



شرکت دلتا درمان پارت
سیستم های آزمایشگاهی و مواد مصرفی



دفتر مرکزی: تهران، میدان آرژانتین، خیابان الوند، خیابان سی و پنجم، پلاک ۱۳، طبقه پنجم

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| | | | |
|-------------------------|-------|---------------------|-----------|
| Data Information | | Calibration | |
| Unit | U/l | Type | LINEAR 1 |
| Decimals | 0 | Standard | |
| Analysys | | Blank 0 | 1 * |
| Type | RATE | 2 | 3 |
| W.Length 1 | 340 | 4 | 5 |
| W.Length 2 | 700 | Normal Range | |
| Method | LIA | | |
| Corr | | | |
| Slope | Inter | Low | High |
| 1.000 × + | 0 | Serum Male | 13.3 63.9 |
| | | Serum Female | 13.3 63.9 |
| | | Urine (mg/24h) | |

| | | | | | |
|-------------------|---------------|----------------------|-------|------------------|-------|
| Aspiration | | Data Process | | Abs Limit | |
| Kind | Single Double | Read | Low | -0.1 | |
| | Volume | Start | End | High | 3.000 |
| Sample | 20 | Main | 32 | 72 | |
| Reagent 1 | 200 | Sub | | | |
| Reagent 2 | | Endpoint Limit 2 | | | |
| | | Linear Check (%) | | | |
| Third Mix | Off ON | Factor | | | |
| R1 Blank | Water R1 - B | Blank Correction | 1.000 | | |
| Monitor | | Prozone Check | | | |
| 0 level Point | 1 | Start | End | Limit | |
| Span | 3.000 | First | | | |
| | | Second | | | |
| | | Third | | | |

ACE

| | | | |
|-------------------------|------------|---------------------|----------|
| Data Information | | Calibration | |
| Unit | U/L | Type | LINEAR 1 |
| Decimals | 1 | Standard | |
| Analysys | | Blank 0 | 1 * |
| Type | RATE | 2 | 3 |
| W.Length 1 | 546 | 4 | 5 |
| W.Length 2 | 700 | Normal Range | |
| Method | COLOUR/ENZ | | |
| Corr | | | |
| Slope | Inter | Low | High |
| 1.000 × + | 0 | Serum Male | 0 15 |
| | | Serum Female | 0 15 |
| | | Urine (mg/24h) | |

| | | | | | |
|-------------------|---------------|----------------------|-------|------------------|-------|
| Aspiration | | Data Process | | Abs Limit | |
| Kind | Single Double | Read | Low | -0.1 | |
| | Volume | Start | End | High | 3.000 |
| Sample | 5 | Main | 52 | 76 | |
| Reagent 1 | 160 | Sub | | | |
| Reagent 2 | 80 | Endpoint Limit 3% | | | |
| | | Linear Check (%) | | | |
| Third Mix | Off ON | Factor | | | |
| R1 Blank | Water R1 - B | Blank Correction | 1.000 | | |
| Monitor | | Prozone Check | | | |
| 0 level Point | 1 | Start | End | Limit | |
| Span | 3.000 | First | | | |
| | | Second | | | |
| | | Third | | | |

ADA

| | |
|-------------------------|-------|
| Data Information | |
| Unit | G/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | 700 |
| Method | BCG |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|---|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 3.5 | 5 |
| | Female | 3.5 | 5 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 405 |
| W.Length 2 | 660 |
| Method | DGKC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 80 | 306 |
| | Female | 64 | 306 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-----------------|------------------|----|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 32 | 33 |
| Sub | Start End | | |
| Endpoint Limit 3% | | | |
| Linear Check (%) | | | |
| Factor | | Blank Correction | |
| | | 1.000 | |
| Prozone Check | | | |
| | Start End Limit | | |
| First | Start End Limit | | |
| Second | Start End Limit | | |
| Third | Start End Limit | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input type="checkbox"/> Single <input checked="" type="checkbox"/> Double |
| Volume | |
| Sample | 4 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-----------------|------------------|----|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 44 | 68 |
| Sub | Start End | | |
| Endpoint Limit 3% | | | |
| Linear Check (%) | | | |
| Factor | | Blank Correction | |
| | | 1.000 | |
| Prozone Check | | | |
| | Start End Limit | | |
| First | Start End Limit | | |
| Second | Start End Limit | | |
| Third | Start End Limit | | |

