Leibniz Institute for Immunotherapy

NGS CORE

NextSeq2000 Booking Request

Submitter:	Experiment Name:
Email:	Requested run date:

Type of flow cell:

P1 (100 mio reads)		100		300	600	Cycles
P2 (400 mio reads)		100	200	300	600	Cycles
P3 (1200 mio reads)	50	100	200	300		Cycles

Type of experiment:

Bulk   Single cell

RNA	Exome	
ChIP	DNA	Other (please specify):
ATAC		

Species/Genome version:

Human	Hg19	Hg38/GRCh38	Other:	
Primary human material		If yes, Ethics Approval Reference No.:		
Mouse	Mm9	Mm10/GRCm38	Other:	Strain:
Other:		Genome version:		

Kit used for library generation:

Indexing strategy: *(Unique dual indices are recommended!)*

Number of indices:	single	dual		None:
Length index 1 (in bp)	6	8	10	Other:
Length index 2 (in bp)	6	8	10	Other:

Single or paired-end reads:	**Library pooling or quantification:**
Single end	Assistance required?
Paired end	No
If paired end:	Yes
Equal length for both reads:	If yes, please specify:
Different setup (please specify):	

Who receives access to raw data?
Email(s):

Additional comments:

Ort, Datum	Name	Signature