(base) C19DMGD:scripts leyu$ python get\_results.py

This stain is: Tetrapisispora blattae

The number of genes used in the model is: 5377

The number of genes in Uniprot database is: 5385

Reciprocal blast best hits with Pidentity more than 80%: 5056

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This stain is: Naumovozyma castellii

The number of genes used in the model is: 5264

The number of genes in Uniprot database is: 5565

Reciprocal blast best hits with Pidentity more than 80%: 5063

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This stain is: Saccharomyces eubayanus

The number of genes used in the model is: 5043

The number of genes in Uniprot database is: 5354

Reciprocal blast best hits with Pidentity more than 80%: 4629

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This stain is: Dekkera bruxellensis

The number of genes used in the model is: 5600

The number of genes in Uniprot database is: 4849

Reciprocal blast best hits with Pidentity more than 80%: 4310

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This stain is: Lachancea thermotolerans

The number of genes used in the model is: 4808

The number of genes in Uniprot database is: 5093

Reciprocal blast best hits with Pidentity more than 80%: 4650

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This stain is: Kluyveromyces dobzhanskii

The number of genes used in the model is: 4882

The number of genes in Uniprot database is: 4952

Reciprocal blast best hits with Pidentity more than 80%: 4711

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This stain is: Zygosaccharomyces rouxii

The number of genes used in the model is: 4623

The number of genes in Uniprot database is: 4987

Reciprocal blast best hits with Pidentity more than 80%: 4500

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This stain is: Schizosaccharomyces pombe

The number of genes used in the model is: 5134

The number of genes in Uniprot database is: 5141

Reciprocal blast best hits with Pidentity more than 80%: 5122

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This stain is: Pichia pastoris

The number of genes used in the model is: 4831

The number of genes in Uniprot database is: 5073

Reciprocal blast best hits with Pidentity more than 80%: 4600

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This stain is: Lachancea fermentati

The number of genes used in the model is: 4879

The number of genes in Uniprot database is: 5229

Reciprocal blast best hits with Pidentity more than 80%: 4709

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This stain is: Kluyveromyces marxianus

The number of genes used in the model is: 5110

The number of genes in Uniprot database is: 4926

Reciprocal blast best hits with Pidentity more than 80%: 4706

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This stain is: Kluyveromyces lactis

The number of genes used in the model is: 4962

The number of genes in Uniprot database is: 5071

Reciprocal blast best hits with Pidentity more than 80%: 4799

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This stain is: Tetrapisispora phaffii

The number of genes used in the model is: 5295

The number of genes in Uniprot database is: 5238

Reciprocal blast best hits with Pidentity more than 80%: 4994

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This stain is: Eremothecium sinecaudum

The number of genes used in the model is: 4314

The number of genes in Uniprot database is: 4524

Reciprocal blast best hits with Pidentity more than 80%: 4172

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