

DYRK1A-related intellectual disability syndrome: a cohort of Portuguese patients

Raquel Gouveia-Silva^{a*}, João Rodrigues-Alves^b, Juliette Dupont^c, Oana Moldovan^d, Patrícia Dias, Márcia Rodrigues^e, Ana Medeira, and Ana B. Sousa

Serviço de Genética Médica, Departamento de Pediatria, Hospital de Santa Maria, Centro Hospitalar e Universitário Lisboa Norte, Lisbon, Portugal

ORCID: ^a0000-0002-5470-5490; ^b0000-0002-8659-1862; ^c0000-0001-8257-5067; ^d0000-0003-0274-4313; ^e0000-0001-6174-2524

Abstract

Introduction: *DYRK1A* heterozygous pathogenic variants have been shown to cause a syndromic form of intellectual disability (ID) with impaired speech development, features of autism spectrum disorder (ASD), microcephaly, and a recognizable facial gestalt that evolves with age. Patients can also present with gait disturbance or hypertonia, epilepsy, brain imaging, ocular, and foot anomalies. **Methods:** This is a cross-sectional study. Clinical data on patients with *DYRK1A* pathogenic variants identified at the Clinical Genetics Department of Santa Maria Hospital, in Lisbon, were retrospectively collected from medical records using a detailed clinical questionnaire. **Results:** We describe eight unrelated patients, six females and two males, aged 4 to 24. Fetal growth restriction (FGR) was present in 5/8, and microcephaly in 7/8. ID, ranging from mild to severe, and language impairment or absent speech were documented in all patients. ASD and/or stereotypic behavior were reported in 6/8. Five patients presented visual anomalies, most commonly optic disc pallor (in 4). Three main facial features were consistently reported: deep-set eyes, thin upper lip, and micro/retrognathia. Foot and hand anomalies were frequent. **Discussion/Conclusions:** Our cohort illustrates the variable degree of severity of a syndromic form of ID, which includes mild cases. Microcephaly and a typical neurobehavioral phenotype are in accordance with the literature, as well as some common dysmorphisms. Interestingly, optic disc pallor seems to be a frequent finding, highlighting the need for ophthalmological surveillance. Our study adds evidence to the existence of a consistent clinical phenotype of *DYRK1A*-related ID, hopefully contributing to increased awareness and improving the recognition of this entity.

Keywords: *DYRK1A*. Intellectual disability. Portuguese cohort.

Perturbação do desenvolvimento intelectual associada ao gene *DYRK1A*: uma coorte de doentes portugueses

Resumo

Introdução: Variantes patogénicas em heterozigotia no gene *DYRK1A* causam uma forma sindrómica de perturbação do desenvolvimento intelectual (PDI) com comprometimento do desenvolvimento da linguagem, características de perturbação do espectro do autismo (PEA), microcefalia e um gestalt facial reconhecível que evolui com a idade. Os doentes também podem apresentar distúrbios da marcha ou hipertonia, epilepsia, alterações cerebrais estruturais, anomalias oculares e nos pés.

*Correspondence:

Raquel Gouveia-Silva

E-mail: ana.g.f.silva@chln.min-saude.pt

2184-3333 / © 2023 Portuguese Society of Pediatrics. Published by Permalyer. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Received: 03-01-2023

