

# 12 q deletion with oculodentodigital dysplasia -like phenotype

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## Abstract

The increase in microarray usage has led to a parallel increase in the discoveries of new genotypes and phenotypes that are partially similar to known genetic syndromes. OculoDentoDigital Dysplasia (ODDD) syndrome is caused by a mutation in the connexin-43 GJA1 gene on the long arm of chromosome 6. We report a 15 year old girl who presents with a partial 12 q monosomy with ODDD-like features including microphthalmia, microcorneas, microdontia, camptodactyly of the fingers and toes, short stature, developmental and speech delays, and fine motor and hand-eye incoordination. Chromosomal Microarray Analysis showed a 12.987 MB loss of in chromosome 12q21.2 to q21.33, and included more than 20 genes, none of which have any known relation to connexin-43. Our patient did not have any deletions or mutations found on chromosome 6q. Two longer deletions on the long arm of chromosome 12 have been documented in connection to the CardioFacioCutaneous (CFC) syndrome, implicating the region 12q21.2-q22 as a candidate region for the gene or genes causing CFC syndrome. This is the first report of a patient with phenotypic features of ODDD syndrome with a microdeletion on chromosome 12q21. Patients who may have an ODDD like phenotype but are negative for connexin-43 mutation should be evaluated for a 12q21 deletion.

## Introduction

OculoDentoDigital Dysplasia (ODDD) is an uncommon genetically inherited disease described as having a distinctive facial appearance with variable involvement of the eyes, teeth, and digits. It follows an autosomal dominant inheritance from a mutation in the connexin-43 (GJA1) gene on chromosome 6q. Facial features include narrow pinched nose with a narrow nasal bridge, prominent columella, and thin anteverted nares. Ocular features comprise prominent epicanthal folds, eye abnormalities that can lead to vision loss (myopia), micro-ophthalmia, microcornea, cataracts, and glaucoma. Dental abnormalities consist of microdontia, enamel hypoplasia, and caries. Digit malformations include syndactyly and camptodactyly [1].

Monosomy of the long arm of chromosome 12 is a rare laboratory finding with 11 previous reports. Of these previous studies two relatively recent reports by Raun *et al.* in 2000 and 2002 implicated the region from 12q21.2 to q22 as a candidate region for the gene or genes causing CardioFacioCutaneous (CFC) syndrome [2,3]. CFC is a genetic disorder characterized by heart defects, facial anomalies, and cutaneous abnormalities, previously reported as typically resulting from a *de novo* dominant mutation [1].

James *et al.* in 2005 reported another patient with 12q deletion with non-specific features including developmental delays, microcephaly, facial features including a high nasal bridge, fine hair and sparse facial hair, narrow hands and feet, and a hyperkeratotic rash [4]. In 1989 Watson *et al.* described an infant with an interstitial deletion from 12q15 to q21.2 with frontal bossing, small sunken eyes, a beaked nose, low-set ears, thin upper lip, high-arched palate, syndactyly of the 2<sup>nd</sup> and 3<sup>rd</sup> toes of the left foot, cutis marmorata, and developmental delay [5].

With the increased use and application of microarray analysis there has been an increased incidence of patients presenting with phenotypic characteristics of a certain genetic syndrome only to have the microarray results show a different genotype that are not consistent with the literature. Microarray has also helped to characterize patients that present with a phenotypic combination that has either not been previously documented or documented as an atypical presentation of a similar syndrome that is genotypically different. These increasing situations have paved the way for characterization of the “Like” syndromes such as Noonan-like, Prader-Willi-like, and others.

We present a 15 year old female with OculoDentoDigital Dysplasia-Like phenotype with a partial 12 q monosomy.

## Case report

Our patient is a 15 year old female who was brought in by her grandparents because of a previously diagnosed chromosomal abnormality. She was born full term via uncomplicated spontaneous vaginal delivery. The pregnancy was uncomplicated; however, her mother only received prenatal care during the last month of pregnancy. She was evaluated at the age of 4 years because of developmental delays. Grandparents describe her as a healthy young girl with no history of

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hospitalizations or recurrent infections. She was found to have a deletion of the long arm of chromosome 12.

On our initial evaluation she had a short attention span, speech delays, and minor bilateral myopia (R 20/40; L 20/30). She is in Special Education classes and struggles particularly with math. She is small for her age. Weight was 36.5 kg, and height was 150 cm; both below the 3<sup>rd</sup> percentile. Also, the frontal occipital circumference was 51 cm which correlates near the 17<sup>th</sup> percentile. Physical examination also revealed minimal scoliosis, small corneas, small teeth, as well as camptodactyly and partial syndactyly (Figures 1-4).

Maternal grandparents and mother were all without the abnormalities present in our patient. The patient has two maternal half-brothers, one of which is in the lower percentiles for height and weight, but both are developmentally adequate. The father was described as “normal”, but he was unavailable for follow up history and chromosomal studies.



Figure 1. Microcornea.

