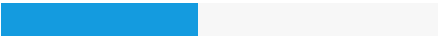


Revisión web de musabase.org

Generado el 26 Marzo 2025 12:47 PM

La puntuación es 41/100



Contenido SEO

	Título	MusaBase Longitud : 8 Preferiblemente, tu título debería contener entre 10 y 70 caracteres (espacios incluidos). Usa esta herramienta gratuita para calcular la longitud del texto.												
	Descripción	Longitud : 0 Muy mal. No hemos encontrado descripción meta en tu página. Usa este generador online gratuito para crear la descripción.												
	Palabras Claves (Keywords)	Muy mal. No hemos encontrado palabras clave (meta keywords) en tu página. Usa este generador de meta tags gratuito para crear tus palabras clave.												
	Propiedades Meta Og	Esta página no usa etiquetas Og. Estas etiquetas permiten a los rastreadores sociales estructurar mejor tu página. Usa este generador de etiquetas og gratuito para crearlas.												
	Titulos	<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	H1	H2	H3	H4	H5	H6	0	2	106	97	0	0
H1	H2	H3	H4	H5	H6									
0	2	106	97	0	0									

Contenido SEO

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

Contenido SEO

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

Contenido SEO

- [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.

Contenido SEO

- [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

Contenido SEO

		• [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
	Imágenes	Hemos encontrado 49 imágenes en esta web. 41 atributos alt están vacíos o no existen. Agrega texto alternativo para que los motores de búsqueda puedan entender las imágenes.
	Ratio Texto/HTML	Ratio : 36% Ideal! El ratio entre texto y código HTML de esta página está entre 25 y 70 por ciento.
	Flash	Perfecto, no se ha detectado contenido Flash en la página.
	Iframe	Genial, no se han detectado Iframes en la página.

Enlaces SEO

	Reescritura URL	Bien. Tus enlaces parecen amigables
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Enlaces SEO

	Guiones bajos en las URLs	Hemos detectado guiones bajos en tus URLs. Deberías usar guiones en su lugar para optimizar tu SEO.
	Enlaces en página	Hemos encontrado un total de 121 enlaces incluyendo 14 enlace(s) a ficheros
	Statistics	Enlaces Externos : noFollow 0% Enlaces Externos : Pasando Jugo 28.1% Enlaces Internos 71.9%

Enlaces en página

Ancla	Tipo	Jugo
order	Interna	Pasando Jugo
MusaBase	Interna	Pasando Jugo
Wizard	Interna	Pasando Jugo
Accessions and Plots	Interna	Pasando Jugo
Organisms	Interna	Pasando Jugo
Progenies and Crosses	Interna	Pasando Jugo
Field Trials	Interna	Pasando Jugo
Genotyping Plates	Interna	Pasando Jugo
Genotyping Data Projects	Interna	Pasando Jugo
Genotyping Protocols	Interna	Pasando Jugo
Accessions Using Genotypes	Interna	Pasando Jugo
Traits	Interna	Pasando Jugo
Markers	Interna	Pasando Jugo
Images	Interna	Pasando Jugo
People	Interna	Pasando Jugo
FAQ	Interna	Pasando Jugo
FTP Data	Interna	Pasando Jugo
User Roles	Interna	Pasando Jugo

Enlaces en página

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

Enlaces en página

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

Enlaces en página

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

Enlaces en página

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

Palabras Clave SEO

	Nube de Palabras Clave	file database plot upload information field genotyping number trial name
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Consistencia de las Palabras Clave

Palabra Clave (Keyword)	Contenido	Título	Palabras Claves (K)	Descripción	Titulos
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Consistencia de las Palabras Clave

			eywords)		
trial	159	✖	✖	✖	✓
file	139	✖	✖	✖	✓
database	107	✖	✖	✖	✓
name	95	✖	✖	✖	✓
field	86	✖	✖	✖	✓

Usabilidad

	Url	Dominio : musabase.org Longitud : 12
	Favicon	Genial, tu web tiene un favicon.
	Imprimibilidad	No hemos encontrado una hoja de estilos CSS para impresión.
	Idioma	No has especificado un idioma. Usa este generador de meta tags gratuito para declarar el idioma de tu sitio.
	Dublin Core	Esta página no usa Dublin Core.

Documento

	Tipo de documento (Doctype)	XHTML 1.0 Transitional		
	Codificación	Perfecto. Has declarado como codificación UTF-8.		
	Validez W3C	Errores : 38 Avisos : 52		
	Privacidad de los Emails	Atención! Hemos encontrado por lo menos una dirección de correo electrónico en texto plano. Usa este protector antispam gratuito para ocultarla de los spammers.		
	HTML obsoleto	<table><tr><th>Etiquetas obsoletas</th><th>Ocurrencias</th></tr></table>	Etiquetas obsoletas	Ocurrencias
Etiquetas obsoletas	Ocurrencias			

Documento



<center>	110
<u>	2

Hemos detectado etiquetas HTML obsoletas que ya no se usa. Es recomendable que cambies esas etiquetas por otras similares.



Consejos de Velocidad

 Atención! Intenta evitar el uso de tablas en HTML.

 Muy mal, tu web está usando estilos embenidos (inline CSS).

 Muy mal, tu página web usa demasiados ficheros CSS (más de 4).

 Muy mal, tu sitio usa demasiados ficheros JavaScript (más de 6).

 Su sitio web se beneficia del tipo de compresión gzip. ¡Perfecto!

Movil

	Optimización Móvil	 Icono para Apple  Etiqueta Meta Viewport  Contenido Flash
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Optimización

	Mapa del sitio XML	No disponible Su sitio web no tiene un mapa del sitio XML. Esto puede traerle problemas. Un mapa del sitio enumera las URL que pueden rastrearse y puede incluir información adicional, como las últimas actualizaciones de su sitio web, la frecuencia de cambios y la importancia de las URL. Esto permite a los motores de búsqueda rastrear el sitio web de una forma más exhaustiva.
	Robots.txt	http://musabase.org/robots.txt ¡Estupendo! Su sitio web tiene un archivo robots.txt.
	Herramientas de Analítica	¡Perfecto! Su sitio web tiene una herramienta de análisis.  Google Analytics

Optimización