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