Pathway Descriptions

pathway_descriptions: Internal script that matches KO genes with pathway whilst displaying the total abundance in the dataset for each pathway. Pathway descriptions are also included.
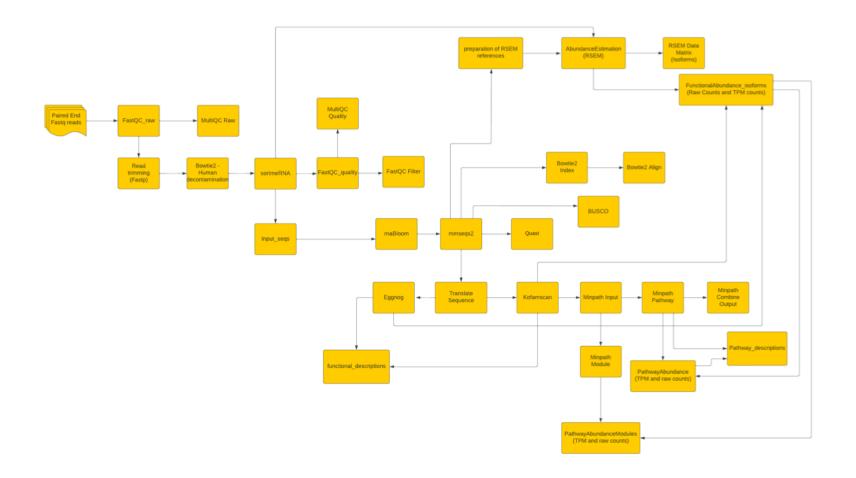

Figure 1: Overview of the Metatranscriptomics pipeline

2   Pipeline Run

Pipeline user: {user-id}

Study name: {study_name}

Study link:
    {pipeline-run-url}

Pipeline: Metatranscriptomics Pipeline

Pipeline version: v1.1