Report/：DifferentialAbundanceAnalysis的图表总结

plots/

* qc/: quality control plots from initial processing e.g. DESeq2
* exploratory/: 样本之间的整体表达丰度的比较
* differential/：contrasts之间的显著差异表达比较
* gprofiler2：显著差异基因的富集分析，gostplot，曼哈顿图

tables/

* annotation/: gene annotation
* differential/: raw & filtered deseq2 results
* gprofiler2/：enriched pathways
* processed\_abundance/: normalized counts & vst counts