Noninvasive screening tools for Down syndrome: a review

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Abstract: Down syndrome is the leading cause of prenatal chromosome abnormalities, accounting for 53% of all reported chromosome conditions. Testing strategies, guidelines, and screening options have expanded from their conception in the 1970s, and now include such options as anatomical ultrasound, maternal serum screening, and noninvasive prenatal testing. This review summarizes all currently available noninvasive diagnostic techniques for the detection of Down syndrome. By understanding fully each technology and the possible alternatives, the physician will be able to provide their patients with all the information necessary to make an informed decision regarding their medical management.

Keywords: Down syndrome, noninvasive screening, diagnostic techniques

Introduction
It has been roughly half a century since Lejeune et al first described Down syndrome in 1959. While the technology employed for prenatal detection of Down syndrome has expanded in leaps and bounds, the primary focus of prenatal care remains the same: to offer women the most thorough risk assessment with the least invasive procedure possible. Down syndrome is the leading cause of prenatal chromosome abnormalities, accounting for 53% of all reported chromosome conditions.

Testing strategies, guidelines, and screening options have expanded from their conception in the 1970s. At that time, any woman aged 35 years or older was considered to be of advanced maternal age, and this was the sole criteria used by the American Congress of Obstetricians and Gynecologists to define pregnancies that should be offered amniocentesis or chorionic villus sampling. As of 2007, the American Congress of Obstetricians and Gynecologists has defined a pregnancy as “high-risk” when any of the following criteria are met: family history of aneuploidy, advanced maternal age, abnormal serum screen, or abnormal ultrasound findings. While multiple screening options are currently available, the only diagnostic tests offered prenatally for Down syndrome are amniocentesis and chorionic villus sampling. This review compares the risks, benefits, and limitations of all currently available prenatal screening methods for detection of Down syndrome.

Established techniques
Anatomical ultrasound
Anatomical ultrasound has been used since the 1980s to provide health care practitioners and expectant mothers with information regarding a pregnancy. With advances in
Maternal age, nuchal translucency ultrasonography, and measurement of maternal serum free beta human chorionic gonadotropin and pregnancy-associated plasma protein A, have been used for screening for Down syndrome. First trimester screening incorporates maternal age-related risk and four maternal serum biomarkers, ie, alpha-fetoprotein, free beta human chorionic gonadotropin, unconjugated estriol, and dimeric inhibin A. Second trimester screening incorporates maternal age, nuchal translucency, and changes detected at ultrasound scan which may be transient and in themselves have little or no pathological significance, but are thought to be more commonly found in fetuses with congenital abnormalities, particularly karyotypic abnormalities. Soft markers that have been linked to Down syndrome include nuchal thickening, echogenic intracardiac focus, echogenic bowel, renal pelvic dilation, shortened long bones, absence of the nasal bone, polydactyly, ventriculomegaly, clinodactyly, and sandal gap toe. In the absence of soft markers, the sensitivity of anatomical ultrasound to detect Down syndrome is relatively low at 50%. However, the presence of one soft marker is associated with an increased risk of Down syndrome, ie, one soft marker increases the risk by two-fold and three or more soft markers increases the risk by 100-fold. Many scoring indices have been created to help maximize sensitivity, while decreasing false-positive rates. These indices incorporate the presence of structural abnormalities and/or soft markers and maternal age to provide physicians with a guideline as to who should be offered more invasive procedures. While these guidelines have assisted in determining the criteria for a positive finding, they remain limited by the quality of the ultrasound and the expertise of the sonographer.

### Maternal serum screening

Beginning in 1984, multiple marker screening provided physicians with a means of offering an individualized risk for Down syndrome without the inherent risk imposed by chorionic villus sampling or amniocentesis (Table 1). This second trimester screening, performed at 15–20 weeks gestation, is often referred to as the quad screen because it incorporates maternal age-related risk and four maternal serum biomarkers, ie, alpha-fetoprotein, free beta human chorionic gonadotropin, unconjugated estriol, and dimeric inhibin A levels. By combining maternal age with the quad screen, the detection rate is roughly 75% for Down syndrome in women younger than 35 years and >80% in women 35 years and older (with a positive screening rate of 5%).

