AlloSeq[™]HCT

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Revolutionizing Chimerism Monitoring

THE LATEST INNOVATION IN CHIMERISM TESTING OFFERS STREAMLINED WORKFLOW WITH BEST-IN-CLASS SENSITIVITY

Chimerism changes after hematopoietic cell transplant can be an early indication of relapse*

Early detection can lead to early intervention

What is AlloSeq HCT?

- A NGS based chimerism testing solution that is able to determine the relative contribution of up to 3 genomes in a single sample i.e. up to two donors per recipient
- A kit based test that offers high quality data with a recommended input of **10 ng of gDNA** and up to **24 samples per batch**
- A one-step multiplex assay that utilizes 202 SNPs across all autosomal chromosomes to differentiate between recipient and donor(s) gDNA
- A highly sensitive, scalable assay with time-saving automated analysis software

AlloSeq HCT is a simple and fast assay with **minimal hands-on time**.

gDNA Sample to Report in Less than 24 hours



• Rashef et al BBMT 2014;20:1758–66 • Tang et al BBMT 2014;20:1139:1144

AlloSeq HCT offers accurate and precise results

72 DNA mixtures simulating post-transplant samples demonstrate test dynamic range of 0.1% - 99.9%

 $R^2 = 0.9999$

AlloSeq HCT accuracy was measured against expected using artificially mixed gDNA samples extracted from fresh whole blood. 72 data points were generated achieving signal linearity with a correlation and slope close to 1. The Limit of Quantification (LOQ) obtained is shown and marked on the graph (for post transplant samples with a single donor)

100

80

AlloSeq HCT is reproducible with high quality data

98%

98%

0.1%

Internal and external validation produces reproducible and precise data

50%

50.7%

0.2%

60

TABLE 1: Internal Validation at CareDx

1%

1.1%

7.7%

10%

10.5%

2.1%

0.4%

0.5%

8.6%

20

40

100

90 80

70

60 50

40

30

20

10

Expected

Fraction Observed

Fraction **Coefficient of**

Variation

AlloSeq HCT result (%)

Sample Number (N)	20	12	11	3	3	4		Sample Number (N)	3	3	3
Table 1 & 2 repr	resent tl	he AlloS	Seq HCT	average r	esult for	artificially	/ mixed	samples across	a numb	er of tech	nnical
replicates. Table	2 shov	vs valida	ation resi	ults at an	external	lab confi	rming tł	he reproducibility	[,] obtaine	ed at Care	eDx.

85%

85.2%

0.5%

Expected (%)

2.00

1.50

1.00

0.50

0.00

0.00

AlloSeg HCT result (%)

TABLE 2: External Validation

Expected Fraction	5%	50%	95%	
Observed Fraction	4.8%	48.2%	94.7%	
Coefficient of Variation	4.8%	0.5%	0.2%	
Sample Number (N)	3	3	3	

Limit of Quantification

1.00

Expected (%)

1.50

2.00

0.50



AlloSeq HCT provides easy and fast results with automated analysis.

The Windows based software is user-friendly and LIMS compatible.

All samples are reported on a single screen

Post-transplant samples

SAMPLE NAME	ANALYSIS DATE	DNA TYPE	SAMPLE QUALITY	RECIPIENT DNA	DONOR-1 DNA	DONOR-2 DNA	DETAILS VIEW
E10R1	NA	gDNA	Pass	0.83%	8.89%	90.27%	•
E10R10	NA	gDNA	Pass	1.01%	8.58%	90.41%	Θ
E10R11	NA	gDNA	Pass	0.89%	8.85%	90.26%	Θ

Recipient-only

SAMPLE NAME	ANALYSIS DATE	DNA TYPE	SAMPLE QUALITY	DETAILS VIEW
Blood3R5	2019-11-21	gDNA	Pass	Θ

Donor-only

SAMPLE NAME	ANALYSIS DATE	DNA TYPE	SAMPLE QUALITY	DETAILS VIEW
Blood1R6	2019-11-21	gDNA	Pass	Ø
Blood2R4	2019-11-21	gDNA	Pass	Θ

seen by clicking on "Details view" Result Details × AlloSeg HCT Sample Analysis Report **é**CareDx⁺ **Detailed %** Analysis output **DNA** display RECIPIENT DNA DONOR-1 DNA DONOR-2 DNA for each 0.83% 8.89% 90.27% [0.66%-1.00%] [8.63%-9.15%] [90.02%-90.52% sample Sample info Sample quality: Pass Sample name E10R1 Mean Coverage: Pas 3252 Threshold: 500 Source type Post-transplant Analysis inputs Uniformity: Pas DNA type aDNA 90% Threshold: 75% DNA amount (ng) 10ng Loci passing filter: Pass Recipient Sample Blood3R5 195 Threshold: 186 Donor1 Sample Blood2R4 Donor2 Sample Blood1R6 Loci within range: NA Analysis information Recipient-only sample filter: Pass Donor-only sample filter: Pass, Pass Analysis date 2019-11-21 15:11:41 Worker version 1.0.0 Other sample metrics Pipeline version 1.2.5.6-hct Informative Loci: 124, 121 Operator XDXINC\cegidio Loci removed from analysis Actions Low coverage Delete Analysis Results Generate Test Report PDF Multiallelic 0 Loci removed from recipient-only and/or donor-only sample(s) Send Results to PDF View Longitudinal Database AlloSeq[®]HCT

"Detailed report by sample can be

PRODUCT	PRODUCT NUMBER	DESCRIPTION		
AlloSeq® HCT	ASHCT.1(24)	Includes all the reagents required to make 24 NGS libraries		

For Research Use Only. Not for use in diagnostic procedures.

Visit <u>www.caredxinc.com/alloseq-hct</u> for more information For inquiries, contact your CareDx representative or reach out to us:

Phone numbers: Americas – <u>orders-US@caredx.com</u> EMEA – <u>orders-at@caredx.com</u> Nordics – orders-aus@caredx.com APAC – orders-se@caredx.com

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