Whose Y is it anyway? Transplantation as a biological cause of noninvasive prenatal testing (NIPT) gender discrepancies

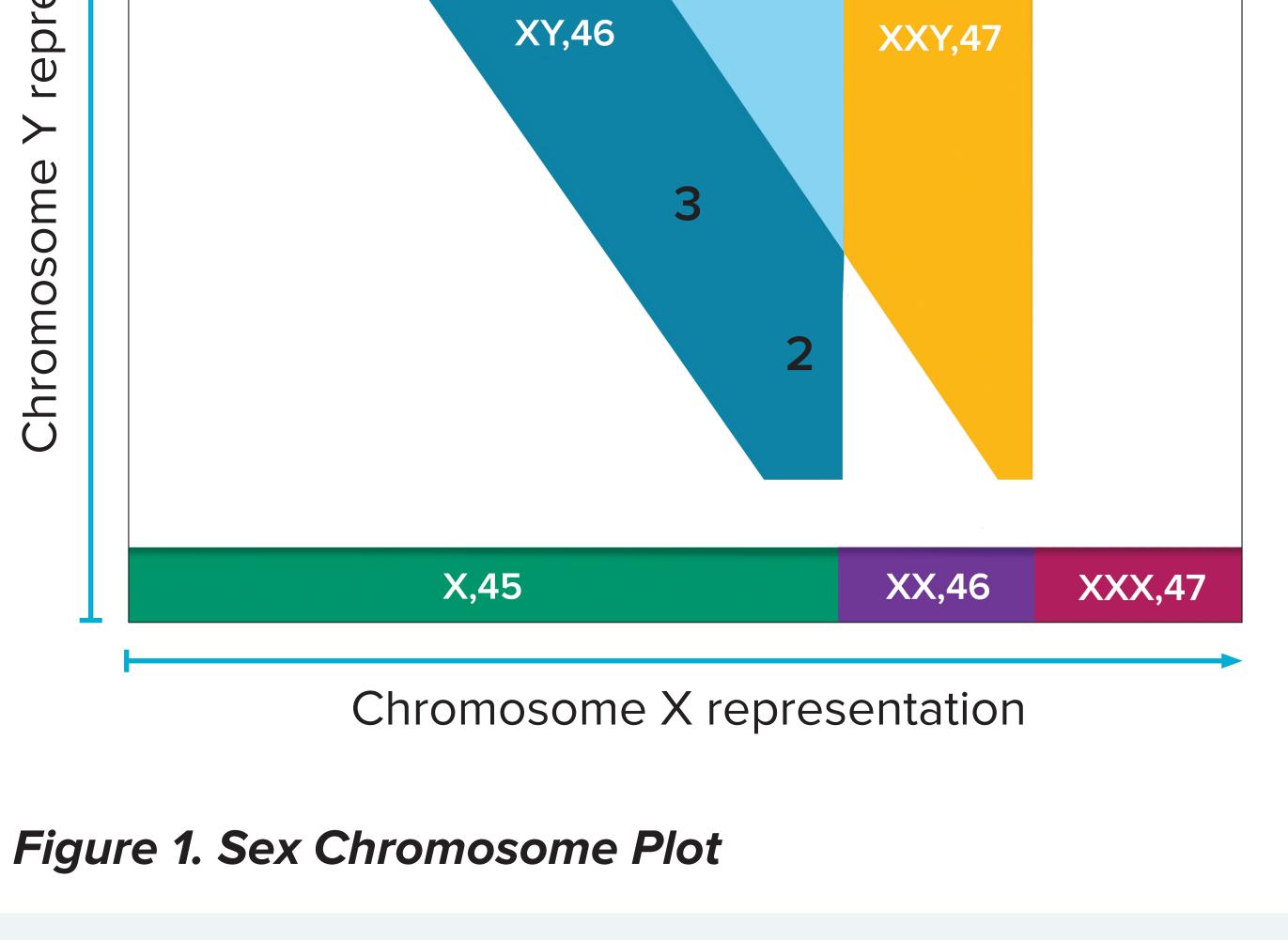
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INTRODUCTION	CASES	
Since the introduction of noninvasive	CASE #1, NIPT result:	
prenatal testing (NIPT) in 2011, more than	Male, negative for aneuploidies.	
400,000 clinical samples have been run	Provider informed us that ultrasound	Justice of the second sec
in Sequenom Laboratories. Discrepant	was consistent with a female fetus. 2^{nd}	NYY,47

biological exresults often aliquot of sample was run. Both samples have planations, including confined placental showed a strong Y signal. Provider later mosaicism, maternal mosaicism, co-twin informed the lab that the patient had a demise, and maternal neoplasm. As cell-free DNA is comprised of maternal and trophoblast placental DNA, tissue from foreign organs can contribute its DNA to the pool. Here we describe three cases of NIPT

liver transplant due to Wilson's disease 16 years prior, from a male donor. CASE#2, NIPT result: Male, aneuploidies. negative for Provider informed us that ultrasound was consistent with a female fetus. Provider later informed the lab that the patient had a bone marrow transplant



CONCLUSIONS

fetal sex discrepancies in patients who had undergone bone marrow and liver transplants.

METHODS

from her brother in '85. Normal female anatomy at birth.

Obtaining a detailed clinical history in cases of NIPT discrepancy can provide unexpected and valuable clues toward resolution and understanding. As NIPT becomes increasingly available to all

populations, pre-test counseling about maternal conditions is essential.

Maternal blood samples submitted to Sequenom Laboratories for MaterniT21[®] PLUS testing were subjected to DNA

trisomies and select microdeletions.²

Male, aneuploidies. negative for

Provider informed us that ultrasound

CASE #3, NIPT result:

was consistent with a female fetus and

extraction, library preparation, and that the patient had a bone marrow transplant from her brother in '02 due parallel massively whole genome sequencing as described by Jensen to aplastic anemia. In reviewing the et al.¹ Sequencing data were analyzed data, the sample showed a strong using a novel algorithm to detect Y signal.

REFERENCES

1. Jensen TJ¹, Zwiefelhofer T, Tim RC, et al. High-throughput massively parallel sequencing for fetal aneuploidy detection from maternal plasma. PLoS One. 2013;8(3):e57381. doi: 10.1371/journal.pone.0057381. Epub 2013 Mar 6.

2. Zhao C, Deciu C, Ehrich M, et al. Detection of fetal subchromosomal abnormalities by sequencing circulating cell-free DNA from maternal plasma. PLoSone. In press.

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