Accepted: 20-06-2023

<https://pjp.spp.pt>

Available online: 07-02-2024

Port J Pediatr. 2024;55(1):13-19

DOI: 10.24875/PJP.M23000130

Métodos: Trata-se de um estudo transversal. Os dados clínicos de doentes com variantes patogénicas no gene *DYRK1A* identificados no Departamento de Genética Clínica do Hospital de Santa Maria, em Lisboa, foram recolhidos retrospectivamente a partir de registos médicos através de um questionário clínico detalhado. **Resultados:** Descrevemos oito doentes não aparentados, seis do sexo feminino e dois do sexo masculino, com idades entre 4 e 24 anos. A restrição do crescimento fetal (RCF) esteve presente em 5/8 e a microcefalia em 7/8. PDI, ligeira a grave, com compromisso da linguagem ou a sua ausência, foi documentada em todos os casos. PEA e/ou comportamento estereotipado foram relatados em 6/8. Cinco pacientes apresentaram anomalias visuais, mais frequentemente palidez do disco óptico (em 4). Três características faciais principais foram consistentemente relatadas: olhos afundados, lábio superior fino e micro/retrognatia. Anomalias nos pés e nas mãos foram frequentes. **Discussão/Conclusões:** A nossa coorte ilustra o grau variável de gravidade de uma forma sindrômica de PDI, que inclui casos ligeiros. A microcefalia e um fenótipo neurocomportamental típico estão de acordo com a literatura, assim como alguns dismorfismos comuns. Curiosamente, a palidez do disco óptico parece ser um achado frequente, destacando a necessidade de vigilância oftalmológica. O nosso estudo acrescenta evidências da existência de um fenótipo clínico consistente de PDI relacionada ao gene *DYRK1A*, contribuindo para aumentar e melhorar o reconhecimento desta entidade.

Palavras-chave: *DYRK1A*. Perturbação intelectual. Coorte portuguesa.

Keypoints

What is known

- *DYRK1A*-related intellectual disability syndrome is a rare condition with variable degrees of developmental delay reported.
- Loss-of-function *DYRK1A* pathogenic variants include disruptive balanced rearrangements, copy number variants, and single nucleotide variants (SNVs).
- No genotype-phenotype correlations could be consistently drawn.

What is added

- We report a new cohort of patients with *DYRK1A*-related intellectual disability syndrome.
- With advances on next-generation sequencing techniques, we expected an earlier diagnosis, raising awareness about common dysmorphic features, delineating a typical gestalt.
- Considering the prevalence of eye problems, namely optic disc pallor, a careful ophthalmological evaluation must be performed.

Introduction

DYRK1A (dual-specificity tyrosine phosphorylation regulated kinase 1A) is a protein kinase encoded on human chromosome 21¹⁻³. The gene is dosage-sensitive with an impact on neurodevelopment⁴. *DYRK1A* is over expressed in Down syndrome (DS) and plays a relevant role in neuronal proliferation, differentiation, plasticity, and aging, making it a primary target for the development of DS therapeutics^{2,5}. Dysregulation of *DYRK1A* levels also occurs in neurodegenerative diseases, such as Alzheimer and Parkinson⁶.

Since 2008, the *DYRK1A* gene has gained attention in the clinical setting, starting with the description of two patients with microcephaly, fetal growth restriction, and febrile seizures. Both had a *de novo* balanced translocation that truncated the *DYRK1A* gene, suggesting *DYRK1A* haplo-insufficiency is associated with brain developmental changes⁷.

In 2011, a *de novo* micro-deletion involving the last three exons of *DYRK1A* was described in a female with microcephaly and intellectual disability⁸. Considering these chromosomal rearrangements, as well as large deletions encompassing additional genes (that may

contribute to extra-*DYRK1A*-related features)^{9,10,11}, van Bon et al. proposed that the heterozygous disruption of this gene causes a distinctive neuro-developmental syndrome, including mild to severe ID, microcephaly, and impaired speech³.

One year later, Courcet et al. described a patient with an Angelman-like phenotype carrying a frameshift variant affecting the N-terminal part of the gene^{12,13}. As more patients with single nucleotide variants in *DYRK1A* were documented, this gene was definitely placed on the list of genes responsible for syndromic ID. Furthermore, it has also been linked to autism spectrum disorder (ASD), with studies showing that *DYRK1A* syndrome accounts for 0.1-0.5% of individuals with ID and/or autism^{3,12,14}.

Core symptoms of *DYRK1A*-related ID syndrome (OMIM#614104) (also known as *DYRK1A*-haploinsufficiency syndrome or autosomal dominant mental retardation 7, MRD7) are ID with impaired speech development of variable degree, features of ASD with anxious and/or stereotypic behavior problems, microcephaly, and a recognizable facial gestalt that evolves with age^{3,15}. Patients can also present with gait disturbance or hypertonía, epilepsy, brain-MRI anomalies, feeding issues, eye problems, and foot anomalies¹⁵⁻¹⁷.

