

转录因子数据库构建实例



PlantRegMap/PlantTFDB v5.0

Plant Transcription Factor Database

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Note:

- The new version PlantTFDB 5.0 has been incorporated into [PlantRegMap](#).

Browse by Species

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Taxonomic Group (165 species) (G)-species with genome sequence

- [Chlorophyta \(16 species\)](#)
- [Charophyta \(1 species\)](#)
- [Marchantiophyta \(1 species\)](#)
- [Bryophyta \(2 species\)](#)
- [Lycopodiophyta \(1 species\)](#)
- [Coniferophyta \(5 species\)](#)
- [Basal Magnoliophyta \(1 species\)](#)
- [Monocots \(38 species\)](#)
- [Eudicots \(100 species\)](#)

Quick links: [Arabidopsis thaliana](#), [Glycine max](#), [Oryza sativa](#), [Populus trichocarpa](#), [Solanum lycopersicum](#), [Zea mays](#)

Couldn't find the species interested? Please try the [extended TF repertoires](#) or identify TFs by the [prediction server](#).

Browse by Family

AP2 (4461)	ARF (4578)	ARR-B (2354)	B3 (10609)	BBR-BPC (1256)
BES1 (1549)	C2H2 (17740)	C3H (9693)	CAMTA (1343)	CO-like (2125)
CPP (1612)	DBB (1651)	Dof (5655)	E2F/DP (1781)	EIL (1234)
ERF (21129)	FAR1 (7527)	G2-like (9874)	GATA (5335)	GRAS (9304)
GRF (1876)	GeBP (1564)	HB-PHD (477)	HB-other (2277)	HD-ZIP (8602)
HRT-like (249)	HSF (4574)	LBD (7216)	LFY (253)	LSD (957)
M-type_MADS (7541)	MIKC_MADS (6918)	MYB (22032)	MYB_related (15369)	NAC (19997)
NF-X1 (403)	NF-YA (2461)	NF-YB (3099)	NF-YC (2446)	NZZ/SPL (109)
Nin-like (2766)	RAV (690)	S1Fa-like (359)	SAP (164)	SBP (4168)
SRS (1327)	STAT (214)	TALE (4433)	TCP (4187)	Trihelix (6256)
VOZ (635)	WOX (2358)	WRKY (14549)	Whirly (530)	YABBY (1719)
ZF-HD (2589)	bHLH (28698)	bZIP (15498)		

网站首页:

[index.html](#) 或 [index.php](#)

功能:

浏览、查询和分析服务

<https://planttfdb.gao-lab.org/>



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(e.g., LFY)

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静态网页

HTML文件在服务器上已存在，
收到请求时直接调用



PlantRegMap/PlantTFDB v5.0

Plant Transcription Factor Database

[Home](#) [TText](#) [BLAST](#) [Prediction](#) [Download](#) [Help](#) [About](#) [Links](#) [PlantRegMap](#) (e.g., LFY)

Transcription Factor Information

[Basic Information](#) | [Signature Domain](#) | [Sequence](#) | [Protein Features](#) | [3D Structure](#) | [Gene Ontology](#) | [Plant Ontology](#) | [Expression](#) | [Function](#) | [Binding Motif](#) | [Cis-element](#) | [Regulation](#) | [Interaction](#) | [Phenotype](#) | [Orthologous Group](#) | [Publication](#)

Basic Information		? help			Back to Top
TF ID	AT5G61850.1				
Common Name	LFY, LFY3, MAC9.18				
Organism	Arabidopsis thaliana				
Taxonomic ID	3702				
Taxonomic Lineage	cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; rosids; malvids; Brassicales; Brassicaceae; Camelineae; Arabidopsis				
Family	LFY				
Protein Properties	Length: 420aa MW: 46582.3 Da PI: 7.1121				
Description	floral meristem identity control protein LEAFY (LFY)				
Gene Model	Gene Model ID	Type	Source	Coding Sequence	
	AT5G61850.1	genome	TAIR	View CDS	

动态网页

HTML文件在服务器上并不存在，收到请求时通过php查询数据库生成相应的html文件

URL: <https://planttfdb.gao-lab.org/tf.php?sp=Ath&did=AT5G61850.1>

用户交互： 表单(form)

```
<form action="search_result.php" method="post">  
  <input type="text" name="input" id="input"><br><br>  
  <input type="submit" value="Submit">  
</form>
```



练习: https://www.w3schools.com/Html/tryit.asp?filename=tryhtml_form_text

相关介绍: https://www.w3schools.com/Html/html_forms.asp

参数传递与接收

- GET 适合传输不敏感、少量数据（如搜索查询），且该数据可公开。
- POST 适合传输大量数据和敏感信息（如用户登录、提交表单等）。

```
$id = $_GET['TF_ID'];
```

```
$id = $_POST['input'];
```

<https://planttfdb.gao-lab.org/tf.php?sp=Ath&did=AT5G61850.1>

```
<form action="search_result.php" method="post">  
  <input type="text" name="input" id="input"><br><br>  
  <input type="submit" value="Submit">  
</form>
```

