

Full name:	Ichthyosis and erythroderma (98 genes) - KUL
Description:	https://firebasestorage.googleapis.com/v0/b/uz-laboboeken.appspot.com/o/GHB_CME%2Fgeno-dermatosen-NGS-specifieke%20genpanels%20(transcripten).pdf?alt=media&token=4ce21463-cf35-4d79-8bb2-c4b58310b259
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Laboratory:	<u>Centrum Menselijke Erfelijkheid - KUL</u>
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Related Diseases

- Acral peeling skin syndrome
- Acral self-healing collodion baby
- Acrodermatitis continua of Hallopeau
- Acrodermatitis enteropathica
- Acute neonatal citrullinemia type I
- Annular epidermolytic ichthyosis
- Arthrogryposis-renal dysfunction-cholestasis syndrome
- Autosomal dominant epidermolytic ichthyosis
- Autosomal dominant focal non-epidermolytic palmoplantar keratoderma with plantar blistering
- Autosomal dominant generalized epidermolysis bullosa simplex, intermediate form
- Autosomal dominant generalized epidermolysis bullosa simplex, severe form
- Autosomal dominant palmoplantar keratoderma and congenital alopecia
- Autosomal recessive epidermolytic ichthyosis
- BRESEK syndrome
- Bathing suit ichthyosis
- Biotinidase deficiency
- CEDNIK syndrome
- CHILD syndrome
- CK syndrome

- Carbamoyl-phosphate synthetase 1 deficiency
- Classic maple syrup urine disease
- Combined immunodeficiency due to partial RAG1 deficiency
- Combined immunodeficiency with granulomatosis
- Congenital ichthyosiform erythroderma
- Congenital ichthyosis-intellectual disability-spastic quadriplegia syndrome
- Congenital reticular ichthyosiform erythroderma
- Cutaneous mastocytoma
- DITRA
- Diffuse palmoplantar keratoderma with painful fissures
- Dowling-Degos disease
- Epidermolysis bullosa simplex with circinate migratory erythema
- Epidermolysis bullosa simplex with mottled pigmentation
- Epidermolytic palmoplantar keratoderma
- Erythrokeratoderma variabilis
- Exfoliative ichthyosis
- Focal palmoplantar keratoderma with joint keratoses
- Generalized pustular psoriasis
- Harlequin ichthyosis
- Hidrotic ectodermal dysplasia
- Holocarboxylase synthetase deficiency
- Hypotrichosis simplex of the scalp
- Ichthyosis follicularis-alopecia-photophobia syndrome
- Ichthyosis hystrix of Curth-Macklin
- Ichthyosis-hypotrichosis syndrome
- Ichthyosis-prematurity syndrome
- Ichthyosis-short stature-brachydactyly-microspherophakia syndrome
- Intermediate maple syrup urine disease
- Intermittent maple syrup urine disease
- Isolated focal non-epidermolytic palmoplantar keratoderma
- KID syndrome
- KRT1-related diffuse nonepidermolytic keratoderma
- Keratoderma hereditarium mutilans
- Keratoderma hereditarium mutilans with ichthyosis
- Keratosis follicularis spinulosa decalvans
- Keratosis linearis-ichthyosis congenita-sclerosing keratoderma syndrome

- Knuckle pads-leukonychia-sensorineural deafness-palmoplantar hyperkeratosis syndrome
- Lamellar ichthyosis
- Late-onset citrullinemia type I
- Localized epidermolysis bullosa simplex
- MEDNIK syndrome
- MEND syndrome
- Menkes disease
- Monilethrix
- Multiple sulfatase deficiency
- Mutilating palmoplantar keratoderma with periorificial keratotic plaques
- Neonatal ichthyosis-sclerosing cholangitis syndrome
- Neonatal inflammatory skin and bowel disease
- Netherton syndrome
- Neutral lipid storage disease with ichthyosis
- Nodular urticaria pigmentosa
- Omenn syndrome
- Pachyonychia congenita
- Palmoplantar keratoderma-deafness syndrome
- Palmoplantar keratoderma-hereditary motor and sensory neuropathy syndrome
- Peeling skin syndrome type A
- Peeling skin syndrome type B
- Pityriasis rubra pilaris
- Plaque-form urticaria pigmentosa
- Progressive symmetric erythrokeratoderma
- Propionic acidemia
- Pustulosis palmaris et plantaris
- Pyruvate dehydrogenase E3 deficiency
- Recessive X-linked ichthyosis
- Refsum disease
- Rhizomelic chondrodysplasia punctata type 1
- Self-improving collodion baby
- Severe combined immunodeficiency due to DCLRE1C deficiency
- Severe combined immunodeficiency due to complete RAG1/2 deficiency
- Severe dermatitis-multiple allergies-metabolic wasting syndrome
- Sjögren-Larsson syndrome
- Striate palmoplantar keratoderma

- [Superficial epidermolytic ichthyosis](#)
- [Syndromic recessive X-linked ichthyosis](#)
- [Thiamine-responsive maple syrup urine disease](#)
- [Trichothiodystrophy](#)
- [Typical urticaria pigmentosa](#)
- [Vitamin B12-unresponsive methylmalonic acidemia type mut-](#)
- [Vitamin B12-unresponsive methylmalonic acidemia type mut0](#)
- [X-linked dominant chondrodysplasia punctata](#)
- [Xeroderma pigmentosum-Cockayne syndrome complex](#)

Related Analytes

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
ABCA12	95.00	0	NM_173076.3
ABHD5	95.00	0	NM_016006.6
ADAM17	95.00	0	NM_003183.6
ALDH3A2	95.00	0	NM_000382.3
ALOX12B	95.00	0	NM_001139.3
ALOXE3	95.00	0	NM_021628.3
AP1B1	95.00	0	NM_001127.4
AP1S1	95.00	0	NM_001283.5
ASPRV1	95.00	0	NM_152792.4
ASS1	95.00	0	NM_000050.4
ATP7A	95.00	0	NM_000052.7
BCKDHA	95.00	0	NM_000709.4
BCKDHB	95.00	0	NM_183050.4