It was not until the late 1990s that first trimester screening was introduced as an earlier screening option for the detection of Down syndrome. First trimester screening incorporates maternal age, nuchal translucency ultrasonography, and

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Note: Table 1 depicts the sensitivity and false positive rates of all serum screening for the detection of Trisomy 21. Adapted based on information obtained from references 9, 12, and 13.
the detection rate from 88% 91% to 94%, it also increases the false-positive rate from 5% to 11%. Thus, it is recommended that women who undergo first-trimester screening for aneuploidy should not also have second-trimester serum screening in the same pregnancy. If a higher detection rate is preferred, an integrated or sequential screening test which combines both first-trimester and second-trimester screening is suggested.

Integrated screening is defined as the process by which a patient’s individualized risk is calculated based on the combination of both the first-trimester and second-trimester screenings. Due to the nature of this screening method, a patient’s first-trimester screening results are not disclosed until second-trimester screening is performed and a combined risk based on both screenings can be calculated. There has been ethical debate regarding integrated screening because the patient’s first-trimester screening results are not disclosed until the conclusion of the second-trimester screening. This precludes patients who are at high risk based on first-trimester screening from being offered chorionic villus sampling and having more options available to them.

In contrast, patients undergoing stepwise sequential screening are provided with their individual risk once the results from the first-trimester screen are available. Patients who screen positive during first-trimester screening are offered genetic counseling and are given the information regarding chorionic villus sampling and amniocentesis. Patients who screen negative are offered second-trimester screening and are provided with an adjusted risk number that incorporates both the first-trimester and second-trimester results. Incorporation of both the first-trimester and second-trimester screenings in this stepwise manner increases the detection rate to 94%–96% while still maintaining a low false-positive rate of 5%–6%.

The final type of sequential screening is known as the contingent screening method. This method is similar to stepwise sequential screening in that scores are calculated based on results from both the first-trimester and second-trimester screening. However, the contingent screening method uses the first-trimester results to classify patients into three subgroups, i.e., screen-positive, screen-negative, and borderline. Second-trimester screening is only offered to patients who fall into either the screen-negative or the borderline group. The detection rate for this method is 94%–95%, with a false-positive rate of about 5%.

**Noninvasive prenatal testing**

It has been known since the mid 1950s that fetal cells are present in the maternal circulation. However, low yield (1 fetal cell/mL of maternal blood), inability to develop an efficient enrichment process, and the tendency for fetal DNA to disintegrate during chromosome extraction have inhibited utilization of these fetal cells for the development of a noninvasive prenatal test. Use of noninvasive prenatal testing targeted to fetal cells was finally abandoned when Bianchi et al demonstrated that fetal cells could remain in the maternal circulation for decades, thus making it impossible to distinguish new fetal cells from those of previous pregnancies.

It was not until 1997 when Lo et al demonstrated the existence of cell-free fetal DNA in the maternal circulation that the scientific community was presented with a new possible target for noninvasive prenatal testing. Fetal cells, theorized to be derived from the placenta, enter the maternal circulation where they undergo apoptosis. In the process, fetal DNA is cleaved into small 150–200 base pairs fragments, which are then released into the maternal bloodstream. These cell-free fetal DNA fragments can be detected as early as the fourth week of gestation and reliably after the seventh week of gestation. Unlike the longevity seen with fetal cells, cell-free fetal DNA has a half-life of only 16 minutes and is cleared from the maternal circulation within 2 hours of delivery. The low false-positive rate as well as the prevalence of cell-free fetal DNA in the maternal circulation (accounting for >10% of all cell-free DNA or 16–80 fetal genomes/mL of maternal blood), made cell-free fetal DNA a desirable target for noninvasive prenatal testing.

