**Data Application**

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| **Basic information of principal investigator** |
| **Name** |  |
| **Affiliation** |  |
| **Post address** |  |
| **Email** |  |
| **Phone number** |  |
| **Application date** |  |

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| **Data to apply** |
| **Project name of the data** |  |
| **Sequencing type of the data** | □ DNA-seq □ RNA-seq □ ATAC-seq □ Methyl-seq □ Etc |
| **Data format** | □ genetic data (fastq) □ genetic data (bam) □ genetic data (vcf) □ Allele frequency data □ Etc |
| **Data quantity** |  |

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| **Plan of data usage** |
| **Purpose of data usage** | □ Commercial use □ Non-commercial (Academic) use □ Etc[Explanation: Commercial use means that some of research done with the shared data may eventually lead to the development of new predictive or diagnostic tests, medicines, or other commercial products. It also means you can make a profit using the data.] |
| **Data usage plan** **(within 200 letters)** |  |
| **Research plan** **(within 200 letters)** |  |
| **Expected outcomes/products using the data** **(within 200 letters)** | \* If you chose “**commercial use**” in question above, please fill in the blanks below. |
| **Period of data usage** |  |

**Researchers (If there are more than five researchers, add an additional table and write all researchers.)**

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| **Number** | **Name** | **Affiliation** | **Email** |
| **1** |  |  |  |
| **2** |  |  |  |
| **3** |  |  |  |
| **4** |  |  |  |
| **5** |  |  |  |

**Application date :**

**Principal investigator : (Signature)**