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| Date of Report | 04-11-2020 |
|----------------|------------|
| PRISCA | 5.0.2.37 |

| | | | | | PRISCA | 5.0.2.37 | |
|--------------------|-----------|------------|---------|---|--|---|--|
| Patient Data | | | | | | | |
| Name | | MRS SONAM | | | Patient ID | 012011030115 | |
| Birthday | | 04-04-1999 | | | Sample ID | 10735219 | |
| Age at delivery | | 21.9 | | Sample Date | 03/11/2020 | | |
| Correction factors | | | | | | | |
| Fetuses | 1 | IVF | | unknown | Previous trisomy 21 | unknown | |
| Weight in kg | 51.2 | Diabetes | | unknown | Pregnancies | unknown | |
| Smoker | Unknown | Origin | | Asian | | | |
| Biochemical Data | | | | Risks at sampl | ling date | | |
| Parameter | Value | Co | orr MoM | Age Risk | | 1:1496 | |
| AFP | 34.4 n | ıg/ml | 0.41 | Neural Tube I | Defect (NTD) Screening | 1:4880 | |
| uE3 | 2.35 n | ıg/ml | 1.18 | Combined Tri | isomy 21 Risk | 1:1181 | |
| hCG | 19538.3 n | nIU/ml | 1.03 | Trisomy 18 | | <1:10000 | |
| Ultrasound Data | | | | | ome Risk (Trisomy 21 S | | |
| Gestational age | 2 | 21+0 | | The calculated risk for Trisomy 21 is below the cut off which indicates a low risk. | | | |
| Method | (| CRL(<>Robi | inson) | After the resul | After the result of the Trisomy 21 test it is expected | | |
| | | | | 9 | 81 women with the same ith a trisomy 21 pregnanc | - | |
| Risk | | | | women with no | ot affected pregnancies. | | |
| Risk | | | | | l risk by PRISCA depend | ds on the accuracy of the physician. Please note that | |
| 1:10 | | | | _ | rovided by the referring p ations are statistical aappr | • | |
| | | | | diagnostic valu | | | |
| | | | | | | | |
| 1:100 | | | | | | | |
| 1:250 | | | Cut off | Trisomy 18 | | | |
| 1:1000 | 7 | | | The calculated | ulated risk for Trisomy 18 is <1:10000, which | | |

area for neural tube defects.

The laboratory can not be held responsible for their impact on the risk assessment! Calculated value has no diagnostic value!

13 15 17 19 21 23 25 27 29 31 33 35 37 39 41 43 45 47 49

indicates a low risk

Neural Tube Defect (NTD) Screening

The corrected MoM for AFP (0.41) is located in the low risk