All Classes

Action
Controller
Dose
IAction
IDose
IMedicine
Medicine

PACKAGE CLASS TREE DEPRECATED INDEX HELP

PREV CLASS NEXT CLASS FRAMES NO FRAMES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Class Action

java.lang.Object Action

All Implemented Interfaces:

IAction

public class Action
extends java.lang.Object
implements IAction

Field Summary

Fields

Modifier and Type Field and Description

private IMedicine medicine

Constructor Summary

Constructors

Constructor and Description

Action()

Method Summary

All Methods Instance Methods	Concrete Methods
Modifier and Type	Method and Description
void	addDose(IDose dose) Adds a new dose.
java.util.ArrayList <java.lang.< td=""><td>ouble> getCurrentConcentration(java.time.LocalTime time) Calculates concentration amount of doses at a specific time.</td></java.lang.<>	ouble> getCurrentConcentration(java.time.LocalTime time) Calculates concentration amount of doses at a specific time.
IMedicine	<pre>getMedicine() Retrieves the Medicine instance.</pre>
<pre>java.util.ArrayList<java.time.< pre=""></java.time.<></pre>	ocalTime> getPeakConcentrationTime(java.lang.Boolean includeTestDoses) Time when the concentration is at its peak.
java.util.ArrayList <java.time.< td=""><td>ocalTime> getWhenToDose(java.lang.Double concentrationDesired) Determines time when the patient should take next dose.</td></java.time.<>	ocalTime> getWhenToDose(java.lang.Double concentrationDesired) Determines time when the patient should take next dose.
void	<pre>loadFile(java.lang.String filename) Load a saved Medicine file.</pre>
void	<pre>newFile(java.lang.String name, java.time.LocalTime tMax, java.time.LocalTime halfLife) Create a new Medicine instance.</pre>
void	<pre>printCurrentConcentration(java.time.LocalTime time)</pre> Prints concentration amount of doses at a specific time.
void	<pre>printDoses() Prints all doses from Medicine's Dose array.</pre>
void	printMedicine() Print the name, tmax, and halfLife of the medicine.
void	<pre>printPeakConcentrationTime(java.lang.Boolean includeTestDoses Print concentration peaks.</pre>

void	<pre>printWhenToDose(java.lang.Double amountDesired) Prints when the patient should take next dose.</pre>
void	removeAllDoses() Delete all doses from the dosages array.
void	<pre>removeDose(int index) Removes a dose from the dose from the doses array.</pre>
void	removeTestDoses() Removes all test doses.
void	<pre>saveFile(java.lang.String filename)</pre> Save Medicine to a file.

Methods inherited from class java.lang.Object

clone, equals, finalize, getClass, hashCode, notify, notifyAll, toString, wait, wait, wait

Field Detail

medicine

private IMedicine medicine

Constructor Detail

Action

public Action()

Method Detail

printDoses

public void printDoses()

$\textbf{Description copied from interface:} \ \mathtt{IAction}$

Prints all doses from Medicine's Dose array.

Specified by:

printDoses in interface IAction

addDose

public void addDose(IDose dose)

Description copied from interface: IAction

Adds a new dose. Added doses can be type Dose (actual dose) or TestDose (what-if dose)

Specified by:

addDose in interface IAction

Parameters:

dose - Dose object.

removeDose

public void removeDose(int index)

Description copied from interface: IAction

Removes a dose from the dose from the doses array.

Specified by:

removeDose in interface IAction

Parameters:

index - Index of the array to be erased

removeAllDoses

public void removeAllDoses()

Description copied from interface: IAction

Delete all doses from the dosages array.

Specified by:

removeAllDoses in interface IAction

getCurrentConcentration

public java.util.ArrayList<java.lang.Double> getCurrentConcentration(java.time.LocalTime time)

Description copied from interface: IAction

Calculates concentration amount of doses at a specific time.

Specified by:

getCurrentConcentration in interface IAction

Parameters:

time - Specific time used to determine concentration amount.

Returns:

ArrayList with the concentration amounts.

printCurrentConcentration

public void printCurrentConcentration(java.time.LocalTime time)

$\textbf{Description copied from interface:} \ \mathtt{IAction}$

Prints concentration amount of doses at a specific time.

Specified by:

printCurrentConcentration in interface IAction

Parameters:

time - Specific time used to determine concentration amount.

getPeakConcentrationTime

public java.util.ArrayList<java.time.LocalTime> getPeakConcentrationTime(java.lang.Boolean includeTestDose

Description copied from interface: IAction

Time when the concentration is at its peak. It allows the user to choose if test doses will be displayed or not.

Specified by:

getPeakConcentrationTime in interface IAction

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

Returns

ArrayList containing peak concentration peak values of each dose.

printPeakConcentrationTime

public void printPeakConcentrationTime(java.lang.Boolean includeTestDoses)

$\textbf{Description copied from interface:} \ \mathtt{IAction}$

Print concentration peaks. Includes dose information and its concentration peak.

