

<b>Copy number variation (CNV) analysis</b>				Document No.	2015-06
				Issue date	2018-10-15
<b>Stem cell Information</b>					
Stem cell line	<b>hUSiPS2</b>	Institute	KSCR		
Cell type	<b>hiPSC</b>	Inspected Date			
Banking status	<b>MCB</b>	Issue date	2018-10-15		
Passage	<b>p25</b>				
Note					
<b>Experiment type</b>					
<b>SNP chip</b>					
<b>Platform</b>	illumina Human Omni 2.5 Exome Beadchip	<b>Analysis program</b>	GenomeStudio, PennCNV		
<b>Reference</b>	hg19	<b>Analysis document</b>	SOP#26-Ver.4		
<b>Statistics</b>					
		<b>Total</b>	<b>Gain</b>	<b>Loss</b>	
the number of total CNVs		13	8	5	
the number of manually filtered CNVs		5	4	1	
the number of CNVs excluded Korean normal CNV DB (KGVDB)		4	3	1	
<b>Result of Data Analysis</b>					
List of CNVs					
	<b>Total</b>	<b>Gain</b>	<b>Loss</b>	<b>Cytoband</b>	
<b>The number of total CNV calls</b>	<b>5</b>	<b>4</b>	<b>1</b>		
The number of Pathogenic CNVs	3	3	0	7q36.2, 16p13.13, 20q11.21	
The number of Recurrent CNVs	1	1	0	20q11.21	
The number of stemness-related CNVs	1	1	0	20q11.21	
The number of Differentiation-related CNVs	1	1	0	20q11.21	
The number of cancer-related CNVs	2	2	0	16p13.13, 20q11.21	
The number of immunogenicity-related CNVs	1	1	0	20q11.21	
*Recurrent CNVs include CNV gain on 1q41, 12p13.31, 17q25.2 and 20q11.21, CNV loss on 10p11.22					
<b>Interpretation</b>					

<b>Copy number variation (CNV) analysis</b>	Document	2015-06
	Issue date	2018-10-15

Stem cell Information			
Stem cell line	<b>hiUSIP52</b>	Institute	KSCR
Cell type	<b>hiPSC</b>	Inspected	
Banking status	<b>MCB</b>	Issue date	2018-10-15
Passage	<b>p25</b>		
Note			

**Result of Data Analysis**

List of CNVs															
chr	Cytoband	chr_start	chr_end	length	Copy Number	CNV	Genes	Pathogenic CNV	Phenotype	Recurrent CNV	Cancer-related	Stemness-related	Differentiation-related	Immunogenicity-related	KGVDB
7	q36.1	149,551,993	149,666,820	114,828	cn=3	Gain	ATPGV0E2,ATPGV0E2-AS1,ZNF862								No
7	q36.2	153,522,051	153,660,141	138,091	cn=3	Gain	PPP6	PPP6	Mental retardation (616311), Ventricular						Yes
15	q11.2	22,410,067	22,512,696	102,630	cn=1	Loss	OR4N3P								No
16	p13.13	10,754,456	11,024,105	269,650	cn=3	Gain	CITAD,EXLN,NUBP1,TEKT5,TVP23A	CITA	Bare lymphocyte syndrome typeII (209920), Rheumatoid		CITA				No
20	q11.21	29,804,016	31,640,095	1,836,080	cn=3	Gain	ABALON,ASXL1,BCL2L1,BPIFB2,BPIFB6,C20orf203,CCND1,COMMMD7,COX4I2,DEFB115,DEFB116,DEFB118,DEFB119,DEFB121,DEFB122,DEFB123,DEFB124,DNMT3B,USP15,FOXO1,HCK,HML3,HML3-AS1,JD1,KIF3B,LINC00028,LOC101929698,LOC149950,MAPRE1,MCTS2P,MIR1825,MIR3193,MIR7641-2,MYLK2,NOL4L,PDZRG1,PLAGL2,POFUT1,REML1,SUN5,TM6SF4,TPX2,TPY26P,TTL9,XKR7	COX4I2, DNMT3B, MYLK2, ASXL1	COX4I2 (Exocrine pancreatic insufficiency) DNMT3B(Immunodeficiency-centromeric instability-facial anomalies syn I) MYLK2 (Cardiomyopathy) ASXL1 (Bohring-Opltz syn, Myelodysplastic syn) SNTA1(LongQT syn)	Yes	ASXL1, BCL2L1	DNMT3B	DNMT3B	DEFB119, DEFB123	No