

## Karyotyp-Report

Einsender DNA ID: NP0012-19iPSC-1  
Interne DNA ID: DE25DIVUKOD100050  
Eingetroffen am: 2015-02-06

Geschlecht  
Angegeben: Unknown  
Aus Chipdaten: Female

## Karyotypisierung

Technologie: Illumina BeadArray  
  
Produkt: HumanOmniExpressExome-8 BeadChip v1.2  
Manifest: HumanOmniExpressExome-8v1-2\_A.bpm  
Clusterfile: HumanOmniExpressExome-8v1-2\_A.egt  
  
ChipID und –Segment: 3999269034 R01C01  
  
Batch und Well: WG0082838-MSA1 A02  
  
Erreichte Callrate: 0,9976457

Typisierung  
Scanner: Illumina iScan, S/N: N234  
Ort des Scans: Life&Brain GENOMICS, Bonn, Deutschland  
Hersteller: Illumina, Inc., San Diego, USA  
Scan-Datum: 2015-02-11-1748

Genotypen-Analyse  
Genome Studio: GenomeStudio V2011.1  
Genotyping Module: Ver. 1.9.4

CNV-Analyse  
Algorithmus: CNV-Partition  
Version: 3.2  
Hersteller: Illumina, Inc., San Diego, USA

## Berichtenswerte Beobachtungen

Keine berichtenswerten, grosseren Events

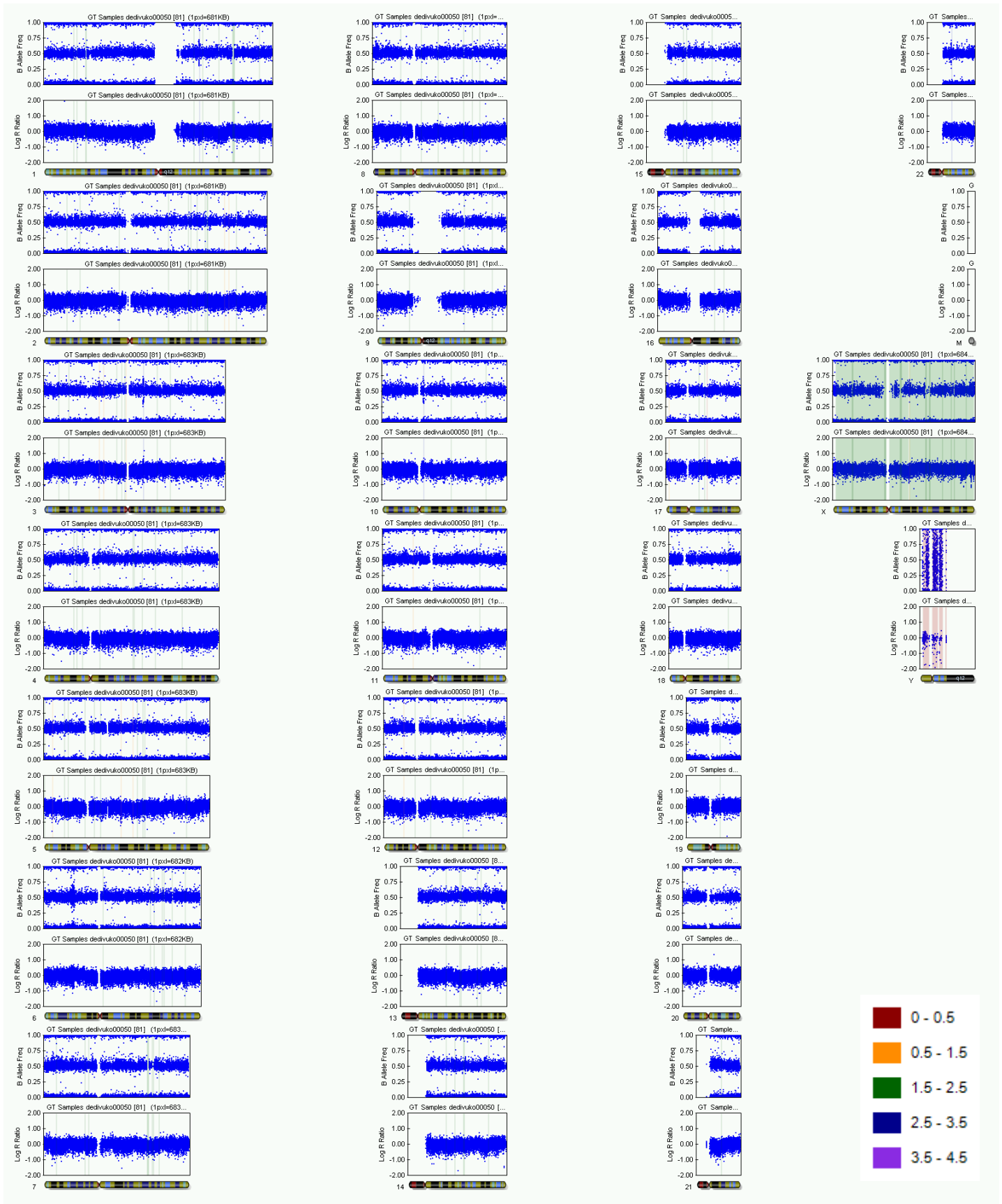
## Auswertender Bioinformatiker

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## Control Dashboard

Sample_ID / Sentrix_Label	Category	Control (BeadType)	Section 1 X	Section 1 Y	State
DEDIVUKO00050 / 3999269034_R01C01	Staining	DNP (High) (27630314)	14559	59	Notable/OK
DEDIVUKO00050 / 3999269034_R01C01	Staining	DNP (Bgnd) (29619375)	190	89	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Staining	Biotin (High) (41666334)	92	6820	OK/Notable
DEDIVUKO00050 / 3999269034_R01C01	Staining	Biotin (Bgnd) (34648333)	114	97	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Extension	Extension (A) (17616306)	34044	269	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Extension	Extension (T) (14607337)	37042	187	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Extension	Extension (C) (12613307)	1112	16467	OK/Notable
DEDIVUKO00050 / 3999269034_R01C01	Extension	Extension (G) (11603365)	1139	16445	OK/Notable
DEDIVUKO00050 / 3999269034_R01C01	Target Removal	Target Removal (31623323)	1069	140	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Hybridization	Hyb (High) (19612319)	1458	16769	OK/Notable