Loss-of-function *DYRK1A* pathogenic variants include disruptive balanced rearrangements, copy number variants, and single nucleotide variants (SNVs)¹⁶. About 500 families are part of the *DYRK1A* Syndrome International Association (DSIA). According to our acknowledge, there are around 112 patients reported in the literature^{1,3,7,12-25} of whom with chromosomal abnormalities, including large deletions, translocations, inversions, inversion/deletions, and other complex rearrangements, and 87 patients harboring a SNV. Most reported SNVs were frameshift variants, followed by nonsense, missense, and splice site variants¹⁶.

With this report, we aim to contribute to further delineating the phenotype of this still under-recognized but increasingly diagnosed condition.

Methods

This is a retrospective and cross-sectional study. Clinical data on patients with *DYRK1A* variants identified at the Genetics Department of Santa Maria Hospital, in Lisbon, were collected from medical records during 2022. We searched in our internal database (Microsoft® Office Access 2019) for patients with this diagnosis followed from January 2010 until December 2022 in our genetic service of a tertiary level hospital. Patient characterization was performed through a detailed questionnaire, completed by the clinician and organized into a standardized form designed in Microsoft® Office Excel 2010. This form was based on international registries and available literature, including information on the demographics, patient and family history, clinical and laboratory findings, genetic studies, and evolution. The molecular diagnosis was possible through whole exome sequencing (WES, CentoXome® from the Centogene laboratory in Germany) in all cases.

Results

A diagnosis of *DYRK1A*-related ID syndrome was made for eight unrelated patients, six females and two males, aged 4 to 24 years old, all representing simplex cases with a pathogenic or likely pathogenic variant in the *DYRK1A* gene. The summary of their main clinical features is shown in [table 1](#).

FGR was present in 5/8. Further growth evaluation showed short stature in 4/8 and microcephaly in 7/8. ID, ranging from mild to severe, and language impairment (6/8) or absent speech (2/8) were documented in all patients. ASD and/or stereotypic behavior were present in 6/8. Only two patients had epilepsy, brain anomalies on MRI were present in 3/7, and 4/8 had gait

disturbance (often ataxic) and hypertonic extremities, with no need to use support devices. Five patients had visual anomalies (strabismus, astigmatism, hypermetropia, optic nerve anomalies), with optic disc pallor present in four of them.

Characteristic facial features consistently reported were deep-set eyes (7/8), thin upper lip (8/8), and micro/retrognathia (5/8). Foot anomalies were described in 5/8 patients. Additional common features included thick eyebrows, prominent nose, large or dysplastic ears, and large mouth ([Fig. 1](#)).

The diagnosis was made on whole exome sequencing (WES) in all cases, but suspected clinically in four. *DYRK1A* variants were classified as pathogenic or likely pathogenic using ACMG criteria. All variants were located between exons four and ten, and 7/8 were loss of function, including three frameshift, two splicing, and two nonsense variants. There was only one missense variant not affecting the kinase domain. Parental segregation studies confirmed all cases were *de novo* ([Table 1](#)).

Patient 6 had an additional pathogenic variant in *PKD2* gene, explaining the presence of renal cysts.

Discussion

The purpose of this study was to characterize the *DYRK1A*-related ID syndrome patients diagnosed in our Medical Genetics Department. No Portuguese cohort has been published so far, and this syndrome is still unknown to many medical professionals dealing with ID/ASD patients. We have attempted to further delineate the phenotype of these patients by comparing them with those already described in the literature harboring a causative variant in *DYRK1A* gene. The clinical findings were consistent with the literature ([Table 1](#)).

