

Free Communications 6: Fat, Metabolism And Obesity 1

FC6.2

Frequency of Bardet-Biedl syndrome variants in a population with early-onset obesity

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Frequency of Bardet-Biedl Syndrome variants in a population with early-onset obesity



Presentation FC6.2

Jesús Argente, MD, PhD, presenting

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DISCLOSURE STATEMENT

Jesús Argente

X I have the following potential conflicts of interest to report:

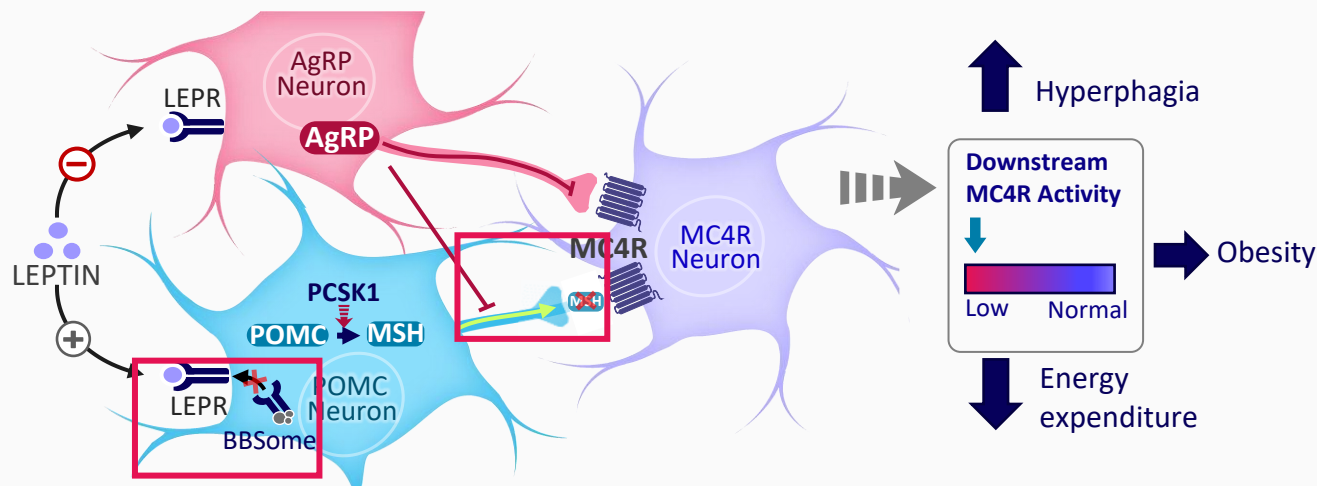
- Research Contracts
- X Consulting
- Employment in the Industry
- Stockholder of a healthcare company
- Owner of a healthcare company
- X Other(s) – speaking engagements for Rhythm Pharmaceuticals, Inc.



The central hypothalamus is a key regulator of energy balance, appetite and bodyweight¹⁻⁴

- The melanocortin-4 receptor (MC4R) pathway in the hypothalamus is critical for the regulation of hunger, energy balance, and body weight¹⁻³
- Individuals who carry disruptive variants in MC4R pathway genes may present with hyperphagia (pathologic, insatiable hunger) and early-onset obesity⁴
- In patients with Bardet–Biedl syndrome (BBS), a rare, genetically heterogeneous, and highly pleiotropic disease, the immotile primary cilia are dysfunctional, leading to impairment of the MC4R pathway

Hypothalamus^{1-4,9,10}



BBSome⁵⁻⁸

The BBSome traffics intracellular proteins to ciliary membranes and other membrane compartments

BBSome genes

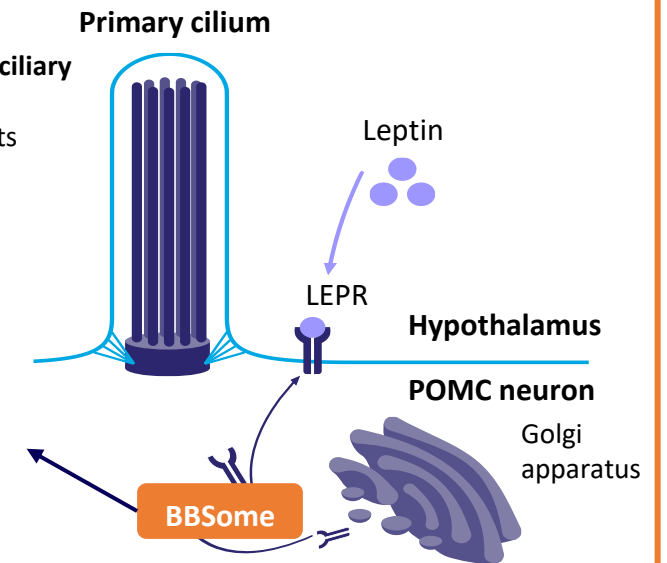
BBS1 *BBS7*
BBS2 *BBS8*
BBS4 *BBS9*
BBS5 *BBS18*