ALB

ALP

| | |
|-------------------------|-------|
| Data Information | |
| Unit | U/L |
| Decimals | 1 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 340 |
| W.Length 2 | 405 |
| Method | IFCC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 40 |
| | Female | 0 | 32 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 405 |
| W.Length 2 | 660 |
| Method | CNPG3 |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 100 |
| | Female | 0 | 100 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> <u>Double</u> <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 20 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 44 | 68 | |
| Sub | | | |
| Endpoint Limit | | 90 | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> <u>Double</u> <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 4 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 24 | 48 | |
| Sub | | | |
| Endpoint Limit | | 3 | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

ALT

AMY

| | |
|-------------------------|-------|
| Data Information | |
| Unit | IU/ML |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 200 |
| | Female | 0 | 200 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | U/L |
| Decimals | 1 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 340 |
| W.Length 2 | 405 |
| Method | IFCC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 37 |
| | Female | 0 | 31 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 3 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start End |
| | 76 77 |
| Sub | |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 20 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start End |
| | 44 68 |
| Sub | |
| Endpoint Limit | |
| Linear Check (90%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

ASO

AST

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 546 |
| W.Length 2 | 700 |
| Method | DMSO |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 0.3 |
| | Female | 0 | 0.3 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 546 |
| W.Length 2 | 700 |
| Method | DMSO |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0.1 | 1.2 |
| | Female | 0.1 | 1.2 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input type="radio"/> Single <input checked="" type="radio"/> Double |
| Volume | |
| Sample | 25 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <input type="radio"/> Off <input type="radio"/> ON |
| R1 Blank | <input type="radio"/> Water <input type="radio"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input type="radio"/> Single <input checked="" type="radio"/> Double |
| Volume | |
| Sample | 6 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <input type="radio"/> Off <input type="radio"/> ON |
| R1 Blank | <input type="radio"/> Water <input type="radio"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

BILI.D

BILI.T

| | | | |
|-------------------------|-------|---------------------|------------|
| Data Information | | Calibration | |
| Unit | MG/DL | Type | Logit.Log2 |
| Decimals | 0 | Standard | |
| Analysys | | Blank 0 | 1 cal/16 |
| Type | END | 2 cal/8 | 3 cal/4 |
| W.Length 1 | 340 | 4 cal/2 | 5 cal |
| W.Length 2 | 700 | Normal Range | |
| Method | LIA | Low | High |
| Corr | | Serum Male | 90 180 |
| Slope | Inter | Serum Female | 90 180 |
| 1.000 × + | 0 | Urine (mg/24h) | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|------------------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 75 | 76 |
| Sub | Start End | 34 | 35 |
| Factor | Endpoint Limit 2 | | |
| Blank Correction | Linear Check (%) | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| First | Start | End | Limit |
| Second | | | |
| Third | | | |

C3

| | | | |
|-------------------------|-------|---------------------|------------|
| Data Information | | Calibration | |
| Unit | MG/DL | Type | Logit.Log2 |
| Decimals | 1 | Standard | |
| Analysys | | Blank 0 | 1 cal/16 |
| Type | END | 2 cal/8 | 3 cal/4 |
| W.Length 1 | 340 | 4 cal/2 | 5 cal |
| W.Length 2 | 700 | Normal Range | |
| Method | LIA | Low | High |
| Corr | | Serum Male | 10 40 |
| Slope | Inter | Serum Female | 10 40 |
| 1.000 × + | 0 | Urine (mg/24h) | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 4 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|------------------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 75 | 76 |
| Sub | Start End | 34 | 35 |
| Factor | Endpoint Limit 2 | | |
| Blank Correction | Linear Check (%) | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| First | Start | End | Limit |
| Second | | | |
| Third | | | |