Specified by:

printPeakConcentrationTime in interface IAction

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

getWhenToDose

public java.util.ArrayList<java.time.LocalTime> getWhenToDose(java.lang.Double concentrationDesired)

Description copied from interface: IAction

Determines time when the patient should take next dose.

Specified by:

getWhenToDose in interface IAction

Parameters:

concentrationDesired - Dose amount desired sought.

Returns

Time when the patient have to take next dose.

printWhenToDose

public void printWhenToDose(java.lang.Double amountDesired)

Description copied from interface: IAction

Prints when the patient should take next dose. Includes dose information and time to take next dose.

Specified by:

printWhenToDose in interface IAction

Parameters:

amountDesired - Dose amount desired sought.

saveFile

public void saveFile(java.lang.String filename)

Description copied from interface: IAction

Save Medicine to a file. By default, save directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Specified by:

saveFile in interface IAction

Parameters:

filename - Name of save file to be saved.

loadFile

public void loadFile(java.lang.String filename)

Description copied from interface: IAction

Load a saved Medicine file. By default, load directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Specified by:

loadFile in interface IAction

Parameters:

filename - Name of the file to be loaded

newFile

Description copied from interface: IAction

Create a new Medicine instance.

Specified by:

newFile in interface IAction

Parameters:

name - Name of medicine

tMax - TMax of medicine.

halfLife - Half life of medicine

printMedicine

Specified by:

public void printMedicine()

Description copied from interface: IAction Print the name, tmax, and halfLife of the medicine.

printMedicine in interface IAction

removeTestDoses

public void removeTestDoses()

Description copied from interface: IAction

Removes all test doses.

Specified by:

removeTestDoses in interface IAction

getMedicine

public IMedicine getMedicine()

Description copied from interface: IAction

Retrieves the Medicine instance.

Specified by:

getMedicine in interface IAction

Returns:

Medicine instance

PACKAGE CLASS TREE DEPRECATED INDEX HELP

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ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Class Action

java.lang.Object Action

All Implemented Interfaces:

IAction

public class Action
extends java.lang.Object
implements IAction

Field Summary

Fields

Modifier and Type Field and Description

private IMedicine medicine

Constructor Summary

Constructors

Constructor and Description

Action()

Method Summary

All Methods	Instance Methods	Concrete Method	s
Modifier and Typ	pe	Meth	nod and Description
void			Dose(IDose dose) Is a new dose.
java.util.Ar	rrayList <java.lang< td=""><td></td><td>CurrentConcentration(java.time.LocalTime time) culates concentration amount of doses at a specific time.</td></java.lang<>		CurrentConcentration(java.time.LocalTime time) culates concentration amount of doses at a specific time.
IMedicine		_	Medicine() rieves the Medicine instance.
java.util.Ar	rrayList <java.time< td=""><td></td><td>PeakConcentrationTime(java.lang.Boolean includeTestDoses) the when the concentration is at its peak.</td></java.time<>		PeakConcentrationTime(java.lang.Boolean includeTestDoses) the when the concentration is at its peak.
java.util.Ar	rrayList <java.time< td=""><td></td><td>WhenToDose(java.lang.Double concentrationDesired) ermines time when the patient should take next dose.</td></java.time<>		WhenToDose(java.lang.Double concentrationDesired) ermines time when the patient should take next dose.
void			dFile(java.lang.String filename) d a saved Medicine file.
void		new	File(java.lang.String name, java.time.LocalTime tMax,

<pre>void</pre>
Prints all doses from Medicine's Dose array. void printMedicine()
void <pre>printPeakConcentrationTime(java.lang.Boolean includeTestDoses</pre> Print concentration peaks.
void <pre>printWhenToDose(java.lang.Double amountDesired) Prints when the patient should take next dose.</pre>
void removeAllDoses() Delete all doses from the dosages array.
void removeDose(int index) Removes a dose from the dose from the doses array.
void removeTestDoses() Removes all test doses.
void saveFile(java.lang.String filename) Save Medicine to a file.

Methods inherited from class java.lang.Object

clone, equals, finalize, getClass, hashCode, notify, notifyAll, toString, wait, wait, wait

Field Detail

medicine

private IMedicine medicine

Constructor Detail

Action

public Action()

Method Detail

printDoses

public void printDoses()

Description copied from interface: IAction

Prints all doses from Medicine's Dose array.

Specified by:

printDoses in interface IAction

addDose

public void addDose(IDose dose)

Description copied from interface: IAction

Adds a new dose. Added doses can be type Dose (actual dose) or TestDose (what-if dose)

Specified by:

addDose in interface IAction

Parameters:

dose - Dose object.

removeDose

public void removeDose(int index)

Description copied from interface: IAction

Removes a dose from the dose from the doses array.

Specified by:

removeDose in interface IAction

Parameters:

index - Index of the array to be erased

removeAllDoses

public void removeAllDoses()

Description copied from interface: IAction

Delete all doses from the dosages array.