Chaperonin complex

BBS6 *BBS12*
BBS10

BBSome trafficking

BBS3 *BBS17*



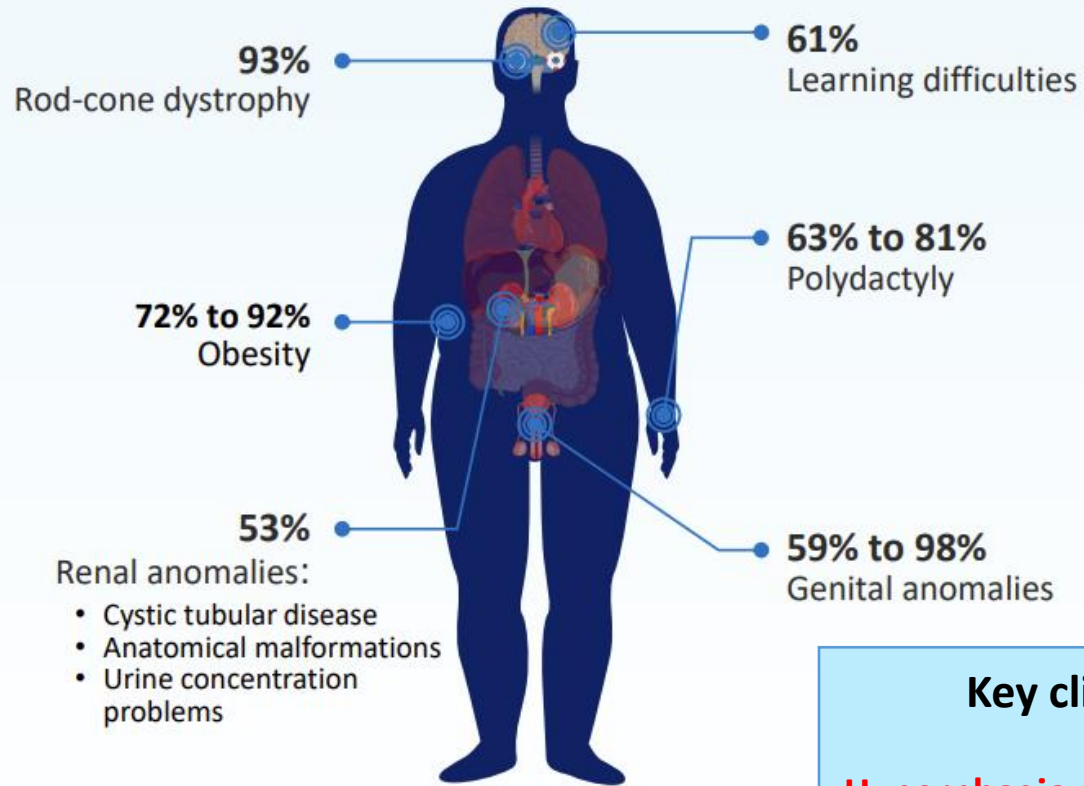
AgRP, agouti-related peptide; LEPR, leptin receptor; MC4R, melanocortin-4 receptor; MSH, melanocyte-stimulating hormone; PCSK1, proprotein convertase subtilisin/kexin type 1; POMC, proopiomelanocortin

1. Huvenne et al. *Obes Facts*. 2016;9:158-173.
2. Yazdi et al. *PeerJ*. 2015;3:e856.
3. Yang et al. *Nat Commun*. 2019;10:1718.
4. Revelli et al. *Obesity (Silver Spring)*. 2011;19:1010-1018.
5. Vaisse et al. *Cold Spring Harb Perspect Biol* 2016;9:a028217;
6. Guo and Rahmouni. *Trends Endocrinol Metab*. 2011;22:286-293;
7. Seo et al. *Hum Mol Genet*. 2009;18:1323-1331;
8. Wang et al. *J Clin Invest*. 2021;131:e146287;
9. Doche et al. *J Clin Invest*. 2012;122:4732-4736.
10. Ghamari-Langroudi et al. *Sci Adv*. 2018;4:eaat0866.

