vos images.
	Ratio texte/HTML	Ratio : 36% Idéal! le ratio de cette page texte/HTML est entre 25 et 70 pour cent.
	Flash	Parfait, aucun contenu FLASH n'a été détecté sur cette page.
	Iframe	Génial, il n'y a pas d'Iframes détectés sur cette page.

Liens

	Réécriture d'URLs	Bien. Vos liens sont optimisés!
	Tiret bas dans les URLs	Nous avons détectés des soulignements dans vos URLs. Vous devriez plutôt utiliser des tirets pour optimiser votre référencement.
	Liens dans la page	Nous avons trouvé un total de 121 lien(s) dont 14 lien(s) vers des fichiers
	Statistics	Liens externes : noFollow 0% Liens externes : Passing Juice 28.1% Liens internes 71.9%

Liens dans la page

Texte d'ancree	Type	Juice
order	Interne	Passing Juice
MusaBase	Interne	Passing Juice
Wizard	Interne	Passing Juice
Accessions and Plots	Interne	Passing Juice
Organisms	Interne	Passing Juice
Progenies and Crosses	Interne	Passing Juice
Field Trials	Interne	Passing Juice
Genotyping Plates	Interne	Passing Juice
Genotyping Data Projects	Interne	Passing Juice
Genotyping Protocols	Interne	Passing Juice
Accessions Using Genotypes	Interne	Passing Juice
Traits	Interne	Passing Juice
Markers	Interne	Passing Juice
Images	Interne	Passing Juice
People	Interne	Passing Juice
FAQ	Interne	Passing Juice

Liens dans la page

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

Liens dans la page

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

Liens dans la page

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

Liens dans la page

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

Mots-clefs

	Nuage de mots-clefs	information number database genotyping upload file name field trial plot
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Cohérence des mots-clefs

Mot-clef	Contenu	Titre	Mots-clefs	Description	Niveaux de titre
trial	159	✖	✖	✖	✓
file	139	✖	✖	✖	✓
database	107	✖	✖	✖	✓
name	95	✖	✖	✖	✓
field	86	✖	✖	✖	✓

Ergonomie

	Url	Domaine : musabase.org Longueur : 12
	Favicon	Génial, votre site web dispose d'un favicon.
	Imprimabilité	Aucun style CSS pour optimiser l'impression n'a pu être trouvé.
	Langue	Vous n'avez pas précisé la langue. Utilisez ce générateur gratuit de balises META en ligne pour préciser la langue de votre site
	Dublin Core	Cette page ne profite pas des métadonnées Dublin Core.

Document

	Doctype	XHTML 1.0 Transitional
	Encodage	Parfait. Votre charset est UTF-8.
	Validité W3C	Erreurs : 38 Avertissements : 52
	E-mail confidentialité	Attention! Au moins une adresse e-mail a été trouvée en texte clair. Utilisez une protection anti-spam gratuite pour cacher vos e-mails aux spammeurs.

Document

	HTML obsolètes	Tags obsolètes	Occurrences
		<center>	110
		<u>	2
		Les balises HTML obsolètes sont des balises qui ne sont plus utilisés. Il est recommandé de supprimer ou de remplacer ces balises HTML, car elles sont désormais obsolètes.	

	Astuces vitesse	 Attention! Essayez d'éviter les tableaux imbriqués au format HTML.
		 Mauvais, votre site web utilise des styles css inline.
		 Mauvais, votre site web contient trop de fichiers CSS (plus de 4).
		 Mauvais, votre site web contient trop de fichiers javascript (plus de 6).
		 Parfait : votre site tire parti de gzip.

Mobile

	Optimisation mobile	 Icône Apple	
		 Méta tags viewport	
		 Contenu FLASH	

Optimisation

	Sitemap XML	Manquant
		Votre site web ne dispose pas d'une sitemap XML, ce qui peut poser problème. La sitemap recense les URLs que les moteurs de recherche peuvent indexer, tout en proposant d'éventuelles informations supplémentaires (comme la date de dernière mise à jour, la fréquence des changements, ainsi que leur niveau d'importance). Ceci permet aux moteurs de recherche de parcourir le site de façon plus efficace.
	Robots.txt	http://musabase.org/robots.txt Votre site dispose d'un fichier robots.txt, ce qui est optimal.
	Mesures d'audience	Votre site web dispose d'une outil d'analytics, ce qui est optimal.

Optimisation



Google Analytics