Specified by:

removeAllDoses in interface IAction

getCurrentConcentration

public java.util.ArrayList<java.lang.Double> getCurrentConcentration(java.time.LocalTime time)

Description copied from interface: IAction

Calculates concentration amount of doses at a specific time.

Specified by:

getCurrentConcentration in interface IAction

Parameters:

time - Specific time used to determine concentration amount.

Returns:

ArrayList with the concentration amounts.

printCurrentConcentration

public void printCurrentConcentration(java.time.LocalTime time)

Description copied from interface: IAction

Prints concentration amount of doses at a specific time.

Specified by:

printCurrentConcentration in interface IAction

Parameters:

time - Specific time used to determine concentration amount.

getPeakConcentrationTime

public java.util.ArrayList<java.time.LocalTime> getPeakConcentrationTime(java.lang.Boolean includeTestDoses)

Description copied from interface: IAction

Time when the concentration is at its peak. It allows the user to choose if test doses will be displayed or not.

Specified by:

getPeakConcentrationTime in interface IAction

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

Returns:

ArrayList containing peak concentration peak values of each dose.

printPeakConcentrationTime

public void printPeakConcentrationTime(java.lang.Boolean includeTestDoses)

Description copied from interface: IAction

Print concentration peaks. Includes dose information and its concentration peak.

Specified by:

printPeakConcentrationTime in interface IAction

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

getWhenToDose

public java.util.ArrayList<java.time.LocalTime> getWhenToDose(java.lang.Double concentrationDesired)

Description copied from interface: IAction

Determines time when the patient should take next dose.

Specified by:

getWhenToDose in interface IAction

Parameters:

concentrationDesired - Dose amount desired sought.

Returns:

Time when the patient have to take next dose.

printWhenToDose

public void printWhenToDose(java.lang.Double amountDesired)

Description copied from interface: IAction

Prints when the patient should take next dose. Includes dose information and time to take next dose.

Specified by:

printWhenToDose in interface IAction

Parameters:

amountDesired - Dose amount desired sought.

saveFile

public void saveFile(java.lang.String filename)

Description copied from interface: IAction

Save Medicine to a file. By default, save directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Specified by:

saveFile in interface IAction

Parameters:

filename - Name of save file to be saved.

loadFile

public void loadFile(java.lang.String filename)

Description copied from interface: IAction

Load a saved Medicine file. By default, load directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Specified by:

loadFile in interface IAction

Parameters:

filename - Name of the file to be loaded

newFile

Description copied from interface: IAction

Create a new Medicine instance.

Specified by:

newFile in interface IAction

Parameters:

```
name - Name of medicine

tMax - TMax of medicine.

halfLife - Half life of medicine
```

printMedicine

public void printMedicine()

Description copied from interface: IAction

Print the name, tmax, and halfLife of the medicine.

Specified by:

printMedicine in interface IAction

removeTestDoses

public void removeTestDoses()

Description copied from interface: IAction

Removes all test doses.

Specified by:

removeTestDoses in interface IAction

getMedicine

public IMedicine getMedicine()

Description copied from interface: IAction

Retrieves the Medicine instance.

Specified by:

getMedicine in interface IAction

Returns:

Medicine instance

PACKAGE CLASS TREE DEPRECATED INDEX HELP

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PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Class Controller

java.lang.Object Controller

public class Controller extends java.lang.Object

Field Summary

Fields

Modifier and Type	Field and Description
(package private) static Action	action
(package private) static java.util.Scanner	userInput

Constructor Summary

Constructors

Constructor and Description

Controller()

Method Summary

All Methods	Static Methods	Concrete Methods	
Modifier and Ty	ре	Method and Description	
private sta	tic void	<pre>checkIfInteger() Validates if user input is an integer only.</pre>	
private sta	tic void	checkIfValidNumber() Validates if user input is a number, either Integer or Double	le.
private sta	tic void	clear() Clears the console screen.	
private sta	tic IDose	<pre>createDose()</pre>	

	Creates Dose from user input.
private static java.time.LocalTime	Generates LocalTime () Generates LocalTime instance from user input.
private static void	<pre>createMedicine()</pre> Creates a Medicine instance from user input.
static void	<pre>main(java.lang.String[] args)</pre>
private static void	Pause scrolling fot the console screen until user hits the Enter key.
private static void	removeDose() Removes dose using an index selected by the user.
private static void	selectAction() Display the list of actions that can be performed by the application.
private static void	start() Starts the program.

Methods inherited from class java.lang.Object

clone, equals, finalize, getClass, hashCode, notify, notifyAll, toString, wait,
wait, wait

Field Detail

userInput

static java.util.Scanner userInput

action

static Action action

Constructor Detail

Controller

public Controller()

Method Detail

createLocalTime

private static java.time.LocalTime createLocalTime()

Generates LocalTime instance from user input. Validates the user input to minimize application crash due to unexpected errors.