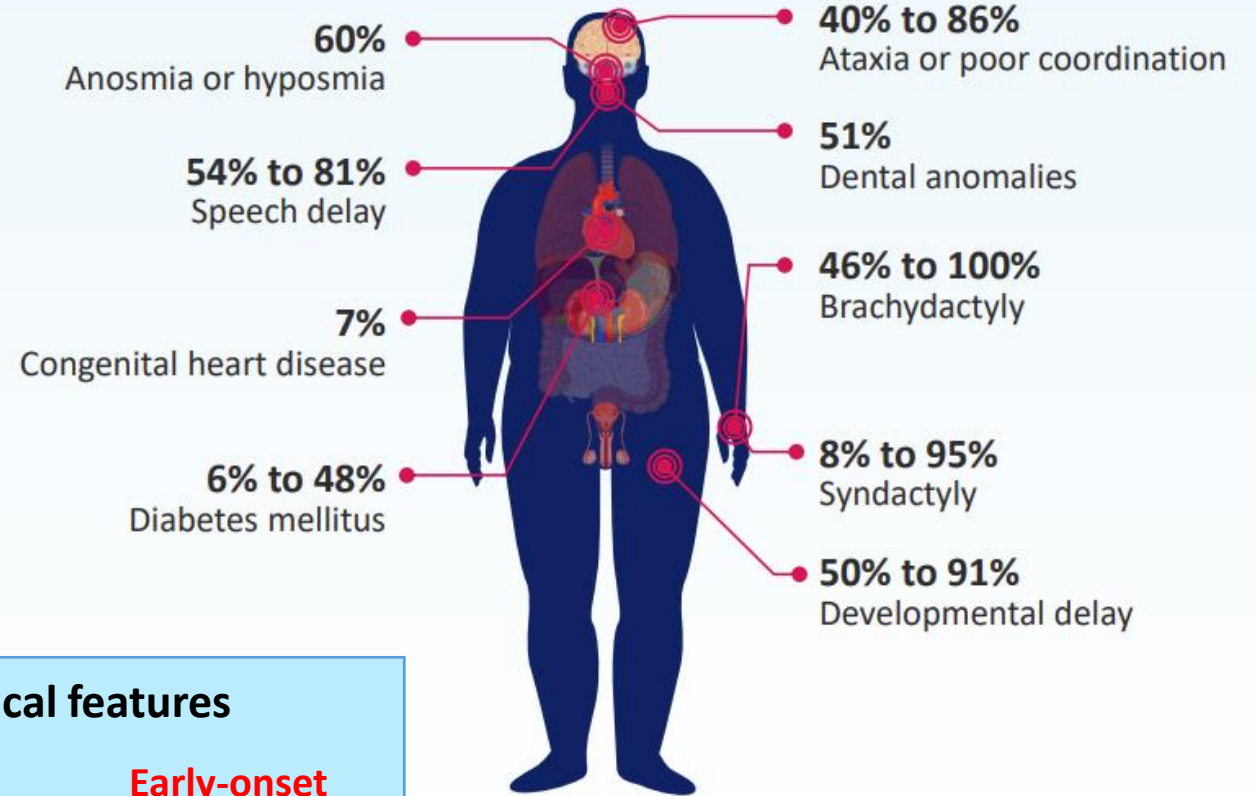
The presence and severity of clinical features are highly variable in BBS¹⁻⁴

Onset of clinical manifestations is variable making diagnosis challenging, as not all symptoms might be present during early testing

Primary clinical features



Secondary clinical features



Key clinical features

Hyperphagia

Early-onset obesity

Early identification of patients with BBS is essential for optimal disease management¹⁻⁴

Early and routine genetic testing can...

Improve identification and diagnosis of individuals with hyperphagia and early-onset obesity caused by rare genetic variants¹⁻⁵

Inform specialized management strategies or eligibility for clinical trials¹⁻⁵

Hyperphagia and obesity caused by rare genetic variants are likely underdiagnosed due to low prevalence and limited access to genetic testing for individuals with obesity^{7,8}



BBS, Bardet-Biedl syndrome.

1. Gregoric et al. *Front Endocrinol (Lausanne)*. 2021;12:689387. 2. Styne et al. *J Clin Endocrinol Metab*. 2017;102:709-757. 3. van der Valk et al. *Obes Rev*. 2019;20:795-804. 4. Zorn et al. *Mol Cell Pediatr*. 2020;7:15. 5. Huvenne et al. *Obes Facts*. 2016;9:158-173; 6. Dollfus et al. *Eur J Hum Genet*. 2024;10.1038/s41431-024-01634-7; 7. Ayers et al. *J Clin Endocrinol Metab*. 2018;103:2601-2612; 8. Clément et al. *Physiol Behav*. 2020;227:113134.