C4

| | |
|-------------------------|--------------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 660 |
| W.Length 2 | 700 |
| Method | ARSENAZO III |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | |
|---------------------|----------|
| Calibration | |
| Type | LINEAR 1 |
| Standard | |
| Blank 0 | 1 * |
| 2 | 3 |
| 4 | 5 |
| Normal Range | |
| | Low High |
| Serum Male | 8.5 10.5 |
| Serum Female | 8.5 10.5 |
| Urine (mg/24h) | |

| | |
|-------------------------|----------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 505 |
| W.Length 2 | 660 |
| Method | CHOD-PAP |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | |
|---------------------|----------|
| Calibration | |
| Type | LINEAR 1 |
| Standard | |
| Blank 0 | 1 * |
| 2 | 3 |
| 4 | 5 |
| Normal Range | |
| | Low High |
| Serum Male | 0 200 |
| Serum Female | 0 200 |
| Urine (mg/24h) | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | <input type="radio"/> Single <input type="radio"/> Double |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="radio"/> Off <input type="radio"/> ON |
| R1 Blank | <input type="radio"/> Water <input type="radio"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 32 33 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | <input type="radio"/> Single <input type="radio"/> Double |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="radio"/> Off <input type="radio"/> ON |
| R1 Blank | <input type="radio"/> Water <input type="radio"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

CA

CHOL

| | |
|-------------------------|---------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 340 |
| W.Length 2 | 405 |
| Method | KINETIC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 24 | 195 |
| | Female | 24 | 170 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|---------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 340 |
| W.Length 2 | 405 |
| Method | KINETIC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 24 |
| | Female | 0 | 24 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 8 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start 52 End 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 16 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start 52 End 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

CK NAC

CK-MB

| | |
|-------------------------|-------|
| Data Information | |
| Unit | UG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 570 |
| W.Length 2 | 700 |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 70 | 140 |
| | Female | 80 | 155 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 505 |
| W.Length 2 | 660 |
| Method | JAFFE |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0.7 | 1.4 |
| | Female | 0.6 | 1.3 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 10 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start End |
| | 32 33 |
| Sub | |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input type="checkbox"/> Single <input checked="" type="checkbox"/> Double |
| Volume | |
| Sample | 20 |
| Reagent 1 | 100 |
| Reagent 2 | 100 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start End |
| | 41 53 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

COPPER

CREATININE JAFFE

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/L |
| Decimals | 1 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|---|
| Calibration | | | |
| Type | SPLINE | | |
| Standard | | | |
| Blank 0 | 1 cal/8 | | |
| 2 cal/4 | 3 cal/2 | | |
| 4 cal | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 6 |
| | Female | 0 | 6 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | UG/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 570 |
| W.Length 2 | 800 |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|------------|----|-----|
| Calibration | | | |
| Type | Logit.Log2 | | |
| Standard | | | |
| Blank 0 | 1 cal/16 | | |
| 2 cal/8 | 3 cal/4 | | |
| 4 cal/2 | 5 cal | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 30 | 220 |
| | Female | 20 | 110 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 76 | 77 | |
| Sub | | | |
| Endpoint Limit 2 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 15 |
| Reagent 1 | 180 |
| Reagent 2 | 60 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 75 | 76 | |
| Sub | 44 | 45 | |
| Endpoint Limit 2 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

CRP

FERRITIN

| | |
|-------------------------|---------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 405 |
| W.Length 2 | 660 |
| Method | CARBOXY |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 11 | 50 |
| | Female | 7 | 32 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|---------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 505 |
| W.Length 2 | 660 |
| Method | GOD-PAP |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 70 | 115 |
| | Female | 70 | 115 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 20 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start 52 End 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start 75 End 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

