Complications and Outcomes of Pregnancies Screening Positive for Microdeletions 22q11.2, 15q11.2, 1p36, 4p, or 5p

Myriad genetics

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Background

 Pregnancy complication types and rates have not been well described among those who screened positive for certain microdeletions.

OBJECTIVE: We aimed to characterize ultrasound findings and pregnancy complications and outcomes in pregnancies screening positive for microdeletions 22q11.2, 15q11.2, 1p36, 4p, or 5p on a prenatal cell-free DNA (pcfDNA) test using insurance claims data.

Methods

- Patients who received Prequel¹ (Myriad Genetics, Inc.), a whole genome sequencing-based pcfDNA screen that incorporates fetal fraction amplification, between July 2020 and January 2023 were linked to the Komodo Healthcare Map™ insurance claims database² using Datavant tokenization³ (Figure 1).
- Included patients were ≥18 years with singleton pregnancies and no other known abnormal pcfDNA screening results (Figure 1).
- Diagnosis codes, procedure codes, and pharmacy fills during pregnancy were used to assess outcomes.
- Outcomes of microdeletion screen-positive (MDS+) patients were compared to microdeletion screen-negative (MDS-) patients using logistic regression with adjustment for health insurance type and history of high-risk pregnancy.

Conclusion

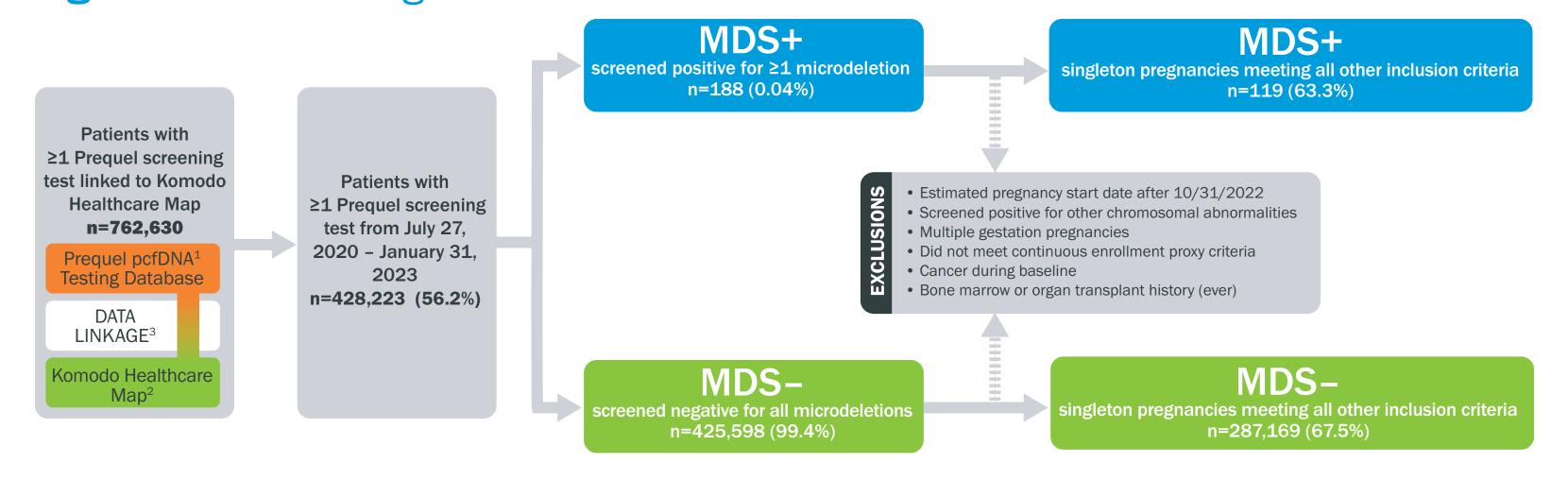
Microdeletion screen-positive pregnancies had higher rates of ultrasound abnormalities, including polyhydramnios and intrauterine growth restriction, and experienced higher rates of pregnancy loss and preterm birth compared to screen-negative pregnancies.

References: 1. Myriad Genetics, Inc., Salt Lake City, UT. Prequel[®] Prenatal Screen. https://myriad.com/womens-health/patient-prequel/. **2.** Komodo Health, Inc. https://www.komodohealth.com/healthcare-map. **3.** Datavant. https://www.datavant.com/.

Disclosures: All authors were employees of Myriad Genetics, Inc. at the time of this study and received salaries and stock as compensation.

• The study population included 119 MDS+ patients and 287,169 MDS- patients (Figure 1; Table 1); the frequency of each microdeletion is shown in Figure 2.