Over the next 15 years, scientists would work on developing and refining methods of detecting pregnancies at risk for aneuploidy. In October, 2011, Sequenom Inc (San Diego, CA, USA) was the first company to make a noninvasive prenatal test commercially available for the detection of Down syndrome. To date, three companies (Verinata Health [Redwood, CA, USA], Ariosa Diagnostics [San Jose, CA, USA], and Sequenom Inc) offer a noninvasive prenatal test for the detection of trisomies 13 and 18 and Down syndrome, with a fourth company (Natera, San Carlo, CA, USA) expected to have a commercially available noninvasive prenatal test within the next few months (Figure 1).

Figure 1 describes the various strategies employed by the four companies to obtain and analyze maternal and/or fetal DNA from the maternal serum sample. Two companies, Sequenom Inc and Verinata Health, utilize massively parallel shotgun sequencing, commonly referred to as shotgun sequencing while the other two companies, Natera and Ariosa Diagnostics, utilize targeted sequence analysis.
The main difference between these two technologies is that shotgun sequencing results in amplification of all genetic information, while targeted sequencing results in amplification of only the genetic information of interest, ie, chromosome(s). Targeted sequencing enables companies such as Natera and Ariosa Diagnostics to perform noninvasive prenatal testing on smaller sample sizes compared with Sequenom Inc and Verinata Health. Massively parallel shotgun sequencing technology requires more “reads” (or DNA fragments) than those required by targeted sequencing to ensure that there are enough fragments from the chromosomes of interest to provide accurate results. A major disadvantage of targeted sequencing is the inability to rule out other chromosomal conditions, such as microduplications and deletions, because they have not been selected for. While this does not seem to be a major limiting factor at this time, due to the fact that companies are not currently reporting nonvalidated findings, it may become a shortcoming in the future as companies aim to expand noninvasive prenatal testing to include other chromosomal abnormalities, such as microdeletions and microduplications.

All four companies currently use variations of two methodologies for analyzing genetic data, ie, quantitative read counting or single nucleotide polymorphism detection. Three of the companies (Sequenom Inc, Verinata Health, and Ariosa Diagnostics) use quantitative read counting, while Natera uses bioinformatic algorithms to analyze single nucleotide polymorphism data obtained by next-generation aneuploidy testing using single nucleotide polymorphisms.

Quantitative read counting analyzes the number of chromosome fragments present from each chromosome or chromosomes of interest. Because the amount of genetic material from each chromosome is directly proportional to the chromosome size, companies can use a known euploid reference sample to calculate the expected proportion of genetic information from each chromosome given a euploid pregnancy.31,32,37–39 For example, chromosome 21 generally accounts for 1.5% of the human genome.31,37–39 Any change in the actual proportions of genetic information from chromosome 21 that is 2.5–3.1 standard deviations from the mean is determined to be an aneuploidy.30,31,40 The benefit of this technology is that it does not require any differentiation between maternal and fetal genetic information. However, this is also a limitation resulting from the fact that, in a fetus with Down syndrome, the contribution of genetic information from chromosome 21 increases from 1.5% to 2.25%, for an overall change of 0.75%. When taking into account the fact that fetal DNA only represents 10% of the total DNA in the sample, the overall change in the amount of chromosome 21 cell-free-DNA would only increase from 1.5% to 1.575%. When the fetal fraction is below 10%, it can result
in inconclusive results, or a no-call. One company using quantitative read counting, Ariosa Diagnostics, has aimed to address this issue by creating the FORTE (fraction optimized risk of trisomy evaluation) assay.\textsuperscript{33,34} This assay utilizes polymorphic and nonpolymorphic regions that are known to differ between fetal and maternal DNA to determine the fetal fraction as well as the overall proportional representation of chromosomal fragments.