创建数据库连接

```
<?php
```

```
// 参数
```

```
$host = 'localhost';
```

```
//数据库服务器IP
```

```
$user = 'username';
```

```
//用户名
```

```
$passwd = 'password';
```

```
//密码
```

```
$db = 'database';
```

```
//数据库名称
```

```
// 创建数据库连接
```

```
$conn = mysqli_connect($host, $user, $passwd, $db);
```

```
// 数据库连接是否成功
```

```
if (!$conn) {
```

```
    die('Connection failed: ' . mysqli_connect_error());
```

```
}
```

```
// 关闭数据库连接
```

```
mysqli_close($conn);
```

```
?>
```

数据库检索

```
<?php
$TF_ID = $_GET['TF_ID']; // 获取 GET 请求中的 TF_ID 参数

$sql = "SELECT * FROM Ath_TF_list WHERE TF_ID = '$TF_ID'"; // 创建 SQL 查询语句

$result = mysqli_query($conn, $sql); // 执行查询

?>
```

数据: https://planttfdb.gao-lab.org/download/TF_list/Ath_TF_list.txt.gz

表格: Ath_TF_list

TF_ID	Gene_ID	Family
AT3G25730.1	AT3G25730	RAV
AT1G68840.1	AT1G68840	RAV
AT1G68840.2	AT1G68840	RAV
.....		

使用预处理语句防止SQL注入: <https://www.php.net/manual/en/mysqli.prepare.php>

检索结果展示

```
<?php
// 检查查询是否失败
if (!$result) {
    die('Query error: ' . mysqli_error($conn));
}
// 检查结果是否为空
elseif (mysqli_num_rows($result) == 0) {
    die('Invalid ID');
}
// 如果查询成功并且有数据
else {
    // 输出表头
    echo '<tr><th>TF ID</th><th>Family</th></tr>';

    // 遍历结果并输出
    while ($row = mysqli_fetch_assoc($result)) {
        echo '<tr>';
        echo '<td>' . $row['TF_ID'] . '</td>';
        echo '<td>' . $row['Family'] . '</td>';
        echo '</tr>';
    }
}
?>
```

预测分析服务

输入与提交 页面示例

Input protein or nucleic acid sequences in FASTA format (Max file size: 75M):

Or load it from disk: No file chosen

Link: ☐ Best hit in *Arabidopsis thaliana*

输入与提交 代码示例

```
<form action="prediction_result.php?task_id=<?php echo($task_id); ?>" method="post" enctype="multipart/form-data">
```

Input protein or nucleic acid sequences in FASTA format (Max file size: 75M):

<textarea id="input_seq" name="input_seq" rows="10" cols="100"></textarea>

 <!-- 输入框 -->

Or load it from disk:

 <!-- 上传文件 -->

Link:

☐ <!-- 复选框 -->

Best hit in *Arabidopsis thaliana*

<!-- 重置按钮 --><!-- 提交按钮 -->

创建任务号

//定义创建随机字符串的函数

```
function create_random($length) {  
    $chars = 'abcdefghijklmnopqrstuvwxyzABCDEFGHIJKLMNOPQRSTUVWXYZ0123456789'; //字符集  
    $id = "";  
    for ($i = 0; $i < $length; $i++) {  
        $id .= $chars[rand(0, strlen($chars) - 1)]; //从字符集中随机选择字符  
    }  
    return $id;  
}
```

```
$task_id = create_random(10);
```

任务接收与后台处理

```
<?php
$tmp_name = $_GET['task_id'];
$tmp_path = "TF_prediction/tmp/" . $tmp_name;
$tmpf = $tmp_path . '/input_seq.fas';

if (!is_dir($tmp_path)) {
    mkdir($tmp_path); // 创建临时文件夹

    if (!empty($_FILES["input_file"]["name"])) { // 检查是否有文件上传
        move_uploaded_file($_FILES["input_file"]["name"], $tmpf);
    } else {
        $seq = $_POST['input_seq']; // 检查是否有序列输入
        if (isset($seq[0]) && $seq[0] == ">") { // 检查序列格式
            $safe_seq = escapeshellarg($seq); // 对字符串中的特殊字符进行转义，防止命令注入
            $cmd = "echo $safe_seq > $tmpf"; // 准备执行的系统命令
            exec($cmd); // 执行命令
        }
    }
}
?>
```

展示结果文件

```
<?php
$result_file = "results.txt"; // 结果文件

if (!file_exists($result_file)) {
    die("结果文件不存在! ");
}

// 打开文件读取内容
$lines = file($filepath, FILE_IGNORE_NEW_LINES | FILE_SKIP_EMPTY_LINES);
if (!$lines) {
    die("无法读取文件内容! ");
}

//遍历每一行并输出
foreach ($lines as $line) {
    $columns = explode("\t", $line); // 使用制表符分隔

    echo "<tr>";
    foreach ($columns as $col) {
        echo "<td>" . htmlspecialchars($col) . "</td>";
    }
    echo "</tr>";
}
?>
```