Returns:

LocalTime instance.

createMedicine

private static void createMedicine()

Creates a Medicine instance from user input.

createDose

private static IDose createDose()

Creates Dose from user input.

Returns:

A Dose instance

removeDose

private static void removeDose()

Removes dose using an index selected by the user.

checklfInteger

private static void checkIfInteger()

Validates if user input is an integer only. If input is not an integer, the program stops.

checklfValidNumber

private static void checkIfValidNumber()

Validates if user input is a number, either Integer or Double. If input is not an number, the program stops.

clear

private static void clear()

Clears the console screen.

pause

private static void pause()

Pause scrolling fot the console screen until user hits the Enter key.

start

private static void start()

Starts the program. Display a welcome screen which allows the user to select whether to create a file or open an existing one.

selectAction

private static void selectAction()

Display the list of actions that can be performed by the application. The user must enter the corresponding number to execute the action.

main

public static void main(java.lang.String[] args)

PACKAGE CLASS TREE DEPRECATED INDEX HELP

PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

PREV CLASS NEXT CLASS

FRAMES NO FRAMES

ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD

DETAIL: FIELD | CONSTR | METHOD

Class Dose

java.lang.Object Dose

All Implemented Interfaces:

IDose, java.io.Serializable

public class Dose
extends java.lang.Object
implements IDose, java.io.Serializable

See Also:

Serialized Form

Field Summary

Fields

	Modifier and Type	Field and Description
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private double amount

private boolean isTestDose

private java.time.LocalTime timeTake

Constructor Summary

Constructors

Constructor and Description

Dose()

Dose(java.time.LocalTime timeTake, double amount)

Method Summary

All Methods

Instance Methods

Concrete Methods

Modifier and Type

Method and Description

void	<pre>createDose(java.time.LocalTime timeTake, double amount) Creates a Dose object.</pre>
double	<pre>getAmount()</pre> Concentration amount of the dose taken by patient.
java.time.LocalTime	getTimeTake() Time when the dose is taken by patient.
boolean	isTestDose() Identifies dose type.
void	setTestDose() Sets dose to test dose.
java.lang.String	toString()

Methods inherited from class java.lang.Object

clone, equals, finalize, getClass, hashCode, notify, notifyAll, wait, wait, wait

Field Detail

timeTake

private java.time.LocalTime timeTake

amount

private double amount

isTestDose

private boolean isTestDose

Constructor Detail

Dose

public Dose()

Dose

Method Detail

isTestDose

public boolean isTestDose()

Description copied from interface: IDose

Identifies dose type.

Specified by:

isTestDose in interface IDose

Returns:

True = TestDose (what-if dose); False = Dose (actual dose);

createDose

Description copied from interface: IDose

Creates a Dose object.

Specified by:

createDose in interface IDose

Parameters:

timeTake - Time when the dose is taken by patient.

amount - Concentration amount of the dose taken by patient.

getAmount

public double getAmount()

Description copied from interface: IDose

Concentration amount of the dose taken by patient.

Specified by:

getAmount in interface IDose

Returns:

Concentration amount of Dose.

getTimeTake

public java.time.LocalTime getTimeTake()

Description copied from interface: IDose

Time when the dose is taken by patient.

Specified by:

getTimeTake in interface IDose

Returns:

LocalTime Time of Dose.

setTestDose

public void setTestDose()

Description copied from interface: IDose

Sets dose to test dose. Used for the what-if feature of the program.

Specified by:

setTestDose in interface IDose

toString

public java.lang.String toString()

Overrides:

toString in class java.lang.Object

PACKAGE CLASS TREE DEPRECATED INDEX HELP

PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Interface IAction

All Known Implementing Classes:

Action

public interface **IAction**

Method Summary

All Methods	Instance Methods	Abstract Methods
Modifier and Typ	oe e	Method and Description
void		addDose(IDose dose) Adds a new dose.
java.util.Ar	rayList <java.lang< td=""><td>.Double> getCurrentConcentration(java.time.LocalTime time)</td></java.lang<>	.Double> getCurrentConcentration(java.time.LocalTime time)
IMedicine		getMedicine() Retrieves the Medicine instance.
java.util.Ar	rayList <java.time< td=""><td>.LocalTime> getPeakConcentrationTime(java.lang.Boolean includeTestDoses</td></java.time<>	.LocalTime> getPeakConcentrationTime (java.lang.Boolean includeTestDoses
java.util.Ar	rrayList <java.time< td=""><td>.LocalTime> getWhenToDose(java.lang.Double concentrationDesired) Determines time when the patient should take next dose.</td></java.time<>	.LocalTime> getWhenToDose (java.lang.Double concentrationDesired) Determines time when the patient should take next dose.
void		<pre>loadFile(java.lang.String filename) Load a saved Medicine file.</pre>
void		<pre>newFile(java.lang.String name, java.time.LocalTime tMax, java.time.LocalTime halfLife) Create a new Medicine instance.</pre>
void		<pre>printCurrentConcentration(java.time.LocalTime time) Prints concentration amount of doses at a specific time.</pre>
void		<pre>printDoses() Prints all doses from Medicine's Dose array.</pre>
void		<pre>printMedicine()</pre> Print the name, tmax, and halfLife of the medicine.
void		<pre>printPeakConcentrationTime(java.lang.Boolean includeTestDos Print concentration peaks.</pre>
void		<pre>printWhenToDose(java.lang.Double amountDesired) Prints when the patient should take next dose.</pre>
void		removeAllDoses() Delete all doses from the dosages array.