Unlike quantitative read counting, the next-generation aneuploidy testing using single nucleotide polymorphisms system created by Natera utilizes the Parental Support\textsuperscript{TM} statistical algorithm to analyze the sequenced fetal cell-free DNA together with maternal genetic information, paternal genetic information, and HapMap technology.\textsuperscript{35,36} This algorithm provides a series of possible “hypotheses” of fetal genotypes (eg, monosomy, disomy, trisomy) based on known common crossover points on the chromosome(s) of interest, parental information, fetal fraction, and fetal chromosome copy number. The sample-specific confidence interval for each hypothesis is calculated, and a “hypothesis” is considered confirmed when the confidence interval is above 98%.\textsuperscript{36}

While the results seen from these four companies are promising (Table 2), it is important to remember that noninvasive prenatal testing is for screening and is not a diagnostic tool.\textsuperscript{30,32,35,36,41} Noninvasive prenatal test results group patients into three possible categories, ie, low risk of aneuploidy, high risk of aneuploidy, or no-call (undeterminable). The clinical implications for low-risk and high-risk populations are the same for noninvasive prenatal testing as they are for first-trimester screening and multiple marker screening. Individuals who are determined to be in the low-risk category would not require any further diagnostic evaluation, while those in the high-risk category are recommended to undergo diagnostic testing (ie, chorionic villus sampling or amniocentesis). Individuals who fall into the third category of “no-call” would require a repeat sample to be drawn. For these reasons, noninvasive prenatal testing is not intended to serve as a replacement for amniocentesis or chorionic villus sampling, but rather a methodology that would enable fewer unnecessary diagnostic procedures by detecting patients who are at high risk for aneuploidy more accurately, when compared with the detection rates for first-trimester screening and multiple marker screening. Physicians should ensure that their patients understand that these tests are not designed to circumvent an unwanted diagnostic procedure, and in actuality may result in the recommendation of such a diagnostic procedure should test results reveal a high risk for aneuploidy.

While noninvasive prenatal tests do have an increased specificity and sensitivity compared with first-trimester screening or multiple marker screening, limitations do exist. False-positive results may be present. Possible causes for false-positive results could be placental mosaicism, vanishing twin syndrome, or an unidentified maternal condition, such as mosaicism or cancer. Secondly, the majority of noninvasive prenatal tests are currently only offered to women with singleton pregnancies who are at high risk for Down syndrome because of family history of aneuploidy, advanced maternal age, an abnormal serum screen, or abnormal ultrasound findings. With the exception of Ariosa Diagnostics, which has conducted validation studies for Harmony\textsuperscript{®} in the low-risk population, it is important to note that these tests have not been validated in low-risk or multiple gestation populations, and their accuracy is unknown at this time.\textsuperscript{42}

**Future directions**

It has long been a goal of prenatal genetic diagnosis to develop a noninvasive test that would allow for detection of aneuploidy and eliminate the need for invasive testing, such as chorionic villus sampling or amniocentesis. While this is still not currently possible, developments in noninvasive

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**Notes:** Table 2 depicts the outcome from the validation studies and current no call rates from the four commercially developed NIPTs. \textsuperscript{c} denotes a no call rate that incorporates all aneuploidies analyzed by Natera (Trisomy 13, 18, 21, and sex chromosome abnormalities). This data was adapted based on information obtained from references 31, 32, 36, 37, and 42.
prenatal testing are the initial steps in this direction. While the scope of this review does not include other aneuploidy conditions, all of the companies discussed do offer screening for trisomy 13 and 18, with varying degrees of success. The noninvasive prenatal tests marketed by Verinata Health and Natera also include a screen for sex chromosome abnormalities, such as Turner syndrome, Kleinfelter syndrome, and 47, XYY syndrome. Validation studies are currently underway for noninvasive prenatal testing in the detection of fetal aneuploidies in the general population as well as multiple gestations. Furthermore, the future of noninvasive prenatal testing may expand beyond the common aneuploidies to include other chromosomal conditions, such as microdeletions and microduplications. With the ever-expanding testing options that are now available to the expectant mother, it is imperative that physicians remain up to date on these technologies and have a clear understanding of the risks, benefits, and limitations of these technologies. By fully understanding each technology and the possible alternatives, physicians can provide their patients with all the information necessary to make an informed decision regarding medical management.

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Disclosure
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References


