



Avaliação do site musabase.org

Gerado a 26 de Março de 2025 12:47 PM

O resultado é de 41/100



Conteúdo SEO

	Título	MusaBase Cumprimento : 8 Idealmente, o Título deve conter entre 10 e 70 caracteres (incluindo espaços).												
	Descrição	Cumprimento : 0 Mau. Não encontrámos nenhuma Descrição META na sua página.												
	Palavras-chave	Mau. Não detetámos palavras-chave META na sua página.												
	Propriedades Og Meta	Esta página não tira vantagens das propriedades Og.												
	Cabeçalhos	<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 • [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	H1	H2	H3	H4	H5	H6	0	2	106	97	0	0
H1	H2	H3	H4	H5	H6									
0	2	106	97	0	0									

Conteúdo SEO

- [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
- [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

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- [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
- [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.

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- [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
- [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

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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
- [H4] Fuzzy Matches
- [H4] Found Accessions
- [H4] Accessions Saved
- [H4] Upload Crosses

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		• [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
	Imagens	Encontrámos 49 imagens nesta página. 41 atributos ALT estão vazios ou em falta. É recomendado adicionar texto alternativo de modo a que os motores de busca identifiquem melhor o conteúdo das suas imagens.
	Rácio Texto/HTML	Rácio : 36% Ideal! O rácio de texto para código HTML desta página está entre 25 e 70 por cento.
	Flash	Perfeito, não foi encontrado conteúdo Flash nesta página.
	Iframe	Excelente, não foram detetadas Iframes nesta página.

Ligações SEO

	Reescrita de URL	Perfeito. As ligações aparentam ser limpas!
	Underscores (traços inferiores) nas URLs	Detetámos 'underscores' (traços inferiores) nas suas URLs. O uso hífen é mais eficiente em termos de otimização SEO.

Ligações SEO

	Ligações para a própria página	Encontrámos um total de 121 ligações incluindo 14 ligações a ficheiros
	Statistics	Ligações externas : noFollow 0% Ligações externas : Passa sumo 28.1% Ligações internas 71.9%

Ligações para a própria página

Âncoras	Tipo	Sumo
order	Internas	Passa sumo
MusaBase	Internas	Passa sumo
Wizard	Internas	Passa sumo
Accessions and Plots	Internas	Passa sumo
Organisms	Internas	Passa sumo
Progenies and Crosses	Internas	Passa sumo
Field Trials	Internas	Passa sumo
Genotyping Plates	Internas	Passa sumo
Genotyping Data Projects	Internas	Passa sumo
Genotyping Protocols	Internas	Passa sumo
Accessions Using Genotypes	Internas	Passa sumo
Traits	Internas	Passa sumo
Markers	Internas	Passa sumo
Images	Internas	Passa sumo
People	Internas	Passa sumo
FAQ	Internas	Passa sumo
FTP Data	Internas	Passa sumo
User Roles	Internas	Passa sumo
Breeding Programs	Internas	Passa sumo
Locations	Internas	Passa sumo

Ligações para a própria página

Accessions	Internas	Passa sumo
Seed Lots	Internas	Passa sumo
Crosses	Internas	Passa sumo
Field Trials	Internas	Passa sumo
Genotyping Plates	Internas	Passa sumo
Tissue Samples	Internas	Passa sumo
Field Book App	Internas	Passa sumo
Phenotyping	Internas	Passa sumo
Barcodes	Internas	Passa sumo
Label Designer	Internas	Passa sumo
NIRS	Internas	Passa sumo
Markerset	Internas	Passa sumo
Download	Internas	Passa sumo
Upload	Internas	Passa sumo
ODK Data Collection	Internas	Passa sumo
Identifier Generation	Internas	Passa sumo
Stored Analyses	Internas	Passa sumo
Compare Trials	Internas	Passa sumo
Graphical Filtering	Internas	Passa sumo
Selection Index	Internas	Passa sumo
Genomic Selection	Internas	Passa sumo
Accession Usage	Internas	Passa sumo
Mixed Models	Internas	Passa sumo
Heritability	Internas	Passa sumo
Stability AMMI	Internas	Passa sumo
GWAS	Internas	Passa sumo
BoxPlotter	Internas	Passa sumo
Image Analysis	Internas	Passa sumo

Ligações para a própria página

BLAST	Internas	Passa sumo
Ontology Browser	Internas	Passa sumo
Compose a New Trait	Internas	Passa sumo
Musa acuminata genome	Internas	Passa sumo
Nematode screening	Internas	Passa sumo
Weevil screening	Internas	Passa sumo
Sigatoka resistance screening	Internas	Passa sumo
Xvm resistance screening	Internas	Passa sumo
FOC-R1 resistances screening (Glasshouse)	Internas	Passa sumo
Phenotyping for FOC-R1	Internas	Passa sumo
Phenotyping for Sigatoka	Internas	Passa sumo
Phenotyping for BXW	Internas	Passa sumo
Sigatoka and Fusarium Collection	Internas	Passa sumo
About	Internas	Passa sumo
Contact	Internas	Passa sumo
Cite Musabase	Internas	Passa sumo
Manual	Externas	Passa sumo
Video tutorials	Externas	Passa sumo
Database statistics	Internas	Passa sumo
Forum	Internas	Passa sumo
Twitter	Externas	Passa sumo
Facebook	Externas	Passa sumo
Design and create breeding trials	Internas	Passa sumo
Upload accessions	Internas	Passa sumo
Make crosses	Internas	Passa sumo
Manage trials tutorials @ SGN	Externas	Passa sumo
Use search & list tutorials @ SGN	Externas	Passa sumo
MGIS	Externas	Passa sumo