Clinically, it is worth emphasizing the variable degree of severity of ID^{3,6,15}, which includes two mild cases in this cohort. Language impairment is seen in a significant proportion of patients^{15,17}, with completely absent speech in two of ours. ASD, stereotypes, anxious behavior, hyperactivity, and sleep disturbances have also been frequently observed^{3,15}, and were reported in all but two individuals in this cohort. Conversely, only two patients had epilepsy, which is lower than in the literature^{12,17}. Seizures of the atonic, absence, and generalized myoclonic types have all been previously described^{1,3,12,13}. FGR was present in 5/8 patients and can be a non-specific clue for the diagnosis of *DYRK1A*-related ID syndrome in the prenatal setting^{3,13,20}, but more reports are needed to reinforce this finding. Head circumference at birth is known to be between -1 and -4 SD, and abnormally slow head growth causes the

Table 1. Clinical characteristics of our cohort of Portuguese patients with *DYRK1A*-related intellectual disability syndrome and literature comparison

	Patient 1	Patient 2	Patient 3	Patient 4	Patient 5	Patient 6	Patient 7	Patient 8	Total n = 8 (%)	Patients in literature (%) 1, 3, 7, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24
Age (years)	14	19	12	24	4	18	5	7	-	-
Genotype	c. 361C>T, p. (Gln121*)	c. 1405C>T, p. (Gln469*)	c. 665-9_ 665-5del	c. 562A>G, p.(Lys188Glu)	c. 336_348del, p.(Ala113Glufs*33)	c. 358del, p. (Gln120A snfs*30)	c. 951+4_ 951+7del	c. 894del, p. (Lys299Asnfs*7)	-	-
Type of variant	Nonsense*	Nonsense	Splicing*	Missense	Frameshift	Frameshift	Splicing*	Frameshift	-	-
Classification	Likely pathogenic	Likely pathogenic	Pathogenic	Likely pathogenic	Likely pathogenic	Likely pathogenic	Pathogenic	Likely pathogenic	-	-
Inheritance	De novo	De novo	De novo	De novo	De novo	De novo	De novo	De novo	-	-
Sex	F	F	F	M	F	M	F	F	-	-
FGR	-	+	+	-	+	+	+	-	5/8 (63%)	33-64%
Microcephaly	-	+	+	+	+	+	+	+	7/8 (88%)	68-98%
Short stature	-	+	+	+	-	-	+	-	4/8 (50%)	10-41%
DD/ID	Mild-Moderate	Moderate	Mild	Severe	Moderate	Severe	Moderate	Moderate	8/8 (100%)	95-100%
Speech impairment	Impaired	Impaired	Impaired	Absent speech	Impaired	Impaired	Absent speech	Impaired	8/8 (100%)	96-100%
ASD/ Stereotypic behaviour	-	+	+	+	+	-	+	+	6/8 (75%)	29-50%
Feeding issues	-	-	-	+	-	-	+	+	3/8 (38%)	48-82%
Gait disturbance and/or hypertonia	-	-	-	+	+	-	+	+	4/8 (50%)	NA
Epilepsy	-	-	-	+	-	+	-	-	2/8 (25%)	20%-67%

(Continues)

Table 1. Clinical characteristics of our cohort of Portuguese patients with *DYRK1A*-related intellectual disability syndrome and literature comparison (*continued*)

	Patient 1	Patient 2	Patient 3	Patient 4	Patient 5	Patient 6	Patient 7	Patient 8	Total n = 8 (%)	Patients in literature (%) 1, 3, 7, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24
Brain MRI anomalies	Megacisterna magna, hypoplasia of optic nerve and chiasma, corpus callosum agenesis	-	-	Cerebellar atrophy	-	-	NA	Bilateral ventricular enlargement	3/7 (43%)	31%-59%
Eye anomalies	Strabismus, astigmatism, hypermetropia, optic disc pallor, optic nerve bilateral atrophy	-	Strabismus, optic disc pallor	Strabismus, optic disc pallor	Temporal optic disc pallor	Strabismus	-	-	5/8 (63%)	46%-58%
Hand/feet anomalies	+	+	-	+	+	+	-	-	5/8 (63%)	37-50%
Deep set eyes	+	+	+	+	+	+	-	+	7/8 (88%)	10-29%
Large ears	+	+	+	+	-	-	-	-	4/8 (50%)	50-58%
Dysplastic ears	-	-	+	-	-	+	-	+	3/8 (38%)	NA
Long/flat filtrum	-	+	-	-	+	-	-	+	3/8 (38%)	NA
Thin upper lip	+	+	+	+	+	+	+	+	8/8 (100%)	NA
Micro/ retrognathia	+	-	-	+	+	+	-	+	5/8 (63%)	37%-54%
Additional features	Long eyelashes, bulbous nose tip, prominent supraciliary arch	Bitemporal narrowing, thick eyebrows, hypotelorism, malar hypoplasia, large mouth	Metopic prominence	Thick eyebrows, long nose, malar hypoplasia, large mouth		Renal cysts due to <i>PKD2</i> heterozygous pathogenic variant				

*Previously reported variants. ASD: autism spectrum disorder; DD: developmental delay; F: female; FGR: fetal growth restriction; ID: intellectual disability; M: male; NA: not available.



