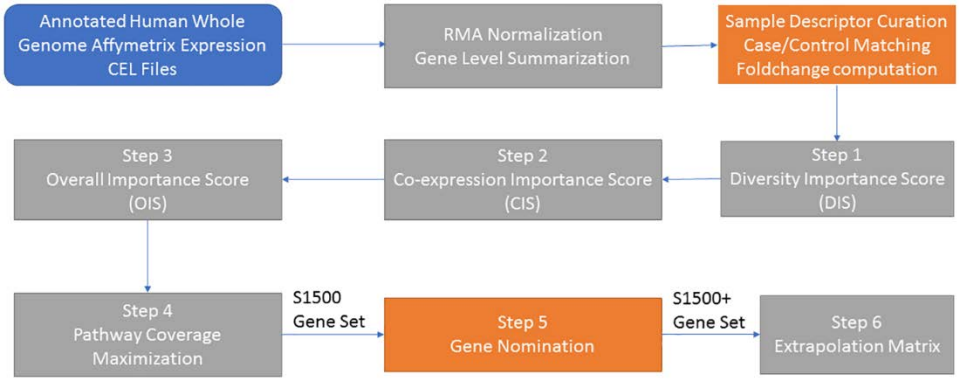
Annotated Human Whole Genome Affymetrix Expression CEL Files
RMA Normalization Gene Level Summarization
Sample Descriptor Curation Case/Control Matching Foldchange computation
Step 1 Diversity Importance Score (DIS)
Step 2 Co-expression Importance Score (CIS)
Step 3 Overall Importance Score (OIS)
Step 4 Pathway Coverage Maximization
S1500 Gene Set
Step 5 Gene Nomination
S1500+ Gene Set
Step 6 Extrapolation Matrix