

Design Summary

Design name	T2685V2hg38
Species	H. sapiens (H. sapiens, hg38, p13, GRCh38, Dec. 2013)
Platform	BGI/Illumina
Date	2025-06-13 15:04:45

Target and Probe Summary

Region size	4.75 kbp
Probe coverage size	10.93 kbp
Coverage predict	100.0000%
Tiling density	3.2854 x

Predict region coverage in base level sorted by coverage rate []

Chromosome	Start	End	Length (bp)	Covered (bp)	Missed (bp)	CoverageRate (%)	PredictCovered (bp)	PredictMissed (bp)	PredictCoverageRate (%)
chr1	97078975	97079146	171	171	0	100.00%	171	0	100.00%
chr1	97082329	97082470	141	141	0	100.00%	141	0	100.00%
chr1	97098488	97098632	144	144	0	100.00%	144	0	100.00%
chr1	97193068	97193248	180	180	0	100.00%	180	0	100.00%
chr1	97234851	97234994	143	143	0	100.00%	143	0	100.00%

"report.txt"

loci.bed / loci-gene.bed
BED loci-gene.bed //Notepad++, WPSWord

probeCov.bed
BED

probeCov.predict.bed
BEDreads

probeMissed.bed / probeMissed.anno.xls
BED probeMissed.anno.xls //

probeMissed.predict.bed / probeMissed.predict.anno.xls
BED probeMissed.predict.anno.xls //

report.txt / report.anno.xls
report.anno.xls //Notepad++WPSExcel