Figure 1. Images depicting similar and key dysmorphic features of presented cases, namely deep set eyes, thin upper lip and micro/retrognathia. **1A:** patient 1 at 11 months. **1B:** 5 years. **1C:** 10 years. **2A:** patient 2 at 3 years. **2B:** 11 years. **2C:** 13 years. **3A:** patient 3 at 1 year. **3B:** 3 years. **3C:** 6 years. **4A:** patient 4 at 13 years. **4B and C:** at 19 years. **5A:** patient 5 at 1 month. **5B:** 6 months. **5C:** 2 years. **6A:** patient 6 at 13 years. **6B and C:** 18 years.

deviation to further increase over time to -2 to -5 SD in the majority of individuals^{3,15,17}. Accordingly, microcephaly was present in all but one patient in this report.

Gait disturbances, broad-based or stiff, has been reported in a smaller fraction of individuals¹⁷, with about

half of the cases presenting this characteristic, very similar to what happens in our cohort.

Brain anomalies were described in 3/7 patients who had an MRI, including ventricular enlargement, sub-ependymal cysts, hypoxic-ischemic-like anomalies, corpus callosum agenesis, and cerebellar atrophy. Brain MRI anomalies are not well established in the literature, involving from grey matter to cerebral spinal fluid and ventricle anomalies^{25,26}.

Ocular anomalies were reported in five patients in our cohort, in line with Méjécase et al., who describe ocular features in 60.3% of patients with *DYRK1A* SNVs, predominantly refractive errors¹⁶. Optic nerve anomalies, including optic disc pallor, have been reported in 22.9% of patients¹⁶, and were present in half of our cases.

Several publications refer to a typical gestalt or common dysmorphisms^{1,3,13,15,18,20}. All patients in this report had a thin upper lip, and deep-set eyes were described in all but one (Fig. 1). Other frequent features include micro/retrognathia and large ears. In adulthood, the nasal bridge may become high and the alae nasi underdeveloped, giving the nose a more prominent appearance³, as shown in patient 6 (Figs. 1-6 A, B, C).

Hand and feet anomalies were also frequent in our cohort and in previously reported patients^{15,17}, namely long splayed fingers, clinodactyly, mild cutaneous syndactyly of toes, hallux valgus, and short fifth toe.

The majority of variants elicited in our cohort were truncating, namely frameshift (Table 1), which is in accordance with previous reports¹⁶. Only three missense variants were previously reported in the literature^{3,26}. No genotype-phenotype correlations could be consistently drawn. All cases were *de novo*.

From the above, it is clear that all *DYRK1A*-related syndrome ID patients need guidance for educational and behavioral problems, and should be regularly monitored for growth parameters and nutritional status, as well as have lifelong specialist follow up for issues affecting particular organs and systems^{6,16,23}. Early motor intervention is also relevant regarding gait disorders and hypertonia, in order to avoid the use of a mobility device in the future¹⁷. Of note, taking into account the prevalence of eye problems, multidisciplinary care must include a careful ophthalmological evaluation.

All in all, national and international registries²⁷, and other forms of data sharing, are paramount in acquiring new insights into the pathophysiology and genetic mechanisms underlying this condition, which will ultimately contribute to improve patient management. In line with this endeavor, we hope this clinical review contributes to raise awareness of *DYRK1A*-related ID syndrome, namely among Portuguese doctors, so as to prompt the diagnosis of more patients.

Acknowledgments

The Genetics Service at Hospital de Santa Maria collaborates with the European Reference Network on Congenital Malformations and Rare Intellectual Disability (ERN-ITHACA) at the date of publication.