void	<pre>removeDose(int index) Removes a dose from the dose from the doses array.</pre>
void	removeTestDoses() Removes all test doses.
void	<pre>saveFile(java.lang.String filename) Save Medicine to a file.</pre>

Method Detail

printDoses

void printDoses()

Prints all doses from Medicine's Dose array.

addDose

void addDose(IDose dose)

Adds a new dose. Added doses can be type Dose (actual dose) or TestDose (what-if dose)

Parameters:

dose - Dose object.

removeDose

void removeDose(int index)

Removes a dose from the dose from the doses array.

Parameters:

index - Index of the array to be erased

removeAllDoses

void removeAllDoses()

Delete all doses from the dosages array.

getPeakConcentrationTime

java.util.ArrayList<java.time.LocalTime> getPeakConcentrationTime(java.lang.Boolean includeTestDoses)

Time when the concentration is at its peak. It allows the user to choose if test doses will be displayed or not.

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

Returns:

ArrayList containing peak concentration peak values of each dose.

printPeakConcentrationTime

void printPeakConcentrationTime(java.lang.Boolean includeTestDoses)

Print concentration peaks. Includes dose information and its concentration peak.

Parameters:

includeTestDoses - Determines whether test doses will be displayed.

getWhenToDose

java.util.ArrayList<java.time.LocalTime> getWhenToDose(java.lang.Double concentrationDesired)

Determines time when the patient should take next dose.

Parameters:

concentrationDesired - Dose amount desired sought.

Returns:

Time when the patient have to take next dose.

printWhenToDose

void printWhenToDose(java.lang.Double amountDesired)

Prints when the patient should take next dose. Includes dose information and time to take next dose.

Parameters:

amountDesired - Dose amount desired sought.

saveFile

void saveFile(java.lang.String filename)

Save Medicine to a file. By default, save directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Parameters:

filename - Name of save file to be saved.

loadFile

```
void loadFile(java.lang.String filename)
```

Load a saved Medicine file. By default, load directory is same directory as the application. Dose and Medicine MUST implement Serializable for this feature to work.

Parameters:

filename - Name of the file to be loaded

newFile

Create a new Medicine instance.

Parameters:

name - Name of medicine

tMax - TMax of medicine.

halfLife - Half life of medicine

printMedicine

void printMedicine()

Print the name, tmax, and halfLife of the medicine.

getCurrentConcentration

java.util.ArrayList<java.lang.Double> getCurrentConcentration(java.time.LocalTime time)

Calculates concentration amount of doses at a specific time.

Parameters:

time - Specific time used to determine concentration amount.

Returns:

ArrayList with the concentration amounts.

printCurrentConcentration

void printCurrentConcentration(java.time.LocalTime time)

Prints concentration amount of doses at a specific time.

Parameters:

time - Specific time used to determine concentration amount.

removeTestDoses

void removeTestDoses()

Removes all test doses.

getMedicine

IMedicine getMedicine()

Retrieves the Medicine instance.

Returns:

Medicine instance

PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

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ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Interface IDose

All Known Implementing Classes:

Dose

public interface IDose

Method Summary

All Methods Instance	Methods Abstract Methods
Modifier and Type	Method and Description
void	<pre>createDose(java.time.LocalTime timeTake, double amount) Creates a Dose object.</pre>
double	getAmount() Concentration amount of the dose taken by patient.
java.time.LocalTime	getTimeTake() Time when the dose is taken by patient.
boolean	isTestDose() Identifies dose type.
void	setTestDose() Sets dose to test dose.

Method Detail

isTestDose

boolean isTestDose()

Identifies dose type.

Returns:

True = TestDose (what-if dose); False = Dose (actual dose);

createDose

Creates a Dose object.

Parameters:

timeTake - Time when the dose is taken by patient.

amount - Concentration amount of the dose taken by patient.

getTimeTake

java.time.LocalTime getTimeTake()

Time when the dose is taken by patient.

Returns:

LocalTime Time of Dose.

getAmount

double getAmount()

Concentration amount of the dose taken by patient.

Returns:

Concentration amount of Dose.

setTestDose

void setTestDose()

Sets dose to test dose. Used for the what-if feature of the program.

