

# ORSO Lipid Example 1 - DSPC Bilayer Analysed Using a Standard Layers Model.

Start by making an empty instance of the project definition class:

```
problem = projectClass('Orso lipid example')
```

```
problem =  
  experimentType: 'Standard Layers with (d,rho,rough) layers'  
  experimentName: 'Orso lipid example'  
  Geometry: 'air/substrate'
```

Parameters: -----

p	Name	Min	Value	Max	Fit?
1	"Substrate Roughness"	1	3	5	true

Layers: -----

Name	Thickness	SLD	Roughness	Hydration	Hydrate with
0	0	0	0	0	0

Bulk In: -----

p	Name	Min	Value	Max	Fit?
1	"SLD Air"	0	0	0	false

Bulk Out: -----

p	Name	Min	Value	Max	Fit?
1	"SLD D2O"	6.2e-06	6.35e-06	6.35e-06	false

Scalefactors: -----

p	Name	Min	Value	Max	Fit?
1	"Scalefactor 1"	0.02	0.23	0.25	false

Backgrounds: -----

(a) Background Parameters:

p	Name	Min	Value	Max	Fit?
1	"Backs par 1"	1e-07	1e-06	1e-05	false

(b) Backgrounds:

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
---	------	------	---------	---------	---------	---------	---------

```
1      "Background 1"      "constant"      "Backs Par 1"      " "      " "      " "      " "
```

```
Resolutions: -----
```

```
(a) Resolutions Parameters:
```

p	Name	Min	Value	Max	Fit?
1	"Resolution par 1"	0.01	0.03	0.05	false

```
(b) Resolutions:
```

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
1	"Resolution 1"	"gaussian"	"Resolution par 1"	" "	" "	" "	" "

```
Data: -----
```

Name	Data	Data Range	Simulation Range
"Data 1"	"No Data"	"-"	"[ 0.0050 , 0.7000 ]"

```
Constrasts: -----
```

```

p
-----
"name"
"Data"
"Background"
"Bulk in"
"Bulk out"
"Scalefactor"
"Resolution"
"Model"
```

This is the standard class containing the model definition for any project. We add to it using the class methods, which are fairly self explanatory (although more detailed instructions are pending) -

```
methods(problem)
```

```
Methods for class projectClass:
```

addBackground	addLayer	getAllAllowedNames	setBackgroundValue	setBulkOut	setPar
addBacksPar	addLayerGroup	projectClass	setBacksPar	setContrast	setPar
addBulkIn	addParam	removeBacksPar	setBacksParConstr	setData	setPar
addBulkOut	addParamGroup	removeData	setBacksParName	setGeometry	setPar
addContrast	addQzshift	removeParam	setBacksParValue	setLayerValue	setPar
addData	addScalefactor	setBackgroundName	setBulkIn	setParamConstr	setSca

```
Methods of projectClass inherited from handle.
```

In this example, we will use a standard layers model. This defines layers in terms of thickness, SLD, roughness and hydration. These are then grouped into layers and the allocated to contrasts in the normal RasCAL way. For this sample, we have a DSPC bilayer on a Silicon substrate, which will be coated by an oxide layer. Ususally there is a hydration layer between the substrate and the lipid, although this depends on the exact conditions

of the measurement. Start by defining the parameters for the oxide layer. We need a thickness, SLD and hydration. The roughness is assumed to be the bulk substrate roughness.

```
Parameters = {
    %           Name           min           val           max           fit?
    {'Oxide thick',          10,          20,          25,          true   };
    {'Oxide SLD',           3e-6,         3.41e-6,         4e-6,         false  };
    {'Oxide Hydration'       0,           20,           30,           true   }};
```

Add these parameters to the project class, and group them into a layer to demonstrate how this is done:

```
problem.addParamGroup(Parameters)
```

```
ans =
  experimentType: 'Standard Layers with (d,rho,rough) layers'
  experimentName: 'Orso lipid example'
  Geometry: 'air/substrate'
```

Parameters: -----

p	Name	Min	Value	Max	Fit?
1	"Substrate Roughness"	1	3	5	true
2	"Oxide thick"	10	20	25	true
3	"Oxide SLD"	3e-06	3.41e-06	4e-06	false
4	"Oxide Hydration"	0	20	30	true

Layers: -----

Name	Thickness	SLD	Roughness	Hydration	Hydrate with
0	0	0	0	0	0

Bulk In: -----

p	Name	Min	Value	Max	Fit?
1	"SLD Air"	0	0	0	false

Bulk Out: -----

p	Name	Min	Value	Max	Fit?
1	"SLD D2O"	6.2e-06	6.35e-06	6.35e-06	false

Scalefactors: -----

p	Name	Min	Value	Max	Fit?
1	"Scalefactor 1"	0.02	0.23	0.25	false

Backgrounds: -----

(a) Background Parameters:

p	Name	Min	Value	Max	Fit?
---	------	-----	-------	-----	------

1	"Backs par 1"	1e-07	1e-06	1e-05	false
---	---------------	-------	-------	-------	-------

