







# **Developers Meeting**

Models2Pathways

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#### Models2Pathways

#### Purpose:

A tool to find significant BioModels, matching to Reactome Pathways, based on the Results of the Reactome Analysis Tool, during Release Process.





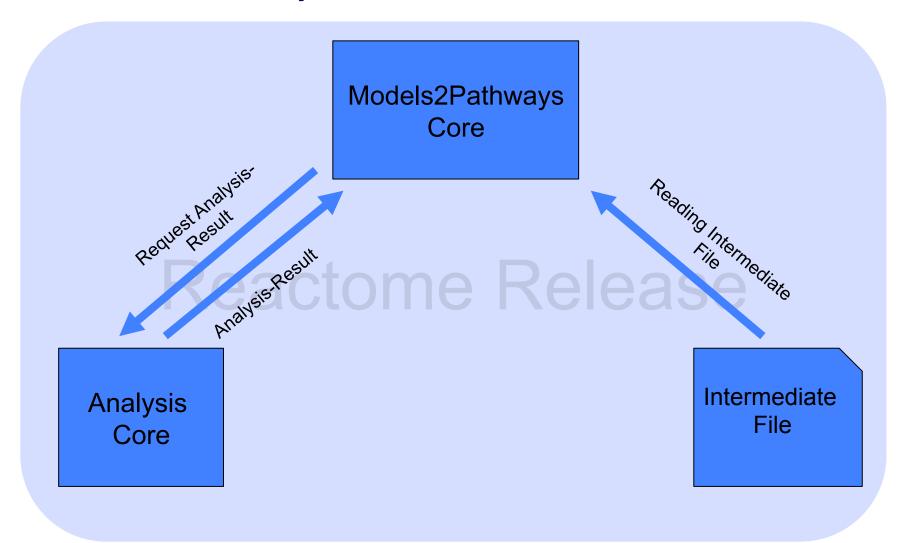
## Reactome Release

Analysis Core Produces Analysis Intermediate File

Intermediate File



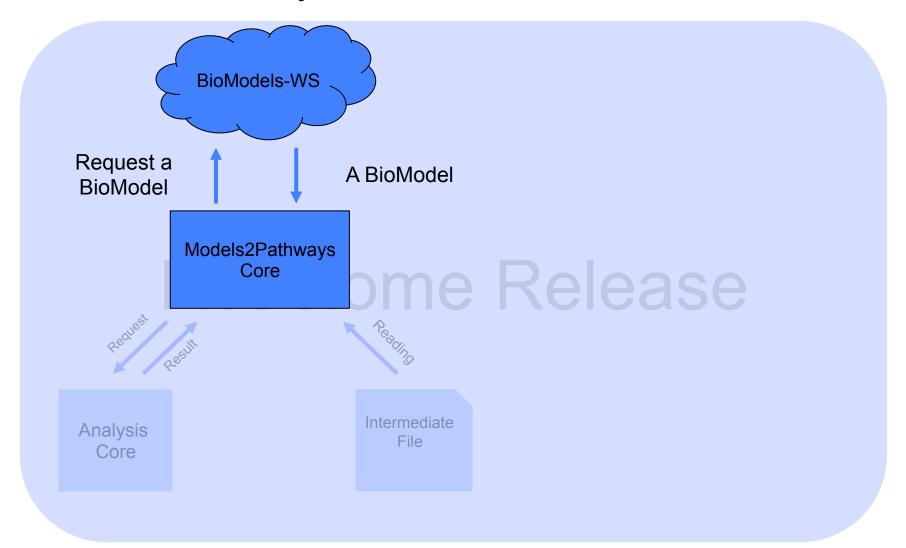






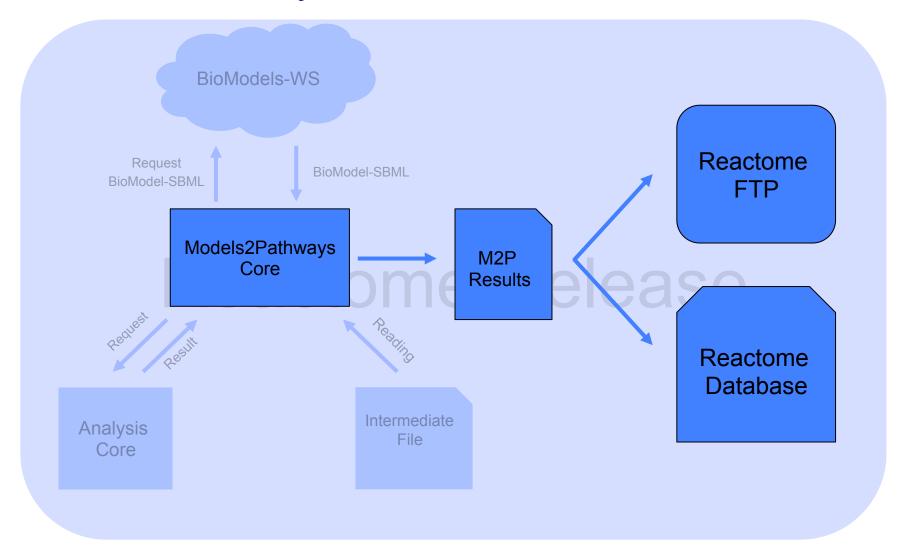
















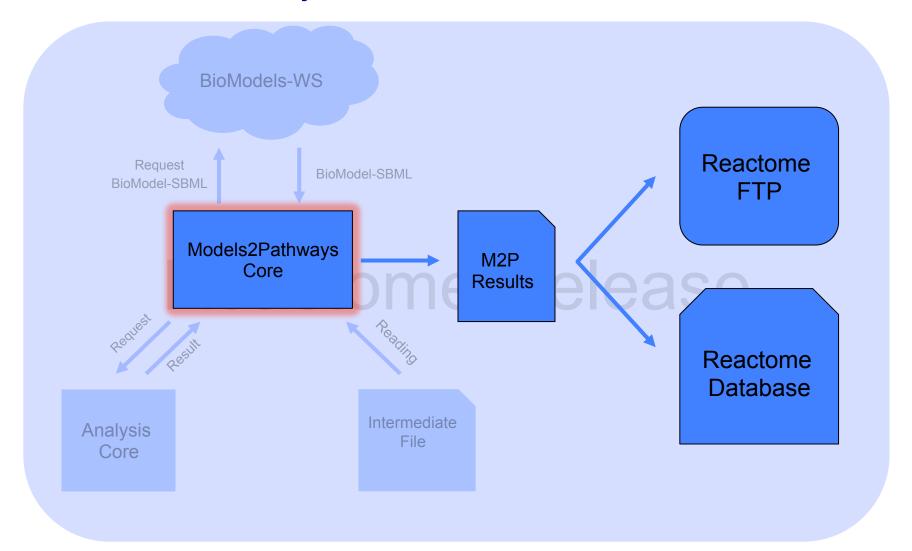