GGT

GLU

| | | | | |
|-----------------|--------|---------------------|-----------------|----------|
| Unit | MG/DL | Calibration | Type | LINEAR 1 |
| Decimals | 1 | | Standard | |
| Analysys | | | Blank 0 | 1* |
| Type | END | | 2 | 3 |
| W.Length 1 | 600 | | 4 | 5 |
| W.Length 2 | 700 | Normal Range | | |
| Method | DIRECT | | Low | High |
| Corr | | Serum | Male | 35 55 |
| Slope | Inter | | Female | 45 65 |
| 1.000 × + | 0 | Urine (mg/24h) | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | <input type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | 2 |
| Reagent 1 | 180 |
| Reagent 2 | 60 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 75 | 76 | |
| Sub | 34 | 35 | |
| Endpoint Limit 3 | | Linear Check (%) | |
| Factor | | Blank Correction | |
| | | 1.000 | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

HDL

| | | | | |
|-----------------|-------|---------------------|-----------------|------------|
| Unit | MG/DL | Calibration | Type | Logit.Log2 |
| Decimals | 0 | | Standard | |
| Analysys | | | Blank 0 | 1 cal/16 |
| Type | END | | 2 cal/8 | 3 cal/4 |
| W.Length 1 | 600 | | 4 cal/2 | 5 cal |
| W.Length 2 | 700 | Normal Range | | |
| Method | LIA | | Low | High |
| Corr | | Serum | Male | 70 400 |
| Slope | Inter | | Female | 70 400 |
| 1.000 × + | 0 | Urine (mg/24h) | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | <input type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | 2 |
| Sample | 160 |
| Reagent 1 | 40 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 75 | 76 | |
| Sub | 34 | 35 | |
| Endpoint Limit 2 | | Linear Check (%) | |
| Factor | | Blank Correction | |
| | | 1.000 | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

IGA

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | 700 |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|------------|-----|------|
| Calibration | | | |
| Type | Logit.Log2 | | |
| Standard | | | |
| Blank 0 | 1 cal/16 | | |
| 2 cal/8 | 3 cal/4 | | |
| 4 cal/2 | 5 cal | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 700 | 1600 |
| | Female | 700 | 1600 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 340 |
| W.Length 2 | 700 |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|------------|----|-----|
| Calibration | | | |
| Type | Logit.Log2 | | |
| Standard | | | |
| Blank 0 | 1 cal/16 | | |
| 2 cal/8 | 3 cal/4 | | |
| 4 cal/2 | 5 cal | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 40 | 230 |
| | Female | 40 | 230 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

IGG

IGM

| | |
|-------------------------|-----------|
| Data Information | |
| Unit | UG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 570 |
| W.Length 2 | |
| Method | FERROZINE |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 35 | 168 |
| | Female | 39 | 149 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/L |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 505 |
| W.Length 2 | 660 |
| Method | LIA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|------|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1 * | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 4.5 | 19.8 |
| | Female | 4.5 | 19.8 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 20 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | Off <input type="checkbox"/> ON <input type="checkbox"/> |
| R1 Blank | Water <input type="checkbox"/> R1 - B <input type="checkbox"/> |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-----------------|------------------|----|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 75 | 76 |
| Sub | Start End | 34 | 35 |
| Endpoint Limit 3 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start End Limit | | |
| First | | | |
| Second | | | |
| Third | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | Off <input type="checkbox"/> ON <input type="checkbox"/> |
| R1 Blank | Water <input type="checkbox"/> R1 - B <input type="checkbox"/> |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-----------------|------------------|----|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start End | 46 | 47 |
| Sub | Start End | | |
| Endpoint Limit 2 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start End Limit | | |
| First | | | |
| Second | | | |
| Third | | | |

IRON

LAC

| | |
|-------------------------|-------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 340 |
| W.Length 2 | 405 |
| Method | DGKC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 530 |
| | Female | 0 | 530 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|--------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 1 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | 700 |
| Method | DIRECT |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 130 |
| | Female | 0 | 130 |
| Urine (mg/24h) | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 5 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | Start 44 End 68 |
| Sub | |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 180 |
| Reagent 2 | 60 |
| Third Mix | <input type="checkbox"/> Off <input checked="" type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low 0.000 |
| | High 3.000 |
| Main | Start 75 End 76 |
| Sub | 34 35 |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