Funding

None.

Conflicts of interest

None.

Ethical disclosures

Protection of human and animal subjects. The authors declare that the procedures followed were in accordance with the regulations of the relevant clinical research ethics committee and with those of the Code of Ethics of the World Medical Association (Declaration of Helsinki).

Confidentiality of data. The authors declare that they have followed the protocols of their work center on the publication of patient data.

Right to privacy and informed consent. The authors have obtained the written informed consent of the patients or subjects mentioned in the article. The corresponding author is in possession of this document.

Use of artificial intelligence for generating text. The authors declare that they have not used any type of generative artificial intelligence for the writing of this manuscript, nor for the creation of images, graphics, tables, or their corresponding captions.

References

- Bronicki LM, Redin C, Drunat S, Piton A, Lyons M, Passemard S, et al. Ten new cases further delineate the syndromic intellectual disability phenotype caused by mutations in *DYRK1A*. *Eur J Hum Genet*. 2015 Nov;23(11):1482-7.
- Duchon A, Herault Y. *DYRK1A*, a Dosage-Sensitive Gene Involved in Neurodevelopmental Disorders, Is a Target for Drug Development in Down Syndrome. *Front Behav Neurosci*. 2016 Jun 3;10:104.
- van Bon BW, Coe BP, Bernier R, Green C, Gerdts J, Witherspoon K, et al. Disruptive de novo mutations of *DYRK1A* lead to a syndromic form of autism and ID. *Mol Psychiatry*. 2016 Jan;21(1):126-32.
- Liu F, Liang Z, Wegiel J, Hwang YW, Iqbal K, Grundke-Iqbal I, et al. Overexpression of *Dyrk1A* contributes to neurofibrillary degeneration in Down syndrome. *FASEB J*. 2008 Sep;22(9):3224-33.
- Meissner LE, Macnamara EF, D'Souza P, Yang J, Vezina G; Undiagnosed Diseases Network, Ferreira CR, Zein WM, Tift CJ, Adams DR. *DYRK1A* pathogenic variants in two patients with syndromic intellectual disability and a review of the literature. *Mol Genet Genomic Med*. 2020;8(12):e1544.
- Arbones ML, Thomazeau A, Nakano-Kobayashi A, Hagiwara M, Delabar JM. *DYRK1A* and cognition: A lifelong relationship. *Pharmacol Ther*. 2019;194:199-221.
- Møller RS, Kübart S, Hoeltzenbein M, Heye B, Vogel I, Hansen CP, et al. Truncation of the Down syndrome candidate gene *DYRK1A* in two unrelated patients with microcephaly. *Am J Hum Genet*. 2008 May;82(5):1165-70.
- van Bon BW, Hoischen A, Hehir-Kwa J, de Brouwer AP, Ruivenkamp C, Gijbbers AC, et al. Intragenic deletion in *DYRK1A* leads to mental retardation and primary microcephaly. *Clin Genet*. 2011;79(3):296-9.
- Fujita H, Torii C, Kosaki R, Yamaguchi S, Kudoh J, Hayashi K, et al. Microdeletion of the Down syndrome critical region at 21q22. *Am J Med Genet A*. 2010;152A:950-3.
- Lyle R, Béna F, Gagos S, Gehrig C, Lopez G, Schinzel A, et al. Genotype-phenotype correlations in Down syndrome identified by array CGH in 30 cases of partial trisomy and partial monosomy chromosome 21. *Eur J Hum Genet*. 2009 Apr;17(4):454-66.
- Matsumoto N, Ohashi H, Tsukahara M, Kim KC, Soeda E, Niikawa N. Possible narrowed assignment of the loci of monosomy 21-associated microcephaly and intrauterine growth retardation to a 1.2-Mb segment at 21q22.2. *Am J Hum Genet*. 1997 Apr;60(4):997-9.