(b) Backgrounds:

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
1	"Background 1"	"constant"	"Backs Par 1"	" "	" "	" "	" "

Resolutions: -----

(a) Resolutions Parameters:

p	Name	Min	Value	Max	Fit?
1	"Resolution par 1"	0.01	0.03	0.05	false

(b) Resolutions:

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
1	"Resolution 1"	"gaussian"	"Resolution par 1"	" "	" "	" "	" "

Data: -----

Name	Data	Data Range	Simulation Range
"Data 1"	"No Data"	"-"	"[ 0.0050 , 0.7000 ]"

Constrasts: -----

p

---

"name"  
 "Data"  
 "Background"  
 "Bulk in"  
 "Bulk out"  
 "Scalefactor"  
 "Resolution"  
 "Model"

```
% Now make the oxide layer
Oxide = { 'Oxide Layer',...           % Name of the layer
          'Oxide thick',...          % Layer thickness
          'Oxide SLD',...            % Layer SLD
          'Substrate Roughness',...  % Layer roughness
          'Oxide Hydration',...      % Oxide hydration (precent)
          'bulk out' };              % Which bulk phase is hydrating the layer

% Add this to the project...
problem.addLayer(Oxide)
```

```
ans =
experimentType: 'Standard Layers with (d,rho,rough) layers'
experimentName: 'Orso lipid example'
Geometry: 'air/substrate'
```

Parameters: -----

p	Name	Min	Value	Max	Fit?
1	"Substrate Roughness"	1	3	5	true
2	"Oxide thick"	10	20	25	true
3	"Oxide SLD"	3e-06	3.41e-06	4e-06	false
4	"Oxide Hydration"	0	20	30	true

Layers: -----

p	Name	Thickness	SLD	Roughness	Hydration	Hy
1	"Oxide Layer"	"Oxide thick"	"Oxide SLD"	"Substrate Roughness"	"Oxide Hydration"	"

Bulk In: -----

p	Name	Min	Value	Max	Fit?
1	"SLD Air"	0	0	0	false

Bulk Out: -----

p	Name	Min	Value	Max	Fit?
1	"SLD D2O"	6.2e-06	6.35e-06	6.35e-06	false

Scalefactors: -----

p	Name	Min	Value	Max	Fit?
1	"Scalefactor 1"	0.02	0.23	0.25	false

Backgrounds: -----

(a) Background Parameters:

p	Name	Min	Value	Max	Fit?
1	"Backs par 1"	1e-07	1e-06	1e-05	false

(b) Backgrounds:

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
1	"Background 1"	"constant"	"Backs Par 1"	" "	" "	" "	" "

Resolutions: -----

(a) Resolutions Parameters:

p	Name	Min	Value	Max	Fit?
1	"Resolution par 1"	0.01	0.03	0.05	false

(b) Resolutions:

p	Name	Type	Value 1	Value 2	Value 3	Value 4	Value 5
1	"Resolution 1"	"gaussian"	"Resolution par 1"	" "	" "	" "	" "

Data: -----

Name	Data	Data Range	Simulation Range
"Data 1"	"No Data"	"-"	"[ 0.0050 , 0.7000 ]"

Constrasts: -----

**p**

"name"  
 "Data"  
 "Background"  
 "Bulk in"  
 "Bulk out"  
 "Scalefactor"  
 "Resolution"  
 "Model"

Now we'll build the layers for the lipid, The bilayer is symmetrical, so we can have mostly the same values for the inner and outer tails and heads. We have to be a bit careful with roughness' however, since we want to have a slightly different roughness for the head/water interface and the head/alkyl interface, and for layer model in this geometry the roughness for each layer refers to the next interface. So we'll make separate layers for inner and outer lipids, but all sharing the same parameters, and just differing in the roughness order.

Define the parameters we will need:

```
% Bilayer parameters:
bilParams = {
    {'Bilayer head thick', 5, 10, 15, true };
    {'Bilayer head sld', 1e-6, 1.47e-6, 1.6e-6, true };
    {'Bilayer heads hydr', 20, 30, 40, true };
    {'Bilayer Heads rough', 5, 7, 15, true };
    {'Bilayer tails thick', 10, 15, 20, true };
    {'Bilayer tails SLD', -5e07, -4e-7, -1e-7, false };
    {'Bilayer tails hydr', 0, 10, 20, true };
    {'Bilayer tails rough', 5, 9, 15, true };
};

% We want the water layer to be matched in SLD to the bulk water phase. So
% we give it an SLD of 0, and a hydration of 100%, and make these
% non-fittable

watParams = {
    {'Water thick', 5, 1, 15, true };
    {'Water SLD', 0, 0, 0, false };
}

}
```