PACKAGE CLASS TREE DEPRECATED INDEX HELP

PREV CLASS NEXT CLASS FRAMES NO FRAMES ALL CLASSES

PREV CLASS NEXT CLASS FRAMES

FRAMES NO FRAMES

ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Interface IMedicine

All Known Implementing Classes:

Medicine

public interface IMedicine

Method Summary

All Methods Instance Methods	Abstract Methods
Modifier and Type	Method and Description
void	<pre>createMedicine(java.lang.String name, java.time.LocalTime tMax, java.time.LocalTime halfLife) Creates a medicine object.</pre>
java.util.ArrayList <idose></idose>	getDoses() Corresponds to all existing doses.
java.time.LocalTime	<pre>getHalfLife()</pre> Half life time of the medicine.
java.lang.String	getName() Name of medicine.
java.time.LocalTime	getTmax() Tmax of medicine

Method Detail

createMedicine

Creates a medicine object.

Parameters:

name - Name of medicine.

tMax - Time when medicine is at its peak concentration.

halfLife - Time required for medicine to decrease by half.

getDoses

java.util.ArrayList<IDose> getDoses()

Corresponds to all existing doses. Includes type Dose and TestDose.

Returns:

An array containing all doses.

getName

java.lang.String getName()

Name of medicine.

Returns:

Return name of the medicine.

getHalfLife

java.time.LocalTime getHalfLife()

Half life time of the medicine.

Returns:

Return the half life time of the medicine.

getTmax

java.time.LocalTime getTmax()

Tmax of medicine

Returns:

Return TMax of medicine.

PACKAGE CLASS TREE DEPRECATED INDEX HELP

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PREV CLASS NEXT CLASS

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ALL CLASSES

SUMMARY: NESTED | FIELD | CONSTR | METHOD DETAIL: FIELD | CONSTR | METHOD

Class Medicine

java.lang.Object Medicine

All Implemented Interfaces:

IMedicine, java.io.Serializable

public class Medicine
extends java.lang.Object
implements IMedicine, java.io.Serializable

See Also:

Serialized Form

Field Summary

Fields

Modifier and Type	Field and Description
<pre>private java.util.ArrayList<idose></idose></pre>	doses
private java.time.LocalTime	halfLife
private java.lang.String	name
private java.time.LocalTime	tmax

Constructor Summary

Constructors

Constructor and Description

Medicine()

Medicine(java.lang.String name, java.time.LocalTime tMax,
java.time.LocalTime halfLife)

Method Summary

All Methods Instance Methods	S Concrete Methods
Modifier and Type	Method and Description
void	<pre>createMedicine(java.lang.String name, java.time.LocalTime tMax, java.time.LocalTime halfLife) Creates a medicine object.</pre>
<pre>java.util.ArrayList<idose></idose></pre>	getDoses() Corresponds to all existing doses.
java.time.LocalTime	getHalfLife() Half life time of the medicine.
java.lang.String	getName() Name of medicine.
java.time.LocalTime	getTmax() Tmax of medicine
java.lang.String	toString()

Methods inherited from class java.lang.Object

clone, equals, finalize, getClass, hashCode, notify, notifyAll, wait, wait, wait

Field Detail

name

private java.lang.String name

tmax

private java.time.LocalTime tmax

halfLife

private java.time.LocalTime halfLife

doses

private java.util.ArrayList<IDose> doses

Constructor Detail

Medicine

```
public Medicine()
```

Medicine

Method Detail

createMedicine

Description copied from interface: IMedicine

Creates a medicine object.

Specified by:

createMedicine in interface IMedicine

Parameters:

```
name - Name of medicine.

tMax - Time when medicine is at its peak concentration.
halfLife - Time required for medicine to decrease by half.
```

getName

```
public java.lang.String getName()
```

Description copied from interface: IMedicine

Name of medicine.

Specified by:

getName in interface IMedicine

Returns:

Return name of the medicine.

getHalfLife

public java.time.LocalTime getHalfLife()

Description copied from interface: IMedicine

Half life time of the medicine.

Specified by:

getHalfLife in interface IMedicine

Returns:

Return the half life time of the medicine.

getDoses

public java.util.ArrayList<IDose> getDoses()

Description copied from interface: IMedicine

Corresponds to all existing doses. Includes type Dose and TestDose.

Specified by:

getDoses in interface IMedicine

Returns:

An array containing all doses.

getTmax

public java.time.LocalTime getTmax()

Description copied from interface: IMedicine

Tmax of medicine

Specified by:

getTmax in interface IMedicine

Returns:

Return TMax of medicine.

toString

public java.lang.String toString()

Overrides:

toString in class java.lang.Object

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PREV PACKAGE NEXT PACKAGE FRAMES NO FRAMES ALL CLASSES

Package <Unnamed>

Interface	Summary
-----------	----------------

Interface	Description	
IAction		
IDose		
IMedicine		

Class Summary

-		
Class	Description	
Action		
Controller		
Dose		
Medicine		

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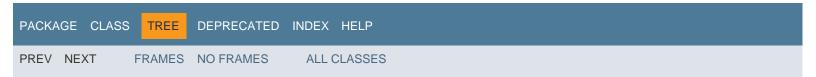
Hierarchy For Package < Unnamed>