LDH

LDL

| | |
|-------------------------|---------|
| Data Information | |
| Unit | U/L |
| Decimals | 0 |
| Analysys | |
| Type | RATE |
| W.Length 1 | 570 |
| W.Length 2 | 700 |
| Method | KINETIC |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 60 |
| | Female | 0 | 60 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|--------------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 546 |
| W.Length 2 | 700 |
| Method | XYLIDYL BLUE |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 1.8 | 2.6 |
| | Female | 1.9 | 2.5 |
| Urine (mg/24h) | | 24 | 244 |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 3 |
| Reagent 1 | 250 |
| Reagent 2 | 50 |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 44 | 60 | |
| Sub | | | |
| Endpoint Limit 3 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

| | |
|-------------------|---|
| Aspiration | |
| Kind | Single <input type="checkbox"/> Double <input checked="" type="checkbox"/> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | | | |
|----------------------|-------|------------------|-------|
| Data Process | | Abs Limit | |
| Read | Low | -0.1 | |
| | High | 3.000 | |
| Main | Start | End | |
| | 32 | 33 | |
| Sub | | | |
| Endpoint Limit 3 | | | |
| Linear Check (%) | | | |
| Factor | | | |
| Blank Correction | 1.000 | | |
| Prozone Check | | | |
| | Start | End | Limit |
| First | | | |
| Second | | | |
| Third | | | |

LIPASE

MG

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/L |
| Decimals | 1 |
| Analysys | |
| Type | END |
| W.Length 1 | 340 |
| W.Length 2 | 700 |
| Method | |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | |
|---------------------|------------|------|
| Calibration | | |
| Type | Logit.Log2 | |
| Standard | | |
| Blank 0 | 1 cal/16 | |
| 2 cal/8 | 3 cal/4 | |
| 4 cal/2 | 5 cal | |
| Normal Range | | |
| | Low High | |
| Serum | Male | 0 30 |
| | Female | 0 30 |
| Urine (mg/24h) | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 340 |
| W.Length 2 | 700 |
| Method | UV |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | |
|---------------------|----------|----------|
| Calibration | | |
| Type | LINEAR 1 | |
| Standard | | |
| Blank 0 | 1* | |
| 2 | 3 | |
| 4 | 5 | |
| Normal Range | | |
| | Low High | |
| Serum | Male | 2.6 4.5 |
| | Female | 2.6 4.5 |
| Urine (mg/24h) | | 300 1000 |

| | |
|-------------------|----------------------|
| Aspiration | |
| Kind | Single <u>Double</u> |
| Volume | |
| Sample | 5 |
| Reagent 1 | 200 |
| Reagent 2 | 50 |
| Third Mix | <u>Off</u> ON |
| R1 Blank | <u>Water</u> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|----------------------|
| Aspiration | |
| Kind | Single <u>Double</u> |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <u>Off</u> ON |
| R1 Blank | <u>Water</u> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 32 33 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

MICRO ALBUMIN

PHOS

| | |
|-------------------------|-------|
| Data Information | |
| Unit | IU/ML |
| Decimals | 1 |
| Analysys | |
| Type | END |
| W.Length 1 | 600 |
| W.Length 2 | |
| Method | UA |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | |
|---------------------|----------|
| Calibration | |
| Type | SPLINE |
| Standard | |
| Blank 0 | 1 Cal/16 |
| 2 Cal/8 | 3 Cal/4 |
| 4 Cal/2 | 5 CAL |
| Normal Range | |
| | Low High |
| Serum Male | 0 20 |
| Serum Female | 0 20 |
| Urine (mg/24h) | |