The Rare Obesity Advanced Diagnosis (ROAD)[®] program

The ROAD[®] genetic testing program aims to encourage and support timely diagnosis by physicians of individuals with suspected rare MC4R pathway diseases

- 79-gene and 1–chromosomal region panel for individuals living in participating regions^a and who meet eligibility criteria
- Testing is conducted by an ISO 15189 accredited clinical laboratory

| Eligibility | | | |
|---|----|---|---|
| ≤ 18 years of age with a BMI | OR | ≥ 19 years of age with a BMI | OR |
| ≥ 97 th percentile | | ≥ 40 kg/m ² and a history of childhood obesity | |
| | |  Immediate family member of select, previously tested patients | OR  Showing clinical symptoms of BBS |

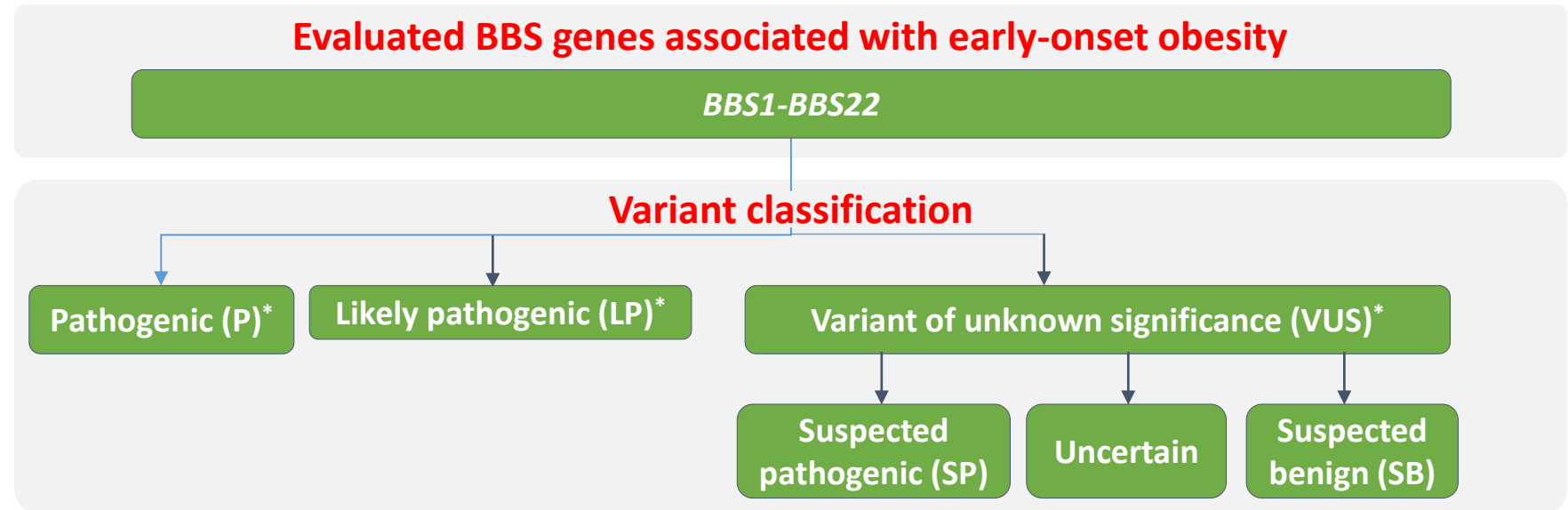
^aGermany, Greece, Ireland, Israel, Italy, Spain, Türkiye, the United Kingdom; Rare Obesity Advanced Diagnosis (ROAD) and its logo are registered trademarks of Rhythm Pharmaceuticals, Inc. BMI, body mass index. ROAD, Rare Obesity Advanced Diagnosis.

BBS gene variant analyses within ROAD[®]

Objective:

- To assess the frequency of BBS genetic variants in individuals with hallmark symptoms of potential underlying genetic causes of early-onset obesity who were sequenced as part of the ROAD[®] genetic testing program

| Genes sequenced | | | |
|--------------------------------|-----------------------------------|------------------------------------|------------------------------------|
| <i>BBS1</i> | <i>BBS7</i> | <i>MKS1</i> (<i>BBS13</i>) | <i>IFT27</i> (<i>BBS19</i>) |
| <i>BBS2</i> | <i>TTC8</i> (<i>BBS8</i>) | <i>CEP290</i> (<i>BBS14</i>) | <i>IFT172</i> (<i>BBS20</i>) |
| <i>ARL6</i> (<i>BBS3</i>) | <i>PTHB1</i> (<i>BBS9</i>) | <i>WDPCP</i> (<i>BBS15</i>) | <i>CFAP418</i> (<i>BBS21</i>) |
| <i>BBS4</i> | <i>BBS10</i> | <i>SDCCAG8</i> (<i>BBS16</i>) | <i>IFT74</i> (<i>BBS22</i>) |
| <i>BBS5</i> | <i>TRIM32</i> (<i>BBS11</i>) | <i>LZTFL1</i> (<i>BBS17</i>) | |
| <i>MKKS</i> (<i>BBS6</i>) | <i>BBS12</i> | <i>BBIP1</i> (<i>BBS18</i>) | |

