

Annex to declaration of accreditation (scope of accreditation)
Normative document: EN ISO/IEC 17025:2017
Registration number: **L 671**

of **Cergentis B.V.**
Services

This annex is valid from: **21-12-2022** to **01-02-2026**

Replaces annex dated: **20-01-2022**

Location(s) where activities are performed under accreditation

Head Office

Yalelaan 62
3584 CM
Utrecht
The Netherlands

Location	Abbreviation/ location code
Yalelaan 62 3584 CM Utrecht The Netherlands	U

No.	Material or product	Type of activity ¹	Internal reference number	Location
1.	Cells isolated from bone marrow, spleen, liver or kidney, White blood cells isolated from fresh blood samples, Cultured T-Cells, ESCs and iPSCs, Cell lines (for example CHO, HELA, HEK293T), Yeast cells, Cells isolated from embryos or whole fish, Vectors	Determination of the integrity of the transgene vector sequence; next generation sequencing (NGS) and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12, SOP25 in house method	U

This annex has been approved by the Board of the Dutch Accreditation Council, on its behalf,

J.A.W.M. de Haas

¹ If there is a referral to a code starting with NAW, NAP, EA or IAF, this concerns a scheme mentioned on [RvA-BR010-lijst](#).
If no date or version number is mentioned for a normative document, the accreditation concerns the most current version of the document or scheme.

of **Cergentis B.V.**
Services

This annex is valid from: **21-12-2022** to **01-02-2026**

Replaces annex dated: **20-01-2022**

No.	Material or product	Type of activity ¹	Internal reference number	Location
2.	Cells isolated from bone marrow, spleen, liver or kidney, White blood cells isolated from fresh blood samples, Cultured T-Cells, ESCs and iPSCs,	Determination of the vector integration site(s) and breakpoint sequences between the vector and genome; next generation sequencing (NGS) and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12 in house method	U
3.	Cell lines (for example CHO, HELA, HEK293T), Yeast cells, Cells isolated from embryos or whole fish	Determination of the presence of structural variants surrounding the vector integration site(s); next generation sequencing (NGS) and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12 in house method	U
4.		Determination of transgene components; next generation sequencing (NGS), comparison with reference libraries of common vector elements and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12 in house method	U
5.		Determination of gene editing events in endogenous loci; next generation sequencing (NGS) and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12 in house method	U
6.	Isolated white blood cells and organoids	Determination of integration site(s) in heterogeneous cell populations; next generation sequencing (NGS) and bio-informatic data analysis	SOP02, SOP04, SOP39, SOP08, SOP12 in house method	U