

```
- <sbml level="2" version="4">  
  - <model id="maiage_bsub_01" name="Metabolism of B. subtilis">  
    + <listOfCompartments></listOfCompartments>  
    + <listOfSpecies></listOfSpecies>  
    + <listOfReactions></listOfReactions>  
  </model>  
</sbml>
```

Standard SBML file, see SBML specifications. Compartments are needed for density constraints. Species and reactions are needed to build the stoichiometry matrix.