Class Hierarchy

- o java.lang.Object
 - Action (implements lAction)
 - Controller
 - **Dose** (implements IDose, java.io.Serializable)
 - Medicine (implements IMedicine, java.io.Serializable)

Interface Hierarchy

- IAction
- IDose
- IMedicine

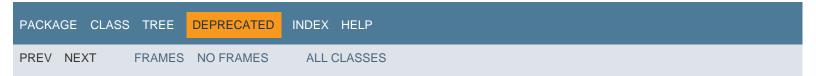


PACKAGE CLASS TREE DEPRECATED INDEX HELP

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Deprecated API

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Α

Action - Class in <Unnamed>

Action() - Constructor for class Action

action - Static variable in class Controller

addDose(IDose) - Method in class Action

addDose(IDose) - Method in interface IAction

Adds a new dose.

amount - Variable in class Dose

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How This API Document Is Organized

This API (Application Programming Interface) document has pages corresponding to the items in the navigation bar, described as follows.

Package

Each package has a page that contains a list of its classes and interfaces, with a summary for each. This page can contain six categories:

- Interfaces (italic)
- Classes
- Enums
- Exceptions
- Errors
- Annotation Types

Class/Interface

Each class, interface, nested class and nested interface has its own separate page. Each of these pages has three sections consisting of a class/interface description, summary tables, and detailed member descriptions:

- Class inheritance diagram
- Direct Subclasses
- All Known Subinterfaces
- · All Known Implementing Classes
- Class/interface declaration
- Class/interface description
- Nested Class Summary
- Field Summary
- Constructor Summary
- Method Summary
- Field Detail
- Constructor Detail
- Method Detail

Each summary entry contains the first sentence from the detailed description for that item. The summary entries are alphabetical, while the detailed descriptions are in the order they appear in the source code. This preserves the logical groupings established by the programmer.

Annotation Type

Each annotation type has its own separate page with the following sections:

- Annotation Type declaration
- Annotation Type description
- Required Element Summary
- Optional Element Summary

Element Detail

Enum

Each enum has its own separate page with the following sections:

- Enum declaration
- Enum description
- Enum Constant Summary
- Enum Constant Detail

Tree (Class Hierarchy)

There is a Class Hierarchy page for all packages, plus a hierarchy for each package. Each hierarchy page contains a list of classes and a list of interfaces. The classes are organized by inheritance structure starting with <code>java.lang.Object</code>. The interfaces do not inherit from <code>java.lang.Object</code>.

- When viewing the Overview page, clicking on "Tree" displays the hierarchy for all packages.
- When viewing a particular package, class or interface page, clicking "Tree" displays the hierarchy for only that package.

Deprecated API

The Deprecated API page lists all of the API that have been deprecated. A deprecated API is not recommended for use, generally due to improvements, and a replacement API is usually given. Deprecated APIs may be removed in future implementations.

Index

The Index contains an alphabetic list of all classes, interfaces, constructors, methods, and fields.

Prev/Next

These links take you to the next or previous class, interface, package, or related page.

Frames/No Frames

These links show and hide the HTML frames. All pages are available with or without frames.

All Classes

The All Classes link shows all classes and interfaces except non-static nested types.

Serialized Form

Each serializable or externalizable class has a description of its serialization fields and methods. This information is of interest to re-implementors, not to developers using the API. While there is no link in the navigation bar, you can get to this information by going to any serialized class and clicking "Serialized Form" in the "See also" section of the class description.

Constant Field Values

The Constant Field Values page lists the static final fields and their values.

This help file applies to API documentation generated using the standard doclet.

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All Classes

Action

Controller

Dose

IAction

IDose

IMedicine

Medicine

PREV NEXT FRAMES NO FRAMES

ALL CLASSES

Serialized Form

Package < Unnamed>

Class Dose extends java.lang.Object implements Serializable

Serialized Fields

timeTake

java.time.LocalTime timeTake

amount

double amount

isTestDose

boolean isTestDose

Class Medicine extends java.lang.Object implements Serializable

Serialized Fields

name

java.lang.String name

tmax

java.time.LocalTime tmax

halfLife

java.time.LocalTime halfLife

doses

java.util.ArrayList<E> doses

PACKAGE CLASS TREE DEPRECATED INDEX HELP

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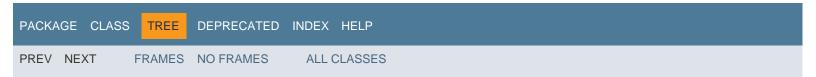
Hierarchy For All Packages

Class Hierarchy

- o java.lang.Object
 - Action (implements lAction)
 - Controller
 - **Dose** (implements IDose, java.io.Serializable)
 - Medicine (implements IMedicine, java.io.Serializable)

Interface Hierarchy

- IAction
- IDose
- IMedicine



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C

checkIfInteger() - Static method in class Controller

Validates if user input is an integer only.

checklfValidNumber() - Static method in class Controller

Validates if user input is a number, either Integer or Double.

clear() - Static method in class Controller

Clears the console screen.

