Questionnaire for Data Processing, Analysis, and Interpretation for

ChIP-seq

## Project Background: Brief description of the project

## Expected deliverables: Please choose the deliverables below and/or specify any additional items/requirements:

Please specify the species below by clicking the checkbox or provide other resource.

|  |  |  |  |
| --- | --- | --- | --- |
| Standard species | Human (Homo sapiens) | hg38 (GRCh38) (the most updated version of human genome) | [ ]  |
| hg19 (old version) | [ ]  |
| Mouse (Mus musculus) | mm10 (GRCm38) | [ ]  |
| Rat (Rattus norvegicus) | Rnor\_6.0 | [ ]  |
| Pig (Sus scrofa) | Sscrofa11.1 | [ ]  |
| Macaque (Macaca mulatta) | Mmul8.0.1 | [ ]  |
| Turkey (Meleagris gallopavo) | UMD2 | [ ]  |
| Fruit fly (Drosophila melanogaster) | DM6 | [ ]  |
| Others |  | [ ]  |

## Specify Input and Replicate details

|  |  |  |
| --- | --- | --- |
| **Sample Information** | **Y/N** | **Description** |
| Input Sample | [ ]  |  |
| Replicates available | ☐ |  |
| Total number of samples |  |

## Standard deliverables:

|  |  |  |
| --- | --- | --- |
| **Analysis Item** | **Y/N** | **If yes, any specific requirements?** |
| QC Statistics | [ ]  |  |
| Sequence alignment | [ ]  |  |
| Summary of alignment quality | [ ]  |  |
| Peak calling | [ ]  |  |
| PCA and sample correlation heatmaps to check sample clustering | [ ]  |  |
| Venn diagram plots of peaks | [ ]  |  |
| Differential bind analysis \* | MA plots | [ ]  |  |
| Volcano plots  |  |
| P-value, FDR, FC table |  |

### Optional deliverables **(each item is additional charges and may affect turnaround time, please request specific pricing from PM team)**:

|  |  |  |
| --- | --- | --- |
| Analysis Item | Y/N | If yes, any specific requirements? |
| DB annotation and GO/KEGG (for species with DB only) | [ ]  |  |
| Combined density profile analysis | [ ]  |  |
| Motif analysis | [ ]  |  |

\*If Differential bind analysis is chosen, please specify the detailed group comparison information below:

For example:

Group A: Sample 1, 2

Group B: Sample 3, 4

Group C: Sample 5, 6

Group D: Sample 7, 8

Group A vs. Group B, Group C vs. Group D

|  |
| --- |
|  |

## If any specific version of genome sequence, annotation file, or software/packages/tools for data analyze is needed, please specify it below:

|  |
| --- |
|  |

## If any other customized data analysis request, please describe below.

Please note that it is not guaranteed to be accepted and may incur additional cost.

|  |
| --- |
|  |

## References: Please attach and or provide links of the relevant references as per your project