Ligações para a própria página

Search accessions	Externas	Passa sumo
Search germplasm collection	Externas	Passa sumo
Taxonomy browser	Externas	Passa sumo
The banana (<i>Musa acuminata</i>) genome and the evolution of monocotyledonous plants	Externas	Passa sumo
Browse the banana genome	Externas	Passa sumo
BLAST search	Externas	Passa sumo
Download reference genome	Externas	Passa sumo
NARO Uganda	Externas	Passa sumo
Mueller lab @BTI	Externas	Passa sumo
For General Help	Externas	Passa sumo
Single Trial Design	Internas	Passa sumo
Multiple Trial Designs	Internas	Passa sumo
Inventory	Externas	Passa sumo
Using Lists	Internas	Passa sumo
Uploading a File	Internas	Passa sumo
data usage policy	Internas	Passa sumo
Zoom Link	Externas	Passa sumo
NARO: New Banana Breed Are Disease Resistant	Externas	Passa sumo
See all news...	Internas	Passa sumo
Genomic Prediction in a Multiploid Crop: Genotype by Environment Interaction and Allele Dosage Effects on Predictive Ability in Banana	Externas	Passa sumo
See all publications...	Internas	Passa sumo
BreedBase Workshop at PAG 32	Externas	Passa sumo
PAG 32	Externas	Passa sumo
See all events...	Internas	Passa sumo
BTRACT	Externas	Passa sumo
Breeding Better Bananas Project	Externas	Passa sumo
Southgreen Banana Genome Hub	Externas	Passa sumo

Ligações para a própria página

MGIS	Externas	Passa sumo
Musapedia	Externas	Passa sumo
Farm Radio	Externas	Passa sumo
IITA banana program	Externas	Passa sumo
Ugandan banana research portal	Externas	Passa sumo
Product profiles	Internas	Passa sumo
Int'l Society for Tropical Root Crops	Externas	Passa sumo
PDF	Internas	Passa sumo
Documentation	Externas	Passa sumo
Videos	Internas	Passa sumo
'+response[i].title+'	Internas	Passa sumo
'+d.seedlot[0]+'	Internas	Passa sumo
'+d.contents[0]+'	Internas	Passa sumo
" + event.title + "	Internas	Passa sumo
" + event.property + "	Internas	Passa sumo
" + event.event_url + "	Internas	Passa sumo
Export	Externas	Passa sumo
directory search	Internas	Passa sumo

Palavras-chave SEO

	Núvem de palavras-chave	name database trial information file plot genotyping field number upload
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Consistência das Palavras-chave

Palavra-chave	Conteúdo	Título	Palavras-chave	Descrição	Cabeçalhos
trial	159	✖	✖	✖	✔
file	139	✖	✖	✖	✔

Consistência das Palavras-chave

database	107	✖	✖	✖	✔
name	95	✖	✖	✖	✔
field	86	✖	✖	✖	✔

Usabilidade

	Url	Domínio : musabase.org Cumprimento : 12
	Favicon	Ótimo, o site tem um favicon.
	Facilidade de Impressão	Não encontramos CSS apropriado para impressão.
	Língua	Não foi declarada nenhuma língua para este site.
	Dublin Core	Esta página não tira vantagens do Dublin Core.

Documento

	Tipo de Documento	XHTML 1.0 Transitional						
	Codificação	Perfeito. O conjunto de caracteres UTF-8 está declarado.						
	Validação W3C	Erros : 38 Avisos : 52						
	Privacidade do Email	Aviso! No mínimo, foi encontrado um endereço de email sob a forma de texto. Isto é um convite para que spammers entupam a caixa de correio deste endereço.						
	HTML obsoleto	<table><tr><th>Etiquetas obsoletas</th><th>Ocorrências</th></tr><tr><td><center></td><td>110</td></tr><tr><td><u></td><td>2</td></tr></table> Etiquetas HTML obsoletas são etiquetas que já não são usadas segundo as normas mais recentes. É recomendado que sejam removidas ou	Etiquetas obsoletas	Ocorrências	<center>	110	<u>	2
Etiquetas obsoletas	Ocorrências							
<center>	110							
<u>	2							

Documento

		substituídas por outras etiquetas atualizadas.
	Dicas de Velocidade	 Atenção! Existem tabelas dentro de outras tabelas em HTML.  Oh não, o site usa estilos CSS nas etiquetas HTML.  Oh, não! O site utiliza demasiados ficheiros CSS (mais que 4).  Oh, não! O site utiliza demasiados ficheiros JavaScript (mais que 6).  Perfeito, o site tira vantagens da compressão gzip.

Dispositivos Móveis

	Otimização para dispositivos móveis	 Icon Apple  Meta Viewport Tag  Conteúdo Flash
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Otimização

	XML Sitemap	Em falta O site não tem um mapa XML do site (sitemap) - isto pode ser problemático. Um mapa do site identifica todas as URLs que estão disponíveis para rastreio, incluindo informação acerca de atualizações, frequência de alterações ou a importancia de cada URL. Isto contribui para uma maior inteligência e eficiência do rastreio.
	Robots.txt	http://musabase.org/robots.txt Perfeito, o seu site tem um ficheiro robots.txt.
	Analytics	Perfeito, o site tem uma ferramenta analítica para a análise de atividade.  Google Analytics