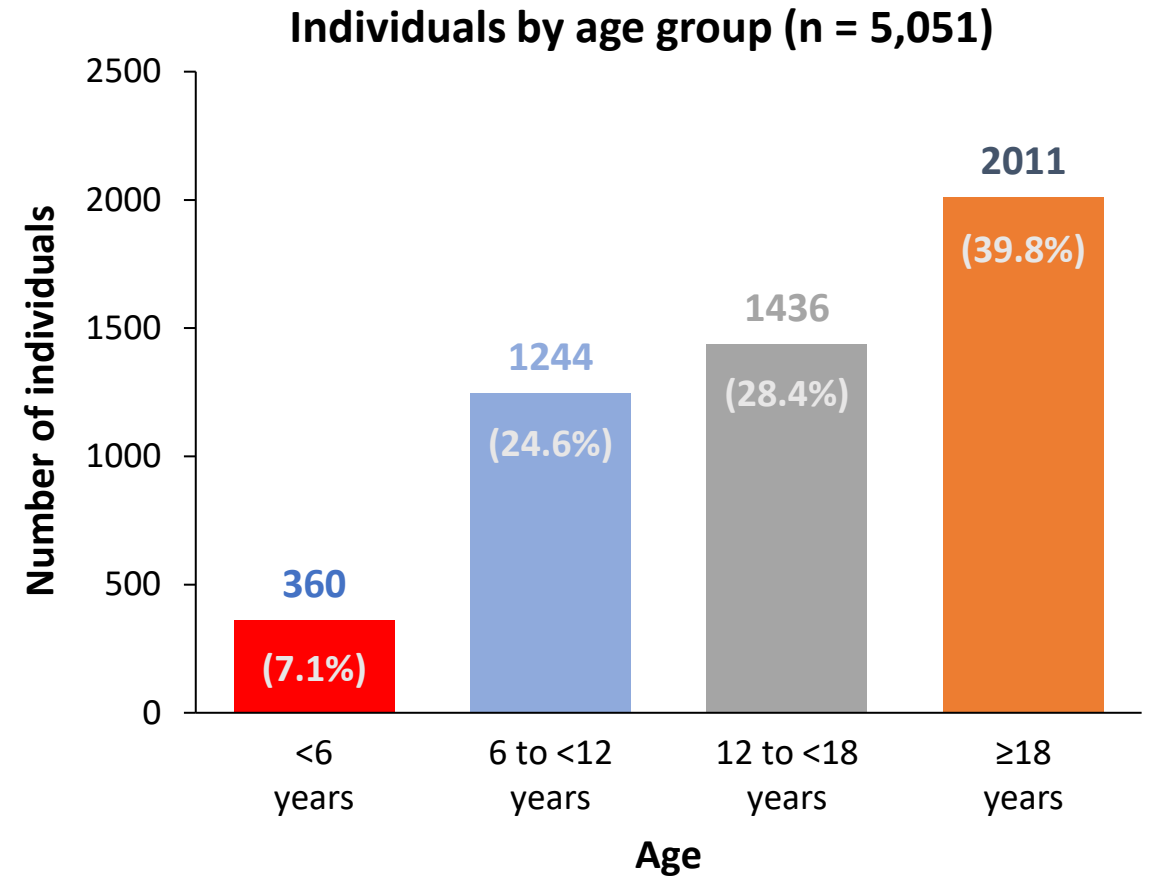


BBS, Bardet-Biedl syndrome; MC4R, melanocortin-4 receptor; ROAD, Rare Obesity Advanced Diagnosis. Rare Obesity Advanced Diagnosis (ROAD) and its logo are registered trademarks of Rhythm Pharmaceuticals, Inc.

*Variants were classified as P/LP/VUS according to American College of Medical Genetics criteria.