- Courcet JB, Faivre L, Malzac P, Masurel-Paulet A, Lopez E, Callier P, et al. The *DYRK1A* gene is a cause of syndromic intellectual disability with severe microcephaly and epilepsy. *Journal of Medical Genetics*. 2012 Dec;49(12):731-736.
- Ji J, Lee H, Argiropoulos B, Dorrani N, Mann J, Martinez-Agosto JA, et al. *DYRK1A* haploinsufficiency causes a new recognizable syndrome with microcephaly, intellectual disability, speech impairment, and distinct facies. *Eur J Hum Genet*. 2015 Nov;23(11):1473-81.
- O'Roak BJ, Vives L, Fu W, Egerton JD, Stanaway IB, Phelps IG, et al. Multiplex targeted sequencing identifies recurrently mutated genes in autism spectrum disorders. *Science*. 2012 Dec 21;338(6114):1619-22.
- Earl RK, Turner TN, Mefford HC, Hudac CM, Gerdts J, Eichler EE, et al. Clinical phenotype of ASD-associated *DYRK1A* haploinsufficiency. *Mol Autism*. 2017 Oct 5;8:54.
- Méjécase C, Way CM, Owen N, Moosajee M. Ocular Phenotype Associated with *DYRK1A* Variants. *Genes (Basel)*. 2021 Feb 5;12(2):234.
- van Bon BW, Coe BP, de Vries BBA, Eichler EE. *DYRK1A* Syndrome. 2015 Dec 17 [Updated 2021 Mar 18]. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. *GeneReviews*® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2022.
- Redin C, Gérard B, Lauer J, Herenger Y, Muller J, Quartier A, et al. Efficient strategy for the molecular diagnosis of intellectual disability using targeted high-throughput sequencing. *J Med Genet*. 2014 Nov;51(11):724-36. doi: 10.1136/jmedgenet-2014-102554.
- Ruaud L, Mignot C, Guét A, Ohl C, Nava C, Héron D, et al. *DYRK1A* mutations in two unrelated patients. *Eur J Med Genet*. 2015 Mar;58(3):168-74.
- Luco SM, Pohl D, Sell E, Wagner JD, Dymeny DA, Daoud D. Case report of novel *DYRK1A* mutations in 2 individuals with syndromic intellectual disability and a review of the literature. *BMC Med Genet*. 2016;17:15.
- Evers JM, Laskowski RA, Bertolli M, Clayton-Smith J, Deshpande C, Eason J, et al. Structural analysis of pathogenic mutations in the *DYRK1A* gene in patients with developmental disorders. *Hum Mol Genet*. 2017 Feb 1;26(3):519-526.
- Murray CR, Abel SN, McClure MB, Foster J 2nd, Walke MI, Jayakar P, et al. Novel Causative Variants in *DYRK1A*, *KARS*, and *KAT6A* Associated with Intellectual Disability and Additional Phenotypic Features. *J Pediatr Genet*. 2017 Jun;6(2):77-83.
- Blackburn ATM, Bekheirnia N, Uma VC, Corkins ME, Xu Y, Rosenfeld JA, et al. *DYRK1A*-related intellectual disability: a syndrome associated with congenital anomalies of the kidney and urinary tract. *Genet Med*. 2019 Dec;21(12):2755-2764.
- Qiao F, Shao B, Wang C, Wang Y, Zhou R, Liu G, et al. A De Novo Mutation in *DYRK1A* Causes Syndromic Intellectual Disability: A Chinese Case Report. *Front Genet*. 2019 Nov 19;10:1194.
- Okazaki T, Yamada H, Matsuura K, Kasagi N, Miyake N, Matsumoto N, et al. Clinical course of epilepsy and white matter abnormality linked to a novel *DYRK1A* variant. *Hum Genome Var*. 2021 Jul 12;8(1):26.
- Frewer V, Gilchrist CP, Collins SE, Williams K, Seal ML, Leventer RJ, et al. A systematic review of brain MRI findings in monogenic disorders strongly associated with autism spectrum disorder. *J Child Psychol Psychiatry*. 2021 Nov;62(11):1339-1352.
- Turnbull C, Scott RH, Thomas E, Jones L, Murugaesu N, Pretty FB, et al. The 100 000 Genomes Project: bringing whole genome sequencing to the NHS. *BMJ*. 2018 Apr 24;361:k1687.