Controller - Class in <Unnamed>

Controller() - Constructor for class Controller

createDose() - Static method in class Controller

Creates Dose from user input.

createDose(LocalTime, double) - Method in class Dose

createDose(LocalTime, double) - Method in interface IDose

Creates a Dose object.

createLocalTime() - Static method in class Controller

Generates LocalTime instance from user input.

createMedicine() - Static method in class Controller

Creates a Medicine instance from user input.

createMedicine(String, LocalTime, LocalTime) - Method in interface IMedicine

Creates a medicine object.

createMedicine(String, LocalTime, LocalTime) - Method in class Medicine

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D

Dose - Class in <Unnamed>

Dose() - Constructor for class Dose

Dose(LocalTime, double) - Constructor for class Dose

doses - Variable in class Medicine

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G

getAmount() - Method in class Dose

getAmount() - Method in interface IDose

Concentration amount of the dose taken by patient.

getCurrentConcentration(LocalTime) - Method in class Action

getCurrentConcentration(LocalTime) - Method in interface IAction

Calculates concentration amount of doses at a specific time.

getDoses() - Method in interface IMedicine

Corresponds to all existing doses.

getDoses() - Method in class Medicine

getHalfLife() - Method in interface IMedicine

Half life time of the medicine.

getHalfLife() - Method in class Medicine

getMedicine() - Method in class Action

getMedicine() - Method in interface IAction

Retrieves the Medicine instance.

getName() - Method in interface IMedicine

Name of medicine.

getName() - Method in class Medicine

getPeakConcentrationTime(Boolean) - Method in class Action

getPeakConcentrationTime(Boolean) - Method in interface | Action

Time when the concentration is at its peak.

getTimeTake() - Method in class Dose

getTimeTake() - Method in interface IDose

Time when the dose is taken by patient.

getTmax() - Method in interface IMedicine

Tmax of medicine

getTmax() - Method in class Medicine

getWhenToDose(Double) - Method in class Action

getWhenToDose(Double) - Method in interface IAction

Determines time when the patient should take next dose.

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Н

halfLife - Variable in class Medicine

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ı

IAction - Interface in <Unnamed>

IDose - Interface in <Unnamed>

IMedicine - Interface in <Unnamed>

isTestDose - Variable in class Dose

isTestDose() - Method in class Dose

isTestDose() - Method in interface IDose Identifies dose type.

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L

loadFile(String) - Method in class Action

loadFile(String) - Method in interface lAction Load a saved Medicine file.

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M

main(String[]) - Static method in class Controller

medicine - Variable in class Action

Medicine - Class in <Unnamed>

Medicine() - Constructor for class Medicine

Medicine(String, LocalTime, LocalTime) - Constructor for class Medicine

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Ν

name - Variable in class Medicine

newFile(String, LocalTime, LocalTime) - Method in class Action

newFile(String, LocalTime, LocalTime) - Method in interface lAction Create a new Medicine instance.

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P

pause() - Static method in class Controller

Pause scrolling fot the console screen until user hits the Enter key. printCurrentConcentration(LocalTime) - Method in class Action

printCurrentConcentration(LocalTime) - Method in interface | Action

Prints concentration amount of doses at a specific time.

printDoses() - Method in class Action

printDoses() - Method in interface IAction

Prints all doses from Medicine's Dose array.

printMedicine() - Method in class Action

printMedicine() - Method in interface IAction

Print the name, tmax, and halfLife of the medicine.

printPeakConcentrationTime(Boolean) - Method in class Action

printPeakConcentrationTime(Boolean) - Method in interface IAction

Print concentration peaks.

printWhenToDose(Double) - Method in class Action

printWhenToDose(Double) - Method in interface IAction

Prints when the patient should take next dose.

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R

removeAllDoses() - Method in class Action

removeAllDoses() - Method in interface IAction

Delete all doses from the dosages array.

removeDose(int) - Method in class Action

removeDose() - Static method in class Controller

Removes dose using an index selected by the user.

removeDose(int) - Method in interface IAction

Removes a dose from the dose from the doses array.

removeTestDoses() - Method in class Action

removeTestDoses() - Method in interface IAction

Removes all test doses.

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S

saveFile(String) - Method in class Action

saveFile(String) - Method in interface IAction

Save Medicine to a file.

selectAction() - Static method in class Controller

Display the list of actions that can be performed by the application.

setTestDose() - Method in class Dose

setTestDose() - Method in interface IDose

Sets dose to test dose.

start() - Static method in class Controller

Starts the program.

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Т

timeTake - Variable in class Dose

tmax - Variable in class Medicine

toString() - Method in class Dose

toString() - Method in class Medicine

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U

userInput - Static variable in class Controller

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PACKAGE CLASS TREE DEPRECATED INDEX HELP

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Constant Field Values

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