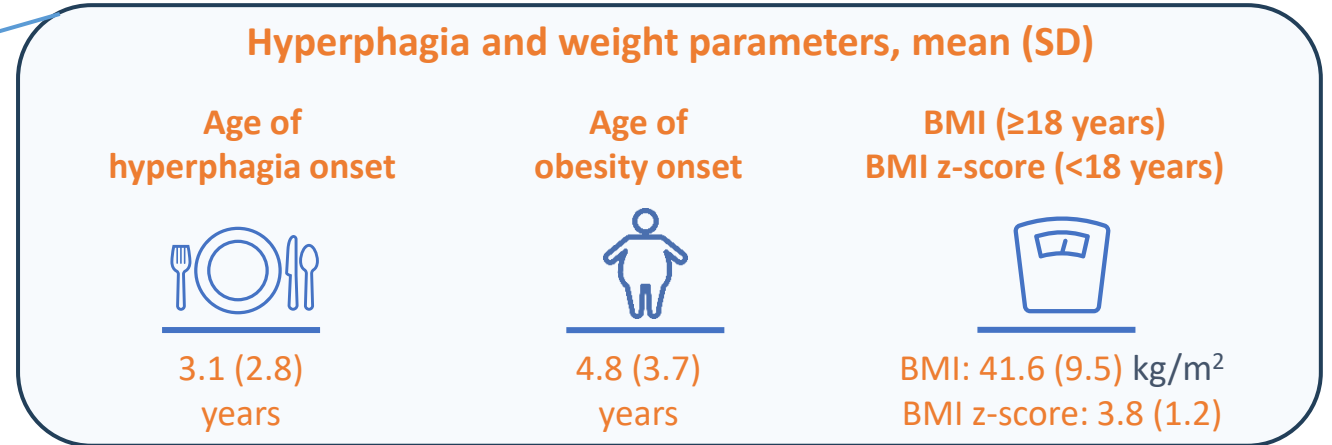
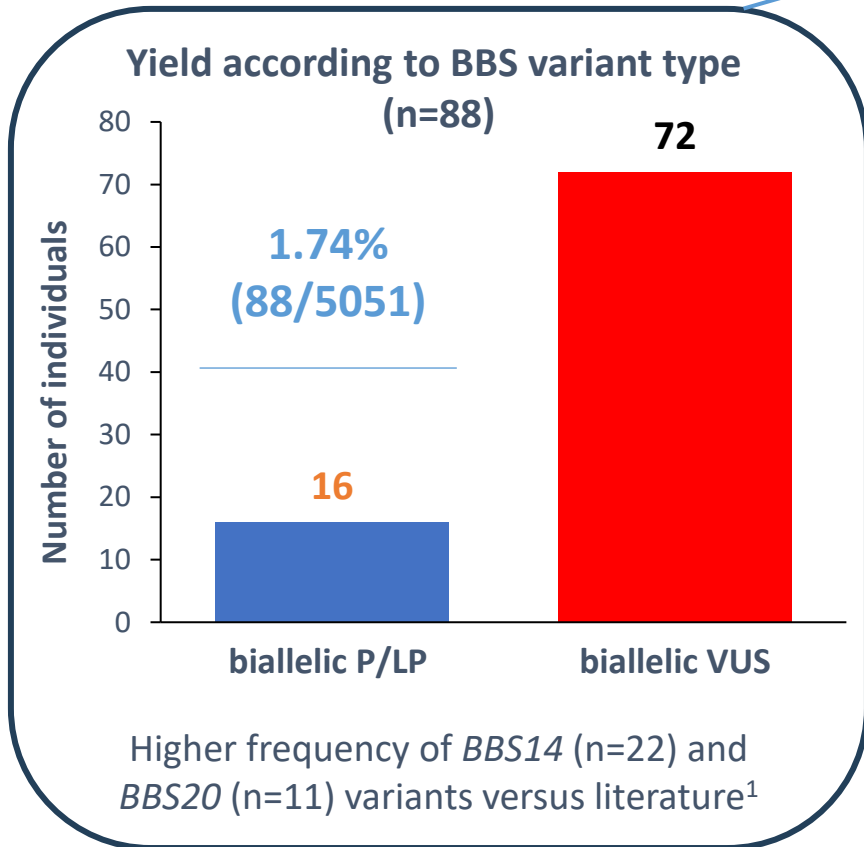
Baseline characteristics of total sequenced individuals

| Parameter | Total (n = 5,051) |
|--|----------------------|
| Sex, n (%) | |
| Female | 2,763 (54.7) |
| Male | 2,285 (45.2) |
| Prefer not to disclose/not provided | 3 (0.1) |
| Age at time of testing, mean (SD) | 23.0 (17.7) |
| Age at onset of obesity (n= 1,554), mean (SD) | 8.0 (3.9) |
| BMI z-score (patients aged <18 years, n= 3026), mean (SD) | 3.6 (1.0) |
| BMI (patients aged ≥18 years, n=1,936), mean (SD), kg/m² | 43.7 (10.1) |
| Location, n (% of tested individuals) | |
| Spain | 1,834 (36.3) |
| Italy | 1,291 (25.6) |
| Türkiye | 653 (12.9) |
| Israel | 501 (9.9) |
| United Kingdom | 445 (8.8) |
| Ireland | 250 (4.9) |
| Germany | 76 (1.5) |



BMI, body mass index; SD, standard deviation.

1.74% of tested individuals carried biallelic variants in one of 22 tested BBS genes and presented with early hyperphagia and obesity onset