Figure 1. Cohort diagram



Other chromosomal abnormalities included monosomy or trisomy results for any autosomal chromosomes (1-22) or any sex chromosome abnormalities. In the MDS+ group, patients screening positive for >1 microdeletion were also excluded. The continuous enrollment proxy required: 1) at least one medical or pharmacy claim from 2 years prior to the estimated pregnancy start date (baseline period) to 8 weeks after the estimated pregnancy start date, 2) at least one medical or pharmacy claim +/- 4 weeks around Prequel test date, and 3) at least one pregnancy-related claim from the estimated pregnancy start date to 42 weeks after the estimated pregnancy start date (follow-up period). This was to ensure most claims occurring during the study period were captured.

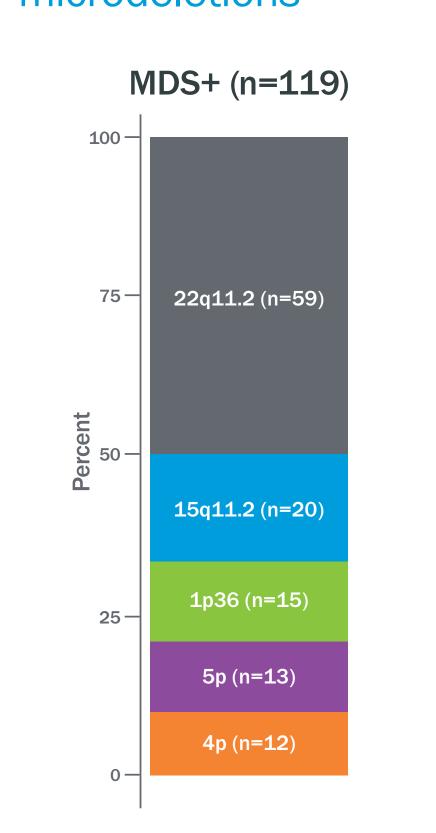
Table 1. Cohort characteristics

Characteristics	MDS-	MDS+
Total N	287,169	119
Maternal age at expected delivery (years)		
Mean (SD)	30.9 (5.8)	30.9 (5.6)
Median (Q1 - Q3)	31.0 (27.0 – 35.0)	32.0 (26.5 - 35.0)
Age ≥35 (n, %)	201,650 (29.4)	34 (28.6)
Payer, n (%)	,	,
Commercial	200,493 (69.8)	>59 (>49.6)
Medicaid	80,742 (28.1)	38 (31.9)
Medicare	1,690 (0.6)	<11 (<9.2)
Other	4,244 (1.5)	<11 (<9.2)
Ethnicity, n (%)	, , ,	,
African or African American	35,868 (12.5)	15 (12.6)
Ashkenazi Jewish	2,386 (0.8)	0 (0)
Asian/Pacific Islander	13,430 (4.7)	<11 (<9.2)
European	28,420 (9.9)	12 (10.1)
French Canadian or Cajun	376 (0.1)	0 (0)
Hispanic	32,529 (11.3)	18 (15.1)
Middle Eastern	2,197 (0.8)	<11 (<9.2)
Native American	1,304 (0.5)	0 (0)
Other/Mixed White	100,811 (35.1)	>31 (>26.1)
Unknown	69,848 (24.3)	21 (17.6)
Pregnancy history, n (%)		
Previous pregnancy	157,750 (54.9)	67 (56.3)
Pregnancy loss	49,506 (31.4)	20 (29.9)
Elective termination	6,110 (3.9)	0 (0)
Preeclampsia	10,647 (6.7)	<11 (<16.4)
Gestational diabetes	12,291 (7.8)	<11 (<16.4)
High-risk pregnancy	133,101 (84.4)	63 (94.0)
IUGR	16,370 (10.4)	11 (16.4)
Previous delivery	110,551 (38.5)	56 (47.1)
Preterm birth	7,917 (7.2)	<11 (<19.6)
Gestational age at pcfDNA testing (weeks)		
Mean (SD)	13.0 (3.7)	14.3 (5.0)
Median (Q1 - Q3)	12.0 (10.9 - 13.4)	12.6 (11.0 - 15.3)
Include microdeletion report results, n (%)	38,960 (13.6)	38 (31.9)

IUGR = intrauterine growth restriction; Q = quartile; SD = standard deviation. Patient numbers <11 were masked to preserve patient privacy.

