This diagram outlines the process of inferring subcellular location from protein sequence data. It is divided into two parts:

**Part I**
- **AC** → GOA Database
- **GO Extraction by searching GOA database**

**Part II**
- **S** → BLAST → Swiss-Prot Database
- **n ACs**
- **GO Vectors Construction**
- **Multi-label SVM + Score Fusion**

The process flow leads to the inference of subcellular location.