| | |
|-------------------------|--------|
| Data Information | |
| Unit | UG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 660 |
| W.Length 2 | 800 |
| Method | DIRECT |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | |
|---------------------|----------|
| Calibration | |
| Type | LINEAR 1 |
| Standard | |
| Blank 0 | 1* |
| 2 | 3 |
| 4 | 5 |
| Normal Range | |
| | Low High |
| Serum Male | 150 550 |
| Serum Female | 150 550 |
| Urine (mg/24h) | |

| | |
|-------------------|----------------------|
| Aspiration | |
| Kind | Single Double |
| Volume | |
| Sample | 3 |
| Reagent 1 | 160 |
| Reagent 2 | 40 |
| Third Mix | Off ON |
| R1 Blank | Water R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 76 77 |
| Sub | |
| Endpoint Limit 2 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|----------------------|
| Aspiration | |
| Kind | Single Double |
| Volume | |
| Sample | 20 |
| Reagent 1 | 200 |
| Reagent 2 | 60 |
| Third Mix | Off ON |
| R1 Blank | Water R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | 34 35 |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

RF

TIBC

| | |
|-------------------------|--------|
| Data Information | |
| Unit | G/DL |
| Decimals | 2 |
| Analysys | |
| Type | END |
| W.Length 1 | 546 |
| W.Length 2 | 700 |
| Method | BIURET |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|-----|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 6.6 | 8.8 |
| | Female | 6.6 | 8.8 |
| Urine (mg/24h) | | | |

| | |
|-------------------------|-------|
| Data Information | |
| Unit | MG/DL |
| Decimals | 0 |
| Analysys | |
| Type | END |
| W.Length 1 | 505 |
| W.Length 2 | 660 |
| Method | |
| Corr | |
| Slope | Inter |
| 1.000 × + | 0 |

| | | | |
|---------------------|----------|---|-----|
| Calibration | | | |
| Type | LINEAR 1 | | |
| Standard | | | |
| Blank 0 | 1* | | |
| 2 | 3 | | |
| 4 | 5 | | |
| Normal Range | | | |
| | Low High | | |
| Serum | Male | 0 | 200 |
| | Female | 0 | 200 |
| Urine (mg/24h) | | | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 5 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 32 33 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

| | |
|-------------------|--|
| Aspiration | |
| Kind | <input checked="" type="checkbox"/> Single <input type="checkbox"/> Double |
| Volume | |
| Sample | 2 |
| Reagent 1 | 200 |
| Reagent 2 | |
| Third Mix | <input type="checkbox"/> Off <input type="checkbox"/> ON |
| R1 Blank | <input checked="" type="checkbox"/> Water <input type="checkbox"/> R1 - B |
| Monitor | |
| 0 level Point | 1 |
| Span | 3.000 |

| | |
|----------------------|-----------------|
| Data Process | |
| Read | Low -0.1 |
| | High 3.000 |
| Main | 75 76 |
| Sub | |
| Endpoint Limit 3 | |
| Linear Check (%) | |
| Factor | |
| Blank Correction | 1.000 |
| Prozone Check | |
| | Start End Limit |
| First | |
| Second | |
| Third | |

TOTAL.PRO

TG

Data Information

Unit: MG/DL
Decimals: 1

Analysys

Type: RATE
W.Length 1: 340
W.Length 2: 405
Method: UV

Corr

Slope: Inter
1.000 × +: 0

Calibration

Type: LINEAR 1

Standard

| | |
|---------|-----|
| Blank 0 | 1 * |
| 2 | 3 |
| 4 | 5 |

Normal Range

| | Low | High |
|----------------|-----|------|
| Serum Male | 19 | 44 |
| Serum Female | 15 | 40 |
| Urine (mg/24h) | 26 | 43 |

Data Information

Unit: MG/DL
Decimals: 2

Analysys

Type: END
W.Length 1: 546
W.Length 2: 700
Method: PAP

Corr

Slope: Inter
1.000 × +: 0

Calibration

Type: LINEAR 1

Standard

| | |
|---------|-----|
| Blank 0 | 1 * |
| 2 | 3 |
| 4 | 5 |

Normal Range

| | Low | High |
|----------------|-----|------|
| Serum Male | 3.6 | 8.2 |
| Serum Female | 2.3 | 6.1 |
| Urine (mg/24h) | | |