DEDIVUKO00050 / 3999269034_R01C01	Hybridization	Hyb (Medium) (20636378)	372	9982	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Hybridization	Hyb (Low) (23617335)	1493	3018	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Stringency	String (PM) (32629312)	22534	330	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Stringency	String (MM) (33668307)	1657	107	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non-Specific Binding	NSB (Bgnd) (26619332)	293	9015	OK/Notable
DEDIVUKO00050 / 3999269034_R01C01	Non-Specific Binding	NSB (Bgnd) (27624356)	240	92	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non-Specific Binding	NSB (Bgnd) (25617343)	4249	438	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non-Specific Binding	NSB (Bgnd) (24616350)	253	3445	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non- Polymorphic	NP (A) (34633358)	13691	229	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non- Polymorphic	NP (T) (16648324)	12255	153	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non- Polymorphic	NP (C) (43641328)	581	9888	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Non- Polymorphic	NP (G) (13642359)	607	9508	OK/OK
DEDIVUKO00050 / 3999269034_R01C01	Restoration	Restore (28637363)	251	217	OK/OK

# Karyogramm



## CNV-Analyse

Berichtet werden Kopienzahlvarianten ab einer Größe von 350'000bp.

DGV-Abgleich: Anzahl CNVs in gesunden Kontrollen, die den ermittelten CNV komplett überspannen.

### CNV-Analyse

Algorithmus: CNV-Partition

Version: 3.2

Hersteller: Illumina, Inc., San Diego, USA

Sample ID	Chr	Start	Ende	Type	Länge	SNPs	CNVConf	DGV-Abgleich
DEDIVUKO00050	3	108608767	109011263	3	402496	120	617.8739	
DEDIVUKO00050	23	2700163	58483247	2	55783084	9384	2618.4650	
DEDIVUKO00050	23	61730485	83128216	2	21397731	2056	1096.1520	
DEDIVUKO00050	23	83132635	154892230	2	71759595	10985	1213.8330	
DEDIVUKO00050	24	2657176	10019072	0	7361896	547	5397.6030	
DEDIVUKO00050	24	13281348	19563894	0	6282546	586	10086.1300	
DEDIVUKO00050	24	20828795	24891486	0	4062691	317	22622.0000	
DEDIVUKO00050	24	28374713	28817442	0	442729	20	877.4322	