

## 4.4 - Final Library Preparation

### Parameters

Parameter Name	Description
LIB_FILE_NAME	the file from which fragments are read, and to which new fragments are written

### Overview

During the "Final Library Preparation" step of the Flux Simulator pipeline the cDNA fragments are optionally filtered by gel segregation, and also optionally amplified by a PCR (i.e., polymerase chain reaction).

### Details

#### [4.4.1 - Size Selection](#)

#### [4.4.2 - PCR Amplification](#)