Aspiration

Kind: Single Double
Volume: _____
Sample: 2
Reagent 1: 160
Reagent 2: 40

Third Mix: Off ON
R1 Blank: Water R1 - B

Monitor

0 level Point: 1
Span: 3.000

Data Process

Read: Low -0.1, High 3.000

| | | |
|------|----|----|
| Main | 41 | 53 |
| Sub | | |

Endpoint Limit 3
Linear Check (%)

Factor

Blank Correction: 1.000

Prozone Check

| | Start | End | Limit |
|--------|-------|-----|-------|
| First | | | |
| Second | | | |
| Third | | | |

Aspiration

Kind: Single Double
Volume: _____
Sample: 5
Reagent 1: 100
Reagent 2: 100

Third Mix: Off ON
R1 Blank: Water R1 - B

Monitor

0 level Point: 1
Span: 3.000

Data Process

Read: Low -0.1, High 3.000

| | | |
|------|----|----|
| Main | 60 | 61 |
| Sub | | |

Endpoint Limit 3
Linear Check (%)

Factor

Blank Correction: 1.000

Prozone Check

| | Start | End | Limit |
|--------|-------|-----|-------|
| First | | | |
| Second | | | |
| Third | | | |

UREA

URIC ACID

Data Information

Unit: MG/L
Decimals: 1

Analysys

Type: END
W.Length 1: 505
W.Length 2: 700
Method: BENZYL CHLORIDE

Corr

Slope: Inter
1.000 × +: 0

Calibration

Type: SPLINE

Standard

| | | |
|---------|-------|---------|
| Blank 0 | 1 | cal/16 |
| 2 | cal/8 | 3 cal/4 |
| 4 | cal/2 | 5 cal |

Normal Range

| | Low | High |
|----------------|-----|------|
| Serum Male | 0 | 150 |
| Serum Female | 0 | 150 |
| Urine (mg/24h) | | |

Data Information

Unit: UG/DL
Decimals: 0

Analysys

Type: END
W.Length 1: 570
W.Length 2: 700
Method: LIA

Corr

Slope: Inter
1.000 × +: 0

Calibration

Type: LINEAR 1

Standard

| | |
|---------|----|
| Blank 0 | 1* |
| 2 | 3 |
| 4 | 5 |

Normal Range

| | Low | High |
|----------------|------|------|
| Serum Male | 76.2 | 127 |
| Serum Female | 77 | 114 |
| Urine (mg/24h) | | |

Aspiration

Kind: Single Double
Volume: _____
Sample: 16
Reagent 1: 160
Reagent 2: 80

Third Mix: Off ON
R1 Blank: Water R1 - B

Monitor

0 level Point: 1
Span: 3.000

Data Process

Read: Low -0.1 High 3.000

| | | |
|------|----|----|
| Main | 75 | 76 |
| Sub | 34 | 35 |

Endpoint Limit 2
Linear Check (%)

Factor

Blank Correction: 1.000

Prozone Check

| | Start | End | Limit |
|--------|-------|-----|-------|
| First | | | |
| Second | | | |
| Third | | | |

Aspiration

Kind: Single Double
Volume: _____
Sample: 10
Reagent 1: 200
Reagent 2: _____

Third Mix: Off ON
R1 Blank: Water R1 - B

Monitor

0 level Point: 1
Span: 3.000

Data Process

Read: Low -0.1 High 3.000

| | | |
|------|----|----|
| Main | 32 | 33 |
| Sub | | |

Endpoint Limit 3
Linear Check (%)

Factor

Blank Correction: 1.000

Prozone Check

| | Start | End | Limit |
|--------|-------|-----|-------|
| First | | | |
| Second | | | |
| Third | | | |

URINE PROTEIN

ZINC