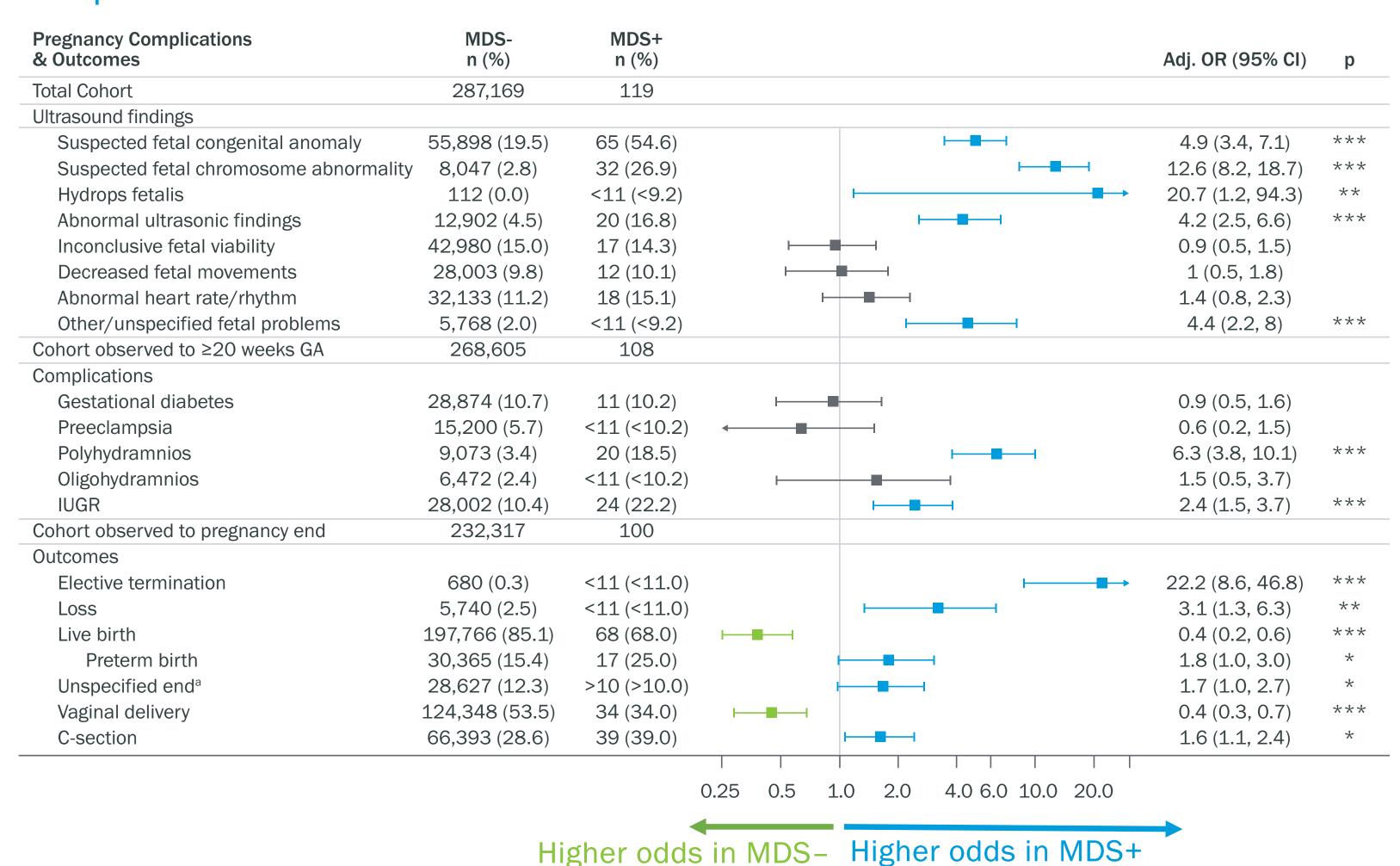
Figure 2. Frequency of microdeletions

Results



- During pregnancy, MDS+ patients were more likely to have abnormal ultrasound findings including polyhydramnios and intrauterine growth restriction compared to MDS- patients, but did not have higher odds of gestational diabetes or preeclampsia (**Figure 3**).
- Among patients with observed pregnancy outcomes, MDS+ patients were significantly more likely to experience pregnancy loss, undergo pregnancy termination, and deliver by caesarean section than MDS- patients.
- Among patients with a live birth, MDS+ patients were more likely to have a preterm delivery.

Figure 3. Association between microdeletion screen status and pregnancy complications and outcomes



Adj. OR=adjusted odds ratio; Cl=confidence interval; IUGR=intrauterine growth restriction. *Indicates p<0.05, ** indicates p<0.01, *** indicates p<0.001. Odds ratios adjusted for payer type and history of high-risk pregnancy. Cell sizes <11 masked to protect patient privacy. Blue ORs and Cls correspond to outcomes with significantly higher odds in MDS+, green ORs and Cls correspond to outcomes with significantly different odds for either group. a. Pregnancies with a delivery code indicating an end to the pregnancy, but without specific evidence of an outcome such as live birth or loss.