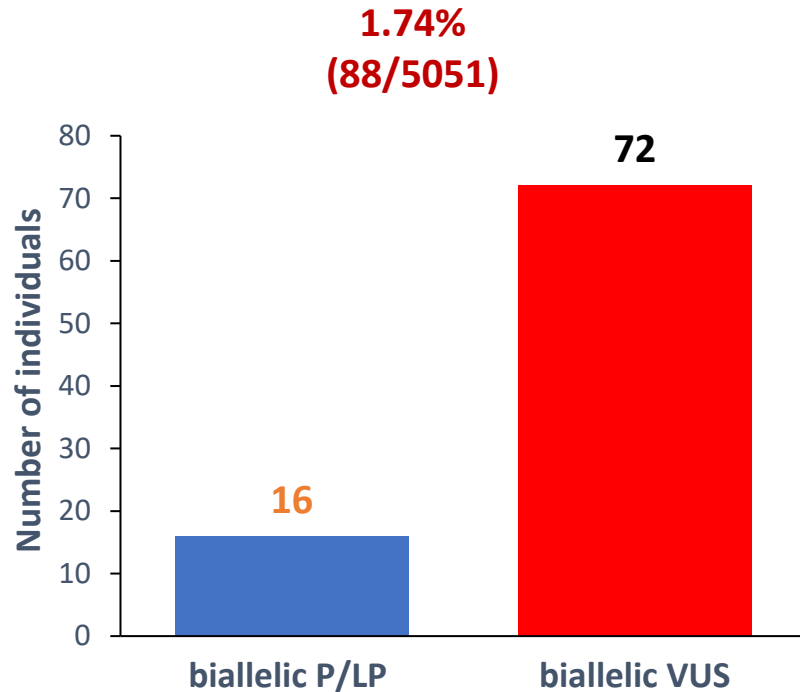


BBS, Bardet-Biedl syndrome; BMI, body mass index; P/LP, pathogenic/likely pathogenic; SD, standard deviation; VUS, variant of unknown significance.

1. Forsythe E, Beales PL. *Eur J Hum Genet.* 2013;21:8-13.

1.74% of tested individuals carried biallelic variants in one of 22 tested BBS genes; BBS was not suspected in most cases

Yield according to BBS variant type (n=88)



Higher frequency of *BBS14* (n=22) and *BBS20* (n=11) variants versus literature

Hyperphagia and weight parameters, mean (SD)

Age of hyperphagia onset



3.1 (2.8) years

Age of obesity onset



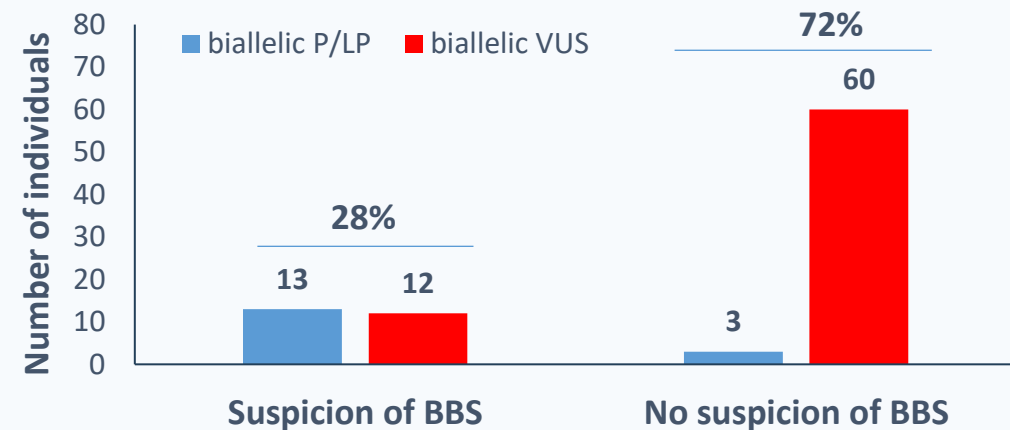
4.8 (3.7) years

BMI (≥18 years)
BMI z-score (<18 years)



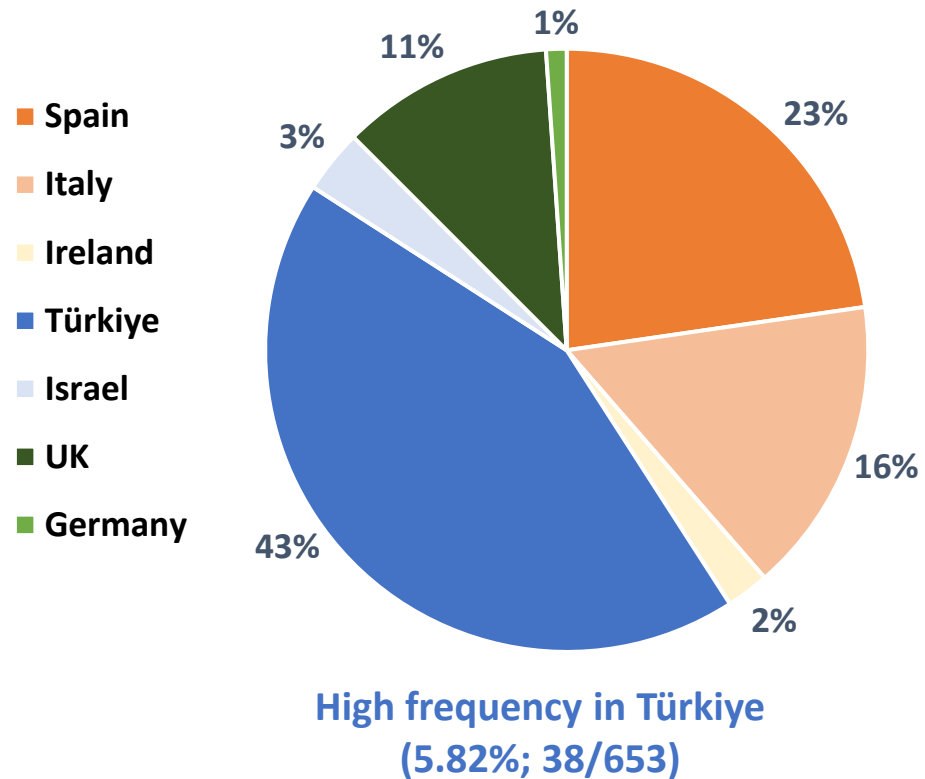
BMI: 41.6 (9.5) kg/m²
BMI z-score: 3.8 (1.2)

