

## ExportableDF

Full name:	Overgrowth (24 genes) - IPG
Version number:	version 3
Laboratory:	Centre de Génétique-Institut de Pathologie et de Génétique (IPG)
Created:	04 Jul 2019 - 10:41
Changed:	10 Oct 2022 - 15:16

### Related Diseases

- [5q35 microduplication syndrome](#)
- [Beckwith-Wiedemann syndrome due to CDKN1C mutation](#)
- [Beckwith-Wiedemann syndrome due to NSD1 mutation](#)
- [CLOVES syndrome](#)
- [Cowden syndrome](#)
- [Deletion 5q35](#)
- [Hemimegalencephaly](#)
- [IMAGe syndrome](#)
- [Intellectual disability-macrocephaly-hypotonia-behavioral abnormalities syndrome](#)
- [Intrauterine growth restriction-short stature-early adult-onset diabetes syndrome](#)
- [Malan overgrowth syndrome](#)
- [Marshall-Smith syndrome](#)
- [Megalencephaly-capillary malformation-polymicrogyria syndrome](#)
- [Megalencephaly-polymicrogyria-postaxial polydactyly-hydrocephalus syndrome](#)
- [Proteus syndrome](#)
- [Segmental outgrowth-lipomatosis-arteriovenous malformation-epidermal nevus syndrome](#)
- [Segmental progressive overgrowth syndrome with fibroadipose hyperplasia](#)
- [Silver-Russell syndrome due to a point mutation](#)
- [Simpson-Golabi-Behmel syndrome](#)
- [Sotos syndrome](#)
- [Tatton-Brown-Rahman syndrome](#)
- [Weaver syndrome](#)

## Related Analytes

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>ADARB2</u>	100.00	1	NM_018702.4
<u>AKT1</u>	100.00	1	NM_001014431.2
<u>AKT3</u>	100.00	1	NM_005465.7
<u>BRWD3</u>	100.00	1	NM_153252.5
<u>CDKN1C</u>	100.00	1	NM_001362474.2
<u>CHD3</u>	100.00	1	NM_001005273.3
<u>CHD8</u>	100.00	1	NM_001170629.2
<u>DIS3L2</u>	100.00	1	NM_152383.4
<u>DNMT3A</u>	100.00	1	NM_175629.2
<u>EED</u>	100.00	1	NM_003797.5
<u>EZH2</u>	100.00	1	NM_004456.5
<u>GPC3</u>	100.00	1	NM_001164617.2
<u>H1-4</u>	100.00	1	NM_005321.3
<u>KMT5B</u>	100.00	1	NM_017635.5
<u>MTOR</u>	100.00	1	NM_004958.4
<u>NFIX</u>	100.00	1	NM_002501.4
<u>NSD1</u>	100.00	1	NM_022455.4
<u>PDGFRB</u>	100.00	1	NM_002609.4
<u>PIK3CA</u>	100.00	1	NM_006218.4

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>PPP2R5C</u>	100.00	1	NM_001161725.1
<u>PPP2R5D</u>	100.00	1	NM_006245.4
<u>PTEN</u>	100.00	1	NM_000314.8
<u>SETD2</u>	100.00	1	NM_014159.6
<u>TCF20</u>	100.00	1	NM_005650.3