



.periodicos.

- Tutorial de **Acesso** .



. Engineering Village .



Olá! Eu sou o Carlos e, a partir de agora,
você descobrirá como aproveitar a
plataforma Engineering Village.
Vamos lá?



A **Engineering Village** é uma plataforma de busca poderosa, essencial para pesquisadores, docentes e estudantes, e que usa um amplo leque de recursos que inclui publicações, procedimentos de conferência, dissertações, **publicações especializadas, patentes, relatórios governamentais** e outros.

Pesquise e acesse informações importantes relacionadas a **engenharia** – resultados dentro de um contexto - de maneira conveniente, rápida e confiável, usando a plataforma líder de busca de informações de engenharia.

A **Engineering Village** é validada e indexada por especialistas da área. Dessa forma, os pesquisadores podem ser mais produtivos e facilitar de maneira rápida as descobertas importantes.

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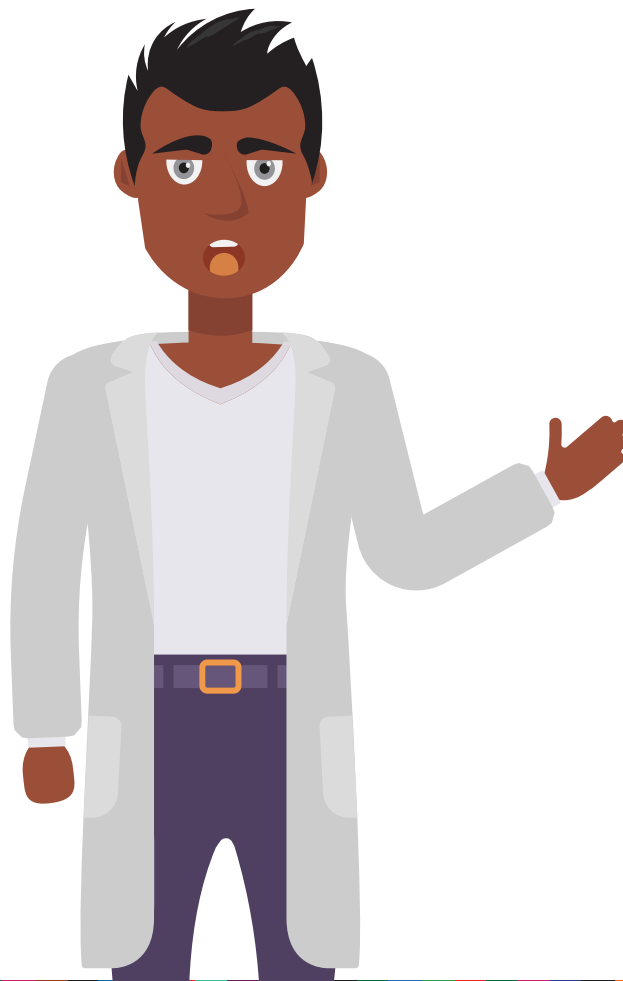
Search

Pesquise por uma frase exata usando aspas ou colchetes:

“rocket propulsion laboratory”
{rocket propulsion laboratory}

Pesquise dentro de um campo específico usando WN
“wearable technology” wn TI and
video wn AB





Booleano e conectores

AND - quando os termos coexistem em um documento ou campo. "AND" restringe o número de resultados obtidos de documentos.

OR - cada termo pode existir separadamente em um documento ou campo. "OR" expande o número de resultados obtidos de documentos.

NOT - exclui termos de um documento ou campo.

NEAR - pesquisa por termos próximos.

ONEAR especifica a ordem exata dos termos.

NEAR ou **ONEAR** não podem ser usados com truncamentos, caracteres curingas, parênteses, chaves ou aspas.

solar NEAR energy

wind NEAR/3 power

Energy ONEAR/0 policy

Use parênteses para fazer uma pesquisa composta ou termos booleanos agrupados

("jet propulsion" OR "rocket propulsion") AND engine

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Dicas adicionais

A Engineering Village não diferencia maiúsculas de minúsculas. As consultas podem ser inseridas em qualquer capitalização ou mistura de capitalização.

Campos mais usados:

AU – author ST – serial title (journal name)

TI – title KY – subject/title/abstract

AB – abstract CV – controlled term (index/thesaurus term)

AF – author affiliation LA – language

CO – country of publication YR – year

O arquivo Engineering Village Help completo pode ser visualizado no menu “Support” (“Suporte”), na barra de navegação superior.

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[Register](#) | [Login](#) | [End Session](#)

Selecione uma ou mais bases de dados. Somente as bases de dados com uma assinatura institucional serão listadas!

Selected records | Settings | Tags & Groups

Support | Ask an expert

Search Expert Search Thesaurus Search

Search History (3)

Databases | Search Tips

DATABASE ☒ Compendex

SEARCH FOR

AND		in	All fields
		in	All fields
AND		in	All fields

Turn Off AutoSuggest | Add search field

ADVANCED OPTIONS

LIMIT TO

All document types
All treatment types
All Languages
1961 TO 2018
1 Updates

SORT BY

☒ Relevance ☐ Date (Newest)

☐ Autostemming off

Search

Browse Indexes

- Author
- Author affiliation
- Controlled term

Digite as palavras-chave na caixa de pesquisa. Utilize os operadores AND, OR e NOT do menu suspenso para restringir ou ampliar os resultados da pesquisa. Clique nas setas "All fields" para habilitar campos específicos na pesquisa (Author, Author affiliation, Controlled term, Source title, etc.). Clique em "Add search field" para adicionar mais linhas (máximo de 12).

