



# Colesterol HDL Cepas Direto

## ANALYTICAL PARAMETERS

| Analytical Conditions |          | Reanalysis conditions         |        | Multi-Standards Setting |                  |                 |             |
|-----------------------|----------|-------------------------------|--------|-------------------------|------------------|-----------------|-------------|
| R1 volume             | 120      | Serum reac. Smp. Vol (µL)     | 0      | Formula Points          | Linear correc. 2 | Axis conv.      | No converte |
| R2 volume             | 0        | Serum dilut. method. Vol (µL) | None   |                         |                  |                 |             |
| R3 volume             | 40       | Serum reac. Smp. Vol (d)      | 0      |                         | FV               | MEAN            |             |
| R4 volume             | 0        | Serum dilut. method. Vol (d)  | None   | BLK                     | 0.00             | ...             |             |
| R1 diluent volume     | 0        | Standards setting             |        | 1                       | *                | ---             |             |
| R2 diluent volume     | 0        | BLK H                         | 9.9999 |                         |                  |                 |             |
| R3 diluent volume     | 0        | BLK L                         | 9.9999 |                         |                  |                 |             |
| R4 diluent volume     | 0        | STD H                         | 9.9999 |                         |                  |                 |             |
| Serum reac. s.volume  | 2        | STD L                         | 9.9999 |                         |                  |                 |             |
| Serum diluent method  | Standard | FV                            | 9.9999 |                         |                  |                 |             |
|                       |          | Abnml. (serum) H              | 9.9999 |                         |                  |                 |             |
| Reaction Time         | 5 min.   | Abnml. (serum) L              | 9.9999 |                         |                  |                 |             |
| Reagent 1 stir.       | Weak     | Calculation method settings   |        | Reation rate method     |                  |                 |             |
| Reagent 2 stir.       | Weak     | M-DET. P.l                    | 0      | Prozone                 |                  | Cycle           | 3           |
| Reagent 3 stir.       | Weak     | M-DET. P.m                    | 96     | Prozone form            | None             | Factor          | 3.0         |
| Reagent 4 stir.       | Weak     | M-DET. P.n                    | 98     | Prozone limit           | 9999             | Reac. Type      | INC         |
| Sub-analy. conditions |          | M-DET. P.p                    | 44     | Prozone judge           | Upper Limit      | E2. corre.      | Not dot     |
| Name                  | HDL DIR  | M-DET. P.r                    | 46     | Judge limit             | 9999             | Blank (µL)      | 9.9999      |
| Digits                | 2        |                               |        | M-DET.P.m               | 0                | Blank (d )      | -9.999      |
| M-wave. L             | 596      | Check D.P.l                   | 0      | M-DET.P.n               | 0                | Sample (µL)     | 9.9999      |
| S-wave. L.            |          | Limit value                   | 0.003  | S-DET. P.p              | 0                | Sample (d )     | -9.999      |
| Analy. mthd           | EPA      | Variance                      | 10     | S-DET.P.r               | 0                | Endpoint method |             |
| Calc. mtdh            | STD      |                               |        |                         |                  | Re. Absorb (µL) | 9.9999      |
| Qualit. judg          | Not do   |                               |        |                         |                  | Re. Absorb (d ) | -9.999      |