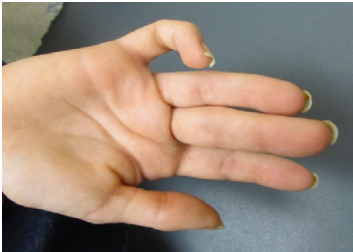


Figure 2. Camptodactyly of fifth digit.



Figure 3. Microdontia.



Figure 4. Camptodactyly and Partial syndactyly of 2<sup>nd</sup> and 3<sup>rd</sup> toes.

## Methods

### Microarray analysis

Chromosomal microarray analysis (CMA – HR + SNP version 9.1.1) was performed at the Kleberg Cytogenetics Laboratory which revealed a 12.987 Mb deletion on chromosome band 12q21.2q21.33 (figures 5). Two smaller deletions were seen on 11q14.1 and 17p13.1. Two small chromosomal gains were seen on Xp11.4 and Xq13.1. No increased blocks of absence of heterozygosity suggestive of uniparental disomy or consanguinity were detected. None of the smaller deletions or gains have been reported to be associated with a clinical phenotype. Maternal FISH analysis showed no evidence of a rearrangement involving chromosome 12 as seen in our patient.

MRI showed eye globes that are smaller than average. The total axial length was 21.1 mm for the left eye, and 22 mm for the right eye. According to Bardakjian *et al.* (2004) this measures as a simple microphthalmia as the lower 2.5% confidence limit for the total axial length is about 21.0 mm [6]. The brain showed no reduction in white matter volume or other white matter abnormalities which is the most common finding in ODDD (Figure 6) [7]. Analysis of chromosome 6 for ODDD mutation was sent to John’s Hopkins DNA Diagnostic Lab showed no evidence of a sequence variation in the coding regions of the *GJA1* gene on chromosome 6.

### Discussion

OculoDentoDigital Dysplasia is a syndrome of a constellation of symptoms and phenotypic findings including characteristic facial features of narrow nose and hypoplastic alae nasi, and the previously mentioned abnormalities of the eyes, teeth, and digits of the hands and feet. Other findings include neurologic problems such as

Change	Chromosome	Min Interval*	Min Size (Mb)	# Probes	Max Interval*	Max Size (Mb)
LOSS	11q14.1	83652737 - 84255911	0.603	86	83641535 - 84267072	0.626
RefSeq Genes: <i>DLG2</i>						
LOSS	12q21.2q21.33	79022885 - 91931022	12.908	1070	78994368 - 91981265	12.987
RefSeq Genes: <i>The region above contains more than 20 genes.</i>						
LOSS	17p13.1	10404962 - 10405873	0.0009	9	10404875 - 10405907	0.001
RefSeq Genes: <i>MYH1</i>						
GAIN	Xp11.4	38493956 - 38565060	0.071	44	38483485 - 38644574	0.161
RefSeq Genes: <i>TSPAN7</i>						
GAIN	Xq13.1	69454420 - 69466846	0.012	29	69438965 - 69478021	0.039
RefSeq Genes: <i>A/WATI</i>						

\* Nucleotide positions based on hg19

Figure 5. Microarray analysis with 12.987Mb deletion.

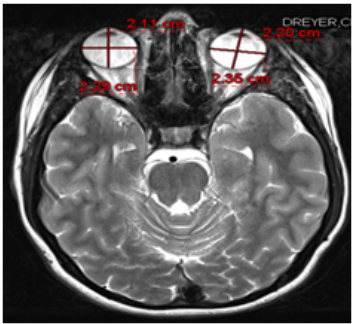


Figure 6. MRI showing small eye globes and negative for white matter abnormalities.

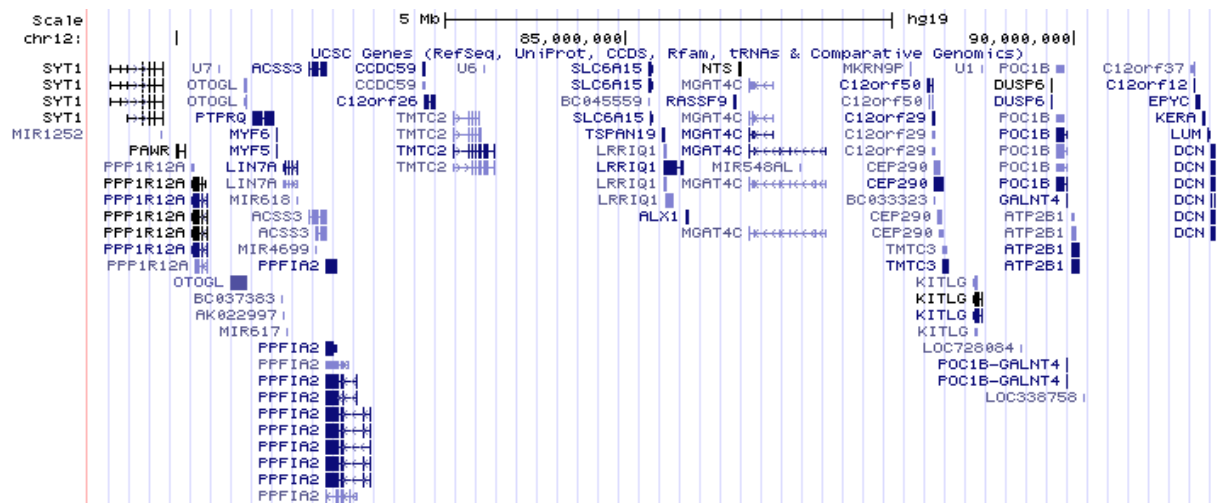


Figure 7. Gene map of deleted region.