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>BTD</u>	95.00	0	NM_001370658.1
<u>BTK</u>	95.00	0	NM_000061.3
<u>CAPN12</u>	95.00	0	NM_144691.4
<u>CARD14</u>	95.00	0	NM_024110.4
<u>CASP14</u>	95.00	0	NM_012114.3
<u>CAST</u>	95.00	0	NM_001042440.5
<u>CDSN</u>	95.00	0	NM_001264.5
<u>CERS3</u>	95.00	0	NM_178842.5
<u>CHST8</u>	95.00	0	NM_001127896.2
<u>CLDN1</u>	95.00	0	NM_021101.5
<u>CPS1</u>	95.00	0	NM_001875.5
<u>CSTA</u>	95.00	0	NM_005213.4
<u>CTSB</u>	95.00	0	NM_001908.5
<u>CYP4F22</u>	95.00	0	NM_173483.4
<u>DBT</u>	95.00	0	NM_001918.5
<u>DCLRE1C</u>	95.00	0	NM_001033855.3
<u>DLD</u>	95.00	0	NM_000108.5
<u>DSG1</u>	95.00	0	NM_001942.4
<u>EBP</u>	95.00	0	NM_006579.3
<u>ELOVL1</u>	95.00	0	NM_001256399.2

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>ELOVL4</u>	95.00	0	NM_022726.4
<u>ERCC2</u>	95.00	0	NM_000400.4
<u>ERCC3</u>	95.00	0	NM_000122.2
<u>FLG</u>	95.00	0	NM_002016.2
<u>FLG2</u>	95.00	0	NM_001014342.3
<u>GBA1</u>	95.00	0	NM_001005741.3
<u>GJA1</u>	95.00	0	NM_000165.5
<u>GJB2</u>	95.00	0	NM_004004.6
<u>GJB3</u>	95.00	0	NM_024009.3
<u>GJB4</u>	95.00	0	NM_153212.3
<u>GJB6</u>	95.00	0	NM_006783.5
<u>GTF2E2</u>	95.00	0	NM_002095.6
<u>GTF2H5</u>	95.00	0	NM_207118.3
<u>HLCS</u>	95.00	0	NM_000411.8
<u>IL36RN</u>	95.00	0	NM_012275.3
<u>KDSR</u>	95.00	0	NM_002035.4
<u>KIT</u>	95.00	0	NM_000222.3
<u>KRT1</u>	95.00	0	NM_006121.4
<u>KRT10</u>	95.00	0	NM_000421.5
<u>KRT14</u>	95.00	0	NM_000526.5

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>KRT16</u>	95.00	0	NM_005557.4
<u>KRT2</u>	95.00	0	NM_000423.3
<u>KRT5</u>	95.00	0	NM_000424.4
<u>KRT6C</u>	95.00	0	NM_173086.5
<u>KRT83</u>	95.00	0	NM_002282.3
<u>KRT9</u>	95.00	0	NM_000226.4
<u>LIPN</u>	95.00	0	NM_001102469.2
<u>LORICRIN</u>	95.00	0	NM_000427.3
<u>MBTPS2</u>	95.00	0	NM_015884.4
<u>MMUT</u>	95.00	0	NM_000255.4
<u>MPLKIP</u>	95.00	0	NM_138701.4
<u>NIPAL4</u>	95.00	0	NM_001099287.2
<u>NSDHL</u>	95.00	0	NM_015922.3
<u>PCCA</u>	95.00	0	NM_000282.4
<u>PCCB</u>	95.00	0	NM_000532.5
<u>PERP</u>	95.00	0	NM_022121.5
<u>PEX7</u>	95.00	0	NM_000288.4
<u>PHYH</u>	95.00	0	NM_006214.4
<u>PIGL</u>	95.00	0	NM_004278.4
<u>PNPLA1</u>	95.00	0	NM_001145717.1

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>POMP</u>	95.00	0	NM_015932.6
<u>RAG1</u>	95.00	0	NM_000448.3
<u>RAG2</u>	95.00	0	NM_000536.4
<u>RNF113A</u>	95.00	0	NM_006978.3
<u>SDR9C7</u>	95.00	0	NM_148897.3
<u>SERPINB7</u>	95.00	0	NM_003784.4
<u>SERPINB8</u>	95.00	0	NM_198833.2
<u>SGPL1</u>	95.00	0	NM_003901.4
<u>SLC25A13</u>	95.00	0	NM_014251.3
<u>SLC27A4</u>	95.00	0	NM_005094.4
<u>SLC30A2</u>	95.00	0	NM_001004434.3
<u>SLC39A4</u>	95.00	0	NM_130849.4
<u>SNAP29</u>	95.00	0	NM_004782.4
<u>SPINK5</u>	95.00	0	NM_006846.4
<u>SREBF1</u>	95.00	0	NM_004176.5
<u>ST14</u>	95.00	0	NM_021978.4
<u>STS</u>	95.00	0	NM_000351.7
<u>SULT2B1</u>	95.00	0	NM_177973.2
<u>SUMF1</u>	95.00	0	NM_182760.4
<u>TBX1</u>	95.00	0	NM_080647.1

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>TGM1</u>	95.00	0	NM_000359.3
<u>TGM5</u>	95.00	0	NM_201631.4
<u>TRPM4</u>	95.00	0	NM_017636.4
<u>VIPAS39</u>	95.00	0	NM_022067.4
<u>VPS33B</u>	95.00	0	NM_018668.5