# POMDPFiles.jl

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Part I

Home

## **Chapter 1**

### **Home**

This is the first page of my documentation.

#### 1.1 Types

POMDPFiles.ContainerNames - Type.

```
ContainerNames

names: Vector{String} - a vector with the names of the actions, states, or observations.

number: Int - the number of actions, states, or observations.

This type was created to store the names and number of actions, states, and observations when 
→ parsed from the preamble of a .pomdp file format.
```

source

 ${\tt POMDPFiles.InitialStateParam-Type.}$ 

```
InitialStateParam was created to store the initial state distribution of a POMDP. The \hookrightarrow distribution can be of any type, but it is stored as a vector of Float64 values.
```

source

POMDPFiles.SFilePOMDP - Type.

```
SFilePOMDP is used whenever the names of the states, actions, and observations are known. It is

→ used to represent a POMDP problem from a file.

dic_states: dictionary with the names of the states
dic_actions: dictionary with the names of the actions
dic_obs: dictionary with the names of the observations
pomdp: FilePOMDP structure
```

source

POMDPFiles.FilePOMDP - Type.

```
FilePOMDP is the main data structure of the package. It is used to represent a POMDP problem

→ from a file.

ns: number of states
na: number of actions
no: number of observations

support_initialstate: support of the initial state distribution
initialstate_distribution: initial state distribution

discount: discount factor

T: transition matrix
0: observation matrix
R: reward matrix
```

source

#### 1.2 Functions

POMDPFiles.test\_if\_probability - Function.

```
test_if_probability(prob::Union{Vector{Float64}, Vector{Nothing}, Nothing};rtol=le-3)

Built-in function that tests whether a vector is a probability distribution. It checks if the elements are between 0 and 1 and if the sum of the elements is approximately 1. The function returns true if the vector is a probability distribution and false otherwise.
```

source

POMDPFiles.read pomdp - Function.

```
Read a .pomdp file following the specfication at http://www.pomdp.org/code/pomdp-file-spec.html \hookrightarrow and returns a FilePOMDP or SFilePOMDP object that can be used within the POMDPs.jl \hookrightarrow interface.
```

source

Base.write - Function.

Writes out the alpha vectors in the .alpha file format

source

 $Write \ out \ a \ . pomdp \ file \ using \ the \ POMDPs. jl \ interface \ Specification: \ http://cs.brown.edu/research/ai/pomdp/examples/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ of \ the \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ version \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ pomdp \ spec: \ https://pomdp.org/code/pomdp-file-spec.html \ A \ more \ recent \ pomdp \ spec: \ https://pomdp-file-spec.html \ A \ more \ recent \ pomdp \ spec: \ https://pomdp-file-spec.html \ A \ more \ recent \ pomdp \ spec: \ https://pomdp-file-spec.html \ pomdp \ spec:$ 

source

POMDPFiles.prob - Function.

```
value returns the a vector with the initial state distribution.
```

source

POMDPFiles.number - Function.

```
number returns the number of actions, states, or observations in the ContainerNames object.
```

source

number returns the number of states **in** the initial state distribution.

source

POMDPFiles.remove comments and white space - Function.

```
remove_comments_and_white_space(file::Vector{String}) is used by read_pomdp to remove comments \hookrightarrow and white spaces from the file. This function allows for some standardization of process of \hookrightarrow parsing files.
```

source

POMDPFiles.read\_alpha - Function.

```
Read a `.alpha` file as generated by pomdp-solve.
Works the same was as `read_pomdp` in `POMDPXFile.jl`.

The `.alpha` file format is recapped here as follows,
see: http://www.pomdp.org/code/alpha-file-spec.html

A set of vectors is the representation use for the value function and each
vector has an action associated with it. The vectors represent the coefficients
of a hyperplane passing through the origin. The format specified here is what is
output from the [pomdp-solve' program and what will be necessary for input to
the [pomdp-solve' program with the []-terminal_values' command line option.

The format is simply:

A
V1 V2 V3 ... VN

A
V1 V2 V3 ... VN

...

Where `A` is an action number and the `V1` through `VN` are real values
```

representing the components of a particular vector that has the associated action. The action number is the  $\theta$ -based index of the action as specificed in the input POMDP file. The vector represents the coefficients of a hyperplane representing one facet of the piecewise linear and convex value function. Note that the length of the lists needs to be equal to the number of states in the POMDP.

To find which action is the "best" for a given set of alpha vectors, the belief state probabilities would be used in a dot product against each alpha vectors' coefficients. The action associated with the vector with the highest value is the best action to take for that belief state given the value function.

source

POMDPs.states - Function.

Implementing the functions required by the POMDP interface. See

→ [POMDPs.jl](https://juliapomdp.github.io/POMDPs.jl/latest/) for more details on the

→ interface.

#### source

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source

POMDPFiles.convert\_to\_data\_structure - Function.

 $\label{local_convert_to_data_structure} convert\_to\_data\_structure(field::String, preamble::Dict{String,String}) \ is used by read\_pomdp \ to \\ \hookrightarrow \ convert \ the \ information \ in \ the \ preamble \ into \ an \ intermidiate \ format \ before \ passing \ it \ into \ a \\ \hookrightarrow \ ContainerNames \ object.$ 

source

POMDPFiles.check preamble fields - Function.

source

POMDPFiles.support - Function.

support returns the support of the initial state distribution.

source

POMDPFiles.process\_preamble - Function.

```
process_preamble(preamble::Dict{String, String}) is used by read_pomdp to process the preamble

→ of the file and check if the fields "discount", "values", "states", "actions", and

→ "observations" have the correct syntax. The output are the discount, values, actions,

→ states, and observations parameters, where actions, states, and observations are converted

→ into ContainerNames objects.
```

#### source

 ${\tt POMDPFiles.order\_of\_transition\_reward\_observation-Function}.$ 

```
order_of_transition_reward_observation(file_lines::Vector{String}, start_line::Int64) is used by \hookrightarrow read_pomdp to find the order of the transition, reward, and observation matrices in the \hookrightarrow file.
```

source