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| Case No. | Detailed clinical indicator(s)＃: | CNV-seq results | Classification | Follow-up |
| 38 | Fetal congenital heart disease | seq[GRCh38]del(22)(q11.21)  NC\_000022.11:g.18892487\_21465711del | Pathogenic | TOP |
| 39 | Maternal serum screening high risk | seq[GRCh38]del(5)(p15.33p15.1)  NC\_000005.10:g.20001\_17939891del | Pathogenic | TOP |
| seq[GRCh38]dup(7)(q34q36.3)  NC\_000007.14:g.141680201\_159335973dup | Pathogenic |
| 40 | Chromosomal abnormalities of pregnant woman | seq[GRCh38]del(1)(p36.33p36.32)  NC\_000001.11:g.884621\_2823435del | Pathogenic | TOP |
| 41 | Maternal serum screening high risk | seq[GRCh38]del(8)(p23.3p23.1)  NC\_000008.11:g.210001\_7082478del | Pathogenic | TOP |
| 42 | Maternal serum screening high risk | seq[GRCh38]del(17)(p12)  NC\_000017.11:g.14196684\_15516686del | Pathogenic | Term birth,  no obvious abnormality |
| 43 | Fetal lateral ventricle enlargement; AMA | seq[GRCh38]del(16)(p13.3)  NC\_000016.10:g.29555975\_30178708del | Pathogenic | TOP |
| 44 | Maternal serum screening high risk | seq[GRCh38]del(16)(p11.2)  NC\_000016.10:g.28799003\_29077303del | Pathogenic | TOP |
| 45 | Bowel echo enhancement, fetal congenital heart disease | seq[GRCh38]del(2)(p16.3)  NC\_000002.12:g.50880234\_51125144del | Pathogenic，maternally inherited | TOP |
| 46 | AMA; Single umbilical artery, left kidney absent | seq[GRCh38]del(15)(q11.2)  NC\_000015.10:g.22595660\_23102647del | Pathogenic | TOP |
| 47 | AMA; NIPT high-risk for other chromosome | seq[GRCh38]dup(15)(q11.2q13.1)  NC\_000015.10:g.23374854\_28294854dup | Pathogenic | TOP |
| 48 | Maternal chromosome abnormalities | seq[GRCh38]del(1)(p36.33p36.32)  NC\_000001.11:g.884621\_2823435del | Pathogenic | After birth，  Obvious abnormality |
| 49 | AMA; Chromosomal microdeletion syndrome of previous fetus | seq[GRCh38]del(9)(p24.3p24.1)  NC\_000009.12:g.200000\_6760000del  seq[GRCh38]dup(20)(p13p12.3)  NC\_000020.11:g.79360\_8139353dup | Pathogenic | TOP |
| 50 | Fetal lateral ventricle enlargement | seq[GRCh38]del(16)(p13.3)  NC\_000016.10:g.35880\_147065del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 51 | Maternal serum screening high risk | seq[GRCh38]dup(22)(q11.21)  NC\_000022.11:g.18892488\_21125711dup | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 52 | Induction of labor due to congenital heart disease of previous fetus | seq[GRCh38]del(22)(q11.21) NC\_000022.11:g.20362586\_214461821del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 53 | AMA | seq[GRCh38]del(X)(p22.33p22.32)  NC\_000023.11:g.2781959\_5541959del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 54 | NIPT high-risk for other chromosome | seq[GRCh38]del(15)(q13.2q13.3)  NC\_000015.10:g.30767797\_32147799del | Pathogenic | TOP |
| 55 | NT 2.8mm; Maternal serum screening high risk | seq[GRCh38]del(16)(p13.11p12.3)  NC\_000016.10:g.15426143\_18086143del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 56 | NIPT high-risk for sex chromosome | seq[GRCh38]del(X)(p22.31)  NC\_000023.11:g.6541959\_8171959del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 57 | NIPT high-risk for sex chromosome | seq[GRCh38]del(X)(p22.31)  NC\_000023.11:g.6541959\_8171959del | Pathogenic，maternally inherited | Term birth,  no obvious abnormality |
| 58 | NIPT high-risk for sex chromosome | seq[GRCh38]del(X)(p22.31)  NC\_000023.11:g.6537110\_8167062del | Pathogenic | Term birth,  no obvious abnormality |
| 59 | Maternal serum screening high risk | seq[GRCh38]dup(16)(p13.11)  NC\_000016.10:g.15026143\_16646143dup | Likely pathogenic | Term birth,  no obvious abnormality |
| 60 | NT 5.0mm | seq[GRCh38]del(X)(p22.31p22.2)  NC\_000023.11:g.8422532\_10917281del | Likely pathogenic | Cleft lip, pleural effusion, TOP |
| 61 | FGR | seq[GRCh38]dup(X)(q25q26.1)  NC\_000023.11:g.129446225\_130254419dup | Likely pathogenic | Term birth,  no obvious abnormality |
| 62 | Absence of nasal bone; Pregnancy with gestational diabetes | seq[GRCh38]del(13)(q13.2q13.3)  NC\_000013.11:g.34643644\_35422587del | Likely pathogenic | TOP |
| 63 | Short long bone | seq[GRCh38]dup(2)(q37.3)  NC\_000002.12:g.236551357\_242077849dup | Likely pathogenic | Term birth,  no obvious abnormality |
| 64 | Nasal bone dysplasia; Maternal serum screening high risk | seq[GRCh38]del(20)(p13)  NC\_000020.11:g.79360\_1119357del | Likely pathogenic | Term birth,  no obvious abnormality |
| 65 | Nasal bone dysplasia; Maternal serum screening high risk | seq[GRCh38]dup(16)(p13.11)  NC\_000016.10:g.14946143\_16206143dup | Likely pathogenic | Term birth,  no obvious abnormality |
| 66 | History of bearing child with chromosome abnormalities | seq[GRCh38]del(16)(p12.2)  NC\_000016.10:g.21928680\_22428679del | Likely pathogenic | Term birth,  no obvious abnormality |