

# Package 'PWMEnrich.Dmelanogaster.background'

April 14, 2020

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**License** GPL-3

**Title** D. melanogaster background for PWMEnrich

**Type** Package

**LazyLoad** yes

**Author** Robert Stojnic

**Description** PWMEnrich pre-compiled background objects for *Drosophila melanogaster* and MotifDb *D. melanogaster* motifs.

**Version** 4.20.0

**biocViews** *Drosophila\_melanogaster\_Data*

**Date** 2015-09-30

**Depends** methods, PWMEnrich

**git\_url** <https://git.bioconductor.org/packages/PWMEnrich.Dmelanogaster.background>

**git\_branch** RELEASE\_3\_10

**git\_last\_commit** 8ec7496

**git\_last\_commit\_date** 2019-10-29

**Date/Publication** 2020-04-14

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*PWMEnrich.Dmelanogaster.background package overview*

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## Description

This package provides a set of pre-compiled genomic background files for the PWMErich package and *D. melanogaster*. The backgrounds were pre-compiled using unique promoters of *Drosophila melanogaster*.

- `MotifDb.Dmel.PFM` - a list of 740 position frequency matrices (PFMs) from MotifDb
- `MotifDb.Dmel` - the corresponding PWMs generated by using the background frequencies of A,C,G,T in a set of 10031 2kb promoters in *D. Melanogaster*.
- `PWMLogn.dm3.MotifDb.Dmel` - pre-compiled threshold-free lognormal background for JASPAR insects PWMs. The lognormal distribution is fitted to 1kb fragments of 10031 2kb *Drosophila Melanogaster* (dm3) promoters. For each gene one promoter is taken (from the first transcript).
- `PWMCutoff4.dm3.MotifDb.Dmel`, `PWMCutoff5.dm3.MotifDb.Dmel` - pre-compiled Z-score background with cutoff of 4 and 5 (base e) for JASPAR insects PWMs. The number of hits above the score cutoff is counted in 10031 2kb *Drosophila Melanogaster* (dm3) promoters. For each gene one promoter is taken (from the first transcript).
- `PWMPvalueCutoff1e2.dm3.MotifDb.Dmel`, `PPWMPvalueCutoff1e3.dm3.MotifDb.Dmel`, `PPWMPvalueCutoff1e4.dm3.MotifDb.Dmel` - pre-compiled Z-score background with P-value cutoff of 0.01, 0.001, and 0.0001 for JASPAR insects PWMs. The number of hits with smaller P-value than cutoff is counted in 10031 2kb *Drosophila Melanogaster* (dm3) promoters. For each gene one promoter is taken (from the first transcript).

## Usage

```
data(MotifDb.Dmel.PFM)
data(MotifDb.Dmel)
data(PWMLogn.dm3.MotifDb.Dmel)
data(PWMCutoff4.dm3.MotifDb.Dmel)
data(PWMCutoff5.dm3.MotifDb.Dmel)
data(PWMPvalueCutoff1e2.dm3.MotifDb.Dmel)
data(PWMPvalueCutoff1e3.dm3.MotifDb.Dmel)
data(PWMPvalueCutoff1e4.dm3.MotifDb.Dmel)
```

## Details

All of these objects were created with the appropriate functions available in the PWMErich package (see Section 'See also'). We recommend using these functions to generate backgrounds for a custom set of background sequences and/or DNA motifs.

This package also contains the 2kb upstream sequences for *Drosophila* genes (`dm3.upstream2000`) which were removed from the `BSgenome.Dmelanogaster.UCSC.dm3` package starting from Bioconductor version 3.0.

## Author(s)

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## See Also

[makeBackground](#), [makePWMLognBackground](#), [makePWMCutoffBackground](#), [makePWMEmpiricalBackground](#).

**Examples**

```
data(PWMLogn.dm3.MotifDb.Dmel)
```

```
res = motifEnrichment(DNAString("TGCATCAAGTGTGTAGTGCATGAATGC"), PWMLogn.dm3.MotifDb.Dmel)
```

```
head(motifRankingForGroup(res))
```

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