Number of patients with or without a suspicion of BBS at testing

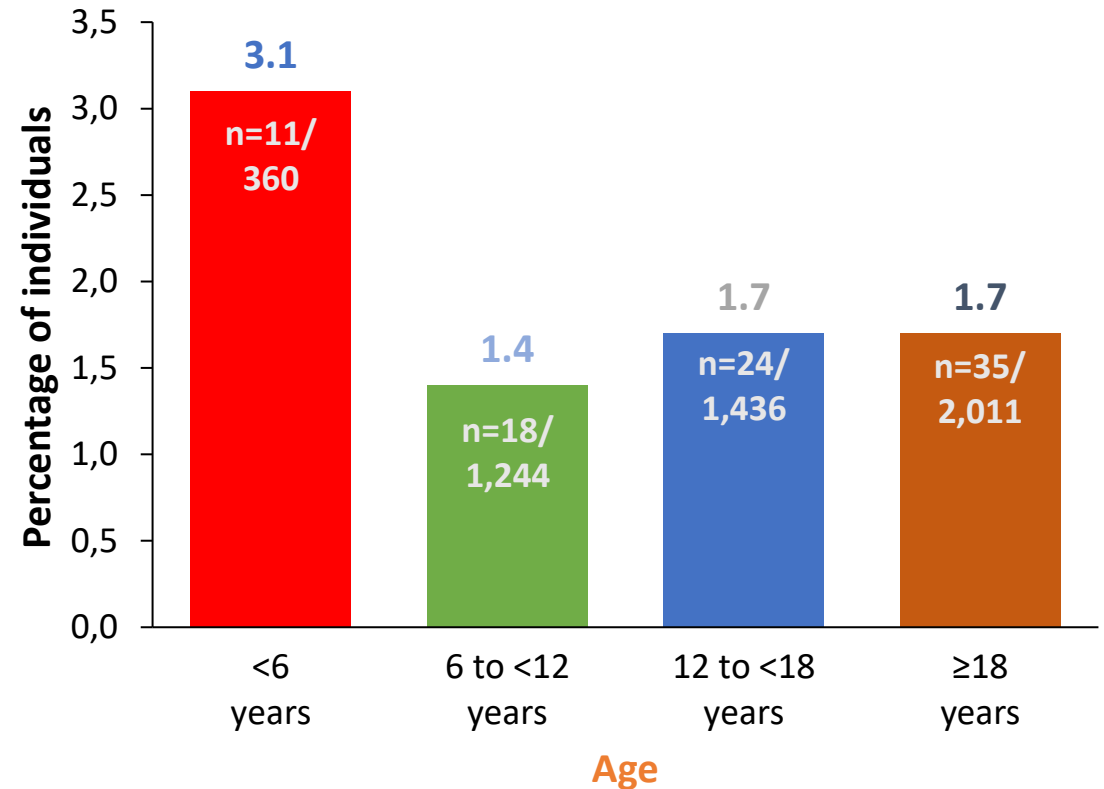


BBS diagnosis in patients <6 years of age

Percentage of individuals with BBS variants across locations (n = 88)



Percentage of individuals with BBS variants by age subgroups of the full cohort (n = 88/5,051)



Summary and conclusions

- The ROAD® testing program encourages and supports timely diagnosis by physicians of individuals with suspected rare genetic causes of obesity in key genes of the MC4R pathway
- Among individuals tested, **1.74%** (88/5,051) carried a biallelic variant in one of 22 tested BBS genes
- **Frequency in Türkiye (5.82%) was highest across locations**
- At diagnosis, not all patients had a clear BBS phenotype indicating that all BBS genes should be included in panels testing for early-onset obesity, not just those investigating suspected syndromic disease

Genetic testing of individuals with early-onset obesity can help improve disease etiology understanding and identify patients who could potentially benefit from specialized care