Table 1. List of some common features seen in ODD or CFC.

Major Clinical Features	OculoDentoDigital Dysplasia (ODDD)	CardioFacioCutaneous (CFC)	Patient's presentation
Camptodactyly	Seen in 90% of patients with ODD [9]	Not seen or very uncommon[10]	Patient has camptodactyly of fifth finger and in toes.
Syndactyly	Seen in 90% of patients with ODD	Not seen or very uncommon	Partial syndactyly of 2 <sup>nd</sup> and 3 <sup>rd</sup> toes
Abnormal white matter on MRI	May be seen in some patients, but data is lacking [11]	May have abnormalities, such as structural changes [12]	Patient had no abnormal white matter seen on MRI
Cariou teeth	Seen in 90% of patients with ODD	Not seen or very uncommon	Patient had enamel hypoplasia and multiple dental carries.
Microcornea	Seen in 90% of patients with ODD	Not seen or very uncommon	Patient has microcorneas and small eye globes seen on MRI
Cognitive impairment	Seen in 50% of patients with ODD	Seen in 90% of patients with CFC	Patient presented with developmental delays and later in life required special education classes

developmental delays, dysarthria, neurogenic bladder disturbances, spastic paraparesis, ataxia, anterior tibial muscle weakness, seizures, conductive hearing loss, and cardiac defects and irregularities of the skin, hair, and nails. It is primarily an autosomal dominant mutation in the human gap junction alpha 1 (*GJA1*) gene encoding for connexin 43, which has a high degree of penetrance but with considerable variability even within families [1,8].

Our patient had ODDD features including long thin nose, microphthalmia, microcorneas, microdontia, syndactyly, camptodactyly, and developmental delays. There are many syndromes with many similar phenotypes but this does not fit any one syndrome exactly. It is like ODDD but not exactly, and with no Ch. 6 abnormality seen. It is also similar to CFC syndrome which has been postulated to have deletions in the same region, however does not exactly fit the CFC syndrome. This is the first report of a patient with phenotypic features *similar to those* of ODDD syndrome with a microdeletion on chromosome 12q21, expanding the phenotype of ODD previously reported.

There have been 11 previous reports of patients with a deletion in or around this 12q21 region all of which have varying presentations but do contain some phenotypic similarities. Watson *et al.* presented a patient in 1989 with a deleted region of 12q15q21 who was not classified with any specific syndrome [5]. The patient had non-specific finding, but did have a long thin nose, syndactyly, and developmental delays. Rauen *et al.* report in 2000 of a patient with a deletion in 12q21.2q22 was more typical of CFC syndrome and included a short nose, cardiac defects, and fine, sparse hair and facial hair [2]. Rauen *et al.* 2002 report was non-specific but did share some features with their 2000 report including the deleted region 12q21.2q22, short

nose, and sparse hair. However, unlike CFC syndrome this patient also had syndactyly and developmental delays. This patient was also born with a head circumference in the 10<sup>th</sup> percentile but later grew into normal ranges [3]. James *et al.* in 2005 presented a patient with a deletion in 12q21.2q22 with non-specific findings of low birth weight in the 3<sup>rd</sup> percentile, microcephaly, developmental and speech delays, strabismus, astigmatism, and narrow hands [4]. Even though these finding were non-specific they can be classified within the spectrum of an ODDD-like phenotype. Common findings in all of these reports were a hyperkeratotic rash, low set ears, and high arched palate. Table 1 lists some common features seen in ODD or CFC.

Of the more than 20 genes contained in the deleted region, there are several genes responsible for the development of cilia including BBS10 and CEP290 genes, and protein components of the Extracellular Matrix and collagen proteins (Figure 7). This can be part of the reason that there have been 11 such deletions in this region with varying phenotypes but with similarities in ectodermal findings. This is a large chromosome area and we cannot be sure the exact location or cause cannot be confirmed at this time. More case reports are required and with the improved use of microarray analysis more cases should be forthcoming.

Patients who may have an ODDD-like phenotype but are negative for connexin-43 mutation should be evaluated for a 12q21 deletion. This report expands the molecular cause for ODDD and CFC, even opening the possibility that an ODDD syndrome with varying presentations and features is present with this chromosomal region containing the genes responsible for the phenotype.

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