Supplementary Data 1. Incidence of clinical features typical for deletion 22q11.2 syndrome in 41 patients enrolled in the study.

 $F - \underline{F}$ emale, $M - \underline{M}$ ale, $R - \underline{R}$ ight, $L - \underline{L}$ eft, $B - \underline{B}$ ilateral, $N - \underline{M}$ ale, $N - \underline{M}$ a

		Con	genital l defect	neart		Craniofacial anomalies											Other anomalies								
Patient	Gender	Heart defect, icluding:	* Conotruncal anomaly	* Fallot's tetralogy	Micro(retro)gnathia	Long nose with broad bridge	Unusually shaped ears	Long slender fingers and hands	Short palpebral fissures	Facial asymmetry	Cleft palate	Velopharyngeal insufficiency	Swallowing difficulty	Learning difficulties / developemtal delay	Epilepsy	Psychiatric disorder	Immunodefficiency / thymic hypoplasia	Hypocalcaemia	Renal abnormalities / Potter sequence	Short stature	Family history				
1	M	+			+		+		+					+	+					+					
2	M	+			+			+					+	+		+									
3	M				+		+							+											
4	M	+	+																						
5	M	+	+	+								+	+						R						
6	M	+				+	+							+											
7	F	+	+	+		+		+					+	+						+					
8	F	+								+															
9	M	+	+			+	+							+											
10	F	+	+	+	+	+								+											
11	F	+	+			+																			
12	M	+			+			+					+	+		+									
13	F	+			+								+	+						+					
14	F	+	+		+		+							+											
15	M				+	+	+		+		+														
16	M	+	+			+	+				+			+						+					
17	F	+	+	+																					
18	F	+											+	+	+				В						

Genomic findings in patients with clinical suspicion of 22q11.2 deletion syndrome.

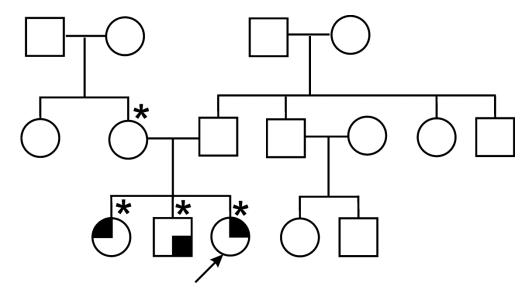
Magdalena Koczkowska, Jolanta Wierzba, Robert Smigiel, Maria Sasiadek, Magdalena Cabala, Ryszard Slęzak, Mariola Iliszko, Iwona Kardas, Janusz Limon, Beata S. Lipska-Zietkiewicz.

19	F	+	+			+	+							+							
20	F	+				+		+		+		+									
21	M	+	+	+		+	+		+	+			+								
22	F	+	+					+					+			+			+		
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24	M	+	+											+			+				l
25	F	+			+			+			+		+	+							
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29	F	+												+	+	+					
30	F	+	+		+		+				+		+	+			nd				
31	M	+	+	+		+															
32	M	+			+		+							+							
33	M	+	+			+	+										+				
34	M	+			+		+		+		+		+	+					В	+	
35	F	+	+		+	+	+		+	+	+			+							<u> </u>
36	M	+			+	+	+		+	+									L		<u> </u>
37	M	+			+	+	+		+	+										+	
38	F	+			+				+				+								—
39	M	+	+														nd	nd	+		igsquare
40	F	+				+					+		+				nd	nd			
41	F				+				+					+							+

Supplementary Data 2.

Co-segregation of the unique 0.65Mb deletion at chromosome 3q26 in Family #41.

A. Family pedigree.



Legend:

* - carriers of arr [hg18] 3q26.31(176,570,887-177,223,154)x1

B. Clinical data.

Affected family members:

Older sister (aged 17) – IUGR, bilateral cleft lip and cleft palate, short stature, congenital hypothyroidism, strabismus (exotropia), normal motor and psychosocial development, recurrent infections; dysmorphic features: long face, deep-set eyes, long nose with broad bridge, short philtrum, full triangle-shaped lips, long slender fingers of hands and feet.

Older brother (aged 16) – IUGR, hemodynamically significant congenital heart defect (VSD+PDA; surgical correction in infancy); normal motor and psychosocial development, dysmorphic features: long face, hypotelorism, long nose with broad bridge, short philtrum, small lips, long slender fingers of hands and feet.

Patient #41 (aged 8) – IUGR, no congenital defects, normal motor development, severe intellectual disability, autism spectrum disorder, dysmorphic features: long face, short palpebral fissures, long nose with broad bridge, short philtrum, micrognathia, long slender fingers of hands and feet.

Mother (aged 39) – no perinatal data, no congenital defects, normal motor and psychosocial development, dysmorphic features: long face, hypotelorism, long nose with broad bridge, short philtrum.

Ethnic background: The family is Kashubian, i.e. comes from a small inbred population of Pomerania region of Poland. Even though the parents negate being related, knowledge of the history of the region suggest relatively high possibility of consanguinity.