#### Models2Pathways - Significant Pathways

public List<PathwaySummary> getReliablePathways( List < PathwaySummary > pathways, Double customFDR. BioModel bioModel. Double reactionCoverage) List < Pathway Summary > reliable Pathways = new ArrayList <> () for (PathwaySummary pathway: pathways) pathway.isLlp() pathway.getEntities().getFdr() <= customFDR pathway.getSpecies() == bimodel.getSpecies() pathway.getEntities().getFound() / pathway.getEntities().getTotal() >= reactionCoverage, Ø Ø reliablePathways.add(pathway) return reliablePathways













## Models2Pathways – Core

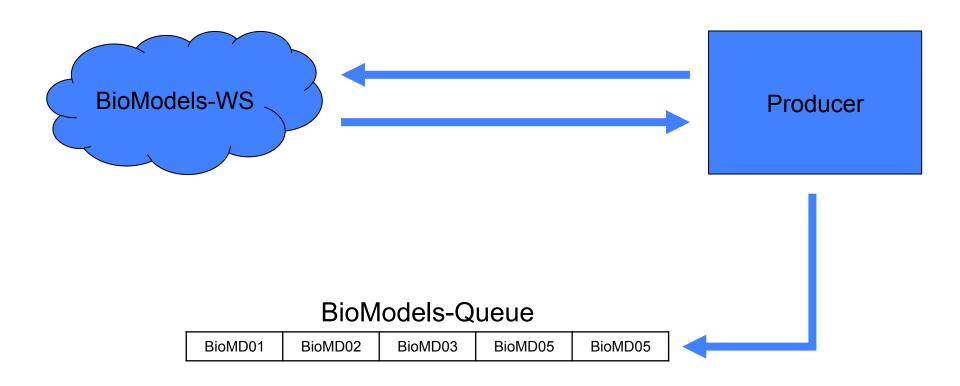
Producer

Consumer





### Models2Pathways – Core







## Models2Pathways – Core