Limitar as pesquisas a um tipo de conteúdo específico dependerá da base de dados selecionada e dos campos usados naquela base de dados. Utilize as setas suspensas para visualizar a lista de limitações para cada campo. Pode-se limitar os campos por tipo de documento (publicações de artigos, procedimentos), tipo de texto (crítica, bibliografia), tipo de disciplina (física, computação) e idioma.

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[Who uses EV?](#)
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Quick Search | [Expert Search](#)

DATABASE SEARCH FOR

☒ Comp

AND

AND

ADVANCED OPTIONS

LIMIT TO

All document types

All treatment types

All Languages

☒ 1961 TO 2011

☐ 1 Updates

SORT BY

☒ Relevance ☐ Date (Newest)

☐ Autostemming off

Search History (3)

Browse Indexes

- Author
- Author affiliation
- Controlled term
- Source title
- Publisher

Latest Resources

More Sources

Interactive Equations and Tools

Powered by Knovel

Uma lista das 50 pesquisas mais recentes feitas durante a seção aparece na caixa "Search history" na parte inferior das páginas Quick Search, Expert Search e Thesaurus Search. As pesquisas podem ser combinadas usando as caixas de seleção e as setas de mover para baixo.

As pesquisas também podem ser editadas, salvas ou usadas para um alerta. Caso uma pesquisa salva ou um alerta sejam criados, o usuário deverá se registrar e fazer login. O histórico de pesquisa é excluído quando o usuário encerra a sessão.

Os resultados podem ser classificados por relevância ou ano de publicação. A função "autostemming" permite a pesquisa por um radical e por palavras formadas com outros possíveis sufixos. Por exemplo, caso o termo "control" seja pesquisado usando essa função, os resultados da lista conterão: controllers, control, controlling, controlled, controls. Este recurs permite a pesquisa de diferentes variações de uma palavra.

Dependendo da base de dados selecionada, a caixa Browse Indexes aparecerá no lado direito da tela. Os usuários podem localizar um autor específico, editor, título do artigo ou termos de índice usados dentro de uma base de dados. Ao pesquisar por termos de índice, os usuários podem localizar e obter resultados altamente precisos.

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Quick Search | Expert Search | Thesaurus Search | Search History (3)

DATABASE SEARCH FOR

☒ **Compindex**

in All fields

in All fields

in All fields

ADVANCED OPTIONS

LIMIT TO

All document types

All treatment types

All Languages

196 TO 201

1 Updates

SORT BY

☒ Relevan

☐ Autoste

Search Reset

Links para fontes de conteúdo de engenharia adicionais disponíveis em seu instituto.

- Browse Indexes**
- Author
 - Author affiliation
 - Controlled term
 - Source title
 - Publisher

Links para materiais de treinamento adicionais, vídeos, notícias e links para feedbacks.

- Latest Resources**
- More Sources**
- Interactive Equations and Tools**
- Powered by Knovel

As ferramentas e equações interativas Powered by Knovel contêm mais de 500 equações interativas de engenharia, além de um conversor de unidades e uma tabela periódica. Assinantes Knovel possuem acesso a todas as mais de 500 equações e os assinantes não Knovel podem usar 125 delas.



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O número de resultados da pesquisa é exibido, seguido por uma lista de bases de dados pesquisadas e um link clicável para a consulta de pesquisa usada.

Iniciar uma nova pesquisa, editar uma pesquisa, salvar uma pesquisa para mais tarde, criar um alerta de pesquisa, definir um feed RSS ou visualizar histórico de pesquisa.

The screenshot shows the Engineering Village search interface. At the top, there's a navigation bar with 'Search', 'Selected records', 'Settings', 'Tags & Groups', 'Support', and 'Ask an expert'. Below this is a search bar with 'transcription factors' entered. The search results are displayed in a list format, showing 22308 articles found in Compendex for 1969-2018. The results are sorted by Relevance. The first result is 'Discovery of WRKY transcription factors through transcriptome analysis and characterization of a novel methyl jasmonate-inducible PqWRKY1 gene from Panax quinquefolius'. The second result is 'Feature-based classification of human transcription factors into hypothetical sub-classes related to regulatory function'. The third result is 'Comparative in Silico evaluation of MYB transcription factors in eucalyptus, sugarcane and rice transcriptomes'. The fourth result is 'GATA transcription factors regulate the expression of the human eosinophil-derived neurotoxin (RNase 2) gene'. The fifth result is 'Functions of heat shock transcription factors involved in response to photooxidative stresses in arabidopsis'. The sixth result is 'Determinants of the DNA binding specificity of class I and class II TCP transcription factors'. The seventh result is 'Are foxO transcription factors implicated in osteoarthritis? Influence of Diacerheind'. The left sidebar contains filters for 'Refine results', 'Add a term', 'Controlled vocabulary', 'Author', 'Author affiliation', 'Classification code', 'Country', 'Document type', 'Language', 'Year', 'Source title', 'Publisher', 'Funding sponsor', and 'Run new search with selected facets'. The URL at the bottom is 'https://www-engineeringvillage-com.ez1.periodicos.capes.gov.br/search#'.

Search | Selected records | Settings | Tags & Groups | Support | Ask an expert

Quick Search | Expert Search | Thesaurus Search | Search History (4)

DATABASE: Compendex
SEARCH FOR: transcription factors in All fields
Turn Off AutoSuggest | Add search field | Search

22308 articles found in Compendex for 1969-2018: (transcription factors) WN All fields

New search | Edit | Save search | Create Alert | RSS feed | Search history

Numeric Filter | Display: 25 results per page

Select: Selected Records (0) | Remove all Selected Records
Email | Print | Download

Sort by: Relevance

1. **Discovery of WRKY transcription factors through transcriptome analysis and characterization of a novel methyl jasmonate-inducible PqWRKY1 gene from Panax quinquefolius**
Sun, Yongzhen (The Key Laboratory of Bioactive Substances and Resources Utilization of Chinese Herbal Medicine, Ministry of Education, Institute of Medicinal Plant Development (IMPLAD), Chinese Academy of Medical Sciences, Peking Union Medical College, Beijing, 100193, China); Niu, Yunyun; Xu, Jiang; Li, Ying; Luo, Hongmei; Zhu, Yingjie; Liu, Mingzhu; Wu, Qiong; Song, Jingyuan; Sun, Chao; Chen, Shilin | Source: *Plant Cell, Tissue and Organ Culture*, v 114, n 2, p 269-277, August 2013
Database: Compendex
Detailed | Show preview | Cited by in Scopus (17) | Full Text
2. **Feature-based classification of human transcription factors into hypothetical sub-classes related to regulatory function**
Ehsani, Rezvan (Norwegian University of Science and Technology, Department of Cancer Research and Molecular Medicine, PO Box 8905, Trondheim; NO-7491, Norway); Bahrami, Shahram; Drabes, Finn | Source: *BMC Bioinformatics*, v 17, n 1, November 14, 2016
Database: Compendex
Detailed | Show preview | Full Text
3. **Comparative in Silico evaluation of MYB transcription factors in eucalyptus, sugarcane and rice transcriptomes**
Soares-Cavalcanti, Nina M. (Genetics Department, Laboratory for Plant Genetics and Biotechnology, Universidade Federal de Pernambuco, Rua Prof. Moraes Rêgo s/n, Recife, PE, Brazil); Wanderley-Nogueira, Ana C.; Belamino, Luis C.; Dos Santos Barros, Petra; Benko-Iseppon, Ana M. | Source: *Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics)*, v 5488 LNBI, p 44-55, 2009, *Computational Intelligence Methods for Bioinformatics and Biostatistics - 5th International Meeting, CIBB 2008, Revised Selected Papers*
Database: Compendex
Detailed | Show preview | Cited by in Scopus (4) | Full Text
4. **GATA transcription factors regulate the expression of the human eosinophil-derived neurotoxin (RNase 2) gene**
Oiu, Zhijun (Laboratory of Allergic Diseases, NIAID, National Institutes of Health, 9000 Rockville Pike, Bethesda, MD 20892, United States); Dyer, Kimberly D.; Xia, Zhihui; Rädinger, Madeleine; Rosenberg, Helene F. | Source: *Journal of Biological Chemistry*, v 284, n 19, p 13099-13109, May 8, 2009
Database: Compendex
Detailed | Show preview | Cited by in Scopus (13) | Full Text
5. **Functions of heat shock transcription factors involved in response to photooxidative stresses in arabidopsis**
Yabuta, Yukinori (Faculty of Agriculture, School of Agricultural, Biological, and Environmental Sciences, Tottori University, Tottori, Japan) | Source: *Bioscience, Biotechnology and Biochemistry*, v 80, n 7, p 1254-1263, 2016
Database: Compendex
Detailed | Show preview | Full Text
6. **Determinants of the DNA binding specificity of class I and class II TCP transcription factors**
Viola, Ivana L. (Instituto de Agrobiotecnología del Litoral (CONICET-UNL), Facultad de Bioquímica Y Ciencias Biológicas, Universidad Nacional del Litoral, CC 242 Paraje El Pozo, 3000 Santa Fe, Argentina); Reinheimers, Renata; Ripoll, Rodrigo; Uberti Manassero, Nora G.; Gonzalez, Daniel H. | Source: *Journal of Biological Chemistry*, v 287, n 1, p 347-356, January 2, 2012
Database: Compendex
Detailed | Show preview | Cited by in Scopus (17) | Full Text
7. **Are foxO transcription factors implicated in osteoarthritis? Influence of Diacerheind**
De Isla, N. (UMR CNRS 7561, Faculté de Médecine, 54500 Vandoeuvre, France); Charif, N.; Stoltz, J.F. | Source: *Bio-Medical Materials and Engineering*, v 20, n 3-4, p 227-233, 2010, *Papers from the 5th Scientific Meeting on Cartilage Engineering, February 2010, Nancy, France*

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O botão “Limit to” restringe os resultados da pesquisa para que inclua apenas termos selecionados a partir das categorias, e o botão “Excluído” elimina os termos das categorias selecionadas. Utilize o botão de pesquisa em “Run new search with selected facets” para ignorar a pesquisa original e usar apenas os termos escolhidos usando as categorias.

Após finalizar uma pesquisa, uma lista de categorias é exibida no lado esquerdo da página de resultados da pesquisa. Cada categoria permite que os usuários modifiquem as consultas de pesquisa. A ordem das caixas pode ser modificada clicando e arrastando a caixa para cima ou para baixo. As categorias permanecerão na nova ordem para os usuários que se registrarem e fizerem logon. As categorias exibidas dependem das bases de dados.

Search | Selected records | Settings | Tags & Groups Support Ask an expert

Quick Search Expert Search Thesaurus Search Search History (4)

DATABASE: Compendex SEARCH FOR: transcription factors in All fields Turn Off AutoSuggest Add search field Search

22308 articles found in Compendex for 1969-2018: ((transcription factors) WN All fields)

Numeric Filter Refine results Limit to Exclude Add a term Controlled vocabulary Genes (7586) Proteins (5909) Transcription (5294) Transcription Factors (5074) Cells (4070) View more

Author Yamamoto, Masayuki (40) Van, Wijnan Andre J. (30) Conway, Ronald C. (28) Stein, Gary S. (28) Shinozaki, Kazuo (28) View more

Author affiliation Classification code Country Document type Language Year Source title Publisher Funding sponsor Limit to Exclude Run new search with selected facets Search

Display: 25 results per page Go to page: 1 of 893 Go

Select: Selected Records (0) Remove all Selected Records Email Print Download Sort by: Relevance

- Discovery of WRKY transcription factors through transcriptome analysis and characterization of a novel methyl jasmonate-inducible PqWRKY1 gene from *Panax quinquefolius*
Sun, Yongzhen (The Key Laboratory of Bioactive Substances and Resources Utilization of Chinese Herbal Medicine, Ministry of Education, Institute of Medicinal Plant Development (IMPLAD), Chinese Academy of Medical Sciences, Peking Union Medical College, Beijing, 100193, China); Niu, Yunyun; Xu, Jiang; Li, Ying; Luo, Hongmei; Zhu, Yingjie; Liu, Mingzhu; Wu, Giong; Song, Jingyuan; Sun, Chao; Chen, Shilin Source: *Plant Cell, Tissue and Organ Culture*, v 114, n 2, p 269-277, August 2013
Database: Compendex
Detailed | Show preview | Cited in Scopus (17) | Full Text
- Feature-based classification of human transcription factors into hypothetical sub-classes related to regulatory function
Ehsani, Rezvan (Norwegian University of Science and Technology, Department of Cancer Research and Molecular Medicine, PO Box 8905, Trondheim; NO-7491, Norway); Bahrami, Shahram; Drables, Finn Source: *BMC Bioinformatics*, v 17, n 1, November 14, 2016
Database: Compendex
Detailed | Show preview | Full Text
- Comparative in Silico evaluation of MYB transcription factors in eucalyptus, sugarcane and rice transcriptomes
Soares-Cavalcanti, Nina M. (Genetics Department, Laboratory for Plant Genetics and Biotechnology, Universidade Federal de Pernambuco, Rua Prof. Moraes Rêgo s/n., Recife, PE, Brazil); Wanderley-Nogueira, Ana C.; Belamino, Luis C.; Dos Santos Barros, Petra; Benko-Iseppon, Ana M. Source: *Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics)*, v 5488 LNBI, p 44-55, 2009, *Computational Intelligence Methods for Bioinformatics and Biostatistics - 5th International Meeting, CIBB 2008, Revised Selected Papers*
Database: Compendex
Detailed | Show preview | Cited in Scopus (4) | Full Text
- GATA transcription factors regulate the expression of the human eosinophil-derived neurotoxin (RNase 2) gene
Qiu, Zhihui (Laboratory of Allergic Diseases, NIAID, National Institutes of Health, 9000 Rockville Pike, Bethesda, MD 20892, United States); Dyer, Kimberly D.; Xie, Zhihui; Rådinger, Madeleine; Rosenberg, Helene F. Source: *Journal of Biological Chemistry*, v 284, n 19, p 13099-13109, May 8, 2009
Database: Compendex
Detailed | Show preview | Cited in Scopus (13) | Full Text
- Functions of heat shock transcription factors involved in response to photooxidative stresses in arabidopsis
Yabuta, Yukinori (Faculty of Agriculture, School of Agricultural, Biological, and Environmental Sciences, Tottori University, Tottori, Japan) Source: *Bioscience, Biotechnology and Biochemistry*, v 80, n 7, p 1254-1263, 2016
Database: Compendex
Detailed | Show preview | Full Text
- Determinants of the DNA binding specificity of class I and class II TCP transcription factors
Viola, Ivana L. (Instituto de Agrobiotecnología del Litoral (CONICET-UNL), Facultad de Bioquímica Y Ciencias Biológicas, Universidad Nacional del Litoral, CC 242 Paraje El Pozo, 3000 Santa Fe, Argentina); Reinheims, Renata; Ripoll, Rodrigo; Uberti Manassero, Nora G.; Gonzalez, Daniel H. Source: *Journal of Biological Chemistry*, v 287, n 1, p 347-356, January 2, 2012
Database: Compendex
Detailed | Show preview | Cited in Scopus (17) | Full Text
- Are for O2 transcription factors implicated in osteoarthritis? Influence of Diacerhermid
De Isla, N. (UMR CNRS 7561, Faculté de Médecine, 54500 Vandoeuvre, France); Charif, N.; Stoltz, J.F. Source: *Bio-Medical Materials and Engineering*, v 20, n 3-4, p 227-233, 2010, *Papers from the 5th Scientific Meeting on Cartilage Engineering, Coimbra, Portugal, 2010*

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Selecione todos os registros da página ou os primeiros 500 registros da lista de resultados usando a caixa “Select” e a seta de mover para baixo. Selecionar os registros permite que os usuários possam usar as opções de Email, Print, Download, ou Save com eles. O número máximo de registros que podem ser selecionados em qualquer momento é de 500.

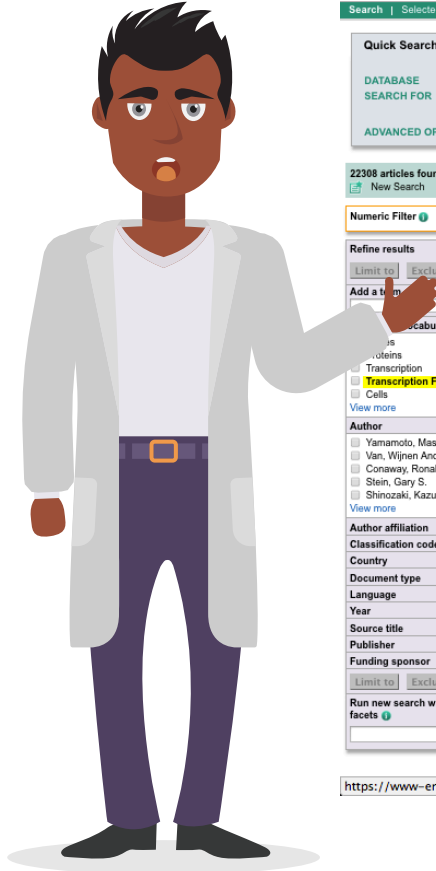
Utilize a caixa “Add a term” para inserir um termo e refinar os resultados, escolhendo:

- O botão “Limit to” para pesquisar o termo.
- O botão “Exclude” para excluir o termo.
- Selecione um ou mais itens nas categorias, clique em “Limit to” ou “Exclude”

The screenshot displays the Engineering Village search results page. At the top, there are navigation links for 'Settings', 'Tags & Groups', 'Support', and 'Ask an expert'. Below these are tabs for 'Search', 'Thesaurus Search', and 'Search History (4)'. A search bar contains the text 'transcription factors' and a 'Search' button. Below the search bar, there are options to 'Turn Off AutoSuggest', 'Add search field', and 'Search history'. The results are displayed in a list format, with a 'Display: 25 results per page' option. On the left side, there is a 'Refine results' panel with a 'Limit to' button and an 'Add a term' input field. Below this, there are categories like 'Controlled vocabulary' and 'Author' with checkboxes and counts. The main results area shows a list of articles, each with a title, author, and source. The first article is 'Discovery of WRKY transcription factors through transcriptome analysis' by Sun, Yongzhen et al. The second article is 'Feature-based classification of human transcription factors into hypothetical sub-classes related to regulatory function' by Ehsani, Rezvan et al. The third article is 'Comparative in Silico evaluation of MYB transcription factors in eucalyptus, sugarcane and rice transcriptomes' by Soares-Cavalcanti, Nina M. et al. The fourth article is 'GATA transcription factors regulate the expression of the human eosinophil-derived neurotoxin (RNase 2) gene' by Qiu, Zhiyun et al. The fifth article is 'Functions of heat shock transcription factors involved in response to photooxidative stresses in arabidopsis' by Yabuta, Yukinori et al. The sixth article is 'Determinants of the DNA binding specificity of class I and class II TCP transcription factors' by Viola, Ivana L. et al. The seventh article is 'Are foxO transcription factors implicated in osteoarthritis? influence of Diacerhein' by De Isla, N. et al.

<https://www-engineeringvillage-com.ez1.periodicos.capes.gov.br/search#>

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Search | Selected records | Settings | Tags & Groups Support Ask an expert

Quick Search | Expert Search | Thesaurus Search | Search History (4)

DATABASE ☒ Compendex
SEARCH FOR transcription factors in All fields
Turn Off AutoSuggest | Add search field

22308 articles found in Compendex for 1969-2018: ((transcription factors) WN All fields)
New Search | Edit | Save Search | Create Alert | RSS feed | Search history

Numeric Filter
Refine results
Add a term
Transcription Factors (5074)
Cells (4070)
Author
Yamamoto, Masayuki (40)
Van, Wijnen Andre J. (30)
Conaway, Ronald C. (28)
Stein, Gary S. (28)
Shinozaki, Kazuo (28)
Author affiliation
Classification code
Country
Document type
Language
Year
Source title
Publisher
Funding sponsor
Run new search with selected facets
Search

Display: 25 results per page
Select: ☒ Selected Records (0)
☒ Email
1. Discovery of WRKY transcription factors through tr
Sun, Yongzhen (The Key Laboratory of Bioactive Substar
College, Beijing, 100193, China); Niu, Yunyun; Xu, Jlang;
Database: Compendex
Detailed | Show preview | Cited by in Scopus (17) | Full Text
2. Feature-based classification of human transcription factors into functional sub-classes related to regulatory function
Ehsani, Razvi
November 14,
Database: Co
Detailed |
3. Comparative
Soares-Cavali
Barros, Petra;
Bioinformatics
Database: Co
Detailed |
4. GATA transcr
Qiu, Zhiyun (L
Chemistry, v 2
Database: Co
Detailed |
5. Functions of
Yabuta, Yukin
Database: Co
Detailed |
6. Determinants
Viola, Ivana L.
Manassero, N
Database: Co
Detailed |
7. Are foxO tra
De Iella, N. (U
Database: Co
Detailed |

Na página de resultados, os usuários podem usar as opções de e-mail, Print, Download, ou Save com os resultados. É necessário se registrar para criar pastas.

É possível baixar os registros diretamente para um computador ou para o Mendeley, Google Drive, DropBox e RefWorks. Os formatos disponíveis incluem texto sem formatação, RefWorks, BibTex, formatos Excel, PDF, RTF ou RIS (o formato RIS é compatível com o EndNote, ProCite e Reference Manager). Caso uma ferramenta de gerenciamento de referências como o Mendeley seja usada, é necessário que ela seja instalada antes de baixar os registros. Os usuários inscritos podem salvar suas preferências de download para cada vez que fizerem login na Engineering Village.

1. Selecione os registros para baixar usando as caixas de seleção próximas a cada registro.
2. Clique em "Save to PC" acima dos resultados de pesquisa. O menu Download Settings é exibido.
3. Selecione o formato e o local para o download.
4. Clique em 'Download'

Go to page: 1 of 893 https://www-engineeringvillage-com.ez1.peric

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Search | Selected records | Settings | Tags & Groups Support | Ask an expert

Quick Search Expert Search Thesaurus Search Search History (4)

DATABASE ☒ Compendex
SEARCH FOR transcription factors in All fields
Turn Off AutoSuggest | Add search field | Search

22308 articles found in Compendex for 1969-2018: ((transcription factors) WN All fields)
New Search | Edit | Save Search | Create Alert | RSS feed | Search history

Display: 25 results per page Go to page: 1 of 893 | Next >

Select: Selected Records (0) | Remove all Selected Records
Email | Print | Download

Sort by: Relevance

1. Discovery of WRKY transcription factors through transcriptome analysis and characterization of a novel methyl jasmonate-inducible PqWRKY1 gene from *Panax quinquefolius*
Wang, Y. (The Key Laboratory of Diagnostic Techniques and Resource Utilization of Chinese Medicinal Materials, Ministry of Education Institute of Medicinal Plant Development (IMPD), Chinese Academy of Medical Sciences, Peking Union Medical College, Beijing, 100193, China); Niu, Yunyun; Xu, Jiang; Li, Ying; Luo, Hongmei; Zhu, Yingjie; Liu, Mingzhu; Wu, Qiong; Song, Jingyuan; Sun, Chao; Chen, Shilin Source: *Plant Cell, Tissue and Organ Culture*, v 114, n 2, p 269-277, August 2013
Database: Compendex
Detailed | Show preview | Cite

2. Feature-based classification of human transcription factors related to regulatory function
Ehsani, Reza; Rezvan (Norwegian University of Science and Technology, Trondheim, Norway); Bahrani, Shahram; Drablos, Finn Source: *BMC Bioinformatics*, v 17, n 1, November 14, 2016
Database: Compendex
Detailed | Show preview | Full Text

3. Comparative in Silico evaluation of MYB transcription factors in eucalyptus, sugarcane and rice transcriptomes
Soares-Cavalcanti, Nina M. (Genetics Department, Laboratory for Plant Genetics and Biotechnology, Universidade Federal de Pernambuco, Rua Prof. Moraes Rêgo s/n., Recife, PE, Brazil); Wanderley-Nogueira, Ana C.; Belarmino, Luis C.; Dos Santos Barros, Petra; Benko-Iseppon, Ana M. Source: *Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics)*, v 5488 LNBI, p 44-55, 2009, *Computational Intelligence Methods for Bioinformatics and Biostatistics - 5th International Meeting, CIBB 2008, Revised Selected Papers*
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4. GATA transcription factors regulate the expression of the human eosinophil-derived neurotoxin (RNase 2) gene
Oliu, Zhijun (Laboratory of Allergic Diseases, NIAID, National Institutes of Health, 9000 Rockville Pike, Bethesda, MD 20892, United States); Dyer, Kimberly D.; Xie, Zhihui; Rådinger, Madeleine; Rosenberg, Helene F. Source: *Journal of Biological Chemistry*, v 284, n 19, p 13099-13109, May 8, 2009
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5. Functions of heat shock transcription factors involved in response to photooxidative stresses in arabidopsis
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6. Determinants of the DNA binding specificity of class I and class II TCP transcription factors
Viola, Ivana L. (Instituto de Agrobiotecnología del Litoral (CONICET-UNL), Facultad de Bioquímica Y Ciencias Biológicas, Universidad Nacional del Litoral, CC 242 Paraje El Pozo, 3000 Santa Fe, Argentina); Reinheimers, Renata; Ripoll, Rodrigo; Uberti Manassero, Nora G.; Gonzalez, Daniel H. Source: *Journal of Biological Chemistry*, v 287, n 1, p 347-356, January 2, 2012
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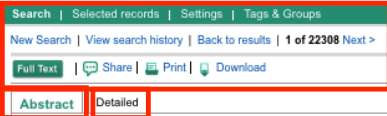
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Source: *Plant Cell, Tissue & Organ Culture* 157; DOI: 10.1007/s11240-013-0323-1; Publisher: Springer Netherlands

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1 The Key Laboratory of Bioactive Substances and Resources Utilization of Chinese Herbal Medicine, Ministry of Education, Institute of Medicinal Plant Development (IMPLAD), Chinese Academy of Medical Science, Beijing, 100193, China
2 College of Horticulture and Landscape Architecture, Southwest University (SWU), Chongqing, 400715, China
3 School of Pharmacy, Guilin Medical College, Guilin, 541004, China

Abstract:
Transcription factors (TFs) are important regulating factors that can mediate many life processes. However, no TF genes have previously been reported in *Panax quinquefolius* (American ginseng), with the exception of some sequence tags. In this study, 753 unigenes (unique sequences) have been annotated in the plant transcription factor database PlnTFDB (version 3.0) by mining a 454 transcriptome dataset of *P. quinquefolius*. After bioinformatics analysis, 45 unigenes were discovered to be annotated as WRKY transcription factors in the public databases. Furthermore, PqWRKY1, one of the WRKY family TF genes that respond to methyl jasmonate, was isolated according to the 454 transcriptome dataset. The cDNA of PqWRKY1 (*P. quinquefolius* WRKY1) encodes a putative protein of 358 amino acids, including a WRKY domain and a zinc finger motif. A subcellular localization assay demonstrated that PqWRKY1 localizes to the nucleus and has strong transcriptional activation activity in transgenic yeast. In comparison to control lines, the PqWRKY1 transgenic *Arabidopsis* line exhibited insensitive phenotypes when exposed to methyl jasmonate. Correspondingly, in transgenic *Arabidopsis* lines, the expression levels of some genes involved in the anti-stress process were relatively higher than those in the control lines. Additionally, genes involved in triterpene biosynthesis were up-regulated in the transgenic *Arabidopsis* line. These results suggest that the transcription factor PqWRKY1 is a positive regulator related to osmotic stress and triterpene biosynthesis in *P. quinquefolius*. © 2013 Springer Science+Business Media Dordrecht.(28 refs)

Main heading: Plants (botany)
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Uncontrolled terms: Expressed sequence tags - Ginsenoside biosynthesis - Osmotic stress - *Panax quinquefolius* - Plant transcription factors - Subcellular localization - Transcription factors

Classification Code: 461 Bioengineering and Biology - 801.2 Biochemistry - 802.3 Chemical Operations
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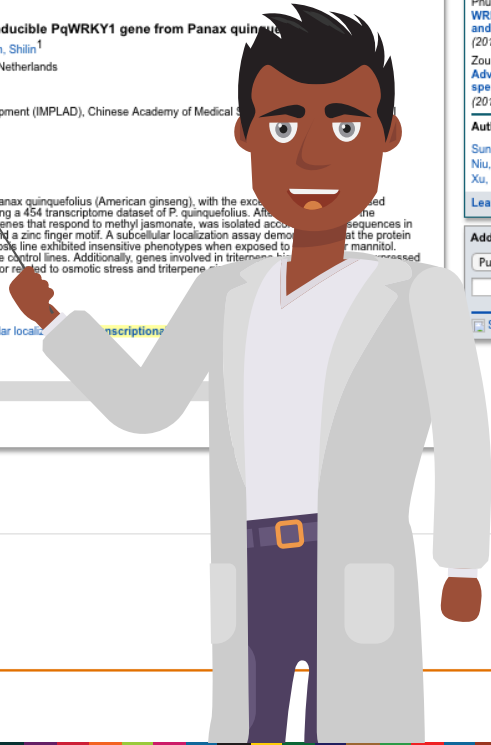
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Sun, Yongzhen¹; Niu, Yunyun¹; Xu, Jiang¹; Li, Ying¹; Luo, Hongmei¹; Zhu, Yingjie¹; Liu, Mingzhu²; Wu, Qiong^{1, 3}; Song, Jingyuan¹; Sun, Chao¹; Chen, Shilin¹

Source: *Plant Cell, Tissue and Organ Culture*, v 114, n 2, p 269-277, August 2013; ISSN: 01676857; DOI: 10.1007/s11240-013-0323-1; Publisher: Springer Netherlands

Author affiliations:

¹ The Key Laboratory of Bioactive Substances and Resources Utilization of Chinese Herbal Medicine, Ministry of Education, Institute of Medicinal Plant Development (IMPLAD), Chinese Academy of Medical Sciences, Peking Union Medical College, Beijing, 100193, China

² College of Horticulture and Landscape Architecture, Southwest University (SWU), Chongqing, 400715, China

³ School of Pharmacy, Guilin Medical College, Guilin, 541004, China

Abstract:

Transcription factors (TFs) are important regulating factors that can mediate many life processes. However, no TF genes have previously been reported in *Panax quinquefolius* (American ginseng), with the exception of a few expressed sequence tags. In this study, 753 unigenes (unique sequences) have been annotated in the plant **transcription factor** database PlnTFDB (version 3.0) by mining a 454 transcriptome dataset of *P. quinquefolius*. After classification of the unigenes, 45 unigenes were discovered to be annotated as **WRKY transcripts** in the public databases. Furthermore, PqWRKY1, one of the WRKY family TF genes that respond to methyl jasmonate, was isolated according to the sequences in the 454 transcriptome dataset. The cDNA of PqWRKY1 (*P. quinquefolius* WRKY1) encodes a putative protein of 358 amino acids, including a WRKY domain and a zinc finger motif. A subcellular localization assay demonstrated that the protein localizes to the nucleus and has strong **transcriptional** activation activity in transgenic yeast. In comparison to control lines, the PqWRKY1 transgenic *Arabidopsis* line exhibited insensitive phenotypes when exposed to high salt or mannitol. Correspondingly, in transgenic *Arabidopsis* lines, the expression levels of some genes involved in the anti-stress process were relatively higher than those in the control lines. Additionally, genes involved in the anti-stress process were relatively higher than those in the control lines. These results suggest that the **transcription factor** PqWRKY1 is a positive regulator related to osmotic stress and trier, *quinquefolius*. © 2013 Springer Science+Business Media Dordrecht.(28 refs)

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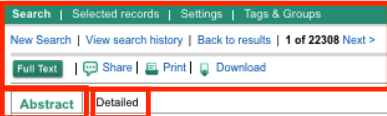
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Sun, Yongzhen¹; Niu, Yunyi²; Zhu, Wu, Qiong^{1,3}; Song, Jingyuan¹; Sun, Chao¹; Chen, Shilin¹
Source: *Plant Cell, Tissue & Organ Culture* 120(1): 1-11, 2018. ISSN: 0167-6369. DOI: 10.1007/s11240-013-0323-1; Publisher: Springer Netherlands

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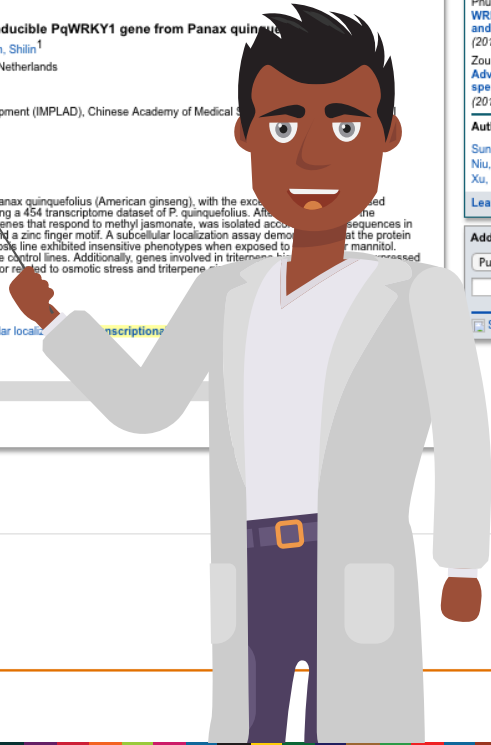
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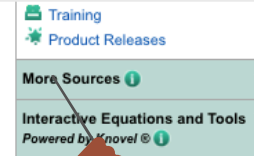
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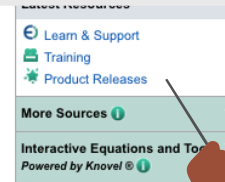
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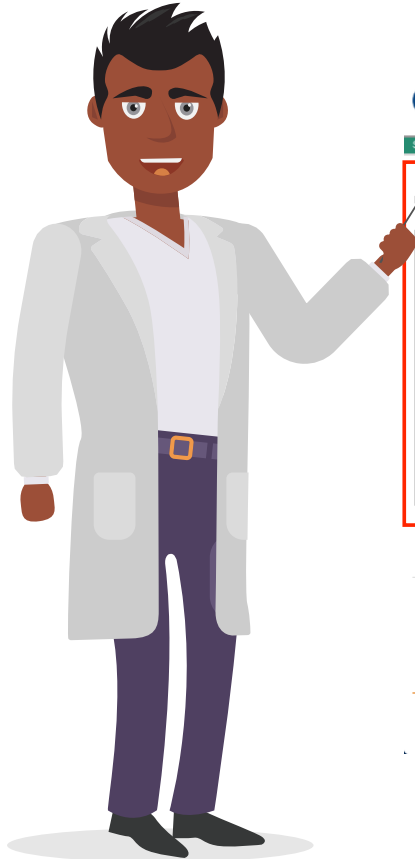
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O ícone Save Search fica próximo da parte superior de qualquer página de resultados de pesquisa da Engineering Village. Para excluir uma pesquisa, clique em Settings na parte superior da página e depois clique em Alerts & Searches. Exclua uma pesquisa específica ou todas as pesquisas salvas clicando em Delete All na parte inferior da caixa My Saved Searches.



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My Email Alerts

No.	Type	Name	Search	Auto- stem	Sort	Results	Year(s)	Database	Date Saved	Current Status	Recently Published	Clear Email Alert	Add Recipients
1. Delete	Quick	((transcription factors) WN All ... Edit	((transcription factors) WN All fields) Edit	On	Relevance	22,308	1969-2018	Compendex	01-Jul-2017 10:31 PM	Enabled	OFF	☐	Cc
2. Delete	Quick	((reservation systems) WN All fi ... Edit	((reservation systems) WN All fields) Edit	On	Relevance	557,574	1969-2018	Compendex	01-Jul-2017 10:32 PM	Enabled	OFF	☐	Cc
Delete All												Clear Email Alerts	

A função Alerts pode ser criada a partir de qualquer página de resultados de pesquisa, clicando no ícone Create Alert na parte superior da página. Para excluir um alerta, clique em Settings na parte superior da página e depois clique em Alerts & Searches. Alertas são fornecidos horas após os novos registros serem carregados dentro das bases de dados da Engineering Village. Os alertas podem ser ativados ou desativados por meio da caixa de gerenciamento de Alert. Alertas desabilitados serão salvos e podem ser reabilitados em uma data posterior.

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Selecione os registros clicando nas caixas de seleção próximas a cada registro, depois clique no ícone Save to Folder na parte superior da página de resultados de pesquisa. Utilize uma pasta existente ou crie uma nova pasta. Para gerenciar as pastas, clique em Settings na parte superior da página e depois em Folders. Os documentos dentro de cada pasta podem ser visualizados ou as pastas podem ser renomeadas ou excluídas. Os registros dentro da pasta podem ser enviados por e-mail, impressos ou baixados. Até dez pastas podem ser criadas contendo até cinquenta registros em cada pasta.

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With your Personal Account, you can create up to ten folders in which to save selected records. Each folder can contain up to 50 records. To create a folder, please enter a folder name:

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Usuários inscritos e que tenham feito login podem salvar as configurações para download, classificação, formato de registro, número de registros exibidos em uma página e cor de destaque. Na barra de navegação superior, utilize o menu Settings para definir a opção My Preferences. Caso um nome de arquivo padrão seja escolhido, o nome do arquivo conterá a data e o horário de download para que arquivos separados sejam criados sempre que acontecer um download.

A screenshot of the 'My Preferences' settings window. The window has a green header bar with the Engineering Village logo and the title 'My Preferences'. Below the header, there are several sections: 'Results' with a 'Download' button and a 'Search' button; 'Number of results per page' with a dropdown set to '25'; 'Facet order' with three 'Select' dropdowns; 'Search terms' with a color selection box (orange) and a 'Text color' label, a checkbox for 'Background highlight', and a note 'Search terms will display with your color choice.'; 'Sorting default' with a dropdown set to 'Relevance'; and 'Show preview' with a 'Hide All' dropdown. At the bottom right, there are 'Save' and 'Cancel' buttons.

My Preferences

Results

Download

Search

Number of results per page

25

Facet order

1. Select

2. Select

3. Select

Search terms

Text color

☐ Background highlight

Search terms will display with your color choice.

Sorting default

Relevance

Show preview

Hide All

Save Cancel

Clique em Register na parte superior da página e preencha o formulário de entrada de dados. As informações inseridas são armazenadas em um perfil que pode ser editado ao clicar em Settings na barra de navegação superior.

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Durante o uso da **Engineering Village**, a opção **Help** (Ajuda) on-line está disponível na barra de navegação no topo da página, clique em **Support** e depois em **Help** para ver instruções ou informações relacionadas.

Clique nos ícones pequenos para ter acesso a informações adicionais de ajuda. A funcionalidade **Help** detecta o contexto e sempre apresenta